1   **Title:**

A high content microscopy screening identifies genes involved in cell width control in *Bacillus subtilis*

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Abstract:

How cells control their size is a fundamental question of biology. In bacteria, cell shape is imposed by

the extracellular cell wall, in particular by the continuous polymer of peptidoglycan (PG) that surrounds

the cell. Thus, bacterial cell morphogenesis results from the coordinated action of the proteins

assembling and degrading the PG shell. Remarkably, during steady-state growth, most bacteria maintain

a defined shape along generations, suggesting that an error-proof mechanism tightly controls the

process. In the rod-shaped model for Gram-positive bacteria, *Bacillus subtilis,* it is well known that the

average cell length varies as a function of growth rate but that cell diameter remains constant throughout

its cell cycle and across growth conditions. Here, in an attempt to shed light on the cellular circuits

controlling bacterial cell width, we developed a screen to identify genetic determinants of cell width in

B. subtilis. Using HCS (high-content screening) fluorescence microscopy and semi-automated

measurement of single-cell dimensions, we screened a library of ~ 4000 single knockout mutants. We

identified 13 mutations significantly altering cell diameter, in genes that belong to several functional

groups. In particular, these results highlight a link between cell width control and metabolism.

Introduction:

The bacterial landscape is surprisingly filled in by a variety of shape that, at the single bacterial species

level, are usually highly conserved (*1*). The rationale behind a specific shape and its selective value

remains speculative in most cases (*1*), as are the molecular mechanisms enabling a specific shape to be

determined and maintain across generations.

The shape of the cells directly depends on that of their cell wall (CW). The CW is composed of a

peptidoglycan (PG) scaffold that forms a rigid shell responsible for the mechanical properties of the cell

envelope and, in Gram-positive (G(+)) bacteria, of PG-linked glycopolymers, the most abundant being

the wall teichoic acids (WTA) (*2*). The PG sacculus is a contiguous matrix of circumferential sugar

strands crosslinked by perpendicular peptide bridges (*3*). Rod-shaped bacteria like *Bacillus subtilis* and

Escherichia coli, the models for G(+) and Gram-negative (G(-)) bacteria respectively, use two different

PG-synthesizing machineries: the divisome and the elongasome (*4, 5*). The divisome is required for cell

division to build the septum that, upon cell separation, will become the new polar caps of the resulting

daughter cells, while the elongasome synthesizes the cylindrical sidewall. The latter comprises two

machineries working independently or semi-independently: some involving aPBPs (class A penicillin

binding proteins), bifunctional enzymes with transpeptidase (TP) and transglycosylase (TG) activity,

and some called "Rod complexes" containing the TG RodA and bPBPs (class B PBPs) carrying mono-

functional TP activity such as PBP2A and PbpH in *B. subtilis* (*5, 6*). The prevailing model postulates

that the Rod complex processively builds the bulk of the organized PG meshwork, while aPBPs would

perform limited and localized, potentially unoriented, strands insertion (*6, 7*). In *B. subtilis*, and

consistent with this model, aPBPs are dispensable (*8, 9*) while most proteins of the Rod complex are

essential such as RodA (*6, 8, 10*) or co-essential as the bPBPs PBP2A and PbpH (*11*). This reflects that

a failure in the proper establishment of the PG mesh compromises the cellular integrity.

In addition to the TP and TG enzymes, the Rod complex includes other essential morphogenetic

proteins: MreC and MreD, presumed regulators of the activity of the complex (*4, 12*), and the actin

homolog MreB and in *B. subtilis* its paralog Mbl, putative coordinators of the complex recently

suggested to orient the direction of the synthetic machineries (*5, 13*). To note that the *B. subtilis* genome

encodes a third *mreB* paralog, *mreBH*, that becomes essential only in absence of the other paralogs,

under stress conditions or at low Mg concentration (*14, 15*). RodZ, a protein of unknown function, is a

suspected member of the rod-complex as well. However, despite its critical role for rod shape

maintenance in both G(-) bacteria *Caulobacter crescentus* and *E. coli,* the protein is essential only in

C. crescentus (*16-19*). Its involvement in shape control and its essentiality are even less clearly

established in G(+) bacteria. Described as essential in *B. subtilis* in an early report (*20*), several

insertional or deletion mutants have been reported since then, displaying minimal shape defects (*21-23*).

It has long been known that rod-shaped bacteria vary their cell size depending on the growth conditions

and in particular on nutrient availability (*24, 25*). Cells growing fast have a larger volume that in slow

growth condition, a relationship often referred to as the (nutrient) growth law (for a review on this topic,

see (*26*) or the very detailed (*27*)). However, while *E. coli* cell width varies greatly (up to 100%) and

concomitantly to its length (*24, 28-30*), *B. subtilis* cells adjust their length but maintain a virtually

constant diameter regardless of the growth conditions (*30-33*). Only limited variations of the width (in

the 50 nm range) have been recently reported, presumably linked to the growth phase (*34*). This

remarkable consistency suggests that cell width is a physiological parameter somehow encrypted in the

genome of *B. subtilis*, and that it must be carefully monitored during growth to correct for potential

deviations to its nominal value. Yet how rod-shaped bacteria check and balance their diameter remains

unclear.

Recently, Garner and co-workers showed how the balance between the Rod and aPBPs elongation

machineries influence cell diameter (*7*). They proposed a model in which the activity of aPBPs introduce

disorganized materials into the PG meshwork, enlarging the tight cylinder imposed by the Rod complex

(*7*). Still, in the absence of PbpA (*B. subtilis* main aPBPs, encoded by *ponA*), albeit thinner, cells remain

perfectly rod-shaped indicating that the 'check & balance' process is still in place (*35, 36*). Conversely,

alteration of the Rod complex leads to increase cell diameter, either because of the raise of the

aPBPs/Rod ratio or because of the rounding of the cells consecutive to the absence of elongation, the

two intertwined causes being difficult to differentiate. Indeed, depletion of each essential (or co-

essential) part(s) of the Rod complex, RodA, MreC, MreD, PBP2A/PbpH or MreB/Mbl/MreBH, leads

to spherical cells following the absence of PG elongation (*11, 37-40*). It is thus not surprising that all

mutants reported to display an enlarged width are involved in CW homeostasis, a phenotype frequently

accompanied by more or less dramatic shape defect such as uneven diameter, bulges, swelling and

frequently lysis. This can be observed for example with the single deletion of the *mreB* or *mbl* paralogs

(*41, 42*), or with the partial depletion of the essential MurB and TagD that catalyze key steps in the

synthesis of the PG and the teichoic acids (TA) precursors, respectively (*34, 43*).

In contrast, a limited number of mutants of *B. subtilis* display a reduced cell diameter to our knowledge.

In addition to the previously mentioned *ponA*, deletion of *ltaS* (*yflE*), *yfnI* (*yetP, ltaSA*) or *yqgS*,

encoding lipoteichoic acid synthases (*44, 45*), or the simultaneous inactivation of the cell division genes

ezrA and *gpsB* (*35*) drive to reduce cell width. Similarly, the absence of MreBH or the PG hydrolase

LytE, two proteins shown to interact in a so-called "LytE-system", was reported to lead to a slightly

reduced diameter (*14, 44, 46*). To note that curiously the simultaneous inactivation of *mreBH* and *lytE*

was reported to widen *B. subtilis* (*44*) while a *lytE* knock out displayed a wild-type width in a separate

study (*47*). Overall, the mechanism(s) at stake for cell width maintenance remains mysterious.

Here, we aimed at identifying at the genome scale level the determinants involved in cell width control

during rapid exponential growth. We screened a complete organized collection of *B. subtilis* deletion

mutants (*22*) using High-Content Screening microscopes (HCSm). Our protocol for mid-throughput

analysis allowed us to uncover several new genes that may work to maintain cell diameter. These are

involved in cell wall synthesis, cell division, metabolism, and translation, suggesting that cell width

homeostasis results from the combined action of several cellular processes. Finally, our analysis

suggests that two main routes are leading to cell width alteration of the cell.

Results

***Bacillus subtilis* displays limited width variability during rapid exponential growth**

It has long been accepted that, in contrast to *E. coli*, the cell diameter of *B. subtilis* cells remains virtually

constant regardless of the growth rate (*30, 31*). We wondered how variable the cell diameter could be in

isogenic *B. subtilis* populations during fast exponential growth. We used MicrobeJ, a FIJI plugin (*48-*

50), to perform cell segmentations and quantify their diameter (see M&M, Sup. Table 1, Sup.

Information). We first compared six independently acquired datasets of wild type *B. subtilis* cells grown

exponentially in rich LB medium (Fig. 1A). We observed that the average width remained remarkably

constant between experiments (below 2 %; Sup. Table 2). Also, the cell-to-cell variability of the

measured width in each population (standard deviation to the mean, sdm) remained low, ranging from

0.071 to 0.089 µm across the different replicates (Sup. Table 2). These variations might reflect potential

-true- differences of cell diameter or just the error of our measurements, but in either case variability

was low. This reproducibility allowed us to expect the detection of potentially small variations of cell

diameter between mutant strains. We next compared the diameter of *B. subtilis* cells exponentially

growing in two different media, rich (LB) and poor (S), and thus supporting different growth rates. In

agreement with previous reports (*30, 31*), we found no significant difference of width between cells

grown in rich and poor media (Fig. 1B). The cell-to-cell variability was similar in the two media,

indicating that this variability is independent of the growth rate. Taken together, these experiments

indicated that *B. subtilis* exerts a tight control over its diameter, whose variability remains below 2 % in

average across conditions and replicates.

HCS microscopy allows screening for small phenotypic variations of cell width

We next defined conditions that would minimize false positives in a microscopy-based screen of a

genome-scale deletion library of *B. subtilis*. First, cells were fixed to obtain snapshots of their

dimensions during exponential growth. Fixation induced a slight reduction of cell width relative to live

cells (Fig. Sup. 1A), but prevents issues resulting from the time required for the preparation of 96-well

plates and the image acquisition inherent to the HCS method. Second, the growth medium was

supplemented with 20 mM of $MgSO_4$ to prevent perturbation of growth and/or the estimation of the cell

width in mutants defective for CW synthesis that might display irregular shapes or lyse. In *B. subtilis,*

high levels of magnesium are known to reduce the activity of PG hydrolases (*51*), limiting the defects

of mutants affected in CW formation and allowing propagation of otherwise lethal mutations (*52-54*).

Importantly, while magnesium rescues most morphological defects, such mutants still present an

abnormal width (*52, 53*). Again, we noticed that the addition of Mg^{2+} to the growth medium slightly

reduced the average width of wild-type cells (Fig. Sup. 1B), as previously reported (*35*). These slight

width differences detected either when cells were fixed (Fig. Sup. 1A) or when grown in high

magnesium (Fig. Sup. 1B) confirmed our ability to detect small variations of width between populations.

Next, to demonstrate the sensitivity of our assay, we tested in our conditions the *mreB* and *ponA* mutants,

known to be wider and thinner, respectively, than wild-type cells (*36, 41*). As shown in Figure 1C, the

altered width of *mreB* and *ponA* mutants was unambiguously detected when cells were grown in high

Mg^{2+}, fixed and observed in either our conventional epifluorescence microscope or the HCSm. The cell-

to-cell variability and the average cell widths noticeably increased when measurements were performed

on HCSm-acquired images but the relative difference of width between the two mutant strains relative

to the wild-type were perfectly conserved (Fig. 1C). These control experiments showed that mutants

affected for the control of width could be identified in our medium-throughput HCSm approach.

Following up our promising preliminary controls, we screened the complete *B. subtilis* kanamycin-

marked ordered deletion library (BKK) (*22*), which contains 3983 single-gene deletion mutants of the

168 strain (GenBank Al009126), thus ~93 % coverage of its genome open reading frames (Fig. 2, see

M&M for details). In order to prevent plate-to-plate fluctuations and to compare the widths of the

mutants across plates, each mutant's width was expressed relatively to the average cell diameter per

plate (ADP (see M&M; Fig. Sup. 2A). The average of the ADPs of the 48 plates and the average cell

width of the wild-type strain grown and imaged in the same conditions showed no significant difference

(Fig. Sup. 2B). For each single mutant, we calculated the delta between its average width and the ADP

of its plate (Sup. Table 3). The 3983 Δwidth obtained present a Gaussian distribution, spreading from -

13.9 to +23.4 % but with 90 % of the values comprised in a narrow +/- 5 % variation to the mean (Fig.

3A). Next, we arbitrarily setup a cutoff of the 1 % most affected strains (0.5 % largest and 0.5 % thinnest)

(Fig. 3A). The 40 mutants selected displayed a difference in diameter > 8 % to that of their ADP (Sup.

Table 4). Using low-throughput epifluorescence microscopy imaging and the wild type *B. subtilis* strain

as a reference, we checked the phenotypes of the selected mutants (see M&M), while the deletion in

each mutant was verified by PCR.

In only half of them the wider or thinner phenotypes could be confirmed when using a low Δwidth cutoff

of 4%, suggesting the presence in our selection of several false positive (Fig. 2 and Sup. Table 4).

Nonetheless, all mutants with a confirmed deletion were kept to the next step and backcrossed into the

wild-type background before attempting further characterization, in order to prevent the possibility that

some phenotypes might be unlinked to the deletions (Fig. 2, Sup. Table 5).

Cell wall and central carbon metabolism genes are linked to cell width control

Next, we carefully measured the diameter of the 40 backcrossed mutants, using low-throughput

epifluorescence imaging (Sup. Table 4). Confirming the trend observed during the previous step, several

mutants presented only a limited width difference with that of the parental strain and were rejected.

Choosing a cautious Δwidth cutoff of 8%, we selected 12 mutants significantly and reproducibly, over

3 independent experiments, wider (7) or thinner (5) (Fig. 3B, Table 1 and Sup. Table 4). We also kept

the Δ*minJ* mutant despite its non-significant Δwidth (-1 %) because it displayed an unusually large

distribution of widths (Fig. 3B, 3D, Sup. Table 4). The width phenotypes were conserved for all 13

mutants when grown without magnesium supplementation (Sup. Table 4), confirming that at the

contrary to the bulging, swelling and lysis observed as consequences of CW synthesis impairment,

magnesium cannot bypass the alteration of width. This might suggest that the width alteration is not a

result from uncontrolled autolysin activities.

Thus, we identified nine new genes affecting cell width of *B. subtilis* and we confirmed the phenotype

of four out of the ten genes previously reported to affect cell width, present in the BKK library and

expected to be found with our method (Sup. Table 6): *rodZ, cwlO, ftsE,* and *ftsX*. To note that despite

not being in the top 1 % of genes retained for further analysis, *ponA* still displayed a significantly reduced

width in the first step of our screen (Sup. Table 3), as expected (Fig. 1C). However, *mreBH, lytE* and

the three recently spotted genes encoding lipoteichoic acid synthases (*ltaS, yqgS, yfnI*) did not display a

significant width difference in our experiment (Sup. Table 3).

The genes identified in our screen belong to different functional categories and, interestingly, a majority

are not directly involved in CW homeostasis (Table 1). In addition to the previously reported *ftsE, ftsX*,

cwlO, and *rodZ*, only *dacA* belong to this group. This gene encodes the bPBP PBP5 involved in PG

maturation (*55, 56*).

The second most represented functional category among our selected width-deficient mutants is the

metabolism (Table 1). One gene, *guaA*, is involved into purine nucleotide synthesis (encoding the GMP

synthetase (*57*)), while the four others are part of the central carbon metabolism (Fig. Sup. 3): *pyk¸*

specifying the pyruvate kinase acting in glycolysis (*58*); *ptsH,* encoding HPr, a component of the sugar

phosphotransferase system (PTS) (*59*); *panD,* involved in coenzyme A biosynthesis (*60*) and *yloR (rpe),*

probably encoding Rpe (ribulose-P-epimerase) of the pentose phosphate pathway.

Noteworthy, while most mutants involved in CW synthesis are larger (Table 1 and Sup. Table 6), four

out of five of them were thinner (Fig. 3B, 3D; Table 1). Only the *rpe* mutant displays a wider diameter

but intriguingly, the phenotype was opposite in the mutant strain from the BKK collection, a point we

will return to below. This observation might suggest that a common mechanism could be at play for

each functional category.

The three remaining genes selected are involved in cell division (*minJ* (*61, 62*)), translation (*yaaA,*

encoding a ribosome assembly factor (*63*)), or have an unknown function (*ybzH).* Using the prediction

HHPRED tool, YbzH was predicted to be a putative transcriptional repressor of the ArsR family (*64*).

ΔminJ and Δrpe display a unique phenotype of diameter instability

The mutant strains selected in our screen displayed a thinner or a larger mean diameter relative to wild-

type cells. Although their mean width differs, these mutants still control their diameter to maintain it

constant over generations. Interestingly, two mutants, Δ*rpe* and Δ*minJ*, displayed a distinctive high

dispersion of width values among their population (Fig 3, Sup. Table 4).

The Δ*rpe* mutant (BKK15790) was first selected based on its slim width (-11.4 % relative to wt) during

the systematic analysis (Sup. Table 4). Surprisingly, once backcrossed into our 168 background, the *rpe*

KO strain (RCL856) displayed the opposite phenotype with an increased average diameter (+11 %)

compared to the wild type (Fig. 4A; Sup. Table 4), nonetheless indicative of a defect of width control.

The most striking difference between the two mutants is however not their average width but their

dispersion (Fig. 4A). While the BKK *rpe-* shows a SD similar to the control strain with cells harboring

thin and regular diameters, the backcrossed mutant presents the broadest width variability (SD 0.179)

of our selection (Fig. 4A, 3B), and arguably the most dramatic loss of control. This mutant also forms

small colonies, indicative of a growth defect, a characteristic not shared with its BKK parent (Fig. Sup.

7). This suggests that the BKK strain might have acquired some suppressor mutation(s), partially

restoring its growth and shape control and reducing the width dispersion to the price of a smaller

diameter.

The second strain with an instable diameter is Δ*minJ*. This mutant of the "Min" system, a complex in

charge of controlling the placement of the divisome, is known to present a reduced septum formation

leading to long cells (*61, 65*). Although its average cell width was marginally affected (-1%), the

standard deviation was unusually large, and extreme widths ranged from 0.6 to 1.5 µm (Fig. 3B). As

observed on Fig. 4B, some filamentous Δ*minJ* cells displayed unusually large and small diameters. This

phenotype affected only a fraction of the population (~15 %), which could explain why it was not

previously reported. A unique trait of this mutant is that the diameter varied within single chains or even

single cells as well (Fig. 4B). This loss of width control was not randomly distributed along the cell

length but presents a polarity, with cells progressively widening or thinning from a given diameter,

suggestive of some kind of gradient between the poles.

RodZ: a non-essential protein involved in cell shape control

We next brought a special attention to *rodZ*, encoding a putative component of the CW machinery, whose mutant appeared wider and longer than the wild type strain in our screen (Fig. 3, Table 1, Sup. Table 4). *rodZ* was reported to be essential in *B. subtilis* (*20*) as it was shown beforehand in G(-) bacteria (*16-19*). But another team reported more recently that the inactivation of the gene led to robust growth and only mild shape defects (*23*). In addition, two collections to our knowledge include a *rodZ* inactivation mutant (*21, 22*). Facing an apparent discrepancy, here we decided to construct by homologous recombination a fresh, independent *rodZ* deletion mutant (CcBs351; see Supplementary Information). Similarly to the backcrossing of the Δ*rodZ* from the BKK library (BKK16910) into the wild-type 168 strain (RCL828) in which countless positive transformants were obtained, the mutant was readily constructed, confirming the non-essentiality of the gene in *B. subtilis* 168.

We assayed the various *rodZ* mutants to our disposal for growth and shape in different conditions. In rich LB medium, *rodZ* KO strains (CcBs351, BKK16910, RCL828) displayed no difference of growth with their parental wild type *B. subtilis* strain (Fig. 5.A). Similarly, their shape could not be easily discriminated (Fig. 5.B), and only the quantification reveals the difference of cell diameter (Fig. 5.C). We confirmed however the presence of minicells (Fig. Sup. 5B), indicative of the perturbation of the division process (*65*). When grown on the slightly poorer MSM, a casamino-acids-based medium, phenotypes of *rodZ* mutant cells worsen: cells widen, frequently divided asymmetrically (Fig. 5.B, D; Fig. Sup. 5A) and shown a solid but slightly reduced growth rate compared to that of the wild type (Fig. 5.A). This reduction of growth in MSM might not be directly linked to the richness of the medium since all strains grew almost identically in the much poorer citrate/glucose-based S medium (Fig. Sup. 6A.). Noticeably, a difference appeared between the mutants, with the BKK16910 displaying a growth lag and a reduced maximum cell density compared to the others. Since the backcrossed deletion of the BKK16910 (RCL828) showed no difference with our independently constructed deletion mutant CcBs351, we inferred that some unknown genetic differences in this strain rather than the deletion itself could be at play.

Pursuing on the idea that the genetic background could affect the phenotype of the *rodZ* mutant, and

realizing that the original mutation attempt from Muchova and coworkers was performed into the

distantly related PY79 wild type *B. subtilis*, we transfered our *rodZ* deletion into the PY79 strain

(CcBs628). However the shape defects (cell widening, asymmetric division) for *rodZ* were similar in

both the 168 and PY79 backgrounds (Fig. Sup. 5A.). The only notable difference was a strong growth

lag correlated to the richness of the medium, for the *rodZ* KO in PY79 (Fig. Sup. 6B).

Taken together, we concluded that *rodZ* is not essential for growth, that it displays shape and division

defects that worsen based on the composition of the medium (which critical factor(s) remain to be

identified) and that the genetic background influences this sensitivity to the medium.

Analysis of the mutants' growth and shape suggest distinct causes are responsible for altered cell

width

During the phenotypic characterization of the selected width-deficient mutants, we noticed that the *guaA*

mutant was unambiguously shorter than the others, while at the opposite the *minJ* mutant formed

distinctive long cells (Fig. 3D). This observation suggests that these mutants might be affected in the

control of the division process. We then systematically measured the length of the selected mutants,

revealing to our surprise that all mutants but *cwlO* displayed an average length significantly greater or

smaller than that of the wild-type cells (Fig. 3C, Table 1). While the wild type strain's average length

variability ranged from 0.1% to 5.6 % across replicates (Sup. Table 7), 11 mutants displayed an average

length >9 % larger or thinner than that of the wild-type cells (Fig. 3C; Table 1). It should be noted that

the minicells formed by Δ*minJ* and Δ*rodZ*, as previously reported (*20, 66*), were not taken into account

in our length quantification.

As shown on Figure 3, the ordering of the strains by width (3B) did not predicted their ordering by

length (3C), indicating the absence of a simple correlation between the two parameters (Fig. Sup. 4A).

However, all metabolism-related mutants but Δ*rpe* were shorter (>12 %) and thinner (>8 %) than the

wild type (Fig. 3B-D and Table 1). In contrast, shorter and wider mutants were all related to CW

metabolism, again to the exception of Δ*rpe*.

In the course of our experiments, we also noticed that the mutants selected were obviously growing at

different pace. Because the length –but not the width- of *B. subtilis* usually correlates with its growth

rate (the "growth law"), we wondered i- if the differences of length observed between the mutants would

reflect their growth and ii- if and how their width would varies regarding their growth. We determined

the generation time of the 12 mutant strains with a significantly affected width during mid exponential

growth (see Material & Method; Fig. Sup. 4B). All the metabolism related mutants, and only these,

presented an altered generation time (GT) with a >75 % increase compared to that of the wild type. For

these, the GT strongly correlates with their length (R^2=0.837) indicating that in such mutants the "growth

law" is conserved (Fig. Sup. 4C), at the contrary to CW homeostasis mutants for which the growth rate

appeared unaffected despite a change in their length (Fig. Sup. 4B-C). However, the GT was not or

poorly correlated with the width for both above-mentioned subcategories suggesting that both

parameters are not connected (Fig. Sup. 4C).

Finally, we calculated the surface area (S) to volume (V) ratio of the mutants, a parameters that was

proposed to be a key values maintained constant in a defined condition, that may be a key determinant

of the cell shape (*67*). Calculation of S/V showed that this ratio was significantly altered in every mutants

(to the exception of *minJ-*) and that the width variation, not the length, was the main contributor to the

S/V variation between mutants (Fig. Sup. 4B). As a consequence, all CW homeostasis mutants presented

a small S/V ratio while the metabolic mutants but not *rpe* displayed a larger S/V. The S/V is remarkably

constant over time for a given conditions, which has been hypothesized to originate from a scaling

between the growth rates of S and V (*68*). Harris & Theriot proposed that this scaling originates from

the rates of surface and volume growth being both functions of V (and not functions of S and V,

respectively) (*67*). A consequence of their model is that a diminishing rate of S growth, for example

when reducing the CW synthesis, both increases cell width and reduces S/V. In the frame of this model,

one could hypothesizes that the increase of width observed in the mutants we have selected may be a

direct consequence of a decreased CW incorporation affecting the S/V homeostasis.

Overall, these results discriminate between two main group of mutants with similar trends: those of

genes involved into cw homeostasis displaying an increased width and reduced S/V but an unaffected

growth rate, and mutants of metabolic genes, strongly affected for growth and presenting a reduced

width and increased S/V. In this dichotomy, *rpe* is stepping out, sharing characteristics of both groups

and suggesting that its phenotypes might reflect a defect in both pathways, which could be a cause for

its instability. Also, based on these observations, it is thus tempting to predict that *ybzH* deletion is

affecting the 'metabolic pathway' while *yaaA* is somehow affecting the 'cw homeostasis' pathway.

Discussion

The cell width is probably one of the most tightly regulated physiological parameter in *B. subtilis* (*33*).

But the mechanism allowing its fine control remains unclear. Our approach aimed at revealing in a

systematic way the (non-essential) genes involved in this process. Reassuringly, we confirmed most of

the non-essential (or non-synthetic) factors previously reported (Sup. Table 6) and identified 9 new

genes presenting a strong effect on *B. subtilis* diameter. Since we arbitrarily setup a cutoff to select the

most drastically affected mutants, it is likely that extra genes could potentially participate in the process,

along with some essential genes or genes acting synthetically. A quick survey of the top 10 % most

affected mutants (Sup. Table 3) shows a few dozen of genes involved in CW (e.g. *walH*, *pbpG*, *lytG*,

yocH, *murQ*, *murE*…), lipid metabolism (*fabI*, *lipL*, *araM*, *fadE*…) or central carbon metabolism (*tkt*,

ywjH, coaA…), albeit to a lesser degree. This complete list should be taken with cautious since the two-

step verification performed on our top 1% selection revealed a significant number of false positive.

Until now, all the genes described to affect the cell width were involved into CW synthesis or affecting

it (Sup. Table 6), and many mutants selected in the present study belongs to this group as well. In

addition to the known *cwlO*, *rodZ*, *ftsX* and *ftsE* we identified *dacA*, encoding the major vegetative DD-

carboxypeptidase PBP5, responsible for the maturation of the PG by trimming the terminal D-Ala of the

pentapeptide (*69*).

However, since the cell shape results from the proper synthesis of the PG meshwork, such mutants are

necessarily overrepresented in our study and were expected. Therefore the underlying question of this

project was: is there proteins not belonging to the CW machinery that could control or contribute to the

tight maintenance of the cell diameter? The answer is yes since most genes reported here have a known

function unrelated to it or have not been characterized yet.

In particular, we identified several known metabolic genes, *guaA*, *ptsH*, *pyk*, *rpe* and *panD*. For two of

them, *guaA* and *ptsH*, a few strings could connect them to the CW synthetic pathway. *guaA*, which

displays the smallest cell diameter of our entire study, encodes a GMP synthetase and therefore belongs

to the purine nucleotide synthetic pathway (*57*). Interestingly, it was shown decades ago that GuaA

inhibition leads to a decreased synthesis and recycling of the PG (*70*). Together, these results suggest a

tight link between purine metabolism and cell shape. The second gene, *ptsH,* encodes Hpr, a component

of the sugar phosphotransferase system (PTS). The complex is responsible for glucose uptake, and as

such is feeding the glycolysis and the PG precursor synthesis pathways (*59*). In addition, mutants of

ptsI, that encodes another piece of the PTS, were previously selected as suppressors of an *mreB* mutant

(*71*). Finally, preliminary results from our lab indicate a connection between the PTS and MreC and

MreD (Keary *et al.,* to be published), strengthening this connection. Thus, one could imagine that Hpr

could be a moon-lightening protein monitoring sugar availability to coordinate its uptake with the

elongation machinery.

Remarkably, previous studies have linked the cell metabolic status with the cell size, although in these

cases the mutants were affected for their length (reviewed in (*26*)). Out of the 17 *B. subtilis* mutants

previously reported, only *pyk,* encoding the pyruvate kinase PykA (that produces pyruvate in the final

step of glycolysis), was identified in our screen (*26*). These studies, aiming at deciphering how the

growth law (which describes the correlation between cell size and growth rate) is encoded, have revealed

how some metabolic genes are encoding enzyme acting as nutrient sensors controlling and coordinating

cell division. It should be pointed out that to a couple of exceptions the mechanisms remain unknown.

Since *B. subtilis* diameter, at the contrary to its length, is usually constant, a similar sensing mechanism

would make little sense. However, Here, one could imagine that the identified metabolic enzymes may

influence one of the two competing elongation machineries (the Rod complex and PBP1). Such effect

could be direct via a moon-lightening activity (as observed for the control of length with UgtP (*32*)) or

indirect if a metabolite influences the activity of a third-party regulatory protein (as observed with PykA

(*72*)).

It is nonetheless striking that all the metabolic genes revealed in the present screen were affected both

for width and length control. In addition, all these mutants were both thinner and shorter as if both

dimensions were somehow linked (to the exception of *rpe* that presents an uncontrolled width, with both

wider and thinner cells). Independently of these metabolic genes, another link between length and width

was found in the course of our study with the selection of the *minJ* and *rodZ* mutants. RodZ is a known

"thinning" factor (Sup. Table 6), member of the Rod complex, and it is suspected to interfere with cell

division because of the presence of anucleated minicells in a depletion mutant, a phenotype we

confirmed here in the KO strains (*20, 65*). This role was proposed to be mediated via its interaction with

MinJ, although the mechanism remains unknown (*65*). MinJ is part of the Min system, composed in

B. subtilis of MinC, MinD, MinJ and DivIVA, that controls the proper positioning of the septum (*61,*

62, 73). The phenotype of the *minJ* knockout in elongation is therefore well known but not its implication

on width. Considering the width, the phenotypes of the two mutants differ slightly, *rodZ-* forming

uniform wider cells, while *minJ-* displays this unique uncontrolled width in a fraction of the population.

Interestingly, two other mutants, encoding parts of the Min system, *divIVA* and *minC¸* presented a

somewhat thinner diameter as well in our primary screen (-6.1 % and -4.9 % respectively) (Sup. Table

3). Since MinJ interacts with both DivIVA and RodZ which have opposite effect on width control, the

mix phenotype of *minJ-* may reflect the composite result of their opposing influences. Nonetheless, it

remains to determine why the perturbation of Min affect the cell diameter. A tempting hypothesis would

be that their primary effector, the divisome, may play a role in cell elongation in such perturbed genetic

background.

Overall, our study have highlighted several new genes involved into a limited number of cellular

pathways that play a role in *B. subtilis* width control. Even though PBP1 and the Rod complex are

certainly important players in this process, these results demonstrate that other factors are required as

well. It will thus be critical to understand the function of the identified genes in hope, in the future, to

decipher the mechanism underlying the setting and the monitoring of the cellular diameter during

bacterial growth.

Materials and methods

General methods and bacterial growth conditions. Methods for growth of *B. subtilis*, transformation, selection of transformants and so on have been described extensively elsewhere (*74*). DNA manipulations were carried out by standard methods. *B. subtilis* strains used in this study are listed in Sup. Table 5. *B. subtilis* strains were grown at 30 or 37 °C in rich lysogeny broth medium (LB), except for assaying growth in poor media were strains were grown in Modified Salt Medium (MSM) supplemented with 10 mM MgSO4 (*52*) and S medium (*31*) with the corrected 1.2 µg/ml of MnSO4. For precultures, media supplements were added at the following final concentrations: $MgSO_4$ 20 mM, and genotype-depending, neomycin 15 µg.ml^{-1}, spectinomycin 100 µg.ml^{-1} or chloramphenicol 5 µg.ml^{-1} (Sup. Table 5). Transformants were selected on LB agar plates supplemented with $MgSO_4$ and neomycin. For the determination of generation time (GT), cells from overnight cultures were diluted to a fixed starting OD_{600nm} of 0.01 in fresh LB medium supplemented with $MgSO_4$ in 96-well cell culture plates (CellStar) and grown in microplate reader at 250 rpm at 37°C. GT was calculated using a Matlab script available at the following link:.

General screening procedure

Screening was performed on the BKK library using HCSm setups (see 'High content screening microscopy'), leading to the selection of 3974 out of the 3983 mutants from this collection (9 clones were absent from the published library or failed to regrow) (*22*). Images were processed and cell diameter was measured (see 'Image processing and cell size quantification'). The 1% most affected strains (40 mutants) were selected and their phenotype confirmed on an epifluorescence widefield microscope (see 'Low throughput epifluorescence microscopy' and 'Image processing and cell size quantification'). Deletions in the selected clones was verified by PCR revealing that 2 out of 40 mutants (*yoqC*, *yorP*) were wild type for the expected locus and discarded for further analysis. The remaining 38 mutants were backcrossed into the wild type 168 strain to be analyzed over triplicates experiments,

using low throughput microscopy. An arbitrary cutoff of 8% Δwidth, obtained by comparison to the wild

type strain width, was chosen and 12 genes were selected.

High content screening microscopy. Cells from overnight cultures, grown in the presence of neomycin

and $MgSO_4$, were diluted $600 \times$ in fresh LB medium supplemented with $MgSO_4$ in 96-well cell culture

plates (CellStar) and grown on an orbital shaker at 250 rpm at 37°C until mid-exponential phase (OD_{600}

~ 0.2). To fix the cells, 150 µL of culture were mixed with 50 µL of fixation solution (0.5 M KPO_4 pH

7, 8 % paraformaldehyde, 0.08 % glutaraldehyde) in 96-well PCR plates and incubated 15 min at room

temperature followed by 15 min on ice. The cells were pelleted by a 5 min centrifugation at 450 g and

the supernatant carefully removed by pipetting. The pellets were washed with 200 µL of washing buffer

(KPO_4 0.1 M pH 7) and centrifuged again. The pellets were resuspended in 20 µL of water containing

3.3 µg/mL FM1-43FX (ThermoFisher, F35355) and incubated 5 min at room temperature. 180 µL of

washing buffer were added and the cells were centrifuged a last time to be concentrated $3.75 \times$ in 40 µL

washing buffer. 96-well (Fisher) or 384-well (Greiner) microscopy plates were treated with 60 µL of

poly-L-lysine 0,01 % and washed with 60 µL of deionized water. 40 µL of cells were put in each well

and discarded after a 1 min incubation. Finally, 40 or 120 µL of deionized water were added into each

well of 96-well or 384-well plates. Imaging was performed either on an ImageXpress micro Confocal

(Molecular Devices) or an INCell 6000 analyzer (GE Healthcare) used in non-confocal mode. The

ImageXpress HCS microscope was equipped with a $60 \times$ Nikon air objective (NA 0.95), a FITC filter

(Ex.488/Em.536) and a Zyla 4.2 Andor sCMOS camera with a final pixel size of 115 nm and controlled

by MetaXpress software package. The INCell 6000 analyzer was equipped with a $60 \times$ water objective

(NA 0.95), a FITC filter (Ex.488/Em.525) and a sCMOS 5.5 Mpixels camera with a final pixel size of

108 nm and controlled by the INCell 6000 Analyzer - Acquisition v.7.1 Software. Images from 4 fields

of view were acquired for each strain.

Low throughput epifluorescence microscopy. Cultures were performed as for HCS microscopy but

in shaking tubes instead of microplates. For live cell imaging, 300 µL of cultures were directly mixed

with FM™ 1-43FX (ThermoFisher) to reach the concentration of 3.3 µg/mL and concentrated 3.75 ×

before 1 µL of the preparation was spotted onto a thin 2 % agarose-LB pad, topped by a coverslip and

immersion oil, and mounted immediately in the temperature-controlled microscope stage. For the

imaging of fixed cells, cells were fixed as described for HCS microscopy except that 300 µL of culture

were mixed with 100 µL of fixation solution and subsequently washed with 300 µL of buffer. Cells were

either loaded on a 2 % agarose-LB pad or loaded on poly-L-lysine treated 96 well microscopy plate. For

the latter, the wells were washed then filled with deionized water. Epifluorescence images of the stained

membranes were acquired on a previously described setup equipped with a $100 \times$ objective (*75*).

Image processing and cell size quantification. The post-acquisition treatment of the images was done

with the Fiji shareware and the measures (mean cell diameter and length) with the MicrobeJ plugin (*48-*

50). In MicrobeJ, the width was calculated as the mean value along the medial axis of the cell. The

parameters used for the MicrobeJ module are listed in Sup. Table 1. Cells aggregates were excluded and

segmentation was manually corrected when necessary.

During high throughput screening, the cell width of each strain was calculated as the mean of 225 cells

(in average). When the four-image set contained less than 30 measurable cells, a new acquisition was

performed. Because the library is devoid of a wild type reference, and to prevent putative plate to plate

variability, the mean cell width of each mutant was compared to the average width of all measured cells

of its 96-well plate, or ADP (average cell diameter per plate) index ($17\text{-}19.10^3$ cells/plate) (Fig. Sup.

2A). Each strain's diameter deviation relatively to this index was calculated, as:

Δwidth (%) = (mean width of the mutant – ADP) × 100 / ADP

From these differences, the 0.5^{st} and 99.5^{th} percentile were calculated and the 99 % of the mutants

between these two values eliminated.

During low-throughput microscopy (for the verification of the BKK candidates and for clones resulting

from the backcross into the 168 strain) the cell width and length of each strain was calculated as the

mean of 245 cells (in average). The calculated Δwidth was expressed by comparison with the wild type

cell size.

Alternative methods for cell width measurement. Cell widths were measured either with the

ChainTracer plugin of the Fiji shareware, or by "manually" determining the width at maximum height

on intensity profiles (*76*) (Fig. S1)**.** For ChainTracer, we used a stack of phase contrast and

epifluorescence images of stained membranes, and only analyzed isolated chains of cells to prevent

segmentation issues. For the measurement using intensity profiles, a line was manually drawn

perpendicularly to the cell's long axis, on epifluorescence images of stained membranes, and a profile

plot of the fluorescence intensity was generated. The cell diameter was determined by measuring the

distance between the two maxima.

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Figure legends

Figure 1. Discrimination of diameter-control deficient *B. subtilis* mutants: a proof of concept.

A. Comparison of cell width distribution of six independent cultures of fixed wild-type *B. subtilis* cells

grown in rich (LB) medium. One-way ANOVA statistical analysis showed no significant differences

between the replicates (Sup. Table 2). **B.** Comparative cell width distributions of fixed wild-type

B. subtilis cells grown in rich (LB) and minimal (S) media. **C.** Qualitative (images) and quantitative

(distribution of measured cell width) comparisons of data acquired on a wide-field epifluorescence

microscope and a confocal HCSm, using the wild-type, Δ*mreB* and Δ*ponA* mutant strains of *B. subtilis*.

Fluorescent images were acquired on cells grown to mid exponential phase ($0.2<OD_{600\ nm}<0.3$), fixed

and stained with the FM1-43fx membrane dye. Discrete fluorescent foci result from cell fixation. Scale

bar, 5 µm. Width distributions are displayed as violin plots with the broken line indicating the mean.

Statistical analyses were performed using (A) a one-way ANOVA or (B, C) nested t-tests. For each

statistical test, **** = $P<0.0001$, *** = $0.0001<P<0.001$, ** = $0.001<P<0.01$, * = $0.01<P<0.05$, and ns

= $P>0.05$. When significant, the difference between the means, expressed as a percent, is indicated on

the plots. B and C are compilations of at least two independent experiments.

Figure 2. Protocol summary of the screening process.

Screening of the *B. subtilis* BKK collection arrayed in 96-well plates was based on automated image

acquisition using an HCSm. The MicrobeJ plugin of FIJI was used for cell segmentation and width

measurements. Average cell diameter of each mutant was compared to the average diameter of all cells

on the plate (ADP). Candidates where confirmed by measuring their diameter on images acquired with

an epifluorescence microscope, relative to the wild-type strain. The selected mutants were then

backcrossed into a wild type background before final width determination over triplicate experiments.

Figure 3. The screen reveals 13 mutants with a cell width variation > 8 % relative to the wild-type.

A. Width difference (Δ width) of each mutant relative to the ADP of its plate. Dotted lines indicate the

cut-offs for the 0.5 % largest (top) and thinnest (bottom) mutants. Red dots mark the 13 mutants with

confirmed diameter defects after deletion were backcrossed into the wild-type strain. **B-C**. Cell width

(B) and cell length (C) distributions of the 13 selected (backcrossed) mutants. Red and green plots

correspond to thinner and wider mutants respectively, compared to their parental wild-type strain (blue).

Broken line: mean. Data are compilations of three independent experiments. The difference between the

means of each mutants and the wild-type is indicated, in %. Statistical significance between the mutant

and the wild-type strain width (B) and length (C) were calculated using nested t-tests (see Table 1 for

P-values). All differences were significant except for the mean width of the Δ*minJ* mutant (B) and the

mean length of the Δ*cwlO* mutant (C). **D.** Phenotypes of the backcrossed mutants segregate into four

classes based on their width and length defects. Δ*minJ* and Δ*rpe* mutants form both thinner (arrowhead,

1) and larger (arrowhead, 2) cells. Displayed are images of FM1-43fx membrane-labelled fixed cells.

Scale bar, 5µm.

Figure 4. Δ*minJ* and Δ*rpe* mutants exhibit unique uncontrolled diameter phenotypes.

A. Images of membrane-labelled strains and corresponding distribution of cell widths for the wild type

strain, Δ*rpe* from the BKK collection and Δ*rpe* backcrossed into 168 wild type background. Broken line:

mean. Data are compilations of two independent experiments and statistical significance between

mutants and the wild type strains was calculated using the Mann-Whitney non parametric test (**** =

P<0.0001; *** = 0.0001<P<0.001; ** = 0.001<P<0.01; * = 0.01<P<0.05; ns = P>0.05). **B.** Images of

membrane-labelled Δ*minJ* mutant, presenting varying width along single cells or chain of cells. This

phenotype is observed in approximately 15% of the population. Percentages indicate the difference of

width compared to that of the wild type strain, at the position of the red marks.

Figure 5. *rodZ* is not essential for *B. subtilis* growth but contributes to cell width control.

A. Typical growth curves of wt and *rodZ* mutants of *B. subtilis* grown in rich LB (blue) and poor MSM

(green) media. Strains are wt (168; solid line), *rodZ-* in 168 background (CcBs351; dashed line), *rodZ-*

from BKK library (BKK16910; open circles) or *rodZ-* from BKK backcrossed into 168 wt (RCL828:

dotted line). **B**. Epifluorescence images of *B. subtilis* strains (same as in A), FM1-43Fx membrane-

labelled, grown to mid exponential phase in LB or MSM media. **C-D**. Distribution of cell width of

strains (same as A) grown to mid exponential phase in rich LB (C) or poor MSM (D) media, in at least

two independent experiments. Cell widths were determined using the manual method. Statistical

analyses were performed using nested t-tests, and **** indicates a *P*-values <0.0001.

Fig. Sup. 1. Comparative cell width distributions of wild type *B. subtilis* cells measured with 3

different methods.

A. Cell width of live and fixed cells were measured using the FIJI plugins MicrobeJ and Chain tracer,

or by manual measurements (see Material and methods). The differences between the means of live and

fixed cells (in %) is specified for each method. **B.** Comparative cell width distribution of fixed wild-

type *B. subtilis* cells grown in LB with and without 20 mM magnesium supplementation. Broken line:

mean. Statistical analysis were performed using nested t-tests, were **** = P<0.0001, *** =

0.0001<P<0.001, ** = 0.001<P<0.01, * = 0.01<P<0.05, and ns = P>0.05. Differences between the

means, expressed as a percent, is indicated on the plots. Data are compilations of at least two independent

experiments.

Fig. Sup. 2. ADPs are constant across plates and equal to the width of the wild-type strain.

A. ADP of the 48 96-well plates containing the BKK library. Each ADP is the mean of all width

(~18 700) on a plate, and error bars are the standard deviations to the mean (SDM). The 48th plate has a

smaller value because the acquisition was performed on the epifluorescence microscope (with 100x

magnification) and not on the HCSm. **B.** Comparison of the average of the ADPs of all 48 plates and

the average width of a wild-type cell population measured with the HCSm. There is no significant

difference between the two values according to the Mann-Whitney non parametric test, indicating ADPs

are similar to the wild type diameter (1.160 µm vs 1.153 µm, respectively). Similarly, the ADP

calculated for the 48th plate is close to the wild-type strain measured in the corresponding microscope

(0.975 µm vs 0.964 µm, respectively). Error bars are SDM.

Fig. Sup. 3. Carbon metabolic pathways involving selected mutants deficient for cell width control.

The four genes selected in our the current screen and involved into carbo metabolism, *panD*, *ptsH*, *pyk*

and *rpe* and encoding enzymes required for the pentothenate, glycolysis and penthose pathway

respectively. PanD converts L-asp into β-alanine, the first step of the pathway leading to Coenzyme A

(Co-A) synthesis. Co-A is used in glycolysis as a substrate for Pyk to form acetyl-CoA. HPr (encoded

by *ptsH*) is a bi-functional protein acting also in glycolysis as a part of the PTS (phosphotransfer system)

required for the import/phosphorylation of sugars, and in the regulation of the carbon catabolite control,

as an allosteric regulator with CcpA. Rpe produces xylulose-5P in the penthose pathway.

Fig. Sup. 4. Relationship between generation time, length, width and surface to volume ratio in

the selected mutants. A. Average cell lengths as a function of average cell widths for each mutant.

Functional categories are color-encoded: orange for metabolism, blue for CW homeostasis and white

for others. Wild type (wt) strains is labelled in green. **B**. Average generation time (GT), length (L),

surface to volume ratio (S/V) and width (W) of cell width deficient mutants compared to the wt. The

dotted line marks the level of the average wt value. Generation time are calculated on populations (see

M&M) and are the average of 4 independent experiments. Length, S/V ratio and width data are

calculated per cell and compiled data from three independent experiments. Statistical significance of the

difference between each mutant and the wild type was calculated using either a Mann-Whitney non

parametric test (generation time) or a nested t-tests (length, S/V ratio and width) and indicated using *P*-

values, as follows: **** = P<0.0001; *** = 0.0001<P<0.001; ** = 0.001<P<0.01; * = 0.01<P<0.05; ns

= P>0.05. **C**. Average length and width as a function of the generation time (GT). Upper panel:

metabolism-related mutant; lower panel: CW homeostasis-related mutant. R² of the linear regressions

(lines) are indicated on the panels. Data are compilations of at least three independent experiments.

Fig. Sup. 5. Cell shape morphologies of several *B. subtilis rodZ* mutants and their cognate parental

wild type strains, grown in LB, MSM or S media.

A. Epifluorescence images of *B. subtilis* cells grown to mid exponential growth phase, colored with

FM1-43Fx membrane dye. Strains are wild type 168 ('wt'; 168), deleted for *rodZ* in 168 background

('351', CcBs351), *rodZ-* from BKK library ('BKK', BKK16910), *rodZ-* from BKK backcrossed into

168 wt ('BC', RCL828), wild type PY79 ('PY', PY79), and deleted for *rodZ* in PY79 background

('628', RCL628). **B**. *rodZ* mutants form minicells during growth in LB medium. Display are images of

CcBs351 grown to mid exponential growth phase in LB medium, colored with FM1-43Fx and imaged

by phase contrast (right panels) or epifluorescence (left panels) microscopy. Arrowheads point to

minicells. Scale bars (A-B): 1 µm.

Fig. Sup. 6. Growth defect of *rodZ* mutants varies depending on growth media and genetic

background.

A-B. Typical growth curves of wt and *rodZ* mutants of *B. subtilis* in rich LB (blue), poor MSM (green)

and S (orange) media. Strains are derivative of the wt 168 strain (A) or PY79 (B), and either wt (plain;

168 or PY79), or deleted for *rodZ-* (dashed; CcBs351 or RCL628). Panel A has two extra curves for

rodZ- from BKK library (circles; BKK16910) and *rodZ-* from BKK backcrossed into 168 wt (dotted;

RCL828).

C. Cell width distributions of the *rodZ* mutant (black) and its parental wild type PY79 strain (red), grown

to mid exponential phase in LB and MSM media. Difference of width between the two strains was

estimated with a nested t-test (**** = *P*-value <0.0001).

Fig. Sup. 7. Backcross of the *rpe* deletion reveals a strong growth defect.

Chromosomal DNA of strain BKK15790, K-O for *rpe*, was transformed into wild type *B. subtilis* 168.

Transformants display a 'small-colony' phenotype indicative of a growth defect. Here are shown isolates

colonies of a typical transformant (RCL856), and its parental wild type (WT; 168) and *rpe* mutant from

the BKK collection (BKK15790), streaked on LB plate and grown for 24h at 37°C.

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Epi.

HCS

WT   ΔponA   ΔmreB

Juillot et al., Sup. Figures

Sup. Fig. 1.

Screen chart

BKK (Bacillus Kanamycin knock-out) organized library

3983 single gene deletion strains on 96-well plates

for each strain

Membrane staining → **HCS microscopy** image acquisitions → **Image analysis** → **Δ width (%) determination**

Fixation

Culture LB + 20 mM MgSO₄

Microbe J

60x magnification 4 images / strain

cell segmentation and cell width measurement

relative to the ADP

for each candidate

Final validation epi. microscope

100X magnification Backcrossed mutant strain compared to 168 WT width

Backcross mutants in 168 WT strain

Phenotype confirmation epi. microscope

100X magnification BKK strain compared to 168 WT width

Data compilation, identification of the 1% most affected strains

− Wild Type width +

Sup. Fig. 2.

A

B

Fig. Sup. 3

Fig. Sup. 7

Fig. Sup. 7. Backcrossing the *rpe* deletion reveals a strong growth defect.
Chromosomal DNA of strain BKK15790, K-O for *rpe*, was transformed into wild type *B. subtilis* 168.
Transformants display a 'small-colony' phenotype indicative of a growth defect. Here are shown isolates colonies of typical transformant (RCL856), and its parental wild type (WT; 168) and *rpe* mutant from the BKK collection (BKK15790), streaked on LB plate and grown for 24h at 37°C.

Fig. 5.

Fig. Sup. 5.

Fig. Sup. 6

LB
MSM
S

WT (PY79)
rodZ- (CcBs628)
<-- alt legends

Fig. Sup. 4.

Table 1. Cellular parameters of the confirmed width-control defficient strains

Gene	Functional category[4]	δ length [1] (%)	P-value[2]	δ width [1] (%)	P-value[2]	δ GT [3] (min)	P-value[2]
minJ	cell division	120.6	****	-1.0	ns	5,6	*
ybzH	unknown (putative TR)	12.9	****	-9.3	****	1,7	*
rodZ	CW homeostasis	10.7	**	12.5	****	-1,1	ns
yaaA	translation	4.4	**	9.7	***	-0,7	ns
wt	-	0,0	-	0,0	-	0,0	
cwlO	CW homeostasis	-5.2	ns	13.1	***	0,3	*
ftsX	CW homeostasis	-8.6	**	10.0	****	-0,7	ns
ftsE	CW homeostasis	-10.9	****	9.4	****	-0,7	ns
dacA	CW homeostasis	-11.7	****	9.3	****	-1,1	ns
ptsH	central C metabolism	-12.2	****	-13.3	****	17,6	*
panD	central C metabolism	-12.5	****	-11.4	****	11,9	*
rpe	central C metabolism	-13.6	***	11.0	****	12,8	*
pyk	central C metabolism	-21.4	****	-8.3	***	36,5	*
guaA	purine metabolism	-36.7	****	-12.7	****	38,1	*

Colors of numbers indicate positively (purple) or negatively (orange) affected value in width or length, or inreased or decreased generation time, compared to the wild type strain
1: difference (in %) relative to wild type, average of 3 independant pooled replicates
2 : statistical significance estimated by nested t-test (δlength, δwidth) or Mann-Whitney test (GT)
P-values are displayed as follow: **** = P<0.0001; *** = 0.0001<P<0.001; ** = 0.001<P<0.01; * = 0.01<P<0.05; ns = P>0.05.
3: difference of generation time (GT) relative to the wild type, averages of 4 independent experiments
4: according to Subtiwiki (Zhu, 2018)

Sup. Table 1. Settings used for MicrobeJ pluggin

Particle selection parameters

Area [μm²]:	3.7-20
Length [μm]:	0-max
Width [μm]:	0.5-2
- Range [μm]:	0-max
- Variation:	0-0.15
Circularity:	0.3-0.7
Curvature:	0-max
Sinuosity:	0-max
Angularity [rad]:	0-0.25
Solidity:	0.85-max
Intensity:	0-max

Segmentation method
Dark, Otsu, median axis

Treatment

Thresholding:	Use ROI (enabled)
Resampling:	Resolution: 0.5, Method: Bilinear
Other:	Include Holes (enabled)
Pre-Processing (enabled):	Subtract Background, rolling=12
Threshold Calculator (enabled):	Binary, Dark, Otsu, offset: 0, area: 1000-max

Sup. Table 2. Average width differences (%) across replicates[1]

	WT 1	WT 2	WT 3	WT 4	WT 5	WT 6
WT 1	-	+1.7 %	+0.8 %	+0.5 %	+0.2 %	+0.1 %
WT 2	-	-	-0.9 %	-1.2 %	-1.5 %	-1.6 %
WT 3	-	-	-	-0.3 %	-0.6 %	-0.7 %
WT 4	-	-	-	-	-0.4 %	-0.4 %
WT 5	-	-	-	-	-	-0.1 %
width (µm)	1.01	1.028	1.012	1.016	1.013	1.012
+/-	0.076	0.080	0.071	0.076	0.089	0.071

1: ordinary one-way ANOVA test between replicates conclude to no significant differences

Sup. Table 3: Cell width of mutants of the BKK collection

BKK name[1]	gene	screening delta[2] (%)	erage width (μ	+/-	nb	ADP
BKK34800	cwlO	23,357	1,418	0,136	301	1,149
BKK12760	xkdW	21,842	1,421	0,125	150	1,166
BKK35260	ftsE	19,51	1,383	0,153	109	1,158
BKK22849	ypzH	15,994	1,276	0,133	401	1,100
BKK00100	dacA	15,94	1,342	0,149	107	1,158
BKK35250	ftsX	15,586	1,329	0,183	286	1,149
BKK20680	yoqC	13,238	1,323	0,180	411	1,168
BKK16825	ymfD	12,668	1,309	0,172	82	1,162
BKK27320	greA	12,654	1,098	0,100	40	0,975
BKK35450	comFC	11,474	1,242	0,102	90	1,114
BKK16910	rodZ	11,399	1,294	0,108	287	1,162
BKK31070	yuaC	11,327	1,293	0,130	238	1,161
BKK12770	xkdX	10,489	1,289	0,121	135	1,166
BKK00030	yaaA	9,941	1,283	0,122	189	1,167
BKK17000	kbl	9,825	1,276	0,164	152	1,162
BKK35440	yvyF	9,578	1,221	0,111	161	1,114
BKK23990	yqiW	9,495	1,263	0,141	142	1,153
BKK35220	minJ	9,381	1,242	0,131	104	1,135
BKK13030	ykhA	9,169	1,273	0,183	109	1,166
BKK02750	natA	9,096	1,311	0,124	127	1,201
BKK19550	yodC	9,083	1,232	0,110	425	1,129
BKK18230	yngG	8,914	1,276	0,084	255	1,172
BKK36280	ywqA	8,891	1,213	0,146	347	1,114
BKK15450	lspA	8,741	1,210	0,108	210	1,113
BKK28800	araA	8,593	1,209	0,100	31	1,113
BKK15730	fmt	8,544	1,058	0,086	72	0,975
BKK08690	ygaD	8,54	1,312	0,134	34	1,208
BKK05344	ydzS	8,47	1,266	0,099	161	1,167
BKK35520	yvhJ	8,434	1,246	0,130	196	1,149
BKK16660	truB	8,397	1,261	0,121	75	1,164
BKK34110	rsbP	8,384	1,230	0,109	124	1,135
BKK36130	ywrA	8,371	1,057	0,089	33	0,975
BKK19440	norM	8,352	1,224	0,130	613	1,129
BKK23520	fur	8,293	1,249	0,140	93	1,153
BKK36269	ywzD	8,241	1,252	0,112	101	1,157
BKK29280	ytnM	8,103	1,203	0,118	155	1,113
BKK12619	ykzL	7,983	1,282	0,140	81	1,187
BKK14750	ylaE	7,862	1,272	0,183	72	1,179
BKK09840	hemZ	7,816	1,269	0,127	140	1,177
BKK27290	yrzA	7,775	1,268	0,128	45	1,177
BKK34760	yvcK	7,743	1,223	0,105	271	1,135
BKK12850	ykaA	7,694	1,252	0,130	271	1,163
BKK19370	odhA	7,657	1,216	0,124	315	1,129
BKK31580	maeN	7,642	1,236	0,099	103	1,148
BKK13280	ykoJ	7,628	1,255	0,166	168	1,166
BKK31570	yufQ	7,616	1,315	0,129	119	1,222
BKK38229	slrA	7,495	1,203	0,112	144	1,119
BKK23250	ribH	7,444	1,275	0,102	339	1,187
BKK18320	ppsC	7,406	1,258	0,113	91	1,172
BKK04410	ydbB	7,404	1,251	0,109	372	1,165
BKK01950	skfE	7,358	1,222	0,097	31	1,138
BKK00330	yobA	7,321	1,265	0,127	430	1,179
BKK18150	xynC	7,264	1,257	0,112	231	1,177
BKK18130	eglS	7,223	1,256	0,116	231	1,172
BKK33710	opuBC	7,139	1,232	0,108	486	1,149
BKK39020	yxjA	7,131	1,293	0,161	103	1,206
BKK14570	ykyA	7,131	1,263	0,180	32	1,179
BKK29730	ytxD	7,116	1,192	0,093	30	1,113
BKK30050	ytgP	7,092	1,309	0,213	112	1,222
BKK22770	mtrB	7,077	1,235	0,143	232	1,153
BKK24930	yqzD	7,045	1,246	0,104	415	1,164
BKK01700	murQ	7,045	1,218	0,146	34	1,138
BKK12660	xkzB	7,034	1,273	0,176	40	1,189
BKK26720	azlB	6,983	1,259	0,109	392	1,177
BKK22620	hisC	6,963	1,177	0,121	304	1,100
BKK05330	aseR	6,954	1,248	0,068	122	1,167
BKK19510	yojB	6,876	1,242	0,141	243	1,162
BKK03040	amyE	6,864	1,208	0,137	60	1,130
BKK03651	bsdD	6,852	1,245	0,102	150	1,165
BKK34130	ganA	6,791	1,212	0,115	30	1,135
BKK02610	ycbR	6,782	1,207	0,128	56	1,130
BKK04770	ydcH	6,741	1,243	0,138	298	1,165
BKK22530	ypjA	6,735	1,216	0,103	76	1,139
BKK28450	sdhC	6,734	1,238	0,097	32	1,160
BKK31370	yugJ	6,723	1,225	0,095	618	1,148
BKK34560	malL	6,715	1,211	0,142	116	1,135
BKK39360	hutU	6,694	1,244	0,152	105	1,166
BKK40060	ycsG	6,653	1,242	0,082	146	1,165
BKK35150	yvzB	6,63	1,210	0,141	62	1,135

BKK name[1]	gene	screening delta[2] (%)	erage width (μ	+/-	nb	ADP
BKK19390	yojN	6,61	1,204	0,116	324	1,129
BKK36380	rapD	6,61	1,188	0,116	769	1,114
BKK10100	yhgC	6,591	1,255	0,115	206	1,177
BKK37580	ywfH	6,58	1,192	0,099	116	1,119
BKK25250	ccpN	6,574	1,240	0,106	98	1,164
BKK19530	yodA	6,542	1,203	0,155	502	1,129
BKK36390	flhP	6,537	1,187	0,123	160	1,114
BKK36860	atpE	6,487	1,222	0,128	253	1,147
BKK25840	phrE	6,454	1,038	0,095	43	0,975
BKK23160	rluB	6,451	1,264	0,105	212	1,187
BKK13840	stoA	6,443	1,242	0,123	39	1,166
BKK31150	uppP	6,429	1,236	0,110	286	1,161
BKK11590	yjbL	6,402	1,265	0,169	516	1,189
BKK26590	blt	6,37	1,300	0,088	72	1,222
BKK14090	ykuI	6,306	1,240	0,133	387	1,166
BKK29400	ytlI	6,306	1,235	0,108	257	1,161
BKK10210	yhfF	6,304	1,251	0,109	116	1,177
BKK05450	ydfK	6,302	1,241	0,116	146	1,167
BKK21750	scuA	6,3	1,226	0,142	215	1,153
BKK15570	cysH	6,219	1,219	0,099	35	1,148
BKK15270	sbp	6,206	1,252	0,143	140	1,179
BKK32080	yuiB	6,197	1,219	0,079	152	1,148
BKK00610	yabQ	6,179	1,244	0,108	94	1,172
BKK13140	ohrA	6,162	1,234	0,146	212	1,163
BKK37350	sboA	6,153	1,188	0,110	116	1,119
BKK40290	yycN	6,108	1,221	0,122	299	1,151
BKK24220	spo0A	6,107	1,257	0,101	161	1,184
BKK15590	sat	6,105	1,239	0,130	241	1,168
BKK09490	yhdJ	6,083	1,249	0,121	86	1,177
BKK23030	ypbB	6,074	1,221	0,124	294	1,151
BKK09620	yhdW	6,056	1,248	0,117	148	1,177
BKK22990	ypbF	6,034	1,167	0,099	570	1,100
BKK05650	ydgH	6,024	1,238	0,126	149	1,167
BKK18400	yoeD	6,008	1,242	0,087	327	1,172
BKK18250	yngI	5,923	1,241	0,117	178	1,172
BKK38480	ywaC	5,912	1,185	0,127	30	1,119
BKK05390	ydfF	5,893	1,236	0,090	391	1,167
BKK04460	dctB	5,887	1,233	0,128	392	1,165
BKK24720	comGB	5,883	1,232	0,124	251	1,164
BKK15780	cpgA	5,86	1,032	0,156	41	0,975
BKK32100	yumB	5,847	1,215	0,095	63	1,148
BKK17890	tkt	5,816	1,229	0,120	550	1,162
BKK34730	yvcN	5,799	1,201	0,124	592	1,135
BKK25700	yqeF	5,798	1,220	0,125	367	1,153
BKK40980	yyaB	5,775	1,253	0,116	97	1,184
BKK22660	trpC	5,766	1,217	0,127	348	1,151
BKK20650	yoqF	5,748	1,221	0,130	194	1,155
BKK03820	yclP	5,726	1,203	0,122	89	1,138
BKK15620	sirB	5,71	1,213	0,092	100	1,148
BKK06220	ydjJ	5,698	1,246	0,111	285	1,179
BKK23460	spoIIAB	5,696	1,275	0,161	93	1,206
BKK13810	ykvS	5,695	1,229	0,136	221	1,163
BKK23120	resD	5,674	1,163	0,099	350	1,100
BKK23510	ripX	5,671	1,254	0,095	148	1,187
BKK05490	ydfO	5,658	1,233	0,115	151	1,167
BKK29870	ytzB	5,644	1,030	0,079	100	0,975
BKK35820	gerBC	5,632	1,214	0,107	408	1,149
BKK31500	yuxK	5,625	1,212	0,147	56	1,148
BKK40050	yyaM	5,597	1,233	0,148	144	1,167
BKK00050	yaaB	5,574	1,245	0,126	178	1,179
BKK25650	yqeI	5,559	1,228	0,099	346	1,164
BKK08820	katA	5,529	1,242	0,124	162	1,177
BKK11300	med	5,512	1,249	0,132	67	1,183
BKK30740	mntD	5,498	1,214	0,123	218	1,151
BKK35670	gtaB	5,489	1,175	0,131	218	1,114
BKK26060	yqbM	5,483	1,241	0,104	292	1,177
BKK34910	hisD	5,475	1,197	0,092	377	1,135
BKK35570	tuaF	5,475	1,212	0,132	504	1,135
BKK37460	rapF	5,469	1,180	0,131	258	1,119
BKK10300	aprE	5,445	1,248	0,141	117	1,183
BKK03490	srfAB	5,444	1,028	0,085	80	0,975
BKK11560	nasE	5,431	1,191	0,145	37	1,130
BKK22700	aroB	5,422	1,249	0,114	132	1,184
BKK02670	lmrB	5,417	1,200	0,124	52	1,138
BKK21030	yonP	5,411	1,216	0,117	398	1,153
BKK27480	yrrC	5,403	1,222	0,105	146	1,160
BKK04130	yczJ	5,388	1,228	0,130	143	1,165
BKK22630	trpA	5,379	1,159	0,107	118	1,100

BKK name[1]	gene	screening delta[2] (%)	erage width (μ	+/-	nb	ADP
BKK39260	bglH	5,364	1,227	0,120	79	1,165
BKK34870	hisF	5,362	1,196	0,100	246	1,135
BKK40579	yyzL	5,346	1,229	0,128	85	1,166
BKK29410	ytkL	5,34	1,223	0,128	410	1,161
BKK40260	yycQ	5,338	1,213	0,115	203	1,151
BKK09340	yhdA	5,327	1,240	0,101	102	1,177
BKK33630	secG	5,322	1,211	0,090	354	1,149
BKK04070	ycsI	5,301	1,227	0,100	92	1,165
BKK02960	yceJ	5,29	1,265	0,148	47	1,201
BKK11580	yjbK	5,282	1,252	0,132	921	1,189
BKK10950	yitD	5,271	1,246	0,123	70	1,183
BKK09990	scoC	5,267	1,239	0,105	67	1,177
BKK19540	yodB	5,241	1,188	0,117	565	1,129
BKK04450	dctS	5,231	1,264	0,094	65	1,201
BKK25380	yqfA	5,211	1,224	0,105	148	1,164
BKK32820	fadE	5,205	1,227	0,114	96	1,166
BKK14810	ylaK	5,201	1,240	0,071	86	1,179
BKK12671	xkdN	5,181	1,251	0,174	77	1,189
BKK33130	lial	5,172	1,193	0,118	62	1,134
BKK03410	bglC	5,146	1,188	0,133	38	1,130
BKK37110	ywjH	5,144	1,176	0,141	115	1,119
BKK19669	yoyF	5,14	1,187	0,133	315	1,129
BKK02880	yceB	5,132	1,188	0,130	67	1,130
BKK04930	yddD	5,113	1,217	0,092	96	1,158
BKK27490	yrrB	5,109	1,219	0,110	148	1,160
BKK39810	csbC	5,096	1,208	0,114	422	1,149
BKK01830	ndhF	5,086	1,219	0,107	274	1,160
BKK05720	ydhE	5,085	1,227	0,081	366	1,167
BKK14510	ktrC	5,078	1,239	0,097	359	1,179
BKK19520	yoja	5,07	1,187	0,116	686	1,129
BKK00170	yaaI	5,06	1,218	0,124	606	1,160
BKK16160	xynD	5,052	1,231	0,115	249	1,172
BKK05350	ydfB	5,036	1,226	0,075	179	1,167
BKK16990	tdh	5,036	1,222	0,105	110	1,164
BKK38440	ywaF	5,015	1,175	0,125	32	1,119
BKK29890	ytnP	5,002	1,169	0,121	39	1,113
BKK11630	prpE	4,999	1,248	0,140	803	1,189
BKK23402	spoVAEB	4,99	1,267	0,187	36	1,206
BKK14500	ykqA	4,976	1,238	0,116	301	1,179
BKK16860	ymfH	4,967	1,221	0,127	82	1,164
BKK22740	hepT	4,961	1,155	0,101	110	1,100
BKK37170	acdA	4,938	1,224	0,116	89	1,166
BKK01800	alkA	4,929	1,217	0,104	324	1,160
BKK09370	lytF	4,921	1,268	0,136	47	1,208
BKK17440	ynbB	4,909	1,221	0,114	160	1,164
BKK06630	yerH	4,908	1,260	0,073	89	1,201
BKK05130	ydeB	4,893	1,224	0,112	366	1,167
BKK28320	ysnD	4,891	1,216	0,103	447	1,160
BKK30170	ytcP	4,889	1,282	0,140	33	1,222
BKK26700	azlD	4,888	1,282	0,078	201	1,222
BKK34880	hisA	4,887	1,191	0,086	270	1,135
BKK24450	efp	4,884	1,245	0,180	402	1,187
BKK29510	ytfI	4,882	1,167	0,110	110	1,119
BKK31740	yuxH	4,882	1,204	0,092	121	1,148
BKK35020	nogBA	4,879	1,190	0,123	126	1,135
BKK35679	yvzH	4,867	1,205	0,122	239	1,149
BKK11700	thiF	4,86	1,247	0,143	929	1,189
BKK11990	yjdB	4,847	1,247	0,180	70	1,189
BKK28750	araN	4,846	1,281	0,078	69	1,222
BKK03810	yclO	4,844	1,260	0,073	113	1,201
BKK22040	ypbQ	4,843	1,154	0,147	35	1,100
BKK21040	yonO	4,833	1,209	0,108	308	1,153
BKK18430	yngN	4,827	1,228	0,086	84	1,172
BKK09900	natB	4,82	1,234	0,091	98	1,177
BKK08920	yhaJ	4,802	1,218	0,120	207	1,163
BKK09965	yhaJ	4,797	1,233	0,131	93	1,177
BKK35390	yvnA	4,79	1,189	0,123	430	1,135
BKK24510	yghO	4,783	1,206	0,112	311	1,151
BKK40000	yxnA	4,779	1,202	0,121	256	1,147
BKK00320	yaaT	4,777	1,235	0,114	214	1,179
BKK29500	ytfJ	4,769	1,166	0,100	181	1,113
BKK16120	topA	4,768	1,241	0,126	110	1,184
BKK34400	padC	4,765	1,189	0,096	68	1,135
BKK24680	comGF	4,754	1,208	0,140	178	1,153
BKK12090	cotT	4,749	1,245	0,203	112	1,189
BKK24500	yghP	4,746	1,206	0,106	404	1,151
BKK33670	rghRB	4,743	1,188	0,139	33	1,134
BKK03050	ldh	4,742	1,184	0,127	46	1,130
BKK21720	ypmT	4,739	1,206	0,132	237	1,151

BKK name[1]	gene	screening delta[2] (%)	erage width (μ	+/-	nb	ADP
BKK10690	gerPD	4,734	1,239	0,107	93	1,183
BKK27580	yrvJ	4,725	1,280	0,064	102	1,222
BKK27710	tgt	4,718	1,214	0,101	352	1,160
BKK29920	ytmP	4,71	1,166	0,128	41	1,113
BKK20690	yoqB	4,7	1,209	0,112	317	1,155
BKK26580	bltR	4,698	1,279	0,110	56	1,222
BKK06076	ydzW	4,693	1,222	0,146	449	1,167
BKK05408	ydzP	4,67	1,193	0,138	32	1,139
BKK12380	uxaB	4,666	1,244	0,170	666	1,189
BKK06200	ydjH	4,662	1,234	0,129	191	1,179
BKK05640	ydgG	4,653	1,222	0,092	289	1,167
BKK27930	spo0B	4,648	1,207	0,145	126	1,153
BKK12600	xkdF	4,635	1,244	0,125	181	1,189
BKK27400	yrzL	4,631	1,213	0,118	464	1,160
BKK14830	glsB	4,626	1,234	0,082	201	1,179
BKK34180	yvfI	4,619	1,188	0,101	230	1,135
BKK09020	yhcB	4,618	1,231	0,109	97	1,177
BKK33180	yvrC	4,613	1,278	0,109	77	1,222
BKK27600	relA	4,606	1,213	0,098	421	1,160
BKK38530	dltD	4,605	1,202	0,166	142	1,149
BKK01845	ybcC	4,599	1,213	0,104	268	1,160
BKK37320	narK	4,597	1,200	0,108	349	1,147
BKK28650	ysgA	4,59	1,164	0,117	121	1,113
BKK18019	ynzL	4,588	1,217	0,119	105	1,164
BKK07480	yfmG	4,582	1,212	0,114	333	1,159
BKK24950	pstBB	4,578	1,262	0,148	150	1,206
BKK34380	slrR	4,578	1,187	0,087	234	1,135
BKK40480	ydbI	4,563	1,256	0,091	47	1,201
BKK28620	rnhC	4,561	1,164	0,099	170	1,113
BKK30390	bceS	4,561	1,278	0,090	74	1,222
BKK13450	sigI	4,549	1,219	0,136	43	1,166
BKK02760	natB	4,547	1,256	0,177	90	1,201
BKK37960	ywdH	4,547	1,200	0,114	436	1,147
BKK20310	yorO	4,516	1,214	0,109	191	1,162
BKK23260	ribAB	4,514	1,241	0,121	254	1,187
BKK17430	ynbA	4,513	1,216	0,109	71	1,164
BKK35410	flgK	4,507	1,165	0,107	1066	1,114
BKK02980	opuAA	4,495	1,255	0,132	69	1,201
BKK06180	pspA	4,493	1,232	0,117	192	1,179
BKK40560	yybP	4,489	1,219	0,123	132	1,166
BKK03840	ycnB	4,477	1,217	0,105	221	1,165
BKK33660	rghRA	4,473	1,019	0,076	55	0,975
BKK09230	yhcV	4,472	1,230	0,096	153	1,177
BKK16000	ylqC	4,461	1,169	0,120	241	1,119
BKK22380	ypmB	4,45	1,149	0,105	218	1,100
BKK14270	moeB	4,448	1,232	0,109	344	1,179
BKK31900	yukD	4,448	1,199	0,111	148	1,148
BKK23680	yqjZ	4,446	1,202	0,112	340	1,151
BKK20700	yoqA	4,444	1,206	0,116	398	1,155
BKK31700	comX	4,443	1,218	0,102	223	1,165
BKK04830	sacV	4,439	1,217	0,091	355	1,165
BKK22230	purT	4,437	1,180	0,123	85	1,130
BKK11670	thiO	4,437	1,242	0,153	883	1,189
BKK04810	inmA	4,435	1,217	0,123	156	1,165
BKK40210	yydC	4,434	1,218	0,093	47	1,166
BKK12080	ctaO	4,427	1,242	0,183	156	1,189
BKK17010	ymcB	4,416	1,215	0,120	146	1,164
BKK06340	yeaD	4,414	1,198	0,134	301	1,147
BKK23270	ribE	4,409	1,239	0,103	363	1,187
BKK27370	yrrL	4,406	1,211	0,105	221	1,160
BKK05290	ydeP	4,398	1,219	0,084	214	1,167
BKK03380	yckB	4,396	1,254	0,130	282	1,201
BKK25460	dnaJ	4,382	1,215	0,126	211	1,164
BKK01900	ybcM	4,373	1,162	0,112	35	1,113
BKK06380	yebC	4,371	1,230	0,144	368	1,179
BKK26259	yqzO	4,358	1,228	0,100	128	1,177
BKK35210	yvkA	4,356	1,200	0,117	376	1,149
BKK37000	prmC	4,352	1,163	0,130	78	1,114
BKK38018	ywzG	4,346	1,217	0,127	71	1,166
BKK28700	ysfE	4,331	1,161	0,078	32	1,113
BKK38850	yxkC	4,329	1,215	0,105	65	1,165
BKK10390	yhfW	4,319	1,234	0,091	176	1,183
BKK03788	yczM	4,309	1,207	0,138	43	1,158
BKK14590	pdhB	4,304	1,230	0,148	301	1,179
BKK02780	ycdA	4,3	1,253	0,212	33	1,201
BKK08860	ssuD	4,292	1,227	0,109	234	1,177
BKK16740	spoVFB	4,285	1,213	0,122	69	1,164
BKK20670	yoqD	4,284	1,204	0,124	286	1,155
BKK34150	ganP	4,283	1,184	0,115	62	1,135

Sup. Table 3: Cell width of mutants of the BKK collection (continued)

BKK name[1]	gene	screening delta[2] (%)	average width (μ)	+/-	nb	ADP
BKK00340	yabB	4,272	1,230	0,152	79	1,179
BKK26150	yqbD	4,272	1,017	0,091	96	0,975
BKK29040	ytbD	4,268	1,274	0,076	114	1,222
BKK39070	bglS	4,253	1,214	0,137	96	1,165
BKK01990	ybdG	4,252	1,178	0,132	69	1,130
BKK35940	rbsA	4,231	1,198	0,130	336	1,149
BKK29050	ytbE	4,217	1,160	0,111	70	1,113
BKK14370	yknZ	4,216	1,229	0,120	77	1,179
BKK27785	yrzF	4,209	1,208	0,099	405	1,160
BKK25900	cwlA	4,196	1,273	0,073	192	1,222
BKK20250	mtbP	4,195	1,211	0,127	199	1,162
BKK13680	motB	4,193	1,217	0,122	296	1,168
BKK31230	tlpB	4,193	1,204	0,131	144	1,156
BKK17330	miaA	4,187	1,212	0,128	109	1,164
BKK34729	yvzJ	4,186	1,198	0,122	289	1,149
BKK22000	sspL	4,185	1,203	0,115	238	1,155
BKK09130	tcyP	4,177	1,226	0,117	77	1,177
BKK10060	ecsC	4,17	1,217	0,120	300	1,168
BKK38400	epr	4,17	1,195	0,112	399	1,147
BKK15370	ylmD	4,169	1,187	0,122	53	1,139
BKK25760	spoIVCB	4,165	1,273	0,102	232	1,222
BKK00810	dusB	4,162	1,208	0,126	361	1,160
BKK31350	pgi	4,162	1,196	0,120	103	1,148
BKK12000	manR	4,158	1,238	0,171	444	1,189
BKK11180	yitZ	4,147	1,232	0,136	118	1,183
BKK24620	tasA	4,144	1,015	0,117	64	0,975
BKK05340	ydfA	4,14	1,015	0,085	47	0,975
BKK19400	sodC	4,136	1,176	0,131	489	1,129
BKK05140	ydzE	4,135	1,208	0,127	633	1,160
BKK20210	yorY	4,135	1,176	0,134	88	1,129
BKK36010	alsS	4,13	1,160	0,114	47	1,114
BKK37430	albG	4,125	1,197	0,114	403	1,149
BKK12400	yjnA	4,11	1,238	0,150	551	1,189
BKK19570	yodE	4,104	1,176	0,098	440	1,129
BKK39600	yxeC	4,103	1,195	0,107	270	1,147
BKK40529	yyzH	4,103	1,214	0,118	105	1,166
BKK30000	ythQ	4,095	1,272	0,174	141	1,222
BKK25110	yqfU	4,094	1,211	0,103	647	1,164
BKK07770	yfkS	4,086	1,216	0,129	357	1,168
BKK12190	yjhB	4,08	1,238	0,095	267	1,189
BKK07860	yfkL	4,073	1,258	0,161	100	1,208
BKK31100	ktrB	4,058	1,208	0,137	223	1,161
BKK02350	nagP	4,056	1,176	0,128	139	1,130
BKK08880	rpsNB	4,054	1,225	0,099	151	1,177
BKK33210	yvrG	4,052	1,214	0,105	121	1,166
BKK29270	ytaF	4,045	1,203	0,099	79	1,156
BKK13819	ykzS	4,043	1,214	0,108	259	1,166
BKK24970	pstA	4,033	1,255	0,156	51	1,206
BKK23210	scpB	4,03	1,232	0,111	79	1,184
BKK02010	ybdK	4,027	1,184	0,128	137	1,138
BKK28310	ilvB	4,024	1,206	0,119	295	1,160
BKK34210 251	yvfG	4,022	1,181	0,108	85	1,135
BKK27910	yufC	4,01	1,206	0,121	200	1,160
BKK12980	ykfB	4,008	1,257	0,101	35	1,208
BKK23320	ypzC	4,001	1,235	0,111	107	1,187
BKK26890	csn	3,998	1,224	0,108	318	1,177
BKK24890	yqgN	3,985	1,210	0,099	280	1,164
BKK40970	parA	3,973	1,232	0,104	130	1,184
BKK32120	yuzG	3,965	1,193	0,110	30	1,148
BKK10009	yhzF	3,952	1,223	0,105	148	1,177
BKK07860	yflH	3,95	1,205	0,119	206	1,159
BKK33830	opuCA	3,95	1,195	0,110	332	1,149
BKK24850	gcvPA	3,945	1,234	0,099	271	1,187
BKK30430	ytrD	3,931	1,207	0,113	280	1,161
BKK30340	ytvA	3,929	1,207	0,114	277	1,161
BKK07750	yflA	3,917	1,214	0,129	253	1,168
BKK09889	yhzD	3,905	1,223	0,126	134	1,177
BKK37560	thrZ	3,9	1,162	0,101	154	1,119
BKK08800	ygaK	3,887	1,204	0,127	394	1,159
BKK14260	mobA	3,883	1,225	0,121	282	1,179
BKK11540	pepF	3,879	1,229	0,113	89	1,183
BKK25289	ygzL	3,878	1,209	0,094	332	1,164
BKK07280	yfnG	3,874	1,225	0,149	108	1,179
BKK10620	addB	3,874	1,229	0,102	114	1,183
BKK30490	ytqB	3,873	1,201	0,104	140	1,156
BKK08210	malQ	3,87	1,255	0,111	31	1,208
BKK20460	yoqZ	3,87	1,173	0,157	67	1,129
BKK18610	yoaH	3,867	1,217	0,113	138	1,172
BKK33430	cysI	3,864	1,013	0,078	41	0,975
BKK06160	gutP	3,856	1,224	0,143	116	1,179
BKK03220	ycgO	3,843	1,173	0,104	114	1,130
BKK35460	comFB	3,826	1,157	0,090	103	1,114
BKK18239	yngHB	3,824	1,216	0,086	342	1,172
BKK15390	sepF	3,82	1,213	0,119	153	1,168
BKK08050	yfjL	3,818	1,254	0,139	74	1,208
BKK26260	yqaM	3,811	1,222	0,111	255	1,177
BKK07250	yetO	3,808	1,202	0,092	78	1,158
BKK08740	ygzB	3,808	1,203	0,110	507	1,159
BKK19670	yodN	3,804	1,172	0,145	185	1,129
BKK04359	ydzK	3,799	1,209	0,091	552	1,165
BKK05630	dinB	3,797	1,212	0,075	188	1,167
BKK04720	rsbW	3,788	1,209	0,106	403	1,165
BKK10310	yhfO	3,786	1,228	0,112	141	1,183
BKK18220	yngF	3,784	1,216	0,099	132	1,172
BKK34840	yvcB	3,784	1,178	0,113	132	1,135
BKK37180	fadF	3,778	1,161	0,130	401	1,119
BKK14550	ykrA	3,775	1,224	0,154	69	1,179
BKK38310	ywbI	3,775	1,161	0,102	43	1,119
BKK06130	ydjC	3,769	1,223	0,141	198	1,179
BKK12230	yjjA	3,761	1,234	0,090	127	1,189
BKK29590	iscS	3,761	1,155	0,127	70	1,113
BKK02990	opuAB	3,756	1,247	0,101	134	1,201
BKK06370	pbuG	3,753	1,223	0,136	384	1,179
BKK28950	thrS	3,749	1,155	0,097	32	1,113
BKK03640	ubiD	3,748	1,209	0,112	170	1,165
BKK13610	mtnB	3,744	1,210	0,145	74	1,166
BKK32660	yurT	3,744	1,177	0,088	90	1,134
BKK24150	mmgC	3,741	1,231	0,103	395	1,187
BKK09760	yheE	3,733	1,221	0,118	133	1,177
BKK12260	yjlA	3,732	1,233	0,065	115	1,189
BKK15970	ylxM	3,729	1,160	0,132	73	1,119
BKK11560	ybl	3,718	1,211	0,114	200	1,168
BKK13750	queF	3,704	1,210	0,133	258	1,166
BKK07760	yfkT	3,698	1,202	0,123	371	1,159
BKK18620	yoaI	3,698	1,215	0,081	250	1,172
BKK26920	yraJ	3,693	1,220	0,110	292	1,177
BKK15260	ylxX	3,681	1,223	0,110	622	1,179
BKK29650	ytrP	3,678	1,154	0,082	59	1,113
BKK40350	rocR	3,667	1,193	0,121	257	1,151
BKK38090	vpr	3,665	1,160	0,125	302	1,119
BKK23010	ypbD	3,663	1,141	0,114	591	1,100
BKK23560	mleN	3,653	1,230	0,104	191	1,187
BKK29820	ytpR	3,65	1,154	0,108	88	1,113
BKK11650	tenA	3,647	1,211	0,129	253	1,168
BKK09720	yheH	3,638	1,252	0,163	44	1,208
BKK33000	htrB	3,629	1,203	0,107	475	1,161
BKK06410	yebG	3,618	1,189	0,111	296	1,147
BKK13509	ykzP	3,604	1,208	0,138	159	1,166
BKK21520	yoIC	3,601	1,196	0,115	408	1,155
BKK40790	yyoO	3,598	1,189	0,121	369	1,147
BKK09480	yhdI	3,591	1,219	0,123	129	1,177
BKK09010	yhcA	3,591	1,204	0,112	245	1,163
BKK16210	fliF	3,591	1,204	0,111	224	1,163
BKK05160	ydeD	3,58	1,209	0,116	65	1,167
BKK39520	yxeK	3,577	1,206	0,115	67	1,165
BKK07900	yfkH	3,576	1,200	0,124	95	1,159
BKK12270	ynzE	3,559	1,213	0,099	191	1,172
BKK26890	yraE	3,559	1,219	0,106	284	1,177
BKK12060	yjdl	3,557	1,231	0,133	340	1,189
BKK10640	sbcD	3,557	1,225	0,100	164	1,183
BKK20910	yopF	3,553	1,196	0,106	336	1,155
BKK11640	yjbQ	3,547	1,231	0,137	824	1,189
BKK07720	yflD	3,544	1,200	0,116	327	1,159
BKK36270	ywqB	3,533	1,208	0,103	95	1,166
BKK38770	cimH	3,533	1,178	0,118	53	1,138
BKK09520	sigM	3,531	1,219	0,150	65	1,177
BKK27390	yrrK	3,521	1,200	0,099	448	1,160
BKK40650	yybG	3,521	1,226	0,116	154	1,184
BKK40990	noc	3,518	1,226	0,100	70	1,184
BKK07630	nosA	3,514	1,200	0,111	433	1,159
BKK06320	yeaB	3,513	1,200	0,114	382	1,160
BKK29780	ytxG	3,513	1,179	0,086	114	1,139
BKK15380	ylmE	3,511	1,221	0,117	47	1,179
BKK31321	mstX	3,511	1,197	0,108	92	1,156
BKK01740	rsiW	3,506	1,244	0,073	33	1,201
BKK08660	sspE	3,503	1,199	0,113	358	1,159
BKK38410	sacX	3,502	1,197	0,106	134	1,157
BKK34420	yveF	3,499	1,175	0,106	150	1,135
BKK09740	yheF	3,49	1,218	0,113	105	1,177
BKK16640	ylxP	3,487	1,204	0,112	110	1,164
BKK37990	ywdE	3,475	1,187	0,110	391	1,147
BKK39480	yxeO	3,474	1,264	0,091	72	1,222
BKK01650	ybbC	3,467	1,243	0,105	34	1,201
BKK12450	yjpA	3,466	1,179	0,128	64	1,139
BKK39320	yxiB	3,458	1,205	0,113	144	1,165
BKK21530	yolB	3,455	1,195	0,128	462	1,155
BKK03280	nasF	3,452	1,169	0,134	64	1,130
BKK08425	mprF	3,446	1,208	0,116	230	1,168
BKK21598	yoyK	3,446	1,191	0,117	243	1,151
BKK23190	dacB	3,446	1,248	0,168	48	1,206
BKK38600	licR	3,442	1,205	0,112	189	1,165
BKK18340	ppsA	3,441	1,212	0,085	43	1,172
BKK24830	yqgT	3,441	1,248	0,165	154	1,206
BKK21160	yonA	3,435	1,195	0,114	404	1,155
BKK07350	yfmT	3,429	1,208	0,124	257	1,168
BKK38380	ywbB	3,425	1,196	0,120	64	1,157
BKK15410	ylmH	3,412	1,219	0,153	72	1,179
BKK00490	spoVG	3,402	1,177	0,124	51	1,138
BKK15820	rpmB	3,397	1,174	0,102	30	1,135
BKK04990	yddJ	3,396	1,199	0,104	98	1,160
BKK06269	ydzJ	3,393	1,219	0,125	242	1,179
BKK31180	yulE	3,387	1,195	0,129	89	1,156
BKK04220	ydaG	3,382	1,204	0,115	274	1,165
BKK06270	ydjO	3,377	1,219	0,115	143	1,179
BKK06730	yefA	3,372	1,219	0,132	263	1,179
BKK22760	hepS	3,371	1,192	0,109	161	1,153
BKK25300	cdd	3,365	1,203	0,102	299	1,164
BKK25470	dnaK	3,365	1,203	0,103	324	1,164
BKK14009	ykzT	3,364	1,229	0,158	168	1,189
BKK20800	yopQ	3,364	1,201	0,110	349	1,162
BKK40140	yydJ	3,363	1,206	0,140	78	1,166
BKK26040	yorB	3,362	1,216	0,094	148	1,177
BKK00700	coaX	3,358	1,178	0,097	106	1,139
BKK01620	yncM	3,358	1,216	0,107	517	1,177
BKK22830	gpsA	3,352	1,137	0,115	44	1,100
BKK20440	yorR	3,336	1,167	0,145	88	1,129
BKK24330	yqhY	3,333	1,189	0,123	307	1,151
BKK30750	mntC	3,328	1,263	0,088	47	1,222
BKK13060	ykjA	3,323	1,205	0,098	44	1,166
BKK09730	yheG	3,32	1,216	0,123	146	1,177
BKK26050	yqdB	3,32	1,192	0,124	242	1,153
BKK11450	oppC	3,318	1,207	0,122	237	1,168
BKK40280	yycO	3,318	1,189	0,114	311	1,151
BKK26180	yceF	3,317	1,216	0,117	139	1,177
BKK09200	yhcS	3,316	1,176	0,111	59	1,138
BKK08970	prkA	3,298	1,216	0,121	163	1,177
BKK28460	yslB	3,296	1,196	0,097	87	1,158
BKK16200	fliI	3,292	1,201	0,109	347	1,163
BKK05730	ydhF	3,291	1,198	0,105	346	1,160
BKK29480	ytxK	3,29	1,150	0,132	82	1,113
BKK37830	spsI	3,286	1,156	0,146	53	1,119
BKK05740	phoB	3,278	1,205	0,083	202	1,167
BKK14450	ampS	3,274	1,218	0,103	66	1,179
BKK39030	yxiT	3,271	1,203	0,133	117	1,165
BKK13799	ykzR	3,256	1,204	0,120	177	1,166
BKK21610	yokF	3,25	1,188	0,114	341	1,151
BKK08270	yfiH	3,247	1,196	0,117	410	1,159
BKK21970	ypeP	3,241	1,192	0,108	486	1,155
BKK11080	pksA	3,24	1,201	0,119	62	1,164
BKK34430	racX	3,238	1,187	0,114	381	1,149
BKK34320	epsF	3,232	1,187	0,109	371	1,149
BKK29270	ytol	3,23	1,149	0,099	186	1,113
BKK34570	malK	3,223	1,172	0,123	31	1,135
BKK22680	trpE	3,222	1,188	0,114	242	1,151
BKK22700	aroB	3,222	1,136	0,094	42	1,100
BKK35470	comFA	3,222	1,199	0,118	398	1,161
BKK07880	yfkJ	3,209	1,206	0,116	272	1,168
BKK10630	addA	3,208	1,221	0,105	125	1,183
BKK14140	ccpC	3,208	1,204	0,117	884	1,165
BKK30700	rpmEB	3,206	1,195	0,104	119	1,158
BKK13970	dacA	3,198	1,200	0,128	215	1,163
BKK32490	pucC	3,195	1,185	0,122	40	1,148
BKK30950	yqzG	3,188	1,193	0,098	122	1,156
BKK37290	arfM	3,17	1,154	0,106	59	1,119
BKK40290	yrdC	3,166	1,202	0,099	156	1,159
BKK26760	yrdC	3,156	1,214	0,110	172	1,177
BKK11410	yjbA	3,155	1,221	0,134	94	1,183
BKK12680	xkdO	3,154	1,227	0,162	83	1,189
BKK35320	fliT	3,153	1,171	0,124	449	1,135
BKK36160	ywqM	3,153	1,149	0,126	92	1,114
BKK15350	thiQ	3,152	1,216	0,152	168	1,179
BKK09330	yhcZ	3,15	1,205	0,118	286	1,168
BKK11010	yitI	3,134	1,220	0,094	130	1,183
BKK39129	yxzl	3,13	1,201	0,116	114	1,165
BKK14072	ykzU	3,128	1,203	0,153	401	1,166
BKK21070	yonJ	3,124	1,191	0,126	185	1,155
BKK40460	yyzB	3,116	1,203	0,105	128	1,166
BKK06210	ydjl	3,115	1,196	0,108	899	1,160
BKK32669	yuzN	3,114	1,170	0,103	160	1,134
BKK24790	yqgX	3,11	1,200	0,100	262	1,164
BKK09560	cueR	3,108	1,246	0,119	96	1,208
BKK07830	yfkO	3,104	1,195	0,120	398	1,159
BKK27620	recJ	3,104	1,196	0,104	151	1,160
BKK35130	yvlA	3,101	1,170	0,104	377	1,135
BKK39830	yxcA	3,097	1,193	0,102	115	1,158
BKK26630	yrdQ	3,093	1,189	0,119	303	1,153
BKK03570	sfp	3,091	1,193	0,107	80	1,158
BKK07840	yfkN	3,09	1,195	0,112	454	1,159
BKK25660	aroD	3,09	1,200	0,101	462	1,164
BKK38360	ywbD	3,085	1,153	0,135	37	1,119
BKK05610	vmlR	3,078	1,195	0,125	425	1,160
BKK40960	parB	3,071	1,221	0,101	118	1,184
BKK28560	lcfA	3,07	1,192	0,110	102	1,156
BKK29000	nrdR	3,063	1,193	0,094	114	1,158
BKK08870	ygaN	3,062	1,213	0,108	128	1,177
BKK10150	yhgD	3,062	1,213	0,103	138	1,177
BKK01550	gerD	3,061	1,173	0,112	35	1,138
BKK06600	pcrB	3,061	1,215	0,113	261	1,179
BKK27420	yrri	3,06	1,189	0,112	290	1,153
BKK14890	ctaC	3,059	1,215	0,109	327	1,179
BKK14770	bipA	3,055	1,215	0,097	259	1,179
BKK09980	yhal	3,054	1,245	0,158	82	1,208
BKK08290	yfiF	3,052	1,194	0,129	108	1,159
BKK28540	fadB	3,049	1,005	0,094	159	0,975
BKK11110	yitS	3,048	1,219	0,099	76	1,183
BKK27809	yrzT	3,048	1,195	0,103	164	1,160
BKK39010	yxjB	3,048	1,200	0,116	101	1,165
BKK27130	rsiV	3,047	1,259	0,067	80	1,222
BKK16180	flgB	3,045	1,204	0,101	348	1,168
BKK37620	rsfA	3,044	1,153	0,107	78	1,119
BKK31020	yuaF	3,043	1,197	0,120	371	1,164
BKK00750	pabA	3,041	1,134	0,116	53	1,100
BKK02920	yceF	3,039	1,238	0,152	67	1,201
BKK38590	licB	3,037	1,200	0,114	83	1,165
BKK08290	yfiJ	3,034	1,245	0,172	39	1,208
BKK13550	mtnA	3,034	1,202	0,126	162	1,166
BKK19450	rsbRC	3,027	1,163	0,118	459	1,129
BKK31710	comQ	3,02	1,133	0,123	52	1,148
BKK22870	fni	3,018	1,133	0,100	251	1,100
BKK05759	citT	3,013	1,203	0,103	261	1,168
BKK17320	ymaF	3,007	1,199	0,096	117	1,164
BKK33680	yvaP	2,985	1,168	0,109	71	1,134
BKK23640	yqkD	2,984	1,222	0,109	241	1,187
BKK00950	mrnC	2,982	1,222	0,099	150	1,187
BKK27670	spoVB	2,966	1,258	0,069	80	1,222
BKK24070	buk	2,959	1,222	0,113	356	1,187
BKK09210	yhcT	2,958	1,212	0,098	147	1,177
BKK20010	yosT	2,957	1,163	0,123	351	1,129
BKK26170	yqbB	2,955	1,211	0,100	159	1,177
BKK05360	ydfC	2,947	1,202	0,089	163	1,167
BKK23930	yqjC	2,942	1,222	0,097	343	1,187
BKK03630	ubiX	2,94	1,199	0,109	77	1,165
BKK10510	yhjH	2,939	1,218	0,093	161	1,183
BKK25520	yqxA	2,937	1,185	0,103	262	1,151
BKK09630	yhdX	2,931	1,211	0,120	85	1,177
BKK35350	yvyC	2,928	1,168	0,106	233	1,135
BKK06520	purH	2,922	1,213	0,122	487	1,179
BKK31959	ybdZ	2,92	1,181	0,107	230	1,148
BKK14200	ykuS	2,919	1,214	0,147	767	1,179
BKK38550	ilvK	2,917	1,191	0,135	85	1,157
BKK20930	yopD	2,914	1,189	0,122	528	1,155
BKK37920	ywdL	2,914	1,191	0,112	67	1,157
BKK06590	yerD	2,912	1,213	0,137	180	1,177
BKK23080	aroC	2,909	1,132	0,097	247	1,100
BKK05500	ydfP	2,908	1,201	0,092	309	1,167
BKK23440	spoVAA	2,906	1,242	0,157	38	1,206

Sup. Table 3: Cell width of mutants of the BKK collection (continued)

BKK name[1]	gene	screening delta[2] (%)	average width (μ)	+/-	nb	ADP
BKK09360	yhdC	2,9	1,211	0,092	94	1,177
BKK29080	mutM	2,9	1,195	0,120	488	1,161
BKK08730	perR	2,899	1,191	0,103	116	1,158
BKK35080	yvmB	2,895	1,168	0,125	76	1,135
BKK03950	ycnJ	2,894	1,199	0,085	135	1,165
BKK39550	yxeH	2,888	1,198	0,101	161	1,165
BKK09460	ctrA	2,887	1,196	0,111	268	1,163
BKK18390	yoeC	2,885	1,195	0,110	289	1,162
BKK18680	yoaO	2,883	1,205	0,101	61	1,172
BKK00080	yaaC	2,872	1,193	0,126	254	1,160
BKK06560	yerA	2,872	1,213	0,104	402	1,179
BKK38640	yxlH	2,869	1,191	0,108	155	1,158
BKK24660	ygzE	2,868	1,197	0,106	173	1,164
BKK06033	ydzT	2,867	1,201	0,088	410	1,167
BKK06880	yesF	2,866	1,213	0,142	279	1,179
BKK31310	yuzB	2,864	1,181	0,138	110	1,148
BKK13950	mcpC	2,857	1,200	0,144	76	1,166
BKK23620	ygkF	2,844	1,186	0,125	296	1,153
BKK32600	frlO	2,843	1,257	0,129	176	1,222
BKK27210	yrhF	2,841	1,210	0,105	358	1,177
BKK06720	dagK	2,84	1,218	0,125	80	1,184
BKK05790	ydhK	2,839	1,200	0,092	148	1,167
BKK13920	splA	2,839	1,199	0,124	135	1,166
BKK28330	ysnE	2,838	1,186	0,124	521	1,153
BKK40820	yyaL	2,836	1,218	0,102	168	1,184
BKK27720	queA	2,831	1,192	0,120	201	1,160
BKK14160	ykuO	2,83	1,213	0,135	1011	1,179
BKK14670	suhB	2,829	1,213	0,125	249	1,179
BKK33040	fumC	2,827	1,166	0,087	48	1,134
BKK28310	ntdC	2,821	1,201	0,107	252	1,168
BKK07000	yesR	2,82	1,212	0,125	227	1,179
BKK01460	ybaE	2,819	1,170	0,116	62	1,138
BKK18830	pps	2,819	1,170	0,090	121	1,138
BKK39500	yxeM	2,818	1,256	0,074	65	1,222
BKK21790	yplQ	2,809	1,186	0,110	344	1,153
BKK06490	purF	2,803	1,212	0,121	358	1,179
BKK18260	yngJ	2,798	1,204	0,116	59	1,172
BKK33950	cggR	2,798	1,166	0,099	107	1,134
BKK19240	yocK	2,796	1,194	0,121	410	1,162
BKK20760	yopU	2,796	1,187	0,105	503	1,155
BKK12300	uxaC	2,795	1,222	0,142	347	1,189
BKK20720	yopY	2,791	1,201	0,124	192	1,168
BKK26730	yrdF	2,79	1,210	0,094	77	1,177
BKK38930	yxjl	2,787	1,197	0,130	46	1,165
BKK18290	yngL	2,786	1,194	0,118	340	1,162
BKK26010	yqbQ	2,786	1,196	0,115	72	1,164
BKK38300	thiM	2,781	1,199	0,110	115	1,166
BKK18420	ftsR	2,779	1,204	0,083	219	1,172
BKK25569	yqzM	2,779	1,196	0,102	333	1,164
BKK38720	yxkO	2,777	1,179	0,110	397	1,147
BKK34990	lgt	2,773	1,181	0,106	535	1,149
BKK01730	sigW	2,77	1,192	0,087	103	1,160
BKK23370	ypuA	2,77	1,185	0,124	247	1,153
BKK08770	yqzA	2,769	1,191	0,122	552	1,159
BKK25950	yqcC	2,766	1,002	0,100	88	0,975
BKK29150	ytwI	2,765	1,193	0,116	466	1,161
BKK22400	dinG	2,757	1,131	0,117	87	1,100
BKK40720	yyaT	2,754	1,217	0,096	113	1,184
BKK09150	yhcO	2,751	1,200	0,117	336	1,168
BKK40710	yybA	2,751	1,217	0,096	171	1,184
BKK04590	ydbS	2,75	1,197	0,091	424	1,165
BKK18669	yozS	2,75	1,194	0,120	196	1,162
BKK20530	yoqS	2,749	1,200	0,105	292	1,168
BKK15830	yloU	2,745	1,145	0,096	102	1,114
BKK16730	spoVFA	2,74	1,240	0,125	96	1,206
BKK24940	yazC	2,732	1,195	0,114	222	1,164
BKK28200	lonA	2,732	1,183	0,120	278	1,151
BKK07930	yfkD	2,729	1,190	0,109	450	1,159
BKK12040	yjdG	2,727	1,221	0,157	285	1,189
BKK12130	yjfC	2,727	1,221	0,157	285	1,189
BKK03020	ycgA	2,726	1,234	0,075	101	1,201
BKK34100	rsbQ	2,726	1,166	0,153	96	1,135
BKK22540	qcrC	2,721	1,130	0,119	419	1,100
BKK15530	pyrK	2,72	1,187	0,103	87	1,156
BKK34010	yvbW	2,719	1,189	0,097	77	1,158
BKK16440	cheW	2,714	1,193	0,106	399	1,161
BKK19930	yotC	2,704	1,193	0,131	428	1,162
BKK13180	metE	2,696	1,170	0,130	39	1,139
BKK07800	treP	2,69	1,190	0,115	388	1,159
BKK30560	pckA	2,685	1,187	0,120	102	1,156
BKK30460	ytrA	2,676	1,187	0,130	108	1,156
BKK03170	ycgK	2,668	1,191	0,111	325	1,160
BKK03700	gerKA	2,667	1,196	0,090	126	1,165
BKK26039	yqbN	2,667	1,208	0,103	545	1,177
BKK13640	spo0E	2,664	1,241	0,155	58	1,208
BKK13860	ykvY	2,664	1,197	0,151	329	1,166
BKK01140	ybaC	2,661	1,195	0,112	228	1,164
BKK32290	yutF	2,652	1,192	0,116	377	1,161
BKK08440	yfiY	2,651	1,240	0,172	67	1,208
BKK09880	yhaR	2,641	1,208	0,103	153	1,177
BKK21300	yomM	2,633	1,181	0,105	141	1,151
BKK09100	cspB	2,629	1,208	0,098	129	1,177
BKK27650	secDF	2,627	1,254	0,084	135	1,222
BKK00180	tadA	2,627	1,208	0,098	98	1,177
BKK31310	yugP	2,625	1,186	0,112	75	1,156
BKK21460	bdbA	2,622	1,181	0,120	280	1,151
BKK30010	ythP	2,621	1,186	0,096	92	1,156
BKK08120	yfjF	2,617	1,189	0,110	158	1,159
BKK04540	ydbO	2,616	1,233	0,097	318	1,201
BKK15120	yllA	2,614	1,210	0,126	188	1,179
BKK03880	yczG	2,606	1,195	0,104	478	1,165
BKK28070	comC	2,606	1,190	0,116	242	1,160
BKK31040	yuaD	2,603	1,186	0,117	118	1,156
BKK07120	lplC	2,602	1,240	0,154	538	1,208
BKK18380	iseA	2,601	1,238	0,146	84	1,206
BKK40570	yybO	2,599	1,197	0,115	141	1,166
BKK22060	pbuX	2,598	1,169	0,122	97	1,139
BKK27200	yrhG	2,596	1,207	0,097	296	1,177
BKK18140	ynfE	2,595	1,202	0,124	213	1,172
BKK44250	yycR	2,594	1,177	0,110	255	1,147
BKK21060	yonK	2,593	1,185	0,113	539	1,155
BKK25180	trmK	2,58	1,181	0,113	344	1,151
BKK10110	pbpF	2,579	1,207	0,110	107	1,177
BKK04320	ydaO	2,576	1,232	0,070	42	1,201
BKK29420	ytkK	2,576	1,000	0,061	39	0,975
BKK39380	hutG	2,576	1,195	0,136	88	1,165
BKK28240	ysoA	2,573	1,189	0,103	185	1,160
BKK29190	pfkA	2,573	1,142	0,099	199	1,113
BKK15850	sdaAB	2,564	1,143	0,077	79	1,114
BKK01870	ybcH	2,559	1,000	0,090	59	0,975
BKK26450	yrkN	2,559	1,183	0,119	334	1,153
BKK04610	ydcA	2,55	1,195	0,101	135	1,165
BKK02380	ybgB	2,546	1,159	0,086	63	1,130
BKK11549	yizD	2,543	1,213	0,106	84	1,183
BKK04100	kipR	2,536	1,194	0,080	126	1,165
BKK17100	pksC	2,535	1,193	0,106	145	1,164
BKK28810	abnA	2,533	1,168	0,104	123	1,139
BKK03190	ycgL	2,532	1,207	0,107	99	1,177
BKK09240	yhcW	2,53	1,159	0,128	72	1,130
BKK24410	spoIIIAC	2,53	1,237	0,122	84	1,206
BKK19710	yodQ	2,527	1,158	0,134	190	1,129
BKK26190	yqaT	2,527	1,180	0,135	334	1,151
BKK05800	ycIN	2,52	1,232	0,152	603	1,201
BKK33700	opuBD	2,515	1,163	0,094	94	1,134
BKK13490	htpX	2,507	1,196	0,129	69	1,166
BKK08400	yfiU	2,499	1,197	0,121	249	1,168
BKK23401	spoVAEA	2,495	1,237	0,158	95	1,206
BKK01540	salA	2,494	1,206	0,108	231	1,177
BKK24060	lpdV	2,494	1,217	0,116	246	1,187
BKK19680	yozE	2,494	1,157	0,124	673	1,129
BKK07370	yfmR	2,485	1,238	0,174	111	1,208
BKK28050	maf	2,485	1,188	0,095	158	1,160
BKK25320	yqfG	2,474	1,194	0,111	400	1,165
BKK29310	ytnJ	2,474	1,141	0,093	67	1,113
BKK21210	yomV	2,463	1,183	0,121	218	1,155
BKK09140	yhcM	2,458	1,206	0,108	110	1,177
BKK14380	fruR	2,458	1,208	0,156	69	1,179
BKK34230	epsA	2,453	1,163	0,114	198	1,135
BKK39140	yxiK	2,453	1,193	0,123	112	1,165
BKK00390	yabD	2,448	1,167	0,137	58	1,139
BKK31990	dhbC	2,448	1,186	0,093	116	1,158
BKK27729	yrzS	2,446	1,188	0,101	429	1,160
BKK15840	yloV	2,437	1,141	0,096	124	1,114
BKK30960	glgD	2,434	1,184	0,098	87	1,156
BKK23890	oxaAB	2,434	1,236	0,193	31	1,206
BKK04440	dctB	2,43	1,193	0,117	267	1,165
BKK35190	yvkC	2,43	1,163	0,094	354	1,135
BKK09750	sspB	2,429	1,206	0,094	76	1,177
BKK11100	nprB	2,429	1,212	0,120	122	1,183
BKK20330	yorM	2,428	1,190	0,102	219	1,162
BKK01520	yboK	2,427	0,999	0,056	30	0,975
BKK36170	ywqL	2,427	1,141	0,139	31	1,114
BKK36630	ywnA	2,427	1,175	0,113	275	1,147
BKK34370	epsA	2,426	1,163	0,083	192	1,135
BKK34480	yvdT	2,426	1,163	0,083	192	1,135
BKK21229	youA	2,423	1,183	0,105	382	1,155
BKK35580	tuaD	2,416	1,177	0,124	516	1,149
BKK14350	yknX	2,415	1,165	0,092	154	1,138
BKK16190	flgC	2,412	1,191	0,106	426	1,163
BKK26770	yrdB	2,411	1,205	0,103	262	1,177
BKK31810	yuzE	2,411	1,176	0,120	271	1,148
BKK20929	yoyI	2,41	1,183	0,112	505	1,155
BKK31460	kapB	2,409	1,176	0,127	31	1,148
BKK21220	yomU	2,407	1,179	0,119	403	1,151
BKK03350	yciB	2,389	1,230	0,104	61	1,201
BKK13940	ykwB	2,385	1,190	0,112	280	1,163
BKK38940	yxjI	2,385	1,192	0,124	88	1,165
BKK00450	sspF	2,384	1,191	0,101	123	1,164
BKK31920	yukF	2,384	1,175	0,126	89	1,148
BKK10790	asnO	2,382	1,211	0,116	85	1,183
BKK13290	ykzD	2,377	1,190	0,118	220	1,163
BKK31430	yugE	2,375	1,175	0,128	61	1,148
BKK26990	yraD	2,374	1,205	0,105	205	1,177
BKK37160	rpoE	2,37	1,145	0,118	331	1,119
BKK07320	yfnC	2,354	1,237	0,153	522	1,208
BKK06980	yesP	2,349	1,165	0,101	30	1,138
BKK08930	cspR	2,349	1,190	0,114	213	1,163
BKK18810	yobA	2,347	1,235	0,138	54	1,206
BKK03440	tlpC	2,344	1,187	0,127	413	1,160
BKK09220	yhcU	2,344	1,205	0,111	165	1,177
BKK37670	ywfl	2,339	1,212	0,113	98	1,184
BKK02660	ycbU	2,334	0,998	0,098	50	0,975
BKK02410	ybgG	2,326	1,156	0,088	126	1,130
BKK38990	scoA	2,326	1,192	0,107	74	1,165
BKK36980	ywlA	2,315	1,184	0,114	126	1,158
BKK07100	lplA	2,314	1,236	0,119	105	1,208
BKK05520	ydzH	2,313	1,194	0,083	333	1,167
BKK38970	yxjF	2,312	1,193	0,107	82	1,166
BKK05470	ydfM	2,31	1,229	0,099	95	1,201
BKK33800		2,309	1,176	0,115	336	1,149
BKK20830	yopN	2,307	1,182	0,122	343	1,155
BKK31279	yxaJ	2,303	1,183	0,112	152	1,156
BKK00410	rnmV	2,3	1,190	0,117	75	1,164
BKK33490		2,298	1,188	0,112	291	1,161
BKK37420	albF	2,296	1,193	0,109	119	1,166
BKK17280	ymaD	2,295	1,190	0,117	96	1,164
BKK12650	xkdK	2,292	1,216	0,152	131	1,189
BKK03300	nasD	2,29	1,156	0,132	58	1,130
BKK25360	yqfC	2,29	1,190	0,103	151	1,164
BKK08110	sspH	2,289	1,185	0,122	149	1,159
BKK24980	pstC	2,289	1,177	0,124	413	1,151
BKK04580	yybN	2,288	1,193	0,136	109	1,166
BKK36510	amtB	2,286	1,176	0,108	331	1,149
BKK12290	ndh	2,285	1,216	0,127	932	1,189
BKK28230	tig	2,284	1,186	0,101	191	1,160
BKK22240	mpr	2,283	1,156	0,113	144	1,130
BKK35230	swrAA	2,282	1,161	0,116	321	1,135
BKK09080	yhcI	2,281	1,195	0,121	420	1,168
BKK00590	yabA	2,271	1,198	0,128	138	1,172
BKK36440	ywoH	2,271	1,184	0,092	98	1,158
BKK37520	ywhH	2,271	1,193	0,109	91	1,166
BKK28400	ysmB	2,268	1,186	0,091	267	1,160
BKK00850	moa	2,264	1,211	0,105	118	1,184
BKK10230	yhfH	2,262	1,204	0,116	112	1,177
BKK38890	yxjN	2,261	1,191	0,111	166	1,165
BKK20610	yoqJ	2,255	1,181	0,135	124	1,155
BKK26650	ytpS	2,254	1,203	0,105	277	1,177
BKK29805	ytpS	2,254	1,250	0,055	36	1,222
BKK36050	cotB	2,252	1,184	0,094	106	1,158
BKK31970	dhbB	2,25	1,174	0,110	112	1,148
BKK06540	yezC	2,248	1,205	0,130	198	1,179
BKK17830	yndM	2,243	1,190	0,110	80	1,164
BKK25780	arsC	2,241	1,177	0,122	361	1,151
BKK07780	yfkR	2,238	1,185	0,122	258	1,159
BKK12140	yjgA	2,234	1,176	0,159	160	1,149
BKK23530	spoIIM	2,231	1,179	0,115	310	1,153
BKK35780	lytD	2,227	1,175	0,123	251	1,149
BKK40700	yybB	2,224	1,211	0,092	101	1,184
BKK34470	yveA	2,223	1,187	0,108	461	1,161
BKK40020	yczH	2,219	1,191	0,079	160	1,165
BKK24600	sinI	2,218	0,997	0,082	103	0,975
BKK07660	yflJ	2,217	1,184	0,131	307	1,159
BKK34930	hisZ	2,217	1,192	0,109	191	1,166
BKK04190	ydaD	2,216	1,191	0,106	383	1,165
BKK10990	yitH	2,214	1,209	0,122	132	1,183
BKK01510	ybaJ	2,211	0,997	0,066	53	0,975
BKK12590	xkdE	2,21	1,215	0,188	155	1,189
BKK15320	sigE	2,207	1,205	0,122	164	1,179
BKK05430	ydfJ	2,205	1,228	0,096	75	1,201
BKK00400	gsiB	2,204	1,191	0,094	188	1,165
BKK32620	yurQ	2,203	1,159	0,109	110	1,134
BKK19380	yojO	2,202	1,154	0,125	293	1,129
BKK22100	kdgA	2,202	1,125	0,123	117	1,100
BKK34330	epsE	2,202	1,192	0,111	214	1,166
BKK17110	pksD	2,2	1,189	0,106	174	1,164
BKK00030	ycsD	2,19	1,185	0,117	345	1,160
BKK31980	dhbE	2,188	1,187	0,105	422	1,161
BKK25770	spoIVCA	2,185	1,249	0,090	124	1,222
BKK37980	ywdF	2,183	1,143	0,114	141	1,119
BKK31510	yufK	2,182	1,164	0,093	67	1,139
BKK12890	ykcC	2,176	1,235	0,151	56	1,208
BKK06580	yerC	2,169	1,204	0,111	334	1,179
BKK13000	ykfD	2,169	1,188	0,117	283	1,163
BKK40574	yyzJ	2,168	1,181	0,098	139	1,156
BKK13880	glcT	2,165	1,192	0,127	241	1,166
BKK38610	yxzF	2,163	1,190	0,099	56	1,165
BKK13480	ykrK	2,162	1,192	0,148	66	1,166
BKK39200	yxzC	2,157	1,190	0,131	96	1,165
BKK37450	ywhK	2,156	1,172	0,113	303	1,147
BKK01450	ybxA	2,154	1,189	0,096	175	1,164
BKK08540	yfhI	2,153	1,234	0,155	165	1,208
BKK08899	ygzD	2,146	1,202	0,086	107	1,177
BKK34940	yvpB	2,146	1,159	0,105	429	1,135
BKK39150	yxiI	2,144	1,190	0,132	66	1,165
BKK11799	yjzK	2,142	1,214	0,086	338	1,189
BKK37550	ywhA	2,139	1,143	0,103	168	1,119
BKK08160	yfjB	2,138	1,184	0,112	284	1,159
BKK37210	ywjC	2,138	1,143	0,110	111	1,119
BKK23950	yqjA	2,137	1,178	0,120	335	1,153
BKK28490	uvrC	2,136	0,996	0,066	43	0,975
BKK24570	gcvT	2,135	1,212	0,095	102	1,187
BKK31410	yugG	2,135	1,172	0,097	63	1,148
BKK10720	gerPA	2,132	1,209	0,099	170	1,183
BKK05180	ydeF	2,123	1,192	0,100	251	1,167
BKK00790	pdaA	2,122	1,183	0,103	419	1,159
BKK20990	yonU	2,12	1,179	0,111	447	1,155
BKK05750	yflT	2,115	1,183	0,114	217	1,159
BKK20390	yorG	2,115	1,186	0,102	481	1,162
BKK18010	yneN	2,113	1,192	0,137	83	1,164
BKK02030	yosR	2,113	1,160	0,073	126	1,129
BKK18970	yobI	2,109	1,186	0,103	478	1,162
BKK04390	ydbA	2,105	1,184	0,114	397	1,160
BKK15520	pyrAB	2,104	1,180	0,106	63	1,156
BKK26160	yqbC	2,1	1,201	0,123	146	1,177
BKK07220	yetL	2,097	0,995	0,086	60	0,975
BKK24730	comGA	2,096	1,178	0,121	248	1,153
BKK06860	yezE	2,092	1,204	0,127	476	1,179
BKK38020	pdxK	2,092	1,181	0,112	101	1,157
BKK23150	resA	2,088	1,232	0,182	124	1,206
BKK37480	ywhH	2,088	1,142	0,096	370	1,119
BKK16220	fliG	2,081	1,187	0,100	283	1,163
BKK38560	licH	2,081	1,189	0,137	35	1,165
BKK40160	yydH	2,079	1,191	0,145	85	1,166
BKK06839	phrH	2,077	1,203	0,113	285	1,179
BKK35480	yviA	2,077	1,137	0,139	71	1,114
BKK23610	nudF	2,075	1,212	0,090	174	1,187
BKK15960	ylqB	2,074	1,142	0,113	31	1,119
BKK31550	yufO	2,072	1,247	0,071	88	1,222
BKK08230	catD	2,071	1,233	0,130	100	1,208
BKK13340	ykoM	2,065	1,190	0,094	172	1,166
BKK02500	ycbG	2,062	1,184	0,113	369	1,160
BKK02160	ybaD	2,059	1,153	0,108	167	1,130
BKK09570	yhdR	2,057	1,201	0,114	78	1,177
BKK00440	veg	2,054	1,187	0,102	171	1,164
BKK40730	yyaS	2,054	1,209	0,101	132	1,184
BKK02470	ycbD	2,053	1,153	0,135	95	1,130

Sup. Table 3: Cell width of mutants of the BKK collection (continued)

BKK name[1]	gene	screening delta[2] (%)	erage width (µ	+/-	nb	ADP
BKK39160	yxil	2,052	1,189	0,110	101	1,165
BKK17120	pksE	2,051	1,231	0,150	206	1,206
BKK16390	flhA	2,049	1,177	0,132	387	1,153
BKK06970	yesO	2,048	1,226	0,081	234	1,201
BKK25820	yqcl	2,046	1,177	0,119	230	1,153
BKK04940	yddE	2,045	1,191	0,118	503	1,167
BKK29530	sppA	2,041	1,163	0,097	191	1,139
BKK12229	yjzl	2,039	1,213	0,089	1324	1,189
BKK13040	hmp	2,038	1,190	0,117	451	1,166
BKK17230	pksS	2,036	1,187	0,115	151	1,164
BKK27070	levD	2,036	1,201	0,093	261	1,177
BKK11680	thiS	2,032	1,213	0,150	755	1,189
BKK35650	lytR	2,031	1,173	0,106	529	1,149
BKK03000	opuAC	2,03	1,226	0,065	146	1,201
BKK05590	ydgD	2,027	1,191	0,077	159	1,167
BKK16080	ylqH	2,026	1,183	0,126	259	1,160
BKK27310	pbpl	2,024	0,995	0,106	123	0,975
BKK30450	ytrB	2,023	1,174	0,117	246	1,151
BKK26910	yraK	2,022	1,200	0,106	204	1,177
BKK04660	ndoA	2,02	1,188	0,092	72	1,165
BKK36110	ywrC	2,013	1,190	0,131	93	1,166
BKK39470	yxeP	2,013	1,188	0,107	95	1,165
BKK29910	ytzH	2,011	1,135	0,135	46	1,113
BKK26250	yqaN	2,007	1,200	0,103	195	1,177
BKK40540	yybR	2,006	1,179	0,105	120	1,156
BKK12710	xkdR	2,005	1,213	0,154	142	1,189
BKK23470	spoIIAA	2,001	1,231	0,157	184	1,206
BKK26790	aadK	1,996	1,200	0,101	286	1,177
BKK39440	yxxB	1,993	1,231	0,109	192	1,206
BKK40660	yybF	1,993	1,190	0,100	407	1,167
BKK20928	yoyH	1,992	1,178	0,110	326	1,155
BKK14440	panE	1,99	1,203	0,116	210	1,179
BKK22030	ypbR	1,99	1,176	0,114	377	1,153
BKK22720	cheR	1,99	1,176	0,120	252	1,153
BKK27810	yrbD	1,98	1,183	0,092	212	1,160
BKK33221	yvrH	1,976	0,994	0,081	41	0,975
BKK33990	yvbU	1,976	1,157	0,126	72	1,134
BKK03590	tcyC	1,974	1,225	0,085	153	1,201
BKK07130	lplD	1,974	1,202	0,118	423	1,179
BKK11590	xynB	1,972	1,186	0,112	205	1,164
BKK21780	yplP	1,97	1,178	0,130	274	1,155
BKK11190	argC	1,966	1,207	0,106	73	1,183
BKK37750	ywfA	1,965	1,172	0,112	349	1,149
BKK31520	yufL	1,964	1,170	0,114	329	1,148
BKK22880	ypfD	1,962	1,176	0,114	318	1,153
BKK15800	thiN	1,96	1,157	0,113	318	1,135
BKK34780	yvcl	1,96	1,157	0,145	339	1,135
BKK30630	ytkD	1,959	1,179	0,090	123	1,156
BKK24000	bmrU	1,958	1,210	0,104	414	1,187
BKK29220	ytsJ	1,958	1,162	0,099	103	1,139
BKK03070	yvfT	1,955	1,172	0,120	289	1,149
BKK34580	yvdI	1,954	1,172	0,102	502	1,149
BKK08850	ssuC	1,95	1,232	0,132	128	1,208
BKK03180	cah	1,948	1,182	0,124	632	1,160
BKK03930	gdh	1,944	1,188	0,086	175	1,165
BKK04830	yyaK	1,942	1,207	0,103	172	1,184
BKK33470	bdbC	1,941	1,189	0,104	160	1,166
BKK17490	ynzG	1,939	1,184	0,107	433	1,162
BKK19110	yobW	1,939	1,194	0,116	89	1,172
BKK10130	hemH	1,938	1,178	0,105	92	1,156
BKK36490	ywoC	1,938	1,189	0,095	81	1,166
BKK14130	ykuL	1,937	1,189	0,142	774	1,166
BKK28360	ysnA	1,937	1,173	0,118	288	1,151
BKK40500	rplI	1,935	1,207	0,099	105	1,184
BKK10020	serC	1,934	1,200	0,119	157	1,177
BKK19030	yobO	1,934	1,194	0,139	98	1,172
BKK30680	ytjA	1,932	1,178	0,113	78	1,156
BKK11120	yitT	1,926	1,185	0,112	223	1,163
BKK21170	yomZ	1,923	1,177	0,104	316	1,155
BKK33580	yvaF	1,922	1,156	0,125	114	1,134
BKK26870	trpB	1,921	1,121	0,096	196	1,100
BKK13770	ykvO	1,919	1,189	0,130	68	1,166
BKK21420	bhlA	1,919	1,173	0,111	216	1,151
BKK21800	ypkP	1,918	1,177	0,117	566	1,155
BKK10010	trpP	1,917	1,232	0,127	46	1,208
BKK23810	yqjN	1,912	1,210	0,102	350	1,187
BKK20590	yoqL	1,91	1,177	0,122	249	1,155
BKK00600	yabP	1,909	1,169	0,107	550	1,147
BKK13690	motA	1,898	1,189	0,102	117	1,166
BKK23540	yqkK	1,898	1,210	0,110	283	1,187
BKK03200	ycgM	1,892	1,151	0,119	36	1,130
BKK10360	yhfT	1,891	1,206	0,101	217	1,183
BKK32630	yurR	1,889	1,156	0,092	150	1,134
BKK18490	rtp	1,888	1,184	0,126	266	1,162
BKK11900	yjcL	1,885	1,184	0,120	287	1,163
BKK19340	yocR	1,884	1,179	0,095	155	1,158
BKK27730	ruvB	1,884	1,181	0,107	314	1,160
BKK14990	ylbF	1,883	1,201	0,131	615	1,179
BKK28250	leuD	1,88	1,181	0,099	224	1,160
BKK35170	uvrB	1,876	1,156	0,117	87	1,135
BKK32990	mrgA	1,873	1,156	0,091	90	1,134
BKK10780	yisN	1,871	1,205	0,130	152	1,183
BKK13780	ykvP	1,871	1,188	0,121	416	1,166
BKK35950	rbsC	1,867	1,188	0,105	116	1,166
BKK24140	prpC	1,865	1,209	0,102	659	1,187
BKK26660	yrdN	1,857	1,199	0,117	403	1,177
BKK38630	katX	1,855	1,186	0,131	34	1,165
BKK04860	ydcQ	1,852	1,186	0,121	95	1,165
BKK29540	ppnKB	1,851	1,161	0,089	140	1,139
BKK06440	purB	1,849	1,201	0,124	179	1,179
BKK19470	yojF	1,847	1,190	0,118	412	1,168
BKK22720	yrhE	1,843	1,198	0,100	228	1,177
BKK14300	moaE	1,838	1,201	0,107	880	1,179
BKK17760	yndE	1,836	1,185	0,113	206	1,164
BKK39270	bglP	1,835	1,186	0,113	79	1,165
BKK01090	rplGB	1,832	1,185	0,109	249	1,164
BKK05420	ydfl	1,824	1,223	0,116	68	1,201
BKK07330	yfnB	1,818	1,200	0,112	548	1,179
BKK13830	ykvU	1,816	1,184	0,115	326	1,163
BKK31390	yugI	1,816	1,169	0,105	232	1,148
BKK00970	yacP	1,816	1,181	0,123	307	1,160
BKK10740	yisJ	1,814	1,205	0,113	75	1,183
BKK02480	gudP	1,81	1,223	0,069	197	1,201
BKK32790	yusG	1,81	1,155	0,095	171	1,134
BKK13800	ykvR	1,804	1,187	0,132	58	1,166
BKK05770	ydhl	1,798	1,188	0,110	439	1,167
BKK02490	gudD	1,797	1,150	0,123	43	1,130
BKK25930	yqcE	1,796	1,172	0,104	256	1,151
BKK31050	gbsB	1,796	1,177	0,126	106	1,156
BKK33480	bdbD	1,796	1,155	0,095	157	1,134
BKK17590	xylR	1,79	1,184	0,113	81	1,164
BKK04520	ydbM	1,788	1,180	0,115	696	1,160
BKK21599	yoyK	1,788	1,176	0,107	459	1,155
BKK04370	yycl	1,785	1,150	0,098	131	1,130
BKK40610	yybK	1,784	1,187	0,121	101	1,166
BKK15560	pyrE	1,783	1,160	0,092	111	1,139
BKK38830	aldY	1,781	1,185	0,118	77	1,165
BKK38950	yxjH	1,78	1,178	0,094	70	1,158
BKK03610	tcyA	1,775	1,168	0,119	422	1,147
BKK16140	codV	1,77	1,205	0,125	65	1,184
BKK39140	yozB	1,768	1,182	0,116	362	1,162
BKK09070	acoB	1,767	1,179	0,118	417	1,159
BKK09310	pgcA	1,767	1,230	0,127	38	1,208
BKK02710	yczC	1,765	1,223	0,067	82	1,201
BKK36150	ywqN	1,764	1,134	0,087	54	1,114
BKK40840	yozJ	1,764	1,188	0,117	192	1,167
BKK02170	ydbC	1,763	1,185	0,100	263	1,165
BKK39280	yfbB	1,758	1,168	0,120	442	1,147
BKK14010	yxxE	1,757	1,185	0,122	80	1,165
BKK04010	cheV	1,754	1,183	0,123	333	1,163
BKK02310	ybfO	1,752	1,150	0,097	86	1,130
BKK08830	ssuB	1,743	1,229	0,167	96	1,208
BKK13530	kinE	1,743	1,188	0,103	359	1,168
BKK18690	yoaP	1,743	1,182	0,114	392	1,162
BKK10610	yhjR	1,74	1,204	0,108	150	1,183
BKK21240	yomS	1,739	1,175	0,123	411	1,155
BKK29430	ytzD	1,739	1,159	0,092	208	1,139
BKK29610	ezrA	1,739	1,132	0,116	65	1,113
BKK03100	ycgG	1,737	1,150	0,107	110	1,130
BKK02000	ybdJ	1,735	1,150	0,097	72	1,130
BKK04490	ydbJ	1,734	1,222	0,098	109	1,201
BKK10890	yisX	1,732	1,204	0,120	111	1,183
BKK05320	ydeS	1,726	1,187	0,122	143	1,167
BKK06700	yerO	1,725	1,199	0,126	373	1,179
BKK06073	ydzW	1,722	1,167	0,098	121	1,147
BKK29700	acuB	1,721	1,178	0,105	142	1,158
BKK24210	yqiG	1,719	1,207	0,145	583	1,187
BKK13120	proB	1,714	1,186	0,123	109	1,166
BKK28740	araP	1,713	1,159	0,102	107	1,139
BKK04700	rsbU	1,71	1,185	0,115	93	1,165
BKK10500	yhjG	1,71	1,204	0,102	100	1,183
BKK27350	yrrN	1,698	1,171	0,123	232	1,151
BKK29490	tpx	1,695	1,132	0,120	128	1,113
BKK40578	yyzK	1,692	1,186	0,114	161	1,166
BKK04170	ydaB	1,687	1,185	0,097	343	1,165
BKK22160	yptA	1,68	1,119	0,097	30	1,100
BKK03060	lctP	1,674	1,149	0,121	51	1,130
BKK19689	yokU	1,674	1,188	0,106	330	1,168
BKK24290	xseB	1,672	1,207	0,112	260	1,187
BKK10600	yhjQ	1,665	1,203	0,120	85	1,183
BKK30940	glgP	1,665	1,175	0,103	132	1,156
BKK09400	spoVR	1,663	1,228	0,112	201	1,208
BKK16950	pbpX	1,661	1,227	0,160	62	1,206
BKK21380	yomF	1,661	1,174	0,125	384	1,155
BKK18410	ggt	1,66	1,157	0,097	89	1,138
BKK20090	yosK	1,656	1,148	0,113	70	1,129
BKK28300	ilvH	1,656	1,179	0,096	313	1,160
BKK04180	ydaC	1,655	1,184	0,087	146	1,165
BKK16170	codY	1,649	1,182	0,138	230	1,163
BKK39350	hutH	1,647	1,184	0,115	102	1,165
BKK09260	yhxA	1,643	1,182	0,104	262	1,163
BKK09120	yhcK	1,636	1,182	0,108	295	1,163
BKK30840	yteA	1,632	1,175	0,116	117	1,156
BKK06890	cotJA	1,631	1,198	0,113	433	1,179
BKK25190	cccA	1,63	1,172	0,111	473	1,153
BKK39680	ioll	1,621	1,166	0,106	353	1,147
BKK06500	purM	1,62	1,198	0,121	369	1,179
BKK36060	cotH	1,619	1,168	0,118	334	1,149
BKK36550	spoIIQ	1,619	1,168	0,118	334	1,149
BKK35690	ggaA	1,617	1,176	0,105	68	1,158
BKK04370	ydaS	1,61	1,184	0,094	466	1,165
BKK08550	sspK	1,601	1,177	0,114	318	1,159
BKK10340	phoE	1,601	1,202	0,115	140	1,183
BKK17460	glnA	1,601	1,182	0,115	59	1,164
BKK24540	yqhL	1,6	1,206	0,110	223	1,187
BKK06330	yeaC	1,596	1,198	0,120	155	1,179
BKK27140	oatA	1,596	1,242	0,068	82	1,222
BKK32860	yusN	1,596	1,152	0,102	110	1,134
BKK10970	yitF	1,591	1,202	0,119	107	1,183
BKK20180	yosB	1,589	1,180	0,114	343	1,162
BKK35840	ywtF	1,589	1,132	0,100	277	1,114
BKK00220	yaaL	1,586	1,202	0,124	98	1,183
BKK17040	mutS	1,584	1,182	0,112	258	1,164
BKK39060	citH	1,584	1,183	0,114	87	1,165
BKK39290	yxxD	1,584	1,185	0,121	66	1,166
BKK17980	sspP	1,582	1,182	0,111	159	1,164
BKK05870	gmuF	1,581	1,186	0,116	431	1,167
BKK22180	gpsB	1,577	1,118	0,115	76	1,100
BKK06048	ydzU	1,574	1,186	0,113	280	1,167
BKK30590	ytmA	1,574	1,174	0,085	59	1,159
BKK31540	yufN	1,571	1,241	0,082	57	1,222
BKK32980	yusZ	1,569	1,180	0,115	395	1,161
BKK09350	yhdB	1,568	1,195	0,105	280	1,177
BKK05700	ydhC	1,558	1,185	0,075	162	1,167
BKK03920	glcU	1,555	1,220	0,082	32	1,201
BKK34680	yvcT	1,554	1,153	0,107	60	1,135
BKK19050	yhoQ	1,551	1,190	0,106	142	1,172
BKK29330	ytmO	1,551	1,130	0,108	791	1,113
BKK12340	uxuA	1,549	1,207	0,092	1324	1,189
BKK11170	yitY	1,547	1,202	0,098	110	1,183
BKK30330	ytvB	1,547	1,174	0,089	111	1,156
BKK23900	yjgF	1,546	1,225	0,164	94	1,206
BKK09650	yhdZ	1,545	1,195	0,110	122	1,177
BKK17990		1,543	1,180	0,124	356	1,162
BKK39740	iolC	1,542	0,990	0,076	90	0,975
BKK28670	yjzL	1,541	1,157	0,096	170	1,139
BKK34250	epsL	1,538	1,153	0,093	142	1,135
BKK07870	yfkK	1,537	1,180	0,120	342	1,163
BKK09420	lytE	1,537	1,227	0,115	39	1,208
BKK16590		1,535	1,179	0,127	326	1,161
BKK22869	ypzl	1,535	1,117	0,088	110	1,100
BKK08650	ydjL	1,533	1,177	0,155	135	1,159
BKK40130	yydK	1,531	1,165	0,095	388	1,147
BKK12960	dppE	1,529	1,155	0,093	140	1,138
BKK09380	nsrR	1,528	1,195	0,104	120	1,177
BKK08029	yfzA	1,524	1,176	0,110	516	1,159
BKK05230	ydeK	1,523	1,157	0,085	95	1,139
BKK27610	apt	1,518	1,177	0,098	142	1,160
BKK02090	ybxl	1,516	1,147	0,129	111	1,130
BKK08760	spo0M	1,514	1,227	0,147	248	1,208
BKK37590	ywgA	1,514	1,136	0,131	72	1,119
BKK25010	yqgE	1,512	1,225	0,116	49	1,206
BKK29840	ytpP	1,511	1,157	0,103	98	1,139
BKK23050	fmnP	1,509	1,117	0,103	148	1,100
BKK00210	recR	1,502	1,165	0,163	503	1,147
BKK21990	ypdQ	1,502	1,172	0,103	484	1,155
BKK06910	cotJC	1,5	1,197	0,115	261	1,179
BKK35420	yvyG	1,5	1,131	0,108	862	1,114
BKK40950	yyaC	1,498	1,202	0,096	139	1,184
BKK04840	ydcO	1,495	1,177	0,131	359	1,160
BKK24610	sinR	1,492	1,202	0,088	162	1,184
BKK11660	tenI	1,49	1,207	0,118	264	1,189
BKK30660	ytkA	1,486	1,240	0,055	32	1,222
BKK12970	ykfA	1,484	1,226	0,139	121	1,208
BKK33450	helD	1,476	1,151	0,085	97	1,134
BKK05620	ydgF	1,475	1,177	0,113	690	1,160
BKK19630	deoD	1,472	1,146	0,119	548	1,129
BKK19430	yojI	1,469	1,146	0,132	269	1,129
BKK20820	yopO	1,465	1,172	0,106	450	1,155
BKK13010	ykgB	1,462	1,183	0,126	555	1,166
BKK10710	gerPB	1,461	1,201	0,117	127	1,183
BKK16800	spoIIIE	1,461	1,224	0,175	62	1,206
BKK26810	yrpC	1,454	1,240	0,084	165	1,222
BKK01660	ybbD	1,451	1,219	0,083	84	1,206
BKK11390	appB	1,451	1,226	0,162	190	1,208
BKK06150	gutB	1,447	1,196	0,126	82	1,179
BKK32160	yutM	1,446	1,178	0,110	434	1,161
BKK33720	opuBB	1,445	1,183	0,114	181	1,166
BKK32500	pucB	1,444	1,183	0,114	156	1,166
BKK32300	yutE	1,441	1,178	0,113	384	1,161
BKK35810	gerBB	1,435	1,130	0,116	31	1,114
BKK25860	yqcG	1,431	1,168	0,107	300	1,151
BKK27280	yrrT	1,431	0,989	0,085	47	0,975
BKK33590	yvaG	1,43	1,151	0,107	178	1,134
BKK26020	yqbP	1,427	1,193	0,122	119	1,177
BKK19120	czrA	1,426	1,188	0,109	157	1,172
BKK40630	yybl	1,426	1,201	0,109	194	1,184
BKK07570	yflS	1,422	1,226	0,140	73	1,208
BKK34240	epsM	1,421	1,151	0,108	130	1,135
BKK08530	yfhH	1,417	1,175	0,106	257	1,159
BKK11220	pksR	1,415	1,180	0,108	157	1,164
BKK28910	lrgA	1,404	1,178	0,108	203	1,161
BKK34080	yvfS	1,404	1,166	0,097	241	1,149
BKK40120	yyzE	1,398	0,989	0,063	46	0,975
BKK06740	yefB	1,394	1,195	0,113	251	1,179
BKK28040	radC	1,394	1,169	0,118	183	1,153
BKK35860	pgdS	1,393	1,165	0,115	76	1,100
BKK34120	ganB	1,391	1,177	0,106	416	1,161
BKK14310	mooD	1,386	1,195	0,107	574	1,179
BKK24430	spoIIIAA	1,386	1,167	0,113	452	1,151
BKK40590	yubM	1,383	1,182	0,107	186	1,166
BKK08950	yhbE	1,382	1,225	0,165	85	1,208
BKK14009	ykzT	1,382	1,182	0,130	124	1,166
BKK09000	yhbJ	1,381	1,193	0,123	131	1,177
BKK27820	yrbC	1,38	1,169	0,114	470	1,153
BKK25370	yqfB	1,378	0,988	0,083	49	0,975
BKK07440	yfmK	1,372	1,175	0,106	308	1,159
BKK32810	yusl	1,366	1,150	0,103	65	1,134
BKK37030	racA	1,362	1,173	0,121	116	1,157
BKK24820	yqgU	1,361	1,179	0,106	322	1,164
BKK03830	yclQ	1,36	1,218	0,091	69	1,201
BKK25050	yqgA	1,36	1,169	0,115	206	1,153
BKK36560	ywnH	1,36	1,182	0,103	162	1,166
BKK05370	ydfD	1,355	1,175	0,128	409	1,160
BKK01590	ybaS	1,353	1,153	0,107	123	1,138
BKK22070	xpt	1,352	1,155	0,112	112	1,139
BKK05250	ycbJ	1,348	1,145	0,089	215	1,130
BKK08980	yhbH	1,347	1,193	0,106	183	1,177
BKK29290	ytnL	1,346	1,155	0,092	133	1,139
BKK05100	yddT	1,34	1,183	0,112	351	1,167
BKK32440	pucK	1,339	1,163	0,110	68	1,148
BKK17600	xylA	1,333	1,179	0,138	65	1,164
BKK26089	yqzN	1,329	1,192	0,118	191	1,177
BKK31830	yueE	1,328	1,163	0,111	251	1,148
BKK19650	yodM	1,325	1,144	0,118	1093	1,129

Sup. Table 3: Cell width of mutants of the BKK collection (continued)

BKK name[1]	gene	screening delta[2] (%)	average width (µ	+/-	nb	ADP
BKK02630	tatAD	1,323	1,217	0,079	43	1,201
BKK21920	ugtP	1,322	1,222	0,162	73	1,206
BKK13230	thiV	1,317	1,224	0,144	59	1,208
BKK11710	thiD	1,316	1,205	0,097	800	1,189
BKK33980	yvbT	1,311	1,177	0,117	449	1,161
BKK35400	flgL	1,309	1,150	0,127	456	1,135
BKK35990	ywrO	1,308	1,129	0,093	92	1,114
BKK26370	yqaC	1,306	1,192	0,098	415	1,177
BKK07190	yezD	1,304	1,194	0,120	279	1,179
BKK14290	mobB	1,304	1,195	0,098	513	1,179
BKK41020	trmE	1,303	1,175	0,113	189	1,160
BKK29860	ytoP	1,301	1,128	0,120	71	1,113
BKK10320	yhfP	1,298	1,178	0,106	390	1,163
BKK22020	ypbS	1,298	1,115	0,131	151	1,100
BKK15440	ylyA	1,297	1,154	0,088	127	1,139
BKK24960	pstBA	1,293	1,222	0,123	132	1,206
BKK05150	ydeC	1,292	1,173	0,117	96	1,158
BKK24530	yqhM	1,29	1,202	0,120	251	1,187
BKK19360	odhB	1,287	1,144	0,121	549	1,129
BKK37090	ywjI	1,282	1,162	0,108	254	1,147
BKK38260	efeN	1,277	1,181	0,110	63	1,166
BKK02250	ybfJ	1,275	1,217	0,068	77	1,201
BKK35310	yvyD	1,272	1,150	0,124	178	1,135
BKK22090	kdgT	1,27	1,222	0,126	196	1,206
BKK30620	ytlD	1,269	1,238	0,081	34	1,222
BKK03530	ycxA	1,267	1,217	0,115	381	1,201
BKK18180	yngB	1,266	1,222	0,167	172	1,206
BKK04040	ycsE	1,264	1,174	0,120	687	1,160
BKK35970	ywsB	1,263	1,128	0,143	32	1,114
BKK24520	mntR	1,262	1,168	0,118	145	1,153
BKK05529	ydzR	1,26	0,987	0,076	37	0,975
BKK15110	ylbQ	1,26	1,194	0,129	310	1,179
BKK15100	ylbP	1,259	1,171	0,107	130	1,156
BKK19060	yobR	1,258	1,199	0,139	119	1,184
BKK05570	ydgB	1,25	1,182	0,127	169	1,167
BKK14960	ylbC	1,249	1,194	0,116	99	1,179
BKK11090	yitR	1,244	1,177	0,102	427	1,163
BKK29620	hisJ	1,242	1,127	0,106	116	1,113
BKK30160	ytcQ	1,242	1,237	0,132	33	1,222
BKK20200	yorZ	1,238	1,176	0,118	242	1,162
BKK21360	yomH	1,238	1,169	0,108	660	1,155
BKK30190	bioI	1,237	1,170	0,111	146	1,156
BKK36940	ywlD	1,237	0,987	0,067	60	0,975
BKK18470	proJ	1,233	1,186	0,107	362	1,172
BKK21560	yokK	1,232	1,168	0,113	374	1,153
BKK06319	ydzX	1,231	1,193	0,111	263	1,179
BKK11740	cotZ	1,229	1,177	0,114	369	1,163
BKK32540	bsn	1,223	1,153	0,085	138	1,139
BKK06120	ydjB	1,222	1,153	0,103	162	1,139
BKK29690	acuA	1,222	1,153	0,091	97	1,139
BKK03070	mdr	1,221	1,216	0,071	60	1,201
BKK32570	frlD	1,218	0,987	0,085	108	0,975
BKK18330	ppsB	1,217	1,186	0,104	265	1,172
BKK03660	yclE	1,216	1,152	0,084	77	1,138
BKK10580	yhjO	1,213	1,223	0,167	120	1,208
BKK35960	rbsB	1,21	1,175	0,108	324	1,161
BKK07390	yfmP	1,206	1,173	0,122	115	1,159
BKK03210	ycgN	1,205	1,144	0,117	102	1,130
BKK12830	spoIISA	1,205	1,182	0,112	386	1,168
BKK23670	yqkA	1,203	1,165	0,111	373	1,151
BKK40780	tetL	1,202	1,161	0,100	434	1,147
BKK37400	albD	1,199	1,180	0,104	91	1,166
BKK02770	yccK	1,193	1,173	0,111	306	1,160
BKK27090	aapA	1,193	1,191	0,095	644	1,177
BKK08450	yfiZ	1,191	0,987	0,077	110	0,975
BKK32390	yunF	1,19	1,162	0,104	91	1,148
BKK33880	yvbJ	1,19	0,987	0,073	46	0,975
BKK23330	ypuC	1,187	1,201	0,097	299	1,187
BKK11550	yjbH	1,185	1,197	0,107	164	1,183
BKK18890	yobF	1,181	1,151	0,087	139	1,138
BKK24110	yqzF	1,181	1,167	0,120	340	1,153
BKK30480	ytqA	1,179	1,170	0,119	97	1,156
BKK31800	yueF	1,174	1,161	0,113	30	1,148
BKK07580	citS	1,172	1,223	0,135	97	1,208
BKK23340	ypuB	1,172	1,201	0,096	492	1,187
BKK19200	desR	1,17	1,185	0,081	187	1,172
BKK13700	clpE	1,166	1,180	0,171	37	1,166
BKK02210	ybfH	1,165	1,143	0,101	173	1,130
BKK36970	iolJ	1,165	1,178	0,100	128	1,165
BKK19690	kamA	1,164	1,142	0,121	516	1,129
BKK30690	ytiB	1,163	1,169	0,101	118	1,156
BKK34450	sacB	1,163	1,148	0,118	344	1,135
BKK18310	ppsD	1,159	1,151	0,074	48	1,138
BKK24420	spoIIIAB	1,155	1,220	0,143	44	1,206
BKK20999	yoyJ	1,154	1,168	0,101	696	1,155
BKK37790	rocG	1,141	1,132	0,104	77	1,119
BKK22580	ypiB	1,14	1,113	0,101	432	1,100
BKK20840	yopM	1,138	1,167	0,109	463	1,153
BKK35120	yvlB	1,138	1,148	0,095	221	1,135
BKK11600	yjbM	1,13	1,202	0,123	874	1,189
BKK08500	yfhE	1,128	1,172	0,107	303	1,159
BKK25610	yqeM	1,125	1,177	0,086	116	1,164
BKK09860	khtT	1,124	1,222	0,149	81	1,208
BKK21230	yomT	1,118	1,168	0,121	384	1,155
BKK02400	ybgF	1,117	1,143	0,092	116	1,130
BKK06077	ydzW	1,114	1,171	0,098	123	1,158
BKK34530	cotR	1,114	1,148	0,101	139	1,135
BKK06300	cotA	1,113	1,192	0,137	231	1,179
BKK39100	yxiO	1,113	1,220	0,122	240	1,206
BKK19040	csaA	1,103	1,185	0,105	258	1,172
BKK29300	rbfK	1,1	1,152	0,102	129	1,139
BKK13090	ykkC	1,099	1,222	0,147	54	1,208
BKK15870	recG	1,094	1,170	0,106	87	1,157
BKK33070	gerAC	1,092	1,147	0,090	128	1,134
BKK19900	yotF	1,09	1,170	0,105	101	1,158
BKK11270	yjlB	1,086	1,202	0,139	1310	1,189
BKK03320	nasB	1,083	1,172	0,122	628	1,160
BKK14080	ykuH	1,083	1,179	0,121	496	1,166
BKK31530	yufM	1,082	1,164	0,123	167	1,151
BKK02160	yosD	1,078	1,141	0,145	94	1,129
BKK18770	cyeA	1,073	1,150	0,093	165	1,138
BKK22230	ypuF	1,071	1,200	0,109	87	1,187
BKK12620	xkdH	1,07	1,202	0,146	325	1,189
BKK32420	pucR	1,068	1,174	0,104	189	1,161
BKK08130	yfjE	1,06	0,985	0,086	35	0,975
BKK27340	yrrO	1,056	1,163	0,119	280	1,151
BKK34090	yvfR	1,056	1,162	0,114	355	1,149
BKK26800	yrpB	1,054	1,189	0,104	219	1,177
BKK30640	ytkC	1,053	1,150	0,100	144	1,138
BKK39930	yxaM	1,051	1,162	0,102	406	1,149
BKK32200	yutJ	1,05	1,151	0,091	155	1,139
BKK03160	ycgJ	1,049	1,172	0,103	409	1,160
BKK21290	yomN	1,047	1,167	0,102	284	1,155
BKK30230	bioK	1,047	1,163	0,122	231	1,151
BKK23730	yqjV	1,044	1,199	0,107	514	1,187
BKK09250	yhcX	1,04	1,189	0,106	161	1,177
BKK18820	yobB	1,038	1,174	0,114	363	1,162
BKK39340	hutP	1,031	1,177	0,110	128	1,165
BKK26210	yqaR	1,027	1,165	0,113	220	1,153
BKK06037	ydzT	1,026	1,179	0,128	384	1,167
BKK39330	abnB	1,022	1,159	0,097	385	1,147
BKK37780	rocA	1,02	1,159	0,119	283	1,147
BKK14520	adeC	1,018	1,191	0,134	117	1,179
BKK21250	yomR	1,016	1,167	0,109	564	1,155
BKK21050	yomN	1,015	1,167	0,111	523	1,155
BKK02120	ybeC	1,014	1,141	0,091	47	1,130
BKK08330	yfiN	1,011	1,171	0,101	487	1,159
BKK17410	cwlC	1,01	1,219	0,172	262	1,206
BKK27080	levR	1,009	1,189	0,102	272	1,177
BKK26380	yqaB	1,008	1,189	0,105	524	1,177
BKK05710	ydhD	1,005	1,214	0,085	97	1,201
BKK39690	iolH	1,004	1,176	0,107	125	1,165
BKK32720	yurZ	1,003	1,146	0,109	150	1,134
BKK02830	ycdF	1,001	1,171	0,097	372	1,160
BKK18670	yoaN	1,001	1,149	0,087	138	1,138
BKK28340	ysnF	1,001	1,171	0,102	308	1,160
BKK31260	mcpB	1	1,173	0,113	392	1,161
BKK27440	glnH	0,995	1,171	0,111	306	1,160
BKK16330	cheY	0,994	1,173	0,112	242	1,162
BKK29390	ytml	0,992	1,173	0,112	387	1,161
BKK04800	ydcL	0,989	1,159	0,119	400	1,147
BKK27840	safA	0,988	1,165	0,122	303	1,153
BKK11210	argB	0,985	1,195	0,107	73	1,183
BKK34190	yvfH	0,985	1,146	0,112	621	1,135
BKK39220	yxxG	0,984	1,176	0,119	107	1,165
BKK27880	nifS	0,982	1,162	0,113	333	1,151
BKK14880	ctaB	0,981	1,191	0,144	304	1,179
BKK18980	yobJ	0,981	1,173	0,107	322	1,162
BKK39210	yxiF	0,98	1,176	0,087	93	1,165
BKK04120	yczI	0,979	1,176	0,071	157	1,165
BKK17530	ynoE	0,978	1,173	0,109	333	1,162
BKK34410	yveG	0,977	1,146	0,127	133	1,135
BKK40060	gntK	0,976	1,179	0,123	198	1,168
BKK05170	ydeE	0,975	0,985	0,090	172	0,975
BKK15470	pyrR	0,975	1,151	0,074	139	1,139
BKK27060	levE	0,973	1,162	0,119	322	1,151
BKK11880	metC	0,972	1,201	0,109	76	1,189
BKK33170	yvrB	0,971	1,145	0,072	109	1,134
BKK05300	ydeQ	0,969	1,171	0,111	442	1,160
BKK24750	yqhB	0,966	1,175	0,101	199	1,164
BKK22150	ypvA	0,965	1,111	0,129	31	1,100
BKK30890	ytxO	0,965	1,167	0,102	119	1,156
BKK37410	albE	0,963	1,161	0,113	302	1,149
BKK09280	glpF	0,962	1,179	0,100	204	1,168
BKK11860	yjcH	0,962	1,200	0,130	1045	1,189
BKK13390	ykoT	0,962	1,178	0,160	133	1,166
BKK25910	yqxH	0,951	1,169	0,093	161	1,158
BKK30510	ytpA	0,951	1,167	0,088	123	1,156
BKK14100	ykuJ	0,949	1,150	0,096	157	1,139
BKK33050	gerAA	0,948	1,145	0,108	98	1,134
BKK27460	glnP	0,944	1,234	0,069	98	1,222
BKK04140	pbpC	0,941	1,213	0,074	38	1,201
BKK35930	rbsD	0,94	1,160	0,109	441	1,149
BKK27540	yrvM	0,937	1,164	0,118	194	1,153
BKK18510	yoxC	0,936	1,183	0,106	142	1,172
BKK34340	epsD	0,935	1,177	0,099	149	1,166
BKK32840	fadN	0,933	1,145	0,109	126	1,134
BKK29990	pbuO	0,932	1,233	0,098	43	1,222
BKK40080	yoxO	0,93	1,179	0,112	275	1,168
BKK40470	yycC	0,929	1,177	0,122	74	1,166
BKK36540	ywnJ	0,925	1,125	0,114	142	1,114
BKK20410	yorE	0,921	1,173	0,108	274	1,162
BKK30500	ytpB	0,921	1,167	0,117	135	1,156
BKK28060	spoIIB	0,918	1,233	0,069	47	1,222
BKK02580	ycbO	0,912	1,140	0,092	111	1,130
BKK08260	yfiG	0,912	1,219	0,123	61	1,208
BKK08460	yoaE	0,912	1,179	0,110	271	1,168
BKK11929	yjzG	0,91	1,173	0,124	238	1,163
BKK15690	ynzO	0,907	1,145	0,113	146	1,134
BKK35500	degS	0,907	1,195	0,096	368	1,184
BKK21080	yonI	0,894	1,165	0,111	263	1,155
BKK38170	qoxA	0,892	1,167	0,111	118	1,157
BKK06230	ioIT	0,891	1,212	0,099	51	1,201
BKK30350	yttB	0,891	1,233	0,104	53	1,222
BKK29600	braB	0,888	1,150	0,097	141	1,139
BKK32220	yutI	0,888	1,158	0,116	96	1,148
BKK07910	yfkF	0,887	1,173	0,111	307	1,161
BKK18720	yoaR	0,887	1,172	0,117	408	1,162
BKK27430	glnQ	0,885	1,233	0,073	63	1,222
BKK33270	yvrD	0,885	1,177	0,082	100	1,166
BKK02850	adcA	0,882	1,212	0,185	62	1,201
BKK28530	etfB	0,88	1,170	0,104	54	1,160
BKK15810	spoVM	0,875	1,217	0,143	136	1,206
BKK18849	yozV	0,872	1,148	0,096	136	1,138
BKK12500	xkdA	0,87	1,199	0,131	581	1,189
BKK21180	yomY	0,87	1,165	0,105	121	1,155
BKK30710	ythA	0,869	1,171	0,110	517	1,161
BKK32380	yunE	0,869	1,233	0,088	187	1,222
BKK30659	ytzI	0,864	1,166	0,099	112	1,156
BKK29630	ytzD	0,863	1,161	0,103	266	1,151
BKK40022	yxaC	0,862	1,159	0,118	436	1,149
BKK29940	ytIR	0,859	1,166	0,091	104	1,156
BKK13620	mtnD	0,858	1,173	0,106	247	1,163
BKK05670	ydaJ	0,856	1,177	0,083	202	1,167
BKK03680	yclG	0,855	1,148	0,083	150	1,138
BKK39540	yxeL	0,853	1,176	0,161	61	1,166
BKK17680	thyA	0,843	1,172	0,102	391	1,162
BKK07735	yfIB	0,842	1,169	0,116	352	1,159
BKK06170	ydjE	0,841	1,169	0,109	242	1,160
BKK11490	ydzN	0,84	1,193	0,108	104	1,183
BKK30990	thiT	0,839	1,232	0,144	449	1,222
BKK18110	ynfP	0,838	1,181	0,097	86	1,172
BKK26490	yrkJ	0,836	1,163	0,118	313	1,153
BKK37280	narD	0,836	1,128	0,095	66	1,119
BKK06480	purL	0,835	1,157	0,105	230	1,147
BKK30135	yteP	0,834	1,167	0,105	100	1,158
BKK32770	yusE	0,831	1,144	0,103	127	1,134
BKK15250	ylxW	0,829	1,178	0,106	230	1,168
BKK29160	ytvI	0,829	0,983	0,092	59	0,975
BKK35050	yvnA	0,828	1,171	0,104	484	1,161
BKK38330	lrgB	0,825	1,176	0,115	146	1,166
BKK17020	ymcA	0,823	1,173	0,121	89	1,164
BKK06240	bdhA	0,822	1,189	0,118	208	1,179
BKK14560	defB	0,822	1,189	0,111	157	1,179
BKK23180	spmA	0,821	1,161	0,110	214	1,151
BKK33310	fhuB	0,821	1,176	0,089	132	1,166
BKK16260	ylxF	0,819	1,172	0,105	385	1,163
BKK19000	yobL	0,818	1,171	0,109	228	1,162
BKK05540	ydfS	0,817	1,157	0,108	443	1,147
BKK16870	ymfI	0,815	1,173	0,127	152	1,164
BKK23960	artR	0,815	1,147	0,126	64	1,138
BKK34440	pbpE	0,815	1,159	0,104	531	1,149
BKK17180	pksJ	0,813	1,173	0,138	53	1,164
BKK03940	ycnI	0,808	1,167	0,103	145	1,158
BKK16940	recA	0,808	1,171	0,136	314	1,162
BKK02440	glnJ	0,805	1,139	0,095	145	1,130
BKK36820	atpG	0,805	1,157	0,108	161	1,147
BKK15490	pyrB	0,801	1,165	0,095	92	1,156
BKK16890	ymfK	0,801	1,173	0,116	110	1,164
BKK38160	qoxB	0,8	1,157	0,106	300	1,147
BKK20940	yopC	0,797	1,164	0,106	347	1,155
BKK23740	yqjU	0,796	1,197	0,101	476	1,163
BKK04260	topB	0,794	1,169	0,124	161	1,160
BKK13070	ykkA	0,794	1,176	0,091	75	1,166
BKK36140	ywqO	0,792	1,123	0,129	34	1,114
BKK38670	yxlE	0,791	1,174	0,100	71	1,165
BKK04970	yddH	0,788	1,167	0,103	187	1,158
BKK15040	ylbK	0,786	1,188	0,099	57	1,179
BKK35980	ywsA	0,785	1,166	0,104	214	1,157
BKK12170	yjgD	0,782	1,198	0,171	663	1,189
BKK05810	gmuB	0,781	1,176	0,070	262	1,167
BKK04550	ydbP	0,78	1,174	0,115	38	1,165
BKK16980	spoVS	0,78	1,171	0,114	489	1,162
BKK39110	deaD	0,78	1,174	0,108	71	1,165
BKK14930	ctoG	0,778	1,188	0,084	340	1,179
BKK38250	ywbO	0,775	1,166	0,128	73	1,157
BKK20570	yoqN	0,773	1,138	0,129	216	1,129
BKK22430	panB	0,77	1,148	0,093	106	1,139
BKK05030	yddM	0,763	1,168	0,114	341	1,160
BKK07210	yetK	0,763	1,177	0,114	305	1,168
BKK21270	yomP	0,759	1,164	0,106	512	1,155
BKK34350	epsC	0,759	1,158	0,105	493	1,149
BKK18000	citB	0,758	1,171	0,116	333	1,162
BKK23200	ypuI	0,758	1,196	0,113	175	1,187
BKK14690	yktD	0,757	1,188	0,123	159	1,179
BKK38220	ywcC	0,755	1,127	0,129	194	1,119
BKK39910	yxnB	0,754	1,173	0,116	165	1,165
BKK02650	pcp	0,751	1,156	0,099	407	1,147
BKK04110	lipC	0,749	1,168	0,123	363	1,160
BKK33150	yvqK	0,749	1,143	0,105	183	1,134
BKK04690	rsbT	0,747	1,174	0,125	153	1,165
BKK17480	ynzF	0,747	1,171	0,107	549	1,162
BKK38760	cydA	0,746	1,173	0,112	80	1,165
BKK28150	hemC	0,744	1,168	0,122	118	1,160
BKK12810	xlyA	0,738	1,171	0,124	192	1,163
BKK37730	bacB	0,738	1,158	0,110	383	1,149
BKK16880	ymfJ	0,737	1,172	0,115	163	1,164
BKK25800	ygcK	0,735	1,162	0,119	297	1,153
BKK20920	yopE	0,734	1,163	0,093	358	1,155
BKK32340	lytH	0,734	1,231	0,068	103	1,222
BKK22670	trpD	0,727	1,108	0,112	210	1,100
BKK34860	hisI	0,727	1,143	0,118	365	1,135
BKK08750	ygxA	0,722	1,171	0,117	280	1,163
BKK07300	yfnE	0,721	0,982	0,107	90	0,975
BKK17310	ymaG	0,72	1,172	0,103	209	1,164
BKK08220	yfiC	0,719	1,217	0,149	146	1,208
BKK20790	yopR	0,719	1,163	0,109	540	1,155
BKK10330	yhfQ	0,717	1,171	0,101	412	1,163
BKK04839	ydzL	0,716	1,173	0,101	120	1,165
BKK21370	yomG	0,713	1,163	0,115	493	1,155
BKK24190	yqjI	0,713	1,215	0,104	36	1,206
BKK39139	yxzJ	0,713	1,173	0,106	60	1,165
BKK10810	yisP	0,711	1,192	0,109	100	1,183
BKK38980	scoB	0,71	1,175	0,111	95	1,166
BKK07310	yfnD	0,706	1,187	0,117	483	1,179

Sup. Table 3: Cell width of mutants of the BKK collection (continued)

BKK name[1]	gene	screening delta[2] (%)	average width (µ	+/-	nb	ADP
BKK22950	ypdA	0,704	1,108	0,122	68	1,100
BKK16280	flgD	0,7	1,176	0,110	238	1,168
BKK19980	yosW	0,699	1,170	0,113	242	1,162
BKK12010	manP	0,698	1,197	0,110	207	1,189
BKK24400	spoIIIAD	0,697	1,215	0,156	73	1,206
BKK00760	pabC	0,694	1,166	0,108	91	1,158
BKK13760	ykvN	0,694	1,171	0,104	317	1,163
BKK19940	yotB	0,694	1,170	0,112	401	1,162
BKK39310	yxiC	0,693	1,173	0,095	154	1,165
BKK20620	yoqI	0,69	1,170	0,123	243	1,162
BKK30060	ytfP	0,688	1,164	0,097	179	1,156
BKK07410	yfmN	0,686	1,167	0,117	156	1,159
BKK12539	ykzK	0,685	1,157	0,127	328	1,149
BKK12600	xkdF	0,684	1,174	0,139	219	1,166
BKK38620	aag	0,684	1,173	0,118	86	1,165
BKK12030	yjdF	0,683	1,217	0,121	62	1,208
BKK27110	yrhO	0,681	1,185	0,110	374	1,177
BKK27230	yrhD	0,681	1,185	0,110	374	1,177
BKK33460	yvgT	0,681	1,142	0,102	219	1,134
BKK31670	yuxO	0,677	1,156	0,143	105	1,148
BKK33910	yxjL	0,677	1,215	0,133	72	1,206
BKK02260	ybfK	0,676	1,167	0,107	479	1,160
BKK02940	yceH	0,672	1,138	0,090	112	1,130
BKK05000	yddK	0,668	1,167	0,120	413	1,160
BKK18050	yneQ	0,668	1,170	0,120	382	1,162
BKK34000	cyeB	0,665	1,157	0,102	296	1,149
BKK21120	yonE	0,662	1,163	0,103	431	1,155
BKK17750	yndD	0,657	1,171	0,099	143	1,164
BKK24020	bmrR	0,655	1,161	0,113	349	1,153
BKK24230	spoIVB	0,651	1,214	0,151	171	1,206
BKK28140	hemD	0,649	1,167	0,115	32	1,160
BKK30670	luxS	0,648	1,164	0,092	146	1,156
BKK12630	xkdI	0,643	1,197	0,115	243	1,189
BKK37850	spsG	0,643	1,126	0,106	156	1,119
BKK01480	truA	0,642	1,167	0,109	987	1,160
BKK03970	ycnL	0,64	1,172	0,075	138	1,165
BKK02700	estA	0,637	1,209	0,069	46	1,201
BKK04690	yybC	0,637	1,192	0,082	150	1,184
BKK25530	spoIIP	0,634	1,230	0,122	228	1,222
BKK39640	yxdL	0,634	1,230	0,123	41	1,222
BKK18460	gltC	0,633	1,145	0,104	86	1,138
BKK09070	yhcG	0,632	1,184	0,119	349	1,177
BKK39570	yxeF	0,632	1,230	0,084	41	1,222
BKK37970	ung	0,628	1,126	0,117	234	1,119
BKK05400	ydfG	0,627	1,175	0,090	283	1,167
BKK10810	yizA	0,627	1,191	0,116	78	1,183
BKK24250	ahrC	0,626	1,194	0,115	224	1,187
BKK06074	ydzW	0,624	1,175	0,131	184	1,167
BKK15650	yloB	0,623	1,141	0,087	176	1,134
BKK22570	yjiF	0,623	1,107	0,115	586	1,100
BKK08190	malR	0,622	1,166	0,120	268	1,159
BKK29570	sspA	0,62	1,120	0,105	74	1,113
BKK14450	yktB	0,618	1,186	0,131	229	1,179
BKK38200	galK	0,615	1,155	0,101	480	1,147
BKK17699	ynzK	0,614	1,171	0,101	59	1,164
BKK21450	yoIJ	0,614	1,158	0,108	327	1,151
BKK40860	yyaH	0,611	1,192	0,115	101	1,184
BKK07450	yfmJ	0,604	1,166	0,110	189	1,159
BKK23390	spoVAF	0,602	1,214	0,144	33	1,206
BKK39490	yxeN	0,597	1,145	0,103	70	1,138
BKK02450	glnL	0,596	1,167	0,118	302	1,160
BKK12050	yjdH	0,596	1,196	0,132	53	1,189
BKK02360	nagBB	0,593	1,137	0,109	196	1,130
BKK08140	yfjD	0,591	1,166	0,104	594	1,159
BKK00420	ksgA	0,585	1,154	0,103	542	1,147
BKK12610	xkdG	0,585	1,196	0,089	288	1,189
BKK16020	rimM	0,584	1,163	0,115	58	1,156
BKK11470	oppF	0,58	1,169	0,115	270	1,163
BKK16299	ylzI	0,576	1,169	0,124	264	1,163
BKK36870	atpB	0,568	1,173	0,101	96	1,166
BKK38300	ywdD	0,568	1,125	0,083	111	1,119
BKK40380	yycI	0,565	1,136	0,096	121	1,130
BKK00890	yacL	0,563	1,154	0,107	520	1,147
BKK30310	ytwF	0,563	1,163	0,099	107	1,156
BKK15880	fapR	0,562	1,125	0,126	861	1,119
BKK38030	ywdA	0,562	1,125	0,095	98	1,119
BKK21400	yomD	0,558	1,161	0,101	255	1,155
BKK30530	ytnA	0,558	1,162	0,103	83	1,156
BKK31340	yugM	0,553	1,144	0,094	147	1,138

BKK name[1]	gene	screening delta[2] (%)	average width (µ	+/-	nb	ADP
BKK14790	ylaI	0,552	1,186	0,101	798	1,179
BKK36930	ywIE	0,551	1,163	0,095	139	1,157
BKK19060	yobR	0,546	1,144	0,096	95	1,138
BKK32550	yurJ	0,544	1,146	0,091	112	1,139
BKK00880	disA	0,541	1,144	0,128	65	1,138
BKK19090	yobU	0,541	1,168	0,116	346	1,162
BKK29950	ytlQ	0,541	1,162	0,090	85	1,156
BKK16780	rnjB	0,54	1,170	0,103	100	1,164
BKK05910	ydiB	0,539	1,174	0,094	138	1,167
BKK17200	pksM	0,538	1,170	0,123	99	1,164
BKK33440	cysJ	0,538	1,140	0,123	146	1,134
BKK22310	recU	0,533	1,106	0,126	173	1,100
BKK06640	yerI	0,531	1,185	0,138	221	1,179
BKK14030	ykuC	0,53	1,213	0,164	91	1,206
BKK35590	tuaC	0,53	1,172	0,112	139	1,166
BKK13250	ykoG	0,529	1,173	0,098	350	1,166
BKK12100	yjeA	0,526	1,169	0,111	403	1,163
BKK24550	gcvPB	0,522	1,193	0,116	252	1,187
BKK04880	ydcS	0,517	1,171	0,133	55	1,165
BKK03750	yclJ	0,515	1,171	0,090	172	1,165
BKK31380	yuzA	0,512	1,154	0,097	152	1,148
BKK36620	ywnB	0,508	1,153	0,102	370	1,147
BKK15069	ylzH	0,506	1,145	0,123	68	1,139
BKK30470	ytzC	0,505	1,162	0,102	111	1,156
BKK39960	yxaI	0,505	1,171	0,101	113	1,165
BKK38050	sacP	0,503	1,124	0,113	151	1,119
BKK11350	yjaZ	0,502	1,168	0,108	254	1,163
BKK13820	ykvT	0,501	1,172	0,104	198	1,166
BKK32760	yusD	0,498	1,167	0,114	491	1,161
BKK36990	ywkF	0,497	1,163	0,124	81	1,157
BKK18908	yozY	0,495	1,144	0,094	152	1,138
BKK26510	yrkH	0,495	1,183	0,103	265	1,177
BKK22560	qcrA	0,492	1,106	0,116	509	1,100
BKK39780	iolS	0,491	1,170	0,099	67	1,165
BKK09640	yhdY	0,488	1,183	0,113	140	1,177
BKK37720	bacC	0,487	1,124	0,097	60	1,119
BKK16370	fliR	0,486	1,167	0,104	307	1,162
BKK37900	spsB	0,485	1,155	0,109	325	1,149
BKK19620	yodJ	0,484	1,212	0,137	277	1,206
BKK16090	sucC	0,482	1,165	0,106	324	1,160
BKK40870	ccpB	0,48	1,153	0,106	296	1,147
BKK18920	phrK	0,479	1,174	0,123	361	1,168
BKK22940	prsW	0,479	1,212	0,113	177	1,206
BKK08700	ygaE	0,475	1,164	0,116	366	1,159
BKK03720	gerKB	0,474	1,170	0,119	187	1,165
BKK31330	yugN	0,474	1,161	0,101	90	1,156
BKK26140	yqbE	0,472	1,157	0,107	509	1,151
BKK23519	yqzK	0,469	1,193	0,103	228	1,187
BKK20050	yosQ	0,467	1,135	0,128	244	1,129
BKK24240	recN	0,462	1,193	0,098	405	1,187
BKK17290	ebrB	0,459	1,212	0,123	103	1,206
BKK12490	yjqC	0,457	1,105	0,132	479	1,113
BKK32710	polYB	0,455	1,192	0,105	309	1,187
BKK04680	rsbS	0,454	1,170	0,106	190	1,165
BKK13720	queC	0,453	1,172	0,115	305	1,166
BKK04920	yddC	0,451	1,153	0,104	382	1,147
BKK18500	yoxD	0,441	1,167	0,116	316	1,162
BKK28680	glcD	0,441	1,144	0,102	258	1,139
BKK14210	ykuT	0,44	1,168	0,122	316	1,163
BKK32350	yunB	0,439	1,144	0,103	309	1,139
BKK39760	mmsA	0,439	1,163	0,085	196	1,158
BKK40030	yxaB	0,439	1,161	0,102	150	1,156
BKK16690	pnpA	0,437	1,167	0,103	296	1,162
BKK13160	ohrB	0,436	1,171	0,144	251	1,166
BKK13200	rsbRB	0,432	1,168	0,113	293	1,163
BKK40110	bglA	0,432	1,173	0,112	233	1,168
BKK02220	ybfI	0,431	1,135	0,092	170	1,130
BKK28510	xsa	0,428	1,165	0,117	121	1,160
BKK32960	yusY	0,423	1,171	0,100	174	1,166
BKK05270	ydzF	0,422	1,165	0,117	427	1,160
BKK32360	ppiB	0,418	1,158	0,109	475	1,153
BKK38320	cidA	0,418	1,154	0,114	449	1,149
BKK23380	lysA	0,417	1,192	0,107	296	1,187
BKK32910	yusS	0,417	1,139	0,099	102	1,134
BKK23850	zwf	0,414	1,158	0,119	278	1,153
BKK39450	yxeR	0,414	1,211	0,143	161	1,206
BKK17170	pksI	0,412	1,168	0,122	55	1,164
BKK19490	gerT	0,408	1,134	0,110	855	1,129
BKK16290	flgE	0,406	1,167	0,105	264	1,163

BKK name[1]	gene	screening delta[2] (%)	average width (µ	+/-	nb	ADP
BKK39410	nupC	0,406	1,211	0,121	169	1,206
BKK39940	yxaL	0,406	1,169	0,120	149	1,165
BKK12320	yjmC	0,402	1,194	0,112	487	1,189
BKK26870	yraN	0,401	1,227	0,077	226	1,222
BKK00540	yabK	0,399	1,168	0,116	119	1,164
BKK23720	yqzH	0,399	1,192	0,111	149	1,187
BKK17640	alrB	0,398	1,211	0,140	126	1,206
BKK20900	yopG	0,397	1,159	0,111	293	1,155
BKK12069	yizH	0,394	1,167	0,121	380	1,163
BKK05580	ydgC	0,389	1,172	0,078	186	1,167
BKK27120	sigW	0,389	1,181	0,111	106	1,177
BKK15500	pyrC	0,388	1,162	0,099	114	1,158
BKK33049	yvzF	0,388	1,139	0,107	137	1,134
BKK24100	bkdR	0,386	1,162	0,101	76	1,158
BKK36800	atpC	0,386	1,119	0,100	114	1,114
BKK20870	yopJ	0,385	1,156	0,121	241	1,151
BKK29250	nrnA	0,383	1,117	0,091	182	1,113
BKK06010	ydiL	0,381	1,172	0,093	276	1,167
BKK14970	ylbD	0,376	1,172	0,101	330	1,168
BKK28720	abfA	0,375	1,144	0,093	167	1,139
BKK34020	yvbX	0,375	1,139	0,089	166	1,134
BKK31360	yugK	0,372	1,152	0,115	238	1,148
BKK17630	yncC	0,371	1,211	0,179	185	1,206
BKK35340	fliD	0,37	1,139	0,112	342	1,135
BKK01600	ybbA	0,369	1,181	0,096	187	1,177
BKK27980	spoIVFA	0,369	1,227	0,103	37	1,222
BKK08470	yfhB	0,363	1,163	0,106	647	1,159
BKK04280	ydaK	0,359	1,169	0,102	90	1,165
BKK17140	pksF	0,358	1,168	0,102	125	1,164
BKK22170	ypsC	0,356	1,143	0,091	185	1,139
BKK22140	kduD	0,355	1,104	0,136	162	1,100
BKK34950	pelC	0,355	1,170	0,118	121	1,166
BKK03520	srfAD	0,349	1,206	0,110	358	1,201
BKK33190	yvrD	0,349	1,138	0,096	120	1,134
BKK39770	iolR	0,349	1,151	0,099	347	1,147
BKK21620	yokE	0,345	1,157	0,112	268	1,153
BKK06400	yebE	0,338	1,162	0,108	155	1,158
BKK14730	ylaC	0,337	1,183	0,164	47	1,179
BKK08350	estB	0,336	1,166	0,107	343	1,163
BKK31060	gbsA	0,336	1,162	0,101	87	1,158
BKK14120	ykzF	0,333	1,170	0,118	709	1,166
BKK03740	yclI	0,327	1,163	0,114	365	1,160
BKK33840	yvbF	0,322	1,138	0,093	94	1,134
BKK22290	psd	0,318	1,151	0,109	486	1,147
BKK04960	yddG	0,314	1,205	0,144	424	1,201
BKK15140	mraW	0,311	1,210	0,144	38	1,206
BKK20000	yosU	0,309	1,133	0,144	350	1,129
BKK23140	resB	0,309	1,191	0,104	848	1,187
BKK36840	atpH	0,309	1,118	0,117	116	1,114
BKK10490	sipV	0,308	1,166	0,109	259	1,163
BKK08610	yfhO	0,306	1,172	0,112	328	1,168
BKK18640	yoaK	0,304	1,141	0,090	121	1,138
BKK20020	yoS	0,303	1,133	0,130	174	1,129
BKK36720	ywmE	0,302	1,151	0,112	230	1,147
BKK08010	yfjP	0,296	1,162	0,113	362	1,159
BKK28550	fadR	0,294	1,141	0,088	128	1,138
BKK16270	fliK	0,293	1,171	0,120	278	1,168
BKK27330	udk	0,291	0,978	0,120	35	0,975
BKK04850	ydcP	0,29	1,163	0,107	520	1,160
BKK37260	nagZ	0,29	1,122	0,102	110	1,119
BKK00840	mcsA	0,287	1,161	0,106	73	1,158
BKK23790	mcsB	0,277	1,190	0,109	238	1,187
BKK08840	ssuA	0,273	1,171	0,111	284	1,168
BKK17420	spoVK	0,273	1,210	0,144	106	1,206
BKK33340	folEB	0,272	1,133	0,087	153	1,130
BKK40359	yyzO	0,271	1,153	0,114	356	1,149
BKK32450	pucL	0,269	1,143	0,082	149	1,139
BKK23550	mleA	0,268	1,190	0,103	198	1,187
BKK08000	yfjQ	0,266	1,162	0,101	373	1,159
BKK33820	opuCB	0,266	1,137	0,091	98	1,134
BKK39610	yxeB	0,266	1,151	0,104	590	1,147
BKK00570	yabM	0,265	1,141	0,131	38	1,138
BKK27830	coxA	0,264	1,163	0,101	115	1,160
BKK28220	clpX	0,263	1,163	0,112	147	1,160
BKK15400	ylmG	0,258	1,182	0,156	135	1,179
BKK32780	yusF	0,257	1,137	0,100	178	1,134
BKK25940	yqcD	0,252	1,154	0,104	313	1,151
BKK24310	folD	0,25	1,154	0,120	106	1,151

BKK name[1]	gene	screening delta[2] (%)	average width (µ	+/-	nb	ADP
BKK23750	yqjT	0,249	0,977	0,082	55	0,975
BKK33390	yvgM	0,249	1,141	0,099	123	1,138
BKK07600	yfIP	0,247	1,162	0,103	326	1,159
BKK19010	yobM	0,246	1,174	0,088	124	1,172
BKK10730	yisI	0,243	1,186	0,121	118	1,183
BKK30020	ytzE	0,243	1,159	0,101	138	1,156
BKK22610	tyrA	0,242	1,103	0,142	69	1,100
BKK33400	yvgN	0,236	1,137	0,097	121	1,134
BKK02370	ybgA	0,235	1,133	0,097	68	1,130
BKK18240	yngHA	0,235	1,174	0,104	136	1,172
BKK05070	yddQ	0,232	1,162	0,107	412	1,160
BKK25340	phoH	0,231	1,166	0,098	211	1,164
BKK09500	yhdK	0,23	1,211	0,198	77	1,208
BKK10030	hinT	0,23	1,180	0,108	124	1,177
BKK31010	yuoG	0,228	1,159	0,103	99	1,156
BKK13410	ykoV	0,227	1,169	0,121	65	1,166
BKK14780	ylaH	0,226	1,182	0,104	302	1,179
BKK17570	xynP	0,226	1,166	0,124	87	1,164
BKK05480	ydfN	0,225	1,170	0,100	142	1,167
BKK26120	yqbG	0,225	1,179	0,090	222	1,177
BKK03790	yclM	0,224	1,168	0,101	695	1,165
BKK13169	ykzN	0,223	1,211	0,118	207	1,208
BKK20350	yorK	0,222	1,142	0,109	41	1,139
BKK09810	yhaZ	0,22	1,165	0,110	260	1,163
BKK04730	sigB	0,219	1,167	0,175	164	1,165
BKK31960	dhbF	0,219	1,169	0,112	184	1,166
BKK37070	rpmEA	0,217	1,160	0,098	107	1,158
BKK02550	ycbL	0,215	1,132	0,088	130	1,130
BKK26560	yrkC	0,215	1,156	0,102	583	1,153
BKK31910	yukE	0,215	1,142	0,101	140	1,139
BKK23650	yqkC	0,214	1,190	0,111	197	1,187
BKK14860	pycA	0,213	1,165	0,110	228	1,163
BKK30910	cotSA	0,213	1,158	0,095	140	1,156
BKK18860	yozH	0,212	1,164	0,110	370	1,162
BKK09550	yhdP	0,211	1,211	0,169	39	1,208
BKK10430	yhxD	0,211	1,186	0,119	104	1,183
BKK05750	ydhG	0,203	1,170	0,075	265	1,167
BKK36920	ywlF	0,203	1,159	0,145	30	1,157
BKK39880	yxbC	0,202	1,150	0,112	397	1,147
BKK40300	rapG	0,202	1,169	0,110	107	1,166
BKK30650	dps	0,201	1,158	0,115	113	1,156
BKK23170	spmB	0,197	1,189	0,090	186	1,187
BKK07700	nagP	0,196	1,161	0,113	434	1,159
BKK13260	ykoH	0,196	1,140	0,115	113	1,138
BKK08560	yfhJ	0,195	1,161	0,112	591	1,159
BKK32090	yuiA	0,195	1,150	0,121	401	1,148
BKK32510	pucA	0,192	1,150	0,098	85	1,148
BKK25030	yqgC	0,189	1,156	0,121	260	1,153
BKK39170	yxzG	0,184	1,167	0,110	78	1,165
BKK11870	metI	0,179	1,191	0,133	619	1,189
BKK28570	yshE	0,178	1,162	0,097	70	1,160
BKK34390	pnbA	0,177	1,137	0,108	195	1,135
BKK33220	yutC	0,175	1,168	0,103	241	1,166
BKK37040	ywkB	0,174	1,151	0,115	428	1,149
BKK33620	yvaK	0,167	1,136	0,096	148	1,134
BKK15080	rpmF	0,166	1,181	0,154	33	1,179
BKK00290	yaaQ	0,16	1,181	0,143	721	1,179
BKK10370	bioY	0,156	1,210	0,136	251	1,208
BKK33730	opuBA	0,155	1,163	0,114	603	1,161
BKK01389	ybzG	0,148	1,165	0,103	312	1,164
BKK03310	nasC	0,147	1,132	0,102	91	1,130
BKK01880	ybcI	0,145	1,140	0,108	157	1,138
BKK38800	yxkH	0,145	1,210	0,131	126	1,208
BKK37390	albC	0,144	1,151	0,103	476	1,149
BKK02050	ybdO	0,142	1,132	0,095	130	1,130
BKK17710	tatAD	0,142	1,165	0,110	135	1,164
BKK04890	ydcT	0,14	1,167	0,107	124	1,165
BKK24180	yqiK	0,138	1,155	0,117	264	1,153
BKK22169	ypzC	0,134	1,141	0,090	131	1,139
BKK16420	cheB	0,132	1,163	0,107	233	1,161
BKK33010	cssR	0,128	1,136	0,093	170	1,134
BKK30520	ytoA	0,126	1,157	0,103	133	1,156
BKK25450	prmA	0,125	1,165	0,086	229	1,164
BKK01970	skfG	0,123	1,141	0,092	231	1,139
BKK13220	thiW	0,123	1,210	0,185	60	1,208
BKK32970	yusV	0,121	1,168	0,104	140	1,166
BKK14740	yloD	0,12	1,169	0,109	459	1,168
BKK11940	yjcP	0,117	1,190	0,102	573	1,189

Sup. Table 3: Cell width of mutants of the BKK collection (continued)

BKK name[1]	gene	screening delta²(%)	average width (μ)	+/-	nb	ADP
BKK22750	menH	0,117	1,102	0,111	167	1,100
BKK01960	skfF	0,116	1,159	0,110	97	1,158
BKK06260	ydjN	0,115	1,180	0,126	106	1,179
BKK02019	ybzI	0,113	1,131	0,081	88	1,130
BKK38350	ywbE	0,113	1,120	0,108	196	1,119
BKK36320	ywpG	0,112	1,158	0,106	104	1,157
BKK27100	yrhP	0,111	1,223	0,067	74	1,222
BKK35330	fliS	0,109	1,136	0,116	72	1,135
BKK14600	pdhC	0,108	0,976	0,067	62	0,975
BKK08940	yhbD	0,101	1,169	0,111	312	1,168
BKK33370	yvgK	0,1	1,162	0,112	358	1,161
BKK32900	yusR	0,099	1,135	0,098	156	1,134
BKK31890	yukC	0,098	1,149	0,122	33	1,148
BKK18660	yoaM	0,094	1,163	0,113	539	1,162
BKK10450	yhjB	0,092	1,169	0,100	396	1,168
BKK21510	yolD	0,092	1,156	0,106	445	1,155
BKK14720	ylaB	0,088	1,169	0,114	205	1,168
BKK30250	ytaP	0,088	1,157	0,101	182	1,156
BKK35890	pgsC	0,088	1,150	0,114	508	1,149
BKK06470	purQ	0,087	1,180	0,117	328	1,179
BKK17090	pksB	0,087	1,208	0,142	95	1,206
BKK08780	ygaJ	0,084	1,160	0,108	183	1,159
BKK05040	yddN	0,08	1,161	0,110	508	1,160
BKK11840	yjcF	0,08	1,190	0,102	239	1,189
BKK21630	yokD	0,078	1,156	0,112	537	1,155
BKK31875	yukB	0,077	1,223	0,085	142	1,222
BKK16230	fliH	0,074	1,163	0,109	309	1,163
BKK09390	ygxB	0,073	1,178	0,105	122	1,177
BKK10870	yisU	0,071	1,184	0,122	89	1,183
BKK08779	ygzC	0,069	0,976	0,068	78	0,975
BKK31220	yuxG	0,068	1,157	0,090	121	1,156
BKK07360	yfmS	0,067	1,209	0,162	277	1,208
BKK20500	ligB	0,065	1,140	0,102	119	1,139
BKK20540	yoqR	0,065	1,140	0,096	140	1,139
BKK35040	yvnB	0,064	1,136	0,099	258	1,135
BKK21940	degR	0,062	1,156	0,106	328	1,155
BKK15090	ylbO	0,061	1,180	0,128	295	1,179
BKK22120	kdgR	0,061	1,154	0,105	309	1,153
BKK24800	yqgW	0,061	1,164	0,100	306	1,164
BKK11260	yjzC	0,059	1,184	0,108	83	1,183
BKK14240	rok	0,057	1,163	0,108	254	1,163
BKK03870	ycnE	0,054	1,166	0,109	252	1,165
BKK34060	yvfU	0,053	1,135	0,093	115	1,134
BKK18850	yobD	0,045	1,139	0,083	33	1,138
BKK38040	sacA	0,045	1,167	0,119	82	1,166
BKK14910	ctaE	0,044	1,180	0,124	362	1,179
BKK30570	ytmB	0,036	1,156	0,095	100	1,156
BKK11270	yjzD	0,034	1,184	0,137	84	1,183
BKK31170	yulF	0,033	1,156	0,105	133	1,156
BKK17920	ynzD	0,032	1,164	0,122	83	1,164
BKK19270	yozN	0,031	1,172	0,105	99	1,172
BKK13110	purU	0,025	1,163	0,122	377	1,163
BKK31190	yuiD	0,025	1,151	0,116	329	1,151
BKK39090	yxiP	0,023	1,165	0,111	167	1,165
BKK24170	mmgA	0,022	0,975	0,086	46	0,975
BKK16410	ylxH	0,021	1,156	0,093	246	1,156
BKK28710	cstA	0,021	1,156	0,093	246	1,156
BKK29960	ytlP	0,021	1,156	0,106	75	1,156
BKK12780	xepA	0,015	1,209	0,116	125	1,208
BKK05600	ydgE	0,014	1,167	0,078	333	1,167
BKK11370	appF	0,012	1,168	0,112	350	1,168
BKK10900	yisY	0,01	1,183	0,109	90	1,183
BKK11240	carB	0,01	1,183	0,099	77	1,183
BKK03260	ycgS	0,008	1,130	0,082	74	1,130
BKK16240	fliI	0,008	1,163	0,102	402	1,163
BKK20170	yosC	0,006	1,162	0,103	302	1,162
BKK06200	ybfG	0,005	1,160	0,111	334	1,160
BKK31280	yugU	0,005	1,158	0,098	182	1,158
BKK35510	yvyE	0,004	1,114	0,115	94	1,114
BKK33570	yvaE	0,003	1,222	0,096	64	1,222
BKK05380	ydfE	0	1,167	0,119	117	1,167
BKK38060	ywcJ	-0,001	1,158	0,104	159	1,158
BKK07380	yfmQ	-0,002	1,159	0,114	290	1,159
BKK04870	ydcR	-0,003	1,147	0,113	421	1,147
BKK13310	tnrA	-0,003	1,163	0,107	316	1,163
BKK11079	yizB	-0,005	1,183	0,106	129	1,183
BKK29340	yqjB	-0,007	1,187	0,107	578	1,187
BKK29460	moaB	-0,008	1,113	0,112	56	1,113
BKK35070	yvmC	-0,008	1,158	0,118	102	1,158
BKK33750	sdpA	-0,01	1,166	0,114	98	1,166
BKK03270	ycgT	-0,013	1,130	0,087	70	1,130
BKK19810	ypqP	-0,013	1,162	0,108	490	1,162
BKK03900	gabT	-0,016	1,159	0,116	439	1,160
BKK08280	yfiI	-0,016	1,159	0,109	485	1,159
BKK13020	ykgA	-0,017	1,166	0,113	222	1,166
BKK21980	ypdP	-0,017	1,155	0,104	410	1,155
BKK19080	yobT	-0,019	1,138	0,096	153	1,138
BKK11400	appC	-0,021	1,162	0,113	319	1,163
BKK09060	yhcF	-0,022	1,177	0,109	213	1,177
BKK40940	yyaD	-0,028	1,167	0,086	1044	1,167
BKK02680	lmrA	-0,031	1,130	0,091	136	1,130
BKK17470	ynxB	-0,038	1,161	0,104	322	1,162
BKK06140	gutR	-0,039	1,178	0,110	197	1,179
BKK10700	gerPC	-0,039	1,183	0,099	144	1,183
BKK13580	mtnE	-0,039	1,166	0,100	307	1,166
BKK29930	amyX	-0,044	1,156	0,096	114	1,156
BKK22010	exoA	-0,046	1,157	0,118	101	1,158
BKK17840	fosB	-0,048	1,163	0,101	79	1,164
BKK25420	ygeW	-0,049	1,137	0,093	161	1,138
BKK07610	citM	-0,052	1,208	0,162	70	1,208
BKK23970	artQ	-0,052	1,186	0,092	375	1,187
BKK29679	ytzK	-0,053	1,150	0,096	183	1,151
BKK06038	ydzT	-0,055	1,157	0,097	121	1,158
BKK10480	yhjE	-0,059	1,208	0,125	156	1,208
BKK39240	yxxF	-0,059	1,206	0,133	120	1,206
BKK02280	ybfM	-0,061	1,129	0,088	66	1,130
BKK08020	yfjO	-0,061	1,158	0,110	459	1,159
BKK21150	yonB	-0,067	1,154	0,116	258	1,155
BKK11020	yitK	-0,069	1,162	0,105	400	1,163
BKK36040	ywrJ	-0,072	1,160	0,105	395	1,161
BKK33160	yvrA	-0,076	1,133	0,087	67	1,134
BKK28280	leuA	-0,077	1,159	0,100	263	1,160
BKK22270	yppE	-0,081	1,099	0,127	308	1,100
BKK40420	purA	-0,082	1,147	0,103	301	1,147
BKK40530	cotF	-0,084	1,164	0,100	110	1,165
BKK23570	ansB	-0,086	1,186	0,099	322	1,187
BKK26110	yqbH	-0,088	1,176	0,101	82	1,177
BKK19230	yocI	-0,09	1,161	0,098	389	1,162
BKK05190	ydeG	-0,094	1,200	0,122	600	1,201
BKK12790	xhlA	-0,096	1,165	0,126	299	1,166
BKK18300	ppsE	-0,096	1,137	0,098	126	1,138
BKK21280	yomO	-0,096	1,154	0,102	187	1,155
BKK18930	yobH	-0,097	1,161	0,108	456	1,162
BKK23240	ypzK	-0,097	1,152	0,111	266	1,153
BKK12900	htrA	-0,103	1,161	0,108	201	1,163
BKK33530	yvaA	-0,104	1,133	0,107	126	1,134
BKK00270	yaaO	-0,105	1,188	0,154	255	1,189
BKK30760	mntB	-0,107	1,155	0,104	87	1,156
BKK17510	ynaC	-0,112	1,162	0,108	108	1,164
BKK02040	ybdN	-0,113	1,158	0,112	519	1,160
BKK12990	ykfC	-0,113	1,207	0,136	89	1,208
BKK25060	yqfZ	-0,113	1,152	0,115	437	1,153
BKK00930	cysE	-0,114	1,099	0,119	344	1,100
BKK30830	menF	-0,114	1,155	0,097	132	1,156
BKK24690	comGE	-0,115	1,150	0,117	411	1,151
BKK28430	sdhB	-0,117	0,974	0,088	37	0,975
BKK11930	yjcO	-0,119	1,188	0,115	502	1,189
BKK33030	yuxN	-0,119	1,133	0,092	114	1,134
BKK16460	cheD	-0,12	1,160	0,105	594	1,161
BKK25100	zur	-0,121	1,162	0,099	222	1,164
BKK03670	dtpT	-0,122	1,137	0,098	191	1,138
BKK11200	argJ	-0,122	1,167	0,101	274	1,168
BKK06770	yeeB	-0,123	1,177	0,128	196	1,179
BKK24870	ycsN	-0,126	1,156	0,110	186	1,158
BKK30360	yorJ	-0,126	1,138	0,109	54	1,139
BKK24090	ptb	-0,127	1,186	0,113	204	1,187
BKK03910	gabD	-0,129	1,137	0,094	125	1,138
BKK10550	ntdA	-0,13	1,167	0,107	302	1,168
BKK02900	yceD	-0,131	1,128	0,092	151	1,130
BKK04340	ydaP	-0,131	1,163	0,102	180	1,165
BKK19250	yocL	-0,131	1,170	0,087	154	1,172
BKK35699	yvzE	-0,132	1,160	0,104	471	1,161
BKK22650	trpF	-0,133	1,099	0,099	205	1,100
BKK23350	ypzD	-0,133	1,185	0,095	555	1,187
BKK02740	natR	-0,134	1,128	0,081	33	1,130
BKK31400	ykoU	-0,134	1,165	0,100	214	1,166
BKK21190	yomX	-0,135	1,150	0,103	417	1,151
BKK35680	ggaB	-0,137	1,160	0,101	475	1,161
BKK17500	ynaB	-0,138	1,160	0,102	456	1,162
BKK03500	comS	-0,14	1,156	0,117	101	1,158
BKK18200	yngD	-0,144	1,170	0,116	40	1,172
BKK08410	yfiV	-0,147	1,157	0,110	450	1,159
BKK25590	comEA	-0,147	1,162	0,108	120	1,164
BKK19600	yodH	-0,149	0,974	0,082	95	0,975
BKK22290	sspM	-0,157	1,099	0,129	223	1,100
BKK11530	coiA	-0,158	1,181	0,118	141	1,183
BKK11160	yitW	-0,163	1,181	0,102	106	1,183
BKK23800	proI	-0,163	1,152	0,110	315	1,153
BKK40800	yyaN	-0,163	1,183	0,110	194	1,184
BKK17660	yncF	-0,165	1,160	0,106	345	1,162
BKK39660	qodI	-0,165	1,163	0,121	126	1,165
BKK29750	aroA	-0,167	1,138	0,091	202	1,139
BKK16380	flhB	-0,168	1,151	0,108	434	1,153
BKK38900	yxjM	-0,169	1,206	0,165	87	1,208
BKK10240	yhfl	-0,177	1,175	0,097	124	1,177
BKK28290	ilvC	-0,179	1,158	0,093	161	1,160
BKK36790	ywmA	-0,179	1,155	0,107	122	1,157
BKK12560	xpf	-0,185	1,187	0,140	193	1,189
BKK33520	csoR	-0,19	1,132	0,095	118	1,134
BKK14820	ylaL	-0,191	1,204	0,122	65	1,206
BKK15000	yisV	-0,192	1,177	0,122	43	1,179
BKK28160	hemX	-0,197	1,157	0,121	145	1,160
BKK39980	qodL	-0,198	1,145	0,110	477	1,147
BKK36340	ywpE	-0,202	1,155	0,133	66	1,157
BKK18570	yodE	-0,206	1,136	0,092	106	1,138
BKK19610	yodI	-0,208	1,127	0,127	787	1,129
BKK02860	adcC	-0,209	1,145	0,105	129	1,147
BKK33640	yvaM	-0,209	1,132	0,083	92	1,134
BKK20480	yoqX	-0,21	1,159	0,112	334	1,162
BKK32859	yuzM	-0,211	1,132	0,098	229	1,134
BKK40070	gntP	-0,211	1,145	0,101	447	1,147
BKK05950	ydiF	-0,212	1,199	0,097	201	1,201
BKK10880	yisV	-0,212	1,181	0,120	113	1,183
BKK17810	yndK	-0,212	1,161	0,104	81	1,164
BKK12700	xkdQ	-0,214	1,137	0,122	51	1,139
BKK33550	yvaC	-0,219	1,155	0,093	106	1,158
BKK19880	yotH	-0,224	1,165	0,111	370	1,168
BKK22080	ypwA	-0,224	1,137	0,089	115	1,139
BKK10350	yhfS	-0,229	1,165	0,112	381	1,168
BKK17670	cotU	-0,229	1,161	0,102	60	1,164
BKK40360	htrC	-0,229	1,148	0,114	206	1,151
BKK08150	yfjC	-0,233	1,165	0,104	219	1,168
BKK27300	yrrS	-0,238	1,174	0,113	99	1,177
BKK11080	yitQ	-0,239	1,180	0,098	89	1,183
BKK22970	mecB	-0,239	1,098	0,119	380	1,100
BKK02140	glpT	-0,24	1,157	0,106	419	1,160
BKK07080	yesZ	-0,24	1,176	0,125	170	1,179
BKK11750	cotY	-0,24	1,160	0,105	250	1,163
BKK12070	yjdI	-0,244	1,137	0,115	128	1,139
BKK33650	yvzC	-0,248	1,132	0,107	106	1,134
BKK06850	yeeK	-0,252	1,176	0,118	284	1,179
BKK33690	yvaQ	-0,252	1,131	0,089	144	1,134
BKK13730	queD	-0,253	1,165	0,113	294	1,168
BKK29580	thiI	-0,258	1,110	0,095	79	1,113
BKK10460	yhjC	-0,259	1,180	0,110	94	1,183
BKK17650	yncE	-0,259	1,161	0,121	77	1,164
BKK12950	dppD	-0,262	1,205	0,221	52	1,208
BKK11570	comEC	-0,262	1,148	0,113	195	1,151
BKK16960	ymdA	-0,267	1,160	0,118	66	1,164
BKK03430	nucA	-0,268	1,157	0,111	483	1,160
BKK14700	nprE	-0,272	1,176	0,130	125	1,179
BKK33080	liaR	-0,273	1,131	0,113	94	1,134
BKK33890	yvbK	-0,278	1,146	0,109	488	1,149
BKK15060	yjbM	-0,279	1,179	0,133	31	1,179
BKK08310	yfiL	-0,28	1,205	0,173	344	1,208
BKK29340	tcyN	-0,281	1,136	0,102	79	1,139
BKK03580	yczE	-0,282	1,135	0,083	161	1,138
BKK21090	ybbK	-0,283	1,150	0,106	441	1,153
BKK01720	ybbK	-0,285	1,156	0,099	107	1,160
BKK09470	ythA	-0,285	1,205	0,154	154	1,208
BKK08180	malA	-0,287	1,159	0,106	297	1,163
BKK11500	spxA	-0,288	1,154	0,108	160	1,158
BKK26400	yrkS	-0,289	1,150	0,104	160	1,153
BKK07290	yfnF	-0,294	0,972	0,101	55	0,975
BKK18950	yozL	-0,295	1,168	0,100	174	1,172
BKK41040	oxaAA	-0,295	1,163	0,156	108	1,166
BKK05110	ydeA	-0,297	1,156	0,100	417	1,160
BKK09320	yhcY	-0,299	1,135	0,095	61	1,138
BKK02690	ansZ	-0,3	1,198	0,073	70	1,201
BKK12930	dppB	-0,301	1,163	0,110	480	1,166
BKK11382	appA	-0,302	1,165	0,106	303	1,168
BKK30970	glgC	-0,303	1,153	0,122	89	1,156
BKK14410	sipT	-0,306	1,176	0,095	649	1,179
BKK30980	glgB	-0,307	1,152	0,086	152	1,156
BKK38130	ywcE	-0,307	1,153	0,109	126	1,157
BKK18360	yoxA	-0,308	1,168	0,118	122	1,172
BKK25500	hemN	-0,313	1,160	0,095	276	1,164
BKK26430	yrkP	-0,313	1,147	0,110	263	1,151
BKK08910	yhbA	-0,315	1,173	0,096	108	1,177
BKK20400	yorF	-0,318	1,158	0,109	391	1,162
BKK38110	nfrA	-0,321	1,144	0,114	420	1,147
BKK18280	yngK	-0,322	1,134	0,096	99	1,138
BKK08990	yhbI	-0,324	1,173	0,107	115	1,177
BKK01630	feuA	-0,329	1,198	0,064	38	1,201
BKK05690	ydhB	-0,329	1,163	0,080	303	1,167
BKK10520	glcP	-0,332	1,179	0,101	80	1,183
BKK31030	yuaE	-0,334	1,152	0,097	92	1,156
BKK02800	ycdC	-0,336	1,152	0,099	127	1,156
BKK39650	yxdK	-0,338	1,161	0,105	67	1,165
BKK12220	yjiC	-0,34	1,159	0,105	418	1,163
BKK39850	yxbF	-0,34	1,144	0,108	397	1,147
BKK32800	yusH	-0,341	1,154	0,096	112	1,158
BKK25000	pbpA	-0,343	0,972	0,075	76	0,975
BKK02540	ycbK	-0,344	1,197	0,064	120	1,201
BKK37340	ywiB	-0,344	1,162	0,113	66	1,166
BKK16470	sigD	-0,345	1,157	0,099	212	1,161
BKK39620	yxeA	-0,35	1,161	0,106	109	1,165
BKK08590	yfhM	-0,354	1,155	0,117	288	1,159
BKK17270	ymaC	-0,356	1,159	0,119	88	1,164
BKK19160	yocC	-0,356	1,153	0,102	125	1,158
BKK34050	yvfV	-0,359	1,130	0,093	74	1,134
BKK17850	lexA	-0,361	1,159	0,144	93	1,164
BKK28600	yshB	-0,362	1,155	0,094	108	1,160
BKK19220	yocI	-0,364	1,158	0,107	360	1,162
BKK10440	yhjA	-0,366	1,179	0,097	76	1,183
BKK28099	yszA	-0,366	1,155	0,096	293	1,160
BKK31420	yugF	-0,369	1,135	0,094	104	1,139
BKK26620	yrdR	-0,371	1,217	0,126	196	1,222
BKK39530	yxeJ	-0,374	1,160	0,101	76	1,165
BKK16799	ylzI	-0,375	1,159	0,110	125	1,164
BKK40330	rocE	-0,38	1,147	0,095	426	1,151
BKK25490	hrcA	-0,381	1,159	0,103	336	1,164
BKK33870	yvbI	-0,386	1,157	0,112	359	1,161
BKK31160	yubA	-0,388	1,152	0,105	64	1,156
BKK31110	yubF	-0,389	1,152	0,092	86	1,156
BKK36670	csbD	-0,389	1,110	0,104	79	1,114
BKK29850	ytoQ	-0,39	1,157	0,109	367	1,161
BKK30810	ytxM	-0,393	1,151	0,113	71	1,155
BKK39190	yxiG	-0,394	1,170	0,104	454	1,147
BKK20730	yopX	-0,395	1,150	0,113	409	1,155
BKK05880	gmuG	-0,398	1,155	0,108	553	1,160
BKK08640	yfhS	-0,398	1,154	0,117	132	1,159
BKK22250	yppG	-0,4	1,096	0,138	56	1,100
BKK10260	yhfK	-0,405	1,172	0,093	116	1,177
BKK14330	yknV	-0,405	1,202	0,148	113	1,206
BKK29360	tcyL	-0,405	1,109	0,092	96	1,113
BKK29260	ytpl	-0,406	1,109	0,140	221	1,113
BKK26540	yrkE	-0,407	1,172	0,104	315	1,177
BKK07170	yeti	-0,408	1,174	0,123	200	1,179
BKK36240	ywqE	-0,41	1,157	0,119	478	1,161
BKK07260	ltaSA	-0,414	1,155	0,104	873	1,160
BKK00730	cysK	-0,415	1,150	0,157	186	1,155
BKK05060	lrpB	-0,415	1,155	0,117	452	1,160
BKK10380	hemAT	-0,416	1,203	0,179	117	1,208
BKK12440	phrA	-0,417	1,163	0,108	296	1,168
BKK38650	yxlG	-0,417	1,161	0,146	88	1,166
BKK05250	ydeM	-0,418	1,153	0,094	211	1,158
BKK02180	ybfE	-0,422	1,155	0,111	629	1,160
BKK22920	ypeB	-0,423	1,149	0,143	254	1,153
BKK06830	rapH	-0,426	1,174	0,112	301	1,179
BKK00230	bofA	-0,428	1,178	0,122	117	1,183
BKK02190	ybfF	-0,429	1,155	0,102	935	1,160
BKK06820	yeeG	-0,429	1,174	0,111	408	1,179
BKK34280	epsJ	-0,431	1,130	0,120	451	1,135
BKK40600	yybL	-0,431	1,161	0,112	83	1,166

[1] BKK name

Sup. Table 3: Cell width of mutants of the BKK collection (continued)

BKK name[1]	gene	screening delta[2] (%)	average width (µ	+/-	nb	ADP
BKK31090	ktrA	-0,432	1,151	0,092	124	1,156
BKK23020	recQ	-0,434	1,096	0,111	512	1,100
BKK18840	xynA	-0,435	1,152	0,108	127	1,157
BKK36850	atpF	-0,436	1,161	0,107	114	1,166
BKK22730	ndk	-0,437	1,153	0,094	91	1,158
BKK02510	garD	-0,438	1,125	0,076	32	1,130
BKK27900	pheA	-0,445	1,154	0,110	78	1,160
BKK04010	sipU	-0,449	1,160	0,119	114	1,165
BKK29100	phoR	-0,449	1,108	0,104	53	1,113
BKK06460	purS	-0,45	1,174	0,111	333	1,179
BKK36810	atpD	-0,453	1,109	0,119	259	1,114
BKK06430	purK	-0,455	1,173	0,148	134	1,179
BKK07060	yesX	-0,456	1,152	0,108	93	1,158
BKK25810	arsR	-0,459	0,971	0,081	89	0,975
BKK02950	ycel	-0,46	1,196	0,139	96	1,201
BKK21870	ilvD	-0,46	1,150	0,108	532	1,155
BKK24050	bkdAA	-0,46	1,182	0,112	134	1,187
BKK38340	ywbF	-0,461	1,114	0,120	229	1,119
BKK29560	ytcl	-0,462	1,108	0,120	58	1,113
BKK04380	ydaT	-0,463	1,154	0,106	433	1,160
BKK14900	ctaD	-0,465	1,174	0,101	488	1,179
BKK06710	swrC	-0,466	1,154	0,107	502	1,160
BKK07140	yetF	-0,469	1,173	0,112	281	1,179
BKK21490	sunI	-0,469	1,148	0,110	595	1,153
BKK39590	yxeD	-0,47	1,159	0,108	161	1,165
BKK03359	yczL	-0,471	1,125	0,081	91	1,130
BKK19830	yotM	-0,474	1,124	0,141	604	1,129
BKK19640	yodL	-0,475	1,124	0,101	1106	1,129
BKK27150	yrhK	-0,479	1,148	0,110	477	1,153
BKK21690	msrA	-0,48	1,148	0,107	495	1,153
BKK24920	yqgL	-0,48	1,158	0,089	136	1,164
BKK28960	glcF	-0,484	1,146	0,109	285	1,151
BKK24840	yqgS	-0,485	1,201	0,157	48	1,206
BKK25090	yqfW	-0,489	1,158	0,099	203	1,164
BKK28960	ytxC	-0,491	1,108	0,096	33	1,113
BKK12910	proG	-0,492	1,161	0,094	429	1,166
BKK07160	yetH	-0,495	1,152	0,086	159	1,158
BKK31210	yulB	-0,497	1,150	0,089	111	1,156
BKK31240	mcpA	-0,497	1,150	0,091	103	1,156
BKK38730	cydD	-0,497	1,159	0,099	176	1,165
BKK21740	ypmR	-0,498	1,200	0,116	101	1,206
BKK33110	liaG	-0,498	1,129	0,109	107	1,134
BKK32849	yuzL	-0,499	1,129	0,089	135	1,134
BKK18630	exlX	-0,501	1,166	0,095	178	1,172
BKK13600	mtnX	-0,502	1,161	0,101	269	1,166
BKK33740	yvaV	-0,502	1,155	0,104	423	1,161
BKK39700	iolG	-0,502	1,159	0,125	128	1,165
BKK01680	ybbF	-0,509	1,154	0,108	425	1,160
BKK04039	yczO	-0,51	1,154	0,112	617	1,160
BKK13510	ykzE	-0,511	1,157	0,110	245	1,163
BKK19100	yobV	-0,512	1,166	0,081	90	1,172
BKK25150	yqfQ	-0,513	1,158	0,103	256	1,164
BKK40139	yyzN	-0,515	1,142	0,101	412	1,147
BKK23310	sipS	-0,516	1,181	0,112	172	1,187
BKK32930	yusU	-0,516	1,128	0,085	121	1,134
BKK33029	yuzO	-0,517	1,128	0,100	133	1,134
BKK12250	yikB	-0,52	1,202	0,179	419	1,208
BKK20110	yosl	-0,52	1,123	0,159	139	1,129
BKK27000	yraB	-0,524	1,171	0,097	137	1,177
BKK04230	ydaH	-0,525	1,152	0,118	155	1,158
BKK25100	yesU	-0,527	1,173	0,126	429	1,179
BKK04160	mtlR	-0,532	1,159	0,110	91	1,165
BKK10180	yhfC	-0,533	1,156	0,105	283	1,163
BKK04650	endB	-0,537	1,159	0,096	107	1,165
BKK28610	yshA	-0,539	1,151	0,108	129	1,158
BKK28470	lysC	-0,54	0,970	0,073	32	0,975
BKK04640	yybH	-0,546	1,178	0,108	186	1,184
BKK07340	yfnA	-0,547	1,151	0,107	68	1,158
BKK02430	glsA	-0,548	1,153	0,096	386	1,160
BKK10830	yisR	-0,548	1,177	0,106	71	1,183
BKK08080	acoC	-0,555	1,152	0,100	334	1,159
BKK40100	ahpF	-0,556	1,162	0,106	417	1,168
BKK03360	yciC	-0,559	1,124	0,099	94	1,130
BKK36400	flhO	-0,559	1,155	0,090	293	1,161
BKK27700	yrbF	-0,56	1,215	0,088	197	1,222
BKK03240	ycgQ	-0,561	1,141	0,110	538	1,147
BKK17520	ynaD	-0,564	1,155	0,105	431	1,162
BKK19070	yobS	-0,564	1,165	0,082	212	1,172
BKK07270	yfnH	-0,565	1,151	0,104	109	1,158
BKK02300	ybfN	-0,569	1,153	0,110	417	1,160
BKK28660	sspl	-0,573	1,133	0,103	135	1,139
BKK25690	sda	-0,574	1,147	0,122	141	1,153
BKK11760	cotX	-0,575	1,182	0,129	720	1,189
BKK21900	bsaA	-0,575	1,148	0,114	230	1,155
BKK01890	ybcL	-0,578	1,131	0,119	141	1,138
BKK12930	dppB	-0,581	1,182	0,131	1017	1,189
BKK30920	coti	-0,585	1,149	0,092	124	1,156
BKK33200	yvrE	-0,585	1,128	0,099	106	1,134
BKK06960	yesN	-0,586	1,172	0,105	370	1,179
BKK31250	tlpA	-0,588	1,149	0,104	117	1,156
BKK32590	frlN	-0,6	1,133	0,085	266	1,139
BKK40240	yycS	-0,602	1,144	0,110	363	1,151
BKK27990	minD	-0,61	1,131	0,093	76	1,138
BKK23070	serA	-0,611	1,151	0,112	147	1,158
BKK03120	ycgl	-0,613	1,153	0,108	731	1,160
BKK29980	ytjP	-0,614	1,149	0,107	92	1,156
BKK01580	ybaR	-0,616	1,194	0,074	45	1,201
BKK04980	yddI	-0,618	1,152	0,105	375	1,160
BKK18030	tlp	-0,621	1,156	0,103	91	1,164
BKK33239	yvrJ	-0,622	1,127	0,105	103	1,134
BKK37240	ywiE	-0,625	1,112	0,110	74	1,119
BKK05200	ydeH	-0,628	1,160	0,112	169	1,167
BKK06034	ydzT	-0,628	1,160	0,122	635	1,167
BKK00990	rpmGB	-0,635	0,969	0,092	74	0,975
BKK31270	tgl	-0,635	1,149	0,096	54	1,156
BKK39920	asnH	-0,638	1,157	0,112	88	1,165
BKK39029	yxiT	-0,642	1,157	0,124	56	1,165
BKK17950	yneJ	-0,648	1,154	0,113	407	1,162
BKK30035	ytzG	-0,65	1,149	0,104	177	1,156
BKK22500	ypjD	-0,651	1,093	0,110	315	1,100
BKK07400	yfmO	-0,654	1,200	0,185	537	1,208
BKK00520	ctc	-0,655	1,156	0,119	116	1,164
BKK00720	yocD	-0,655	1,131	0,104	111	1,138
BKK22910	ypfA	-0,656	1,199	0,165	62	1,206
BKK03770	rapC	-0,658	1,131	0,084	139	1,138
BKK11070	yitP	-0,658	1,176	0,130	112	1,183
BKK22190	ypsA	-0,658	1,093	0,115	234	1,100
BKK25480	grpE	-0,658	1,156	0,094	228	1,164
BKK19590	ctpA	-0,659	1,122	0,140	308	1,129
BKK13430	ykoX	-0,661	1,200	0,116	84	1,208
BKK30180	ytbQ	-0,663	1,148	0,087	125	1,156
BKK07560	pel	-0,667	1,151	0,108	301	1,159
BKK09850	yhaU	-0,671	1,200	0,142	65	1,208
BKK32830	fadA	-0,673	1,127	0,096	160	1,134
BKK09440	citA	-0,68	1,155	0,099	349	1,163
BKK07810	treA	-0,683	1,151	0,119	341	1,159
BKK29240	ytrl	-0,684	1,105	0,107	208	1,113
BKK12640	xkdJ	-0,686	1,160	0,117	308	1,168
BKK38520	dltC	-0,686	1,142	0,117	106	1,149
BKK21410	blyA	-0,688	1,108	0,118	318	1,114
BKK15860	sdaAA	-0,69	1,137	0,119	86	1,114
BKK30110	yteS	-0,691	1,130	0,098	172	1,138
BKK35560	tuaF	-0,692	0,968	0,082	46	0,975
BKK24760	rsbRD	-0,694	1,155	0,095	103	1,164
BKK30420	ytrE	-0,694	1,214	0,099	160	1,222
BKK26640	yrdP	-0,697	1,145	0,098	461	1,153
BKK31820	yuzF	-0,697	1,132	0,096	98	1,139
BKK23580	ansA	-0,698	1,179	0,099	267	1,187
BKK38100	ywcH	-0,7	1,149	0,098	103	1,157
BKK12870	mhqA	-0,703	1,158	0,122	71	1,166
BKK36220	ywqG	-0,704	1,149	0,105	146	1,157
BKK36650	ureB	-0,704	1,149	0,105	146	1,157
BKK24460	yqhT	-0,704	1,143	0,104	402	1,151
BKK32920	yusT	-0,705	0,968	0,072	110	0,975
BKK40930	yyaE	-0,705	1,148	0,106	143	1,156
BKK36740	ywmC	-0,708	1,106	0,103	101	1,114
BKK13460	rsgl	-0,71	1,160	0,110	284	1,168
BKK15300	bpr	-0,712	1,154	0,110	315	1,163
BKK24440	yqhV	-0,713	1,179	0,103	289	1,187
BKK17266	ymzE	-0,714	1,155	0,085	76	1,164
BKK40880	exoAA	-0,716	1,176	0,089	156	1,184
BKK16700	ylxY	-0,721	1,155	0,112	89	1,164
BKK37220	ywjB	-0,724	1,158	0,116	106	1,166
BKK26840	sigZ	-0,727	1,168	0,101	270	1,177
BKK15480	pyrP	-0,727	1,148	0,110	96	1,156
BKK06040	ydiM	-0,733	1,131	0,120	118	1,139
BKK23630	yqkE	-0,733	1,178	0,094	656	1,187
BKK25830	rapE	-0,734	1,155	0,093	64	1,164
BKK17700	cotC	-0,739	1,155	0,094	106	1,164
BKK17740	ynzB	-0,739	1,153	0,107	311	1,162
BKK21820	thyB	-0,743	1,143	0,106	188	1,151
BKK09958	yhzE	-0,744	1,168	0,092	134	1,177
BKK35160	uvrA	-0,745	1,127	0,119	693	1,135
BKK24810	yqgV	-0,747	1,155	0,092	280	1,164
BKK17160	pksH	-0,756	1,155	0,097	112	1,164
BKK29740	ccpA	-0,758	1,153	0,142	154	1,161
BKK17240	ymzB	-0,759	1,153	0,108	396	1,162
BKK32890	yusQ	-0,766	1,126	0,098	134	1,134
BKK01810	adaA	-0,767	1,151	0,096	148	1,160
BKK19750	cgeE	-0,771	1,121	0,138	159	1,129
BKK04500	ydbK	-0,773	1,151	0,117	466	1,160
BKK19479	yoyC	-0,781	1,120	0,107	532	1,129
BKK21200	yomW	-0,781	1,146	0,100	543	1,155
BKK14340	yknW	-0,782	1,170	0,122	261	1,179
BKK12150	yjgB	-0,783	1,180	0,151	191	1,189
BKK31140	cdoA	-0,788	1,147	0,111	87	1,156
BKK11620	yjbO	-0,789	1,130	0,105	59	1,139
BKK22960	gudB	-0,791	1,148	0,108	82	1,158
BKK24300	xseA	-0,791	1,178	0,104	258	1,187
BKK27740	ruvA	-0,791	1,150	0,119	126	1,160
BKK05899	ydhU	-0,792	1,130	0,098	190	1,139
BKK36450	ywoG	-0,792	1,140	0,085	265	1,149
BKK22019	ypzF	-0,794	1,092	0,147	61	1,100
BKK31860	yueB	-0,795	1,139	0,110	505	1,148
BKK18990	yobK	-0,797	1,129	0,089	101	1,138
BKK17930	ccdA	-0,799	1,154	0,133	71	1,164
BKK25440	rsmE	-0,801	1,154	0,099	258	1,164
BKK33230	sigO	-0,801	0,967	0,074	94	0,975
BKK02130	glpQ	-0,805	1,150	0,111	551	1,160
BKK19580	yodF	-0,807	1,120	0,106	819	1,129
BKK29370	tcyK	-0,807	1,129	0,094	64	1,138
BKK12720	xkdS	-0,811	1,130	0,101	226	1,139
BKK30070	opuD	-0,812	1,152	0,107	355	1,161
BKK31080	yuaB	-0,818	1,147	0,102	80	1,156
BKK28100	ysxE	-0,825	1,150	0,101	367	1,160
BKK01020	rplK	-0,826	1,154	0,099	453	1,164
BKK25890	yqxl	-0,828	1,144	0,104	437	1,153
BKK14760	ylaF	-0,829	1,169	0,131	403	1,179
BKK18440	gltB	-0,829	1,158	0,108	258	1,168
BKK37760	rocC	-0,83	1,109	0,140	54	1,119
BKK38070	sacT	-0,834	1,109	0,169	30	1,119
BKK05410	ydfH	-0,835	1,138	0,102	406	1,147
BKK38870	yxkA	-0,836	1,138	0,105	378	1,147
BKK11980	yjdA	-0,837	1,153	0,111	259	1,163
BKK02110	ybyB	-0,843	1,120	0,107	72	1,130
BKK08890	ygaO	-0,847	1,158	0,106	277	1,168
BKK34290	epsl	-0,849	1,156	0,109	145	1,166
BKK38080	ywcl	-0,858	1,109	0,093	102	1,119
BKK26930	yral	-0,859	1,167	0,112	205	1,177
BKK07820	treR	-0,861	1,149	0,112	507	1,159
BKK23100	sigX	-0,863	1,091	0,118	364	1,100
BKK24780	yqgY	-0,863	1,154	0,104	231	1,164
BKK00260	yaaN	-0,864	1,156	0,136	719	1,166
BKK16360	fliQ	-0,865	1,152	0,118	488	1,162
BKK19420	yclD	-0,867	1,128	0,098	144	1,138
BKK23910	yojE	-0,867	1,141	0,127	264	1,151
BKK32410	yojl	-0,867	1,130	0,085	208	1,139
BKK32950	yusW	-0,869	1,128	0,090	114	1,138
BKK10160	yhgE	-0,875	1,152	0,076	97	1,177
BKK10250	yhfJ	-0,875	1,167	0,089	101	1,177
BKK03789	yczN	-0,88	1,155	0,104	309	1,165
BKK20470	yoqY	-0,88	1,152	0,113	368	1,162
BKK18870	yojD	-0,883	1,152	0,114	418	1,162
BKK31200	yulC	-0,889	1,147	0,102	142	1,158
BKK33260	yvrN	-0,889	1,147	0,119	95	1,158
BKK01640	ybbB	-0,89	1,149	0,118	376	1,160
BKK05460	ydfL	-0,892	1,149	0,105	224	1,160
BKK03400	yckD	-0,893	0,966	0,075	74	0,975
BKK14060	fadH	-0,893	1,158	0,119	643	1,166
BKK04090	kipA	-0,897	1,154	0,086	123	1,165
BKK31470	yojG	-0,899	1,138	0,118	264	1,148
BKK06812	yeeF	-0,909	1,149	0,105	430	1,160
BKK27450	glnM	-0,91	1,148	0,094	177	1,160
BKK21260	yomQ	-0,911	1,141	0,115	260	1,151
BKK34650	yvdC	-0,911	1,125	0,094	30	1,135
BKK40340	rocD	-0,911	1,141	0,112	191	1,151
BKK35698	yuzl	-0,914	1,104	0,094	225	1,114
BKK35790	yvyl	-0,914	1,104	0,120	98	1,114
BKK39860	aldX	-0,918	1,154	0,091	94	1,165
BKK03030	ycgB	-0,919	1,120	0,100	159	1,130
BKK16300	fliL	-0,921	1,157	0,104	169	1,168
BKK19740	yodT	-0,924	1,119	0,109	569	1,129
BKK24010	bmr	-0,924	1,147	0,097		1,158
BKK10750	yisK	-0,926	1,152	0,113	418	1,163
BKK34720	yvcP	-0,928	1,139	0,095	317	1,149
BKK27570	yrzK	-0,931	1,149	0,089	55	1,160
BKK35200	yvkB	-0,934	1,124	0,109	345	1,135
BKK12470	yjgA	-0,935	1,178	0,098	823	1,189
BKK15070	ylbN	-0,935	1,152	0,105	238	1,163
BKK18040	yneP	-0,935	1,157	0,116	222	1,168
BKK07020	rhgT	-0,943	1,168	0,114	406	1,179
BKK03390	yckC	-0,946	1,147	0,111	106	1,158
BKK03960	ycnK	-0,946	1,149	0,105	386	1,160
BKK29380	tcyJ	-0,946	1,127	0,085	113	1,138
BKK04200	ydaE	-0,947	1,154	0,109	57	1,165
BKK25350	yqfD	-0,948	1,153	0,103	226	1,164
BKK23328	ypzJ	-0,949	1,140	0,105	342	1,151
BKK40270	yycP	-0,951	1,140	0,100	356	1,151
BKK25990	yqbS	-0,953	1,140	0,115	245	1,151
BKK06750	yefC	-0,956	1,168	0,108	131	1,179
BKK12210	yjiB	-0,956	1,178	0,101	1308	1,189
BKK11850	yjcG	-0,958	1,178	0,110	408	1,189
BKK19700	yodP	-0,961	1,195	0,135	101	1,206
BKK20750	yopV	-0,962	1,144	0,097	498	1,155
BKK40190	fbp	-0,966	1,157	0,108	254	1,168
BKK20970	yonX	-0,97	1,142	0,100	490	1,153
BKK01690	ybbH	-0,973	1,165	0,106	289	1,177
BKK32180	yutK	-0,978	1,210	0,067	54	1,222
BKK02420	glnT	-0,98	1,148	0,097	479	1,160
BKK24630	sipW	-0,981	0,965	0,084	50	0,975
BKK25080	yqfX	-0,982	1,152	0,114	249	1,164
BKK17790	yndH	-0,991	1,156	0,108	352	1,168
BKK19780	cgeA	-0,992	1,146	0,110	134	1,158
BKK09660	yheN	-0,993	1,165	0,104	115	1,177
BKK17620	yncB	-0,994	1,150	0,110	420	1,162
BKK13470	sspD	-0,995	1,155	0,112	115	1,166
BKK09710	yheI	-1	1,196	0,114	55	1,208
BKK26670	gltR	-1,003	1,142	0,099	508	1,153
BKK37650	cysL	-1,004	1,108	0,109	257	1,113
BKK13650	eag	-1,005	1,151	0,108	313	1,163
BKK24670	comGG	-1,009	1,152	0,098	163	1,164
BKK24860	yqgQ	-1,012	1,152	0,100	316	1,164
BKK11870	yneB	-1,014	1,150	0,113	307	1,162
BKK25120	yqfT	-1,014	1,152	0,104	482	1,153
BKK14150	ykuN	-1,018	1,167	0,123	907	1,179
BKK36950	ywlC	-1,02	0,965	0,112	41	0,975
BKK32310	resE	-1,029	1,194	0,094	126	1,206
BKK26830	yrpE	-1,029	1,209	0,067	94	1,222
BKK26900	yraL	-1,032	1,165	0,102	431	1,177
BKK34220	epsO	-1,033	1,123	0,111	96	1,135
BKK40010	yxaD	-1,036	1,136	0,107	433	1,147
BKK35880	pgsA	-1,037	1,138	0,095	548	1,149
BKK36669	ywzF	-1,038	1,136	0,100	159	1,147
BKK27190	yrzl	-1,039	1,146	0,112	86	1,158
BKK02530	rtpA	-1,04	1,136	0,101	428	1,147
BKK34690	yvcS	-1,041	1,137	0,105	563	1,149
BKK33780	sdpl	-1,043	1,122	0,096	107	1,134
BKK06780	yeeC	-1,045	1,167	0,102	601	1,179
BKK37610	ywzC	-1,045	1,107	0,104	320	1,119
BKK38540	dltE	-1,046	0,965	0,084	50	0,975
BKK33060	gerAB	-1,047	1,122	0,095	77	1,134
BKK33300	fhuG	-1,047	1,122	0,099	99	1,134
BKK34710	yvcQ	-1,047	1,137	0,106	440	1,149
BKK04030	ycgP	-1,048	1,118	0,091	41	1,130
BKK26880	yraM	-1,051	1,141	0,102	504	1,153
BKK26420	yrkQ	-1,054	1,209	0,092	161	1,222
BKK35800	gerBA	-1,054	1,103	0,087	38	1,114
BKK23000	ypbE	-1,057	1,089	0,113	607	1,100
BKK18750	yoaT	-1,064	1,159	0,106	218	1,172
BKK18900	yojl	-1,064	1,126	0,093	69	1,138
BKK31590	yufS	-1,064	1,127	0,110	61	1,139
BKK05120	cspC	-1,067	1,147	0,120	380	1,160
BKK10910	yisZ	-1,069	1,171	0,097	92	1,183
BKK12820	spoIISB	-1,069	1,195	0,160	87	1,208

Sup. Table 3: Cell width of mutants of the BKK collection (continued)

BKK name[1]	gene	screening delta[2] (%)	average width (µ	+/-	nb	ADP
BKK33120	liaH	-1,071	1,122	0,084	63	1,134
BKK05260	ydeN	-1,075	1,155	0,097	89	1,167
BKK24160	mmgB	-1,075	1,141	0,105	257	1,153
BKK33320	fhuD	-1,075	1,209	0,105	252	1,222
BKK01710	ybbJ	-1,077	1,164	0,102	133	1,177
BKK27470	yrrD	-1,077	1,147	0,099	194	1,160
BKK18350	dacC	-1,078	1,193	0,135	91	1,206
BKK21550	yokL	-1,079	1,142	0,104	393	1,155
BKK39230	wapA	-1,084	1,193	0,143	195	1,206
BKK19870	yotI	-1,086	1,117	0,135	332	1,129
BKK09770	yheD	-1,091	1,155	0,097	330	1,168
BKK21020	yonR	-1,092	1,141	0,099	548	1,153
BKK36420	spoIIID	-1,094	1,137	0,107	357	1,149
BKK00250	xpaC	-1,095	1,154	0,113	166	1,166
BKK17300	ebrA	-1,095	1,127	0,080	151	1,139
BKK28970	ytxB	-1,096	1,149	0,102	394	1,161
BKK18960	yozM	-1,103	1,149	0,111	410	1,162
BKK40150	yydI	-1,104	1,135	0,095	299	1,147
BKK21320	yomL	-1,106	1,141	0,119	521	1,153
BKK23860	gndA	-1,107	1,174	0,123	247	1,187
BKK15760	prpC	-1,108	1,122	0,105	108	1,134
BKK02330	ybfQ	-1,111	1,117	0,101	90	1,130
BKK19300	yozC	-1,114	1,149	0,115	374	1,162
BKK38960	yxjG	-1,116	1,152	0,112	148	1,165
BKK07180	yetI	-1,117	1,166	0,110	343	1,179
BKK03090	ycgF	-1,125	1,147	0,117	276	1,160
BKK08600	csbB	-1,128	1,149	0,111	343	1,163
BKK22260	yppF	-1,128	1,127	0,099	97	1,139
BKK05440	nap	-1,129	1,145	0,093	126	1,158
BKK31680	comA	-1,13	1,127	0,091	114	1,139
BKK34200	sigL	-1,131	1,122	0,150	50	1,135
BKK07890	yfkI	-1,132	1,146	0,112	300	1,159
BKK10580	gerPE	-1,132	1,155	0,094	302	1,168
BKK34850	yvcA	-1,135	1,136	0,117	519	1,149
BKK06280	ydjP	-1,136	1,134	0,107	302	1,147
BKK10420	comK	-1,136	1,170	0,094	96	1,183
BKK11230	carA	-1,136	1,170	0,126	87	1,183
BKK32530	pucF	-1,138	1,135	0,141	51	1,148
BKK27680	yrbG	-1,143	1,146	0,099	381	1,160
BKK13170	guaD	-1,145	1,153	0,132	208	1,166
BKK06390	yebD	-1,146	1,146	0,108	324	1,160
BKK30200	bioB	-1,148	1,143	0,097	92	1,156
BKK30300	melA	-1,149	1,143	0,096	118	1,156
BKK35060	cypX	-1,149	1,122	0,122	509	1,135
BKK17820	yndL	-1,152	1,150	0,117	119	1,164
BKK00450	ycsF	-1,156	1,134	0,098	508	1,147
BKK13870	ykvZ	-1,157	1,149	0,104	324	1,163
BKK39460	yxeQ	-1,157	1,151	0,090	123	1,165
BKK38820	yxkF	-1,158	1,151	0,100	179	1,165
BKK31450	kinB	-1,16	1,208	0,062	59	1,222
BKK27010	adhA	-1,166	1,163	0,099	387	1,177
BKK20980	yonV	-1,167	1,140	0,101	590	1,153
BKK21660	yokA	-1,168	1,141	0,094	236	1,155
BKK18190	yngC	-1,169	1,192	0,143	65	1,206
BKK05080	yddR	-1,171	1,154	0,114	251	1,167
BKK25730	yqeC	-1,173	1,150	0,116	153	1,164
BKK00120	pdxT	-1,176	1,144	0,111	91	1,158
BKK22850	seaA	-1,178	1,087	0,116	159	1,100
BKK00430	yabG	-1,179	1,150	0,094	141	1,164
BKK13299	ykzO	-1,179	1,087	0,112	133	1,100
BKK32400	yunG	-1,179	1,134	0,128	144	1,148
BKK09530	yhdN	-1,18	1,149	0,118	291	1,163
BKK29970	ytkP	-1,181	1,142	0,114	107	1,156
BKK17350	ymzC	-1,186	1,148	0,109	446	1,162
BKK10170	fabHB	-1,187	1,163	0,097	106	1,177
BKK32030	bioYB	-1,187	1,208	0,068	97	1,222
BKK37310	fnr	-1,187	1,134	0,101	308	1,147
BKK37580	ywgB	-1,188	1,105	0,120	802	1,119
BKK12649	ykzM	-1,189	1,154	0,111	241	1,168
BKK02910	yceE	-1,191	1,117	0,099	166	1,130
BKK13670	mhqR	-1,191	1,152	0,116	231	1,166
BKK15660	yloC	-1,191	1,121	0,102	102	1,134
BKK08520	recX	-1,193	1,145	0,103	280	1,159
BKK33090	liaS	-1,193	1,207	0,126	67	1,222
BKK12180	yjhA	-1,199	1,194	0,161	146	1,208
BKK16160	clpY	-1,2	1,149	0,105	311	1,163
BKK04820	immR	-1,201	1,144	0,095	123	1,158
BKK30090	yteU	-1,218	1,124	0,099	116	1,138
BKK38500	dltA	-1,218	1,135	0,124	115	1,149
BKK08380	yfiS	-1,219	1,145	0,110	285	1,159
BKK35920	rbsK	-1,219	1,101	0,093	72	1,114
BKK08320	yfiM	-1,221	1,145	0,107	270	1,159
BKK30380	bceA	-1,222	1,207	0,080	97	1,222
BKK24650	yqzG	-1,223	1,173	0,107	165	1,187
BKK20850	yopL	-1,224	1,137	0,101	257	1,151
BKK36570	ywnG	-1,224	1,101	0,107	208	1,114
BKK20130	yosG	-1,225	1,148	0,109	236	1,162
BKK26780	yrdA	-1,225	1,139	0,111	450	1,153
BKK04630	ydcC	-1,228	1,187	0,115	558	1,201
BKK20740	yopW	-1,228	1,141	0,097	412	1,155
BKK38840	yxkD	-1,233	1,193	0,123	83	1,208
BKK38580	licC	-1,234	1,152	0,127	75	1,166
BKK14170	ykuP	-1,241	1,148	0,098	319	1,163
BKK07230	yetM	-1,242	1,164	0,111	529	1,179
BKK09090	yhcI	-1,244	1,154	0,112	287	1,168
BKK38880	yxjO	-1,244	1,150	0,102	79	1,165
BKK36700	moaA	-1,245	1,100	0,111	86	1,114
BKK06940	yesL	-1,246	1,153	0,099	342	1,168
BKK07500	yfmE	-1,247	1,144	0,102	257	1,159
BKK38710	yxlA	-1,25	1,150	0,114	148	1,165
BKK03330	nasA	-1,255	1,186	0,074	69	1,201
BKK02870	adcB	-1,26	1,124	0,109	108	1,138
BKK39420	deoC	-1,262	1,150	0,108	126	1,165
BKK35910	rbsR	-1,264	1,152	0,108	135	1,166
BKK20890	yopH	-1,265	1,139	0,091	514	1,153
BKK26680	yrdK	-1,269	1,136	0,100	279	1,151
BKK17450	glnR	-1,272	1,147	0,101	632	1,162
BKK11440	oppB	-1,273	1,193	0,162	80	1,208
BKK20070	nrdIB	-1,276	1,115	0,120	112	1,129
BKK00400	yabE	-1,28	1,125	0,106	65	1,139
BKK37130	spo0F	-1,28	1,135	0,095	525	1,149
BKK17130	acpM	-1,293	1,148	0,120	90	1,164
BKK34610	mdxE	-1,296	1,146	0,117	352	1,161
BKK25390	yqeZ	-1,297	1,148	0,095	182	1,164
BKK35280	yvjA	-1,298	1,120	0,111	324	1,135
BKK34040	yvfW	-1,299	1,120	0,090	138	1,134
BKK11000	yitI	-1,302	1,147	0,117	303	1,163
BKK21880	ypgR	-1,302	1,140	0,097	477	1,155
BKK00370	abrB	-1,303	1,164	0,146	91	1,179
BKK06450	purC	-1,303	1,163	0,119	259	1,179
BKK02280	yorR	-1,303	1,153	0,106	244	1,168
BKK38230	ywcB	-1,303	1,104	0,117	69	1,119
BKK17720	yndA	-1,304	1,147	0,106	409	1,162
BKK12730	xkdT	-1,307	1,173	0,126	110	1,189
BKK26390	spoIIIC	-1,313	1,161	0,109	187	1,177
BKK31400	yugH	-1,315	1,133	0,116	120	1,148
BKK12550	xtrA	-1,317	1,153	0,120	208	1,168
BKK34660	yvdB	-1,318	1,134	0,106	357	1,149
BKK06760	yeeA	-1,321	1,163	0,105	420	1,179
BKK07510	yfmD	-1,321	1,192	0,147	53	1,208
BKK20860	yopK	-1,323	1,138	0,112	512	1,153
BKK03980	ycnD	-1,324	1,122	0,103	386	1,147
BKK05560	ydgI	-1,325	1,142	0,106	80	1,158
BKK23430	spoVAB	-1,33	1,190	0,130	100	1,206
BKK07990	yfjR	-1,331	1,143	0,109	324	1,159
BKK04210	ydoF	-1,333	1,149	0,096	136	1,165
BKK37810	spsL	-1,339	1,134	0,110	405	1,149
BKK06920	yesJ	-1,342	1,163	0,127	285	1,179
BKK39510	ytrF	-1,343	1,140	0,101	163	1,156
BKK04780	ydcI	-1,352	1,149	0,133	181	1,165
BKK01980	skfH	-1,353	1,144	0,101	436	1,160
BKK08360	yfiQ	-1,354	1,143	0,120	230	1,159
BKK02620	phoD	-1,357	1,132	0,109	508	1,147
BKK37380	albB	-1,359	1,104	0,094	61	1,119
BKK24200	yqiH	-1,361	1,190	0,163	31	1,206
BKK38790	yxzE	-1,361	1,149	0,108	112	1,165
BKK18800	penP	-1,364	0,962	0,101	39	0,975
BKK27690	yrzE	-1,364	1,144	0,103	394	1,160
BKK31930	ald	-1,364	1,132	0,120	51	1,148
BKK10960	yitE	-1,373	1,167	0,103	143	1,183
BKK13100	ykkD	-1,373	1,192	0,138	130	1,208
BKK25430	rimO	-1,374	1,148	0,103	365	1,164
BKK34920	hisG	-1,377	1,119	0,113	100	1,135
BKK35430	flgM	-1,377	1,141	0,106	89	1,157
BKK11381	appA	-1,38	1,145	0,117	375	1,161
BKK10760	yisL	-1,387	1,167	0,115	123	1,183
BKK29760	ytxJ	-1,39	1,124	0,095	87	1,139
BKK18520	yoxB	-1,391	1,141	0,105	122	1,157
BKK34520	yvdP	-1,391	1,119	0,104	77	1,135
BKK25740	ugtP	-1,396	1,147	0,112	122	1,164
BKK28940	ysaA	-1,398	1,098	0,120	292	1,113
BKK40480	yycB	-1,399	1,131	0,099	412	1,147
BKK21390	yomE	-1,401	1,137	0,109	434	1,153
BKK39120	yxiM	-1,404	1,148	0,116	30	1,165
BKK12460	xlyB	-1,41	1,172	0,109	914	1,189
BKK33380	yvgL	-1,41	1,205	0,089	94	1,222
BKK06310	gabP	-1,411	1,143	0,104	835	1,160
BKK19920	yotD	-1,414	1,113	0,142	144	1,129
BKK03982	mtlF	-1,419	1,148	0,084	173	1,165
BKK17730	yndB	-1,42	1,145	0,105	310	1,162
BKK21540	yolA	-1,423	1,138	0,114	236	1,155
BKK36640	ureC	-1,424	1,140	0,111	147	1,157
BKK37190	clsB	-1,424	1,103	0,104	64	1,119
BKK06250	ydjM	-1,425	1,162	0,107	316	1,179
BKK30770	mntA	-1,426	1,145	0,105	389	1,161
BKK18540	yooB	-1,428	1,122	0,093	174	1,138
BKK33240	oxdC	-1,428	1,118	0,106	149	1,134
BKK07940	yfkC	-1,432	1,142	0,106	337	1,159
BKK37060	tdk	-1,432	1,098	0,123	30	1,114
BKK30080	yteV	-1,434	1,139	0,106	98	1,156
BKK11130	ipi	-1,437	1,166	0,095	131	1,183
BKK23820	yqjM	-1,437	1,137	0,105	489	1,153
BKK15770	prkC	-1,438	1,151	0,101	374	1,168
BKK39339	rsiX	-1,441	1,131	0,105	357	1,147
BKK29680	acsA	-1,445	1,097	0,115	54	1,113
BKK16430	cheA	-1,446	1,122	0,117	107	1,138
BKK27469	yrzR	-1,447	1,134	0,101	274	1,151
BKK06990	yetA	-1,452	1,191	0,193	175	1,208
BKK35140	yvkN	-1,453	1,119	0,112	363	1,135
BKK36880	yooF	-1,454	1,098	0,135	177	1,114
BKK02570	ycbN	-1,461	1,184	0,071	153	1,201
BKK08390	yfhE	-1,468	1,142	0,113	277	1,159
BKK07090	yetA	-1,471	1,162	0,122	231	1,179
BKK22590	csfB	-1,472	1,084	0,101	235	1,100
BKK05090	yddS	-1,475	1,143	0,116	439	1,160
BKK25230	yqxD	-1,475	1,146	0,092	192	1,164
BKK38920	pepT	-1,475	1,148	0,125	149	1,165
BKK18580	atpI	-1,478	1,121	0,092	73	1,138
BKK11510	yjbE	-1,479	1,166	0,107	141	1,183
BKK00240	csfB	-1,48	1,166	0,114	178	1,183
BKK02840	ycdG	-1,48	1,113	0,095	53	1,130
BKK17610	xylB	-1,481	1,145	0,110	292	1,160
BKK10270	lcfB	-1,482	1,166	0,086	74	1,183
BKK16110	dprA	-1,482	1,142	0,122	183	1,160
BKK08790	thiC	-1,487	1,142	0,103	187	1,159
BKK14430	ykpA	-1,487	1,145	0,098	374	1,161
BKK03710	gerKC	-1,488	1,121	0,090	59	1,138
BKK08480	yfhE	-1,488	1,142	0,120	318	1,159
BKK36100	ywrD	-1,488	1,140	0,095	90	1,157
BKK13630	ykvA	-1,49	0,960	0,089	99	0,975
BKK23950	ykzC	-1,493	1,162	0,162	93	1,179
BKK32070	yuiC	-1,495	1,131	0,082	182	1,148
BKK16310	fliM	-1,501	1,145	0,113	212	1,163
BKK16050	rnhB	-1,503	1,141	0,104	306	1,159
BKK19800	phy	-1,503	1,147	0,096	66	1,165
BKK35770	tagC	-1,504	1,144	0,114	343	1,161
BKK18020	sspN	-1,506	1,144	0,113	69	1,164
BKK32520	pucG	-1,513	1,204	0,064	126	1,222
BKK17190	pksL	-1,514	1,150	0,101	460	1,168
BKK23060	ypzE	-1,514	1,140	0,088	95	1,158
BKK11139	yxzL	-1,517	1,145	0,105	410	1,163
BKK26950	yraG	-1,518	1,159	0,109	169	1,177
BKK18130	alsT	-1,521	1,154	0,123	126	1,172
BKK24380	spoIIAF	-1,521	1,188	0,140	54	1,206
BKK23980	artP	-1,522	1,136	0,113	432	1,153
BKK02390	ilvE	-1,528	1,142	0,099	480	1,160
BKK16930	ciaR	-1,529	1,146	0,103	109	1,164
BKK08300	yfiK	-1,533	1,141	0,116	123	1,159
BKK22190	yhpP	-1,535	1,137	0,102	356	1,155
BKK06078	ydzW	-1,536	1,130	0,102	158	1,147
BKK26440	yrkO	-1,536	1,159	0,108	235	1,177
BKK39630	yxdM	-1,539	1,203	0,082	74	1,222
BKK21760	ypmP	-1,542	1,137	0,112	541	1,155
BKK03620	yclA	-1,543	1,147	0,102	218	1,165
BKK29900	trmB	-1,543	1,143	0,115	339	1,161
BKK30100	yteT	-1,544	1,138	0,084	264	1,156
BKK38450	ywaE	-1,544	1,140	0,095	86	1,158
BKK21930	cspD	-1,545	1,136	0,115	292	1,153
BKK37840	spsI	-1,546	1,148	0,109	96	1,166
BKK41060	rpmH	-1,546	1,138	0,141	37	1,156
BKK22550	qcrB	-1,549	1,083	0,122	393	1,100
BKK23600	yqxK	-1,551	1,169	0,102	227	1,187
BKK02460	ycbC	-1,554	1,112	0,103	106	1,130
BKK16100	sucD	-1,556	1,144	0,103	270	1,163
BKK25330	yqfF	-1,559	1,145	0,095	287	1,164
BKK02790	ycdB	-1,56	1,112	0,101	71	1,130
BKK38610	dltB	-1,561	1,148	0,109	108	1,166
BKK02340	gltP	-1,571	1,112	0,086	113	1,130
BKK13380	ykoS	-1,571	1,150	0,104	463	1,168
BKK31730	yuzC	-1,571	1,122	0,085	67	1,139
BKK11809	yjcZ	-1,572	1,170	0,088	328	1,189
BKK30720	ythB	-1,572	1,138	0,101	112	1,156
BKK40040	glxK	-1,572	1,129	0,101	458	1,147
BKK10120	hemE	-1,574	1,158	0,118	98	1,177
BKK29710	acuC	-1,577	1,133	0,106	278	1,151
BKK11910	yjcM	-1,578	1,170	0,112	321	1,189
BKK33222	rsoA	-1,578	1,116	0,082	107	1,134
BKK35360	hag	-1,58	1,117	0,112	490	1,135
BKK33610	rnr	-1,581	1,116	0,098	95	1,134
BKK11770	yndF	-1,588	1,145	0,107	100	1,164
BKK23090	rsiX	-1,588	1,138	0,103	140	1,156
BKK40680	yybD	-1,588	1,166	0,095	235	1,184
BKK08170	yfjA	-1,59	1,140	0,107	392	1,159
BKK16130	trmFO	-1,592	1,141	0,096	555	1,160
BKK28110	spoVID	-1,593	1,141	0,104	89	1,160
BKK29010	speD	-1,593	1,095	0,102	167	1,113
BKK20450	yorA	-1,596	1,149	0,108	282	1,168
BKK13050	ykzH	-1,597	1,144	0,101	351	1,163
BKK07670	yfII	-1,6	1,141	0,099	508	1,160
BKK19410	cwlS	-1,603	1,143	0,119	379	1,162
BKK03540	ycxB	-1,605	1,112	0,097	128	1,130
BKK20040	yosP	-1,606	1,111	0,146	181	1,129
BKK07850	yfkM	-1,608	1,141	0,100	442	1,160
BKK32640	sspG	-1,609	1,116	0,080	162	1,134
BKK36600	mta	-1,609	1,096	0,102	68	1,114
BKK10560	yhjM	-1,611	1,164	0,102	93	1,183
BKK17940	ynel	-1,612	1,143	0,103	382	1,162
BKK03510	srfAC	-1,613	1,112	0,105	104	1,130
BKK24850	glcK	-1,614	1,145	0,097	126	1,164
BKK11570	yjbI	-1,617	1,189	0,136	144	1,208
BKK09600	crcBA	-1,619	1,158	0,098	143	1,177
BKK05890	ydhU	-1,62	1,139	0,095	125	1,158
BKK08900	yhzB	-1,623	1,158	0,096	73	1,177
BKK06840	yeeI	-1,629	1,160	0,112	307	1,179
BKK25880	yqxI	-1,632	0,959	0,070	80	0,975
BKK01530	cwlD	-1,634	0,959	0,076	49	0,975
BKK25580	comEB	-1,639	1,144	0,109	305	1,164
BKK38270	efeM	-1,641	1,131	0,107	296	1,149
BKK08490	yfhD	-1,643	1,140	0,101	197	1,159
BKK20580	yoqM	-1,644	1,136	0,098	448	1,155
BKK34590	mdxG	-1,644	1,131	0,110	557	1,149
BKK30150	ytdP	-1,647	1,137	0,087	99	1,156
BKK37910	spsA	-1,648	1,138	0,131	166	1,157
BKK33860	yvbH	-1,653	1,142	0,107	283	1,161
BKK39890	yxbB	-1,653	1,138	0,121	98	1,158
BKK32050	pepA	-1,655	1,129	0,097	189	1,148
BKK07620	yflN	-1,658	1,140	0,111	437	1,159
BKK07955	yfkA	-1,658	1,143	0,102	358	1,163
BKK08340	padR	-1,659	1,143	0,103	324	1,163
BKK26550	yrkD	-1,661	1,157	0,102	241	1,177
BKK28590	polX	-1,661	1,140	0,094	41	1,160
BKK14280	moeA	-1,667	1,159	0,105	186	1,179
BKK12420	yjoB	-1,668	1,169	0,115	635	1,189
BKK02070	pssA	-1,669	1,111	0,092	91	1,130
BKK33340	sspJ	-1,67	1,142	0,113	342	1,161
BKK21890	ypgQ	-1,672	1,136	0,106	342	1,155
BKK07710	yflE	-1,674	1,119	0,100	128	1,138
BKK32010	besA	-1,675	1,129	0,126	164	1,148
BKK38680	yxlD	-1,675	1,147	0,106	219	1,166
BKK13520	ykrP	-1,676	1,147	0,103	230	1,164
BKK14470	mreBH	-1,677	1,143	0,105	317	1,163
BKK26480	yrkK	-1,677	1,134	0,097	437	1,153

Sup. Table 3: Cell width of mutants of the BKK collection (continued)

BKK name[1]	gene	screening delta[2] (%)	average width (µm)	+/-	nb	ADP	BKK name[1]	gene	screening delta[2] (%)	average width (µm)	+/-	nb	ADP	BKK name[1]	gene	screening delta[2] (%)	average width (µm)	+/-	nb	ADP	BKK name[1]	gene	screening delta[2] (%)	average width (µm)	+/-	nb	ADP
BKK17030	cotE	-1,679	1,186	0,120	146	1,206	BKK38180	ywzA	-1,84	1,136	0,100	88	1,157	BKK32020	yuiH	-2,006	1,125	0,107	138	1,148	BKK06350	yebA	-2,185	1,153	0,110	460	1,179
BKK29320	ytnI	-1,68	1,094	0,124	219	1,113	BKK03140	tmrB	-1,844	1,109	0,093	113	1,130	BKK35100	yvlD	-2,007	1,112	0,106	409	1,135	BKK13190	ispA	-2,185	1,141	0,085	56	1,166
BKK32240	thrB	-1,681	1,129	0,128	97	1,148	BKK10080	yhfA	-1,844	1,186	0,151	72	1,208	BKK37500	speE	-2,01	1,096	0,135	44	1,119	BKK03550	ycxC	-2,186	1,105	0,097	113	1,130
BKK10040	ecsA	-1,685	1,119	0,106	117	1,138	BKK24580	yqhH	-1,844	1,136	0,093	114	1,158	BKK12740	xkdU	-2,012	1,143	0,120	135	1,166	BKK36070	cotG	-2,187	1,131	0,117	91	1,156
BKK00140	dck	-1,686	1,159	0,111	494	1,179	BKK11820	yjcD	-1,85	1,141	0,099	347	1,163	BKK33140	yvaJ	-2,017	1,197	0,083	140	1,222	BKK17250	ymaE	-2,189	1,136	0,107	575	1,162
BKK21960	ypeQ	-1,686	1,132	0,102	312	1,151	BKK19350	yocS	-1,853	1,184	0,144	83	1,206	BKK27040	levG	-2,019	1,153	0,110	255	1,177	BKK26749	yrdD	-2,189	1,151	0,097	166	1,177
BKK27270	mtnN	-1,686	1,138	0,105	125	1,158	BKK40170	yydG	-1,854	1,146	0,126	284	1,168	BKK06036	ydzT	-2,026	1,136	0,101	346	1,160	BKK32210	yuzD	-2,189	1,123	0,093	38	1,148
BKK08060	acoA	-1,69	1,148	0,100	355	1,168	BKK39750	iolB	-1,855	1,143	0,113	74	1,165	BKK17210	pksN	-2,03	1,140	0,106	81	1,164	BKK05240	ydeL	-2,19	1,114	0,105	58	1,139
BKK29720	ytxE	-1,693	1,120	0,098	134	1,139	BKK36350	ywpD	-1,858	1,140	0,101	406	1,161	BKK20600	yoqK	-2,031	1,131	0,114	250	1,155	BKK01400	rpmJ	-2,192	1,134	0,129	188	1,160
BKK00480	yabJ	-1,696	1,144	0,104	133	1,164	BKK39180	yxiH	-1,859	1,143	0,095	72	1,165	BKK05550	cotP	-2,033	1,124	0,104	346	1,147	BKK11480	yjbB	-2,195	1,182	0,120	87	1,208
BKK16720	ymxH	-1,696	1,144	0,110	151	1,164	BKK39970	yxaH	-1,859	1,128	0,097	551	1,149	BKK20230	yorW	-2,033	1,144	0,114	250	1,168	BKK28580	mutSB	-2,198	1,134	0,107	46	1,160
BKK09780	yheC	-1,699	1,157	0,100	171	1,177	BKK11790	yjcA	-1,861	1,167	0,132	759	1,189	BKK02640	tatCD	-2,034	1,177	0,071	79	1,201	BKK31840	yueD	-2,199	1,123	0,093	145	1,148
BKK39990	yxaF	-1,699	1,145	0,116	71	1,165	BKK01935	skfC	-1,866	1,092	0,095	102	1,113	BKK32140	paiB	-2,035	1,125	0,098	42	1,148	BKK15030	ylbJ	-2,2	1,153	0,138	261	1,179
BKK12280	rex	-1,7	1,143	0,111	267	1,163	BKK16650	rbfA	-1,867	1,142	0,135	97	1,164	BKK20120	yosS	-2,036	1,106	0,120	100	1,129	BKK26220	ygaQ	-2,2	1,151	0,094	408	1,177
BKK33960	araE	-1,7	1,142	0,099	359	1,161	BKK21329	youB	-1,867	1,132	0,099	450	1,153	BKK09190	yhcR	-2,037	1,153	0,104	120	1,177	BKK40090	ahpC	-2,201	1,142	0,109	340	1,168
BKK34970	ppaX	-1,7	1,116	0,123	252	1,135	BKK34460	levB	-1,867	1,114	0,116	246	1,135	BKK12570	xtmA	-2,037	1,165	0,129	436	1,189	BKK26000	ygbR	-2,206	1,126	0,092	459	1,151
BKK34600	mdxF	-1,704	1,116	0,102	143	1,135	BKK37710	bacD	-1,868	1,128	0,095	609	1,149	BKK39900	aslA	-2,041	1,142	0,108	112	1,166	BKK37270	narH	-2,206	1,094	0,108	90	1,119
BKK18710	yozF	-1,707	1,152	0,106	89	1,172	BKK03890	gabR	-1,873	1,143	0,129	557	1,165	BKK24900	rpmGA	-2,043	1,140	0,095	211	1,164	BKK25630	ygeK	-2,213	1,138	0,099	142	1,164
BKK02619	yczK	-1,713	1,111	0,105	61	1,130	BKK06570	yerB	-1,876	1,179	0,088	145	1,201	BKK34550	pgcM	-2,046	1,112	0,109	267	1,135	BKK34830	yvzA	-2,215	1,110	0,127	164	1,135
BKK31560	yufP	-1,713	1,146	0,104	163	1,166	BKK04910	yddB	-1,881	1,126	0,096	417	1,147	BKK00560	spoVT	-2,048	0,955	0,075	44	0,975	BKK11360	appD	-2,22	1,182	0,143	116	1,208
BKK05530	ydfR	-1,715	1,140	0,104	319	1,160	BKK32000	dhbA	-1,881	1,118	0,089	39	1,139	BKK21000	yonT	-2,049	1,130	0,105	585	1,153	BKK25540	gpr	-2,22	1,138	0,093	189	1,164
BKK39840	yxbG	-1,716	0,958	0,091	42	0,975	BKK39080	licT	-1,883	1,143	0,098	69	1,165	BKK31319	yuzI	-2,049	1,132	0,113	69	1,156	BKK04470	dctP	-2,221	1,132	0,095	228	1,158
BKK03690	yczF	-1,718	1,145	0,106	134	1,165	BKK26559	yrzN	-1,884	1,155	0,108	337	1,177	BKK24490	yqhQ	-2,05	1,163	0,094	377	1,187	BKK05329	ydzO	-2,222	1,141	0,107	146	1,167
BKK32560	frlR	-1,72	1,120	0,092	204	1,139	BKK20420	yorD	-1,887	1,140	0,105	457	1,162	BKK28780	araD	-2,051	1,090	0,092	351	1,113	BKK33770	spbC	-2,223	1,135	0,097	572	1,161
BKK30120	yteR	-1,721	1,136	0,089	121	1,156	BKK38140	qoxD	-1,887	1,135	0,089	126	1,157	BKK25920	yqxG	-2,054	1,140	0,130	58	1,164	BKK13150	ohrR	-2,225	1,140	0,125	195	1,166
BKK19890	yotG	-1,724	1,110	0,125	487	1,129	BKK01240	rpmC	-1,888	1,126	0,106	154	1,147	BKK18590	yosG	-2,055	1,115	0,094	104	1,138	BKK01760	ybbR	-2,226	1,134	0,104	558	1,160
BKK11330	fabHA	-1,725	1,163	0,114	77	1,183	BKK07070	yesY	-1,888	1,157	0,135	181	1,179	BKK05840	gmuD	-2,056	1,143	0,077	244	1,167	BKK11220	argD	-2,235	1,142	0,109	390	1,168
BKK24640	yqxM	-1,725	1,131	0,100	200	1,151	BKK32370	yunD	-1,889	1,126	0,118	44	1,148	BKK09110	yhcI	-2,056	1,183	0,142	691	1,208	BKK19150	yocB	-2,239	1,136	0,103	468	1,162
BKK36830	atpA	-1,727	1,095	0,114	135	1,114	BKK37230	ywjA	-1,892	1,126	0,106	474	1,147	BKK27660	comN	-2,058	1,136	0,107	232	1,160	BKK34510	yvdQ	-2,239	1,135	0,101	418	1,161
BKK32460	pucM	-1,728	1,128	0,139	50	1,148	BKK20780	yopS	-1,893	1,140	0,105	463	1,162	BKK20710	yopZ	-2,06	1,131	0,104	381	1,155	BKK40200	yydD	-2,239	1,125	0,102	270	1,151
BKK12940	dppC	-1,729	1,187	0,151	94	1,208	BKK18740	yozG	-1,894	1,149	0,125	113	1,172	BKK21010	yonS	-2,06	1,182	0,106	94	1,206	BKK30440	ytrC	-2,249	1,195	0,088	239	1,222
BKK26860	yraO	-1,73	1,156	0,093	212	1,177	BKK09259	yhzG	-1,897	1,141	0,101	340	1,163	BKK33250	yvrL	-2,06	1,137	0,104	253	1,161	BKK18530	yoaA	-2,25	1,132	0,108	113	1,158
BKK38240	ywcA	-1,732	1,099	0,144	41	1,119	BKK17860	yneA	-1,9	1,141	0,121	79	1,164	BKK02730	natK	-2,062	1,177	0,059	52	1,201	BKK18780	yooW	-2,251	1,145	0,095	267	1,172
BKK25620	yqeL	-1,74	1,143	0,093	205	1,164	BKK25960	yqcB	-1,901	1,141	0,112	132	1,164	BKK30360	yttA	-2,065	1,132	0,101	148	1,156	BKK00680	hprT	-2,252	1,122	0,103	265	1,147
BKK11690	yncM	-1,741	1,142	0,097	467	1,162	BKK22330	ypoC	-1,905	1,129	0,101	491	1,151	BKK05820	gmuA	-2,068	1,124	0,095	395	1,147	BKK11810	spoVIF	-2,255	1,162	0,121	413	1,189
BKK04740	rsbX	-1,745	1,145	0,137	276	1,165	BKK37700	bacE	-1,91	1,127	0,113	257	1,149	BKK26530	yrkF	-2,07	1,130	0,109	398	1,153	BKK37770	rocB	-2,255	1,094	0,123	46	1,119
BKK31440	potB	-1,745	1,128	0,096	187	1,148	BKK02970	yceK	-1,913	1,108	0,089	99	1,130	BKK35239	swrAA	-2,071	1,112	0,123	410	1,135	BKK02890	yceC	-2,256	1,131	0,095	132	1,158
BKK32870	yusO	-1,745	1,115	0,106	105	1,134	BKK01750	ybbP	-1,915	1,137	0,087	209	1,160	BKK37740	bacA	-2,072	1,096	0,098	90	1,119	BKK29140	citZ	-2,261	1,088	0,138	132	1,113
BKK07420	yfmM	-1,75	1,187	0,136	44	1,208	BKK25130	nfo	-1,917	1,141	0,097	250	1,164	BKK15630	sirC	-2,074	1,124	0,112	433	1,148	BKK39560	yxeG	-2,262	1,138	0,092	115	1,165
BKK34140	ganQ	-1,75	1,137	0,087	145	1,158	BKK29450	argG	-1,919	1,129	0,102	330	1,151	BKK13560	mtnK	-2,076	1,138	0,107	430	1,163	BKK34170	ganR	-2,265	1,109	0,121	92	1,135
BKK05990	tatCY	-1,759	1,139	0,110	489	1,160	BKK08960	yhbF	-1,923	1,154	0,096	209	1,177	BKK21130	yonD	-2,079	1,131	0,097	406	1,155	BKK09870	khtS	-2,266	1,181	0,180	175	1,208
BKK17880	ynzC	-1,766	1,141	0,100	341	1,162	BKK26570	yrkB	-1,923	1,131	0,102	355	1,153	BKK30870	ytcB	-2,08	1,197	0,063	445	1,222	BKK25750	nucB	-2,267	0,953	0,064	47	0,975
BKK21640	yokC	-1,773	1,133	0,100	472	1,153	BKK16250	fliI	-1,927	1,146	0,102	344	1,168	BKK12240	yjkA	-2,087	1,183	0,158	459	1,208	BKK26619	yrzO	-2,271	1,150	0,109	114	1,177
BKK39710	iolF	-1,773	1,146	0,099	194	1,166	BKK21590	yokH	-1,929	1,133	0,109	426	1,155	BKK05800	pbuE	-2,088	1,176	0,051	59	1,201	BKK05510	ydfQ	-2,272	1,141	0,106	647	1,167
BKK23830	yqjL	-1,776	1,166	0,112	135	1,187	BKK26075	yqbK	-1,933	1,129	0,105	474	1,151	BKK06000	ydiK	-2,091	1,143	0,098	204	1,167	BKK07640	yffL	-2,272	1,132	0,106	495	1,159
BKK22300	yppC	-1,778	1,081	0,120	153	1,100	BKK06420	purE	-1,938	1,156	0,124	258	1,179	BKK11890	yjcK	-2,094	1,164	0,121	356	1,189	BKK13500	ktrD	-2,274	1,140	0,101	289	1,166
BKK33500	copA	-1,778	1,118	0,101	103	1,138	BKK40310	phrG	-1,938	1,129	0,097	459	1,151	BKK06190	ydjG	-2,096	1,154	0,103	168	1,179	BKK15050	ylbL	-2,275	1,114	0,107	81	1,139
BKK04270	ydaJ	-1,78	1,180	0,070	54	1,201	BKK13790	ykvQ	-1,939	1,144	0,107	90	1,166	BKK05760	ydhH	-2,098	1,143	0,131	139	1,167	BKK00550	mfd	-2,277	1,137	0,113	86	1,164
BKK19999	yojW	-1,781	1,109	0,134	310	1,129	BKK33100	liaF	-1,941	1,112	0,095	174	1,134	BKK24700	comGD	-2,099	1,127	0,104	387	1,151	BKK20810	yopP	-2,28	1,125	0,108	268	1,151
BKK24080	bcd	-1,787	1,166	0,112	291	1,187	BKK27380	yrzB	-1,946	1,129	0,102	372	1,151	BKK39970	araR	-2,101	1,133	0,091	124	1,158	BKK27050	levF	-2,282	1,150	0,094	349	1,177
BKK11970	yjcS	-1,79	1,142	0,111	337	1,163	BKK02150	ybeF	-1,948	1,108	0,113	74	1,130	BKK12840	pit	-2,103	1,183	0,132	80	1,208	BKK07970	yfjT	-2,289	1,132	0,114	252	1,159
BKK30595	ytlA	-1,791	1,200	0,059	243	1,222	BKK13240	thiU	-1,95	1,144	0,117	52	1,166	BKK34740	crh	-2,105	1,111	0,094	379	1,133	BKK02720	yccF	-2,29	1,174	0,082	45	1,201
BKK01040	yonF	-1,793	1,134	0,116	243	1,155	BKK17540	ynaF	-1,95	1,109	0,107	360	1,162	BKK26690	brnQ	-2,114	1,196	0,086	222	1,222	BKK36580	ywnF	-2,291	1,089	0,141	214	1,114
BKK32480	pucD	-1,793	1,145	0,108	84	1,166	BKK00070	csgA	-1,952	1,108	0,093	59	1,130	BKK02320	ybfP	-2,117	1,106	0,103	125	1,130	BKK13980	bbpH	-2,294	1,136	0,107	403	1,163
BKK13990	kinA	-1,798	1,163	0,113	98	1,184	BKK28410	gerE	-1,952	1,137	0,093	132	1,160	BKK37490	speB	-2,117	1,095	0,099	57	1,119	BKK38810	msmX	-2,294	1,181	0,162	257	1,208
BKK05780	ydhJ	-1,799	1,139	0,099	444	1,160	BKK02060	ybxG	-1,953	1,108	0,078	45	1,130	BKK07520	yfmC	-2,119	1,183	0,125	181	1,208	BKK30930	ytaB	-2,296	1,135	0,100	429	1,161
BKK32360	yunC	-1,799	1,127	0,115	101	1,148	BKK30240	bioW	-1,956	1,133	0,098	264	1,156	BKK12480	yjqB	-2,126	1,164	0,147	729	1,189	BKK13930	splB	-2,299	1,140	0,131	47	1,166
BKK03981	mtlA	-1,801	1,144	0,089	125	1,165	BKK14580	pdhA	-1,958	1,156	0,161	118	1,179	BKK36090	ywrE	-2,126	1,091	0,098	458	1,114	BKK33410	yvgO	-2,3	1,108	0,079	85	1,134
BKK10940	yitC	-1,801	1,162	0,112	128	1,183	BKK35540	tuaH	-1,961	1,127	0,100	691	1,149	BKK17470	yfmI	-2,131	1,135	0,110	546	1,160	BKK21650	yokB	-2,301	1,127	0,106	388	1,153
BKK15720	defA	-1,801	1,114	0,115	85	1,134	BKK35630	lytB	-1,961	1,127	0,100	691	1,149	BKK25670	yqeH	-2,131	1,127	0,133	130	1,151	BKK26558	yrzM	-2,302	1,127	0,104	248	1,153
BKK26740	cypA	-1,802	1,155	0,101	376	1,177	BKK19330	sodF	-1,961	1,149	0,112	116	1,172	BKK37800	yweA	-2,137	1,125	0,105	439	1,149	BKK39250	yxiE	-2,304	1,138	0,117	85	1,165
BKK35090	yvmA	-1,802	1,129	0,097	453	1,149	BKK37140	ywjG	-1,968	1,097	0,112	690	1,119	BKK08240	catE	-2,141	1,134	0,109	292	1,159	BKK10470	yhjD	-2,305	1,136	0,106	422	1,163
BKK08100	acoR	-1,805	1,138	0,098	462	1,159	BKK31300	yugS	-1,97	1,138	0,102	296	1,161	BKK28930	tatD	-2,146	1,123	0,098	421	1,147	BKK28930	lytS	-2,308	1,125	0,115	236	1,151
BKK12390	uxaA	-1,805	1,147	0,105	422	1,168	BKK38499	ywzH	-1,972	1,097	0,122	85	1,119	BKK30220	bioF	-2,15	1,131	0,103	114	1,156	BKK10090	yhgB	-2,313	1,150	0,096	126	1,177
BKK39390	hutM	-1,809	1,144	0,107	106	1,165	BKK24280	ispA	-1,974	1,164	0,099	281	1,187	BKK31780	crh	-2,152	1,123	0,107	205	1,148	BKK00040	recF	-2,314	1,140	0,135	688	1,167
BKK22930	sleB	-1,81	1,080	0,128	100	1,100	BKK25040	yggB	-1,974	1,141	0,114	149	1,164	BKK06790	yeeD	-2,157	1,135	0,096	487	1,160	BKK04580	cshA	-2,315	1,133	0,121	225	1,160
BKK07430	yfmL	-1,811	1,139	0,109	434	1,160	BKK36530	bcrC	-1,975	1,092	0,073	51	1,114	BKK21430	yezC	-2,158	1,130	0,103	474	1,155	BKK17960	yneK	-2,316	1,135	0,112	478	1,162
BKK22810	yphF	-1,812	1,185	0,126	147	1,206	BKK03730	yclH	-1,976	1,178	0,116	979	1,201	BKK18450	gltA	-2,159	1,113	0,087	57	1,138	BKK33350	yvsG	-2,316	1,108	0,100	127	1,134
BKK07530	yfmB	-1,815	1,138	0,106	496	1,159	BKK27760	csbX	-1,977	1,137	0,113	408	1,160	BKK10920	yjdB	-2,163	1,158	0,114	121	1,183	BKK01820	adaB	-2,321	1,133	0,101	151	1,160
BKK36660	ureA	-1,815	1,094	0,121	218	1,114	BKK15330	sigG	-1,978	1,117	0,114	66	1,139	BKK38420	sacY	-2,163	1,095	0,113	183	1,119	BKK26230	ygaP	-2,327	1,149	0,086	152	1,177
BKK33810	yvbG	-1,819	0,957	0,071	65	0,975	BKK00350	yazA	-1,978	1,156	0,155	152	1,179	BKK01610	feuC	-2,166	1,151	0,095	232	1,177	BKK39000	yxjC	-2,327	1,138	0,102	117	1,165
BKK14110	ykuK	-1,821	1,141	0,111	232	1,163	BKK14420	ykoA	-1,984	1,156	0,117	450	1,179	BKK38390	ywbA	-2,167	1,132	0,124	61	1,157	BKK26130	ygbF	-2,33	0,952	0,105	50	0,975
BKK34160	cycB	-1,821	1,129	0,113	428	1,149	BKK24710	comGC	-1,984	1,140	0,091	464	1,164	BKK16010	yttD	-2,168	1,132	0,107	126	1,157	BKK10860	yisT	-2,332	1,156	0,106	103	1,183
BKK34820	bmrA	-1,823	1,128	0,094	590	1,149	BKK32940	yusV	-1,985	1,198	0,098	38	1,222	BKK24030	bkdB	-2,168	1,161	0,108	173	1,187	BKK20770	yopT	-2,332	1,135	0,103	225	1,162
BKK25600	comER	-1,825	1,142	0,094	229	1,164	BKK35240	ctpB	-1,988	1,113	0,131	420	1,135	BKK22130	kduI	-2,171	1,076	0,137	144	1,100	BKK17150	pksG	-2,333	1,136	0,105	116	1,164
BKK01060	ybxB	-1,83	1,142	0,097	388	1,164	BKK13130	proA	-1,989	1,143	0,113	173	1,166	BKK29520	yteJ	-2,178	1,089	0,105	72	1,113	BKK38570	licA	-2,335	1,138	0,092	79	1,165
BKK06079	ydzW	-1,832	1,136	0,101	98	1,158	BKK16790	tepA	-1,99	1,140	0,117	96	1,164	BKK09670	dltA	-2,179	1,182	0,127	45	1,208	BKK22860	yphA	-2,337	1,178	0,119	103	1,206
BKK15460	rluD	-1,835	1,093	0,110	129	1,113	BKK27590	dtd	-1,991	1,137	0,095	316	1,160	BKK32060	yuiD	-2,18	1,115	0,109	82	1,139	BKK28920	lytT	-2,346	1,193	0,093	69	1,222
BKK31850	yueC	-1,837	1,127	0,115	237	1,148	BKK08570	yfhK	-1,996	1,136	0,118	187	1,159	BKK25850	yqzI	-2,181	1,138	0,112	104	1,164	BKK39870	yxbD	-2,348	1,137	0,097	99	1,165

Sup. Table 3: Cell width of mutants of the BKK collection (continued)

BKK name[1]	gene	screening delta[2] (%)	erage width (µ	+/-	nb	ADP
BKK33420	nhaK	-2,351	1,134	0,101	490	1,161
BKK14710	yloA	-2,355	1,151	0,113	150	1,179
BKK19990	yosV	-2,356	1,141	0,108	317	1,168
BKK32470	pucE	-2,357	1,113	0,096	129	1,139
BKK01470	ybaF	-2,358	1,136	0,092	172	1,164
BKK05560	ydgA	-2,36	1,130	0,103	167	1,158
BKK12110	yjfA	-2,36	1,161	0,135	210	1,189
BKK40180	yydF	-2,363	1,140	0,113	245	1,168
BKK10280	yhfM	-2,366	1,155	0,091	112	1,183
BKK39820	htpG	-2,366	1,137	0,086	112	1,165
BKK01350	rplO	-2,372	1,136	0,099	135	1,164
BKK38860	galE	-2,375	1,137	0,119	81	1,165
BKK29110	phoP	-2,376	1,087	0,104	34	1,113
BKK29239	ytrH	-2,378	1,087	0,080	70	1,113
BKK16450	cheC	-2,379	1,134	0,102	380	1,161
BKK15540	pyrD	-2,384	1,128	0,122	73	1,156
BKK19259	yoyB	-2,384	1,144	0,108	70	1,172
BKK20220	yorX	-2,386	1,112	0,112	59	1,139
BKK14390	fruK	-2,387	1,151	0,141	131	1,179
BKK20380	yorH	-2,39	1,102	0,124	48	1,129
BKK25980	yqbT	-2,392	0,952	0,072	120	0,975
BKK39580	yxeE	-2,394	1,137	0,089	128	1,165
BKK17050	mutL	-2,398	1,136	0,104	144	1,164
BKK16620	rplGA	-2,399	1,130	0,100	81	1,158
BKK06049	ydzV	-2,404	1,139	0,111	125	1,167
BKK14640	yktA	-2,404	1,135	0,110	295	1,163
BKK30270	msmE	-2,414	1,111	0,096	110	1,138
BKK37300	ywiC	-2,416	1,092	0,110	94	1,119
BKK12920	dppA	-2,417	1,138	0,135	52	1,166
BKK28700	yrzH	-2,42	1,123	0,114	398	1,151
BKK31322	yugO	-2,421	1,128	0,104	48	1,156
BKK27160	cypB	-2,422	1,148	0,110	280	1,177
BKK26610	yrkA	-2,425	1,148	0,121	121	1,177
BKK24740	yqxL	-2,426	1,177	0,128	71	1,206
BKK20960	yopA	-2,427	1,127	0,105	270	1,155
BKK30280	amyD	-2,427	1,110	0,101	68	1,138
BKK07040	yesV	-2,431	1,150	0,111	337	1,179
BKK22800	spoIVA	-2,434	1,112	0,092	55	1,139
BKK25510	lepA	-2,435	1,135	0,103	312	1,164
BKK26750	yrdD	-2,437	1,148	0,105	326	1,177
BKK35110	yvlC	-2,438	1,107	0,103	428	1,135
BKK10590	yhjP	-2,439	1,134	0,100	274	1,163
BKK27870	nadD	-2,441	1,131	0,099	246	1,160
BKK08040	yfjM	-2,442	1,131	0,097	529	1,159
BKK19480	yojE	-2,442	1,102	0,101	828	1,129
BKK12310	exuM	-2,445	1,179	0,120	75	1,208
BKK18880	yobE	-2,448	1,133	0,117	361	1,162
BKK27240	yrhC	-2,448	1,148	0,109	166	1,177
BKK03450	hxlB	-2,454	1,102	0,094	30	1,130
BKK17900	yneE	-2,456	1,135	0,085	105	1,164
BKK12800	xhlB	-2,463	1,138	0,129	184	1,166
BKK10840	degA	-2,473	1,154	0,100	142	1,183
BKK32650	yurS	-2,474	1,137	0,105	138	1,166
BKK34030	yvbY	-2,474	1,106	0,091	88	1,134
BKK13660	kinD	-2,476	1,177	0,160	152	1,206
BKK30210	bioD	-2,479	1,127	0,118	81	1,156
BKK11928	yjzF	-2,488	1,139	0,105	531	1,168
BKK17360	ymzA	-2,491	1,135	0,107	132	1,164
BKK38740	cydC	-2,491	1,172	0,129	459	1,201
BKK14490	kinC	-2,494	1,176	0,151	48	1,206
BKK39300	yxiD	-2,497	1,137	0,094	81	1,166
BKK12350	uxuB	-2,498	1,159	0,141	177	1,189
BKK16350	fliP	-2,498	1,133	0,108	581	1,162
BKK15610	sumT	-2,501	1,119	0,100	165	1,148
BKK30880	ytcC	-2,505	1,132	0,107	301	1,161
BKK40050	gntR	-2,516	1,119	0,119	149	1,147
BKK12360	exuT	-2,521	1,178	0,149	104	1,208
BKK18940	yazK	-2,523	1,142	0,105	362	1,172
BKK19450	yojG	-2,523	1,101	0,120	194	1,129
BKK03850	ycnC	-2,524	1,136	0,098	318	1,165
BKK40750	yyaQ	-2,524	1,119	0,106	545	1,147
BKK26820	yrpD	-2,527	1,147	0,111	280	1,177
BKK39730	iolD	-2,527	1,135	0,108	107	1,165
BKK20150	yosE	-2,528	1,101	0,157	129	1,129
BKK32250	thrC	-2,529	1,119	0,096	75	1,148
BKK28520	etfA	-2,532	1,128	0,107	72	1,158
BKK21340	yomJ	-2,533	1,126	0,096	315	1,155
BKK37020	ywkD	-2,533	1,086	0,081	34	1,114
BKK22900	ypfB	-2,534	1,072	0,109	247	1,100
BKK27009	yrzP	-2,535	1,124	0,106	366	1,153
BKK10670	gerPF	-2,539	1,153	0,097	128	1,183
BKK07790	yfkQ	-2,54	1,129	0,114	366	1,159
BKK15560	yodD	-2,541	1,101	0,129	112	1,129
BKK24260	yqxC	-2,542	1,128	0,090	113	1,158
BKK24470	yqhS	-2,542	1,128	0,090	113	1,158
BKK05409	ydzQ	-2,543	1,118	0,098	434	1,147
BKK26940	yraH	-2,544	1,147	0,098	433	1,177
BKK30739	ytzL	-2,548	1,127	0,091	60	1,156
BKK10000	yhaH	-2,551	1,133	0,101	400	1,163
BKK05310	ydeR	-2,553	1,130	0,114	429	1,160
BKK18898	yozW	-2,555	1,109	0,093	41	1,138
BKK21770	ilvA	-2,556	0,950	0,072	150	0,975
BKK18550	yoaC	-2,562	1,109	0,109	53	1,138
BKK12200	yjiA	-2,563	1,159	0,083	1736	1,189
BKK08030	dusC	-2,566	1,129	0,106	451	1,159
BKK18560	yoaD	-2,568	1,132	0,110	415	1,162
BKK18060	yneR	-2,577	1,134	0,105	133	1,164
BKK27860	nadC	-2,578	1,121	0,115	422	1,151
BKK26600	bltD	-2,579	1,146	0,099	140	1,177
BKK28260	leuC	-2,582	1,130	0,085	298	1,160
BKK36310	ssbB	-2,59	1,085	0,114	158	1,114
BKK40259	yyzG	-2,594	1,120	0,120	232	1,149
BKK16810	ymfC	-2,598	1,133	0,087	86	1,164
BKK00580	yabN	-2,603	1,133	0,093	136	1,164
BKK18170	yngA	-2,612	1,141	0,125	150	1,172
BKK36120	ywrB	-2,614	1,136	0,096	96	1,166
BKK28900	lrgB	-2,618	1,084	0,121	239	1,113
BKK39050	katE	-2,621	1,134	0,094	87	1,165
BKK10770	wprA	-2,626	1,152	0,118	131	1,183
BKK29170	ytzA	-2,626	1,131	0,102	350	1,161
BKK08200	malP	-2,633	1,128	0,098	405	1,159
BKK09910	yhaO	-2,633	1,153	0,102	172	1,184
BKK17550	ynaG	-2,633	1,133	0,151	89	1,164
BKK20630	yoqH	-2,633	1,131	0,103	434	1,162
BKK35900	pgsB	-2,633	1,119	0,103	461	1,149
BKK06530	purD	-2,634	1,148	0,118	479	1,179
BKK40021	yxzK	-2,636	1,119	0,089	601	1,149
BKK21440	bdbB	-2,643	1,124	0,103	548	1,155
BKK06290	yeaA	-2,644	1,148	0,116	159	1,179
BKK35030	yvoA	-2,648	1,135	0,113	78	1,166
BKK01380	mapA	-2,649	1,133	0,104	327	1,164
BKK22050	bcsA	-2,651	1,071	0,103	79	1,100
BKK25140	cshB	-2,654	1,133	0,107	190	1,164
BKK33810	opuCC	-2,656	1,119	0,101	243	1,149
BKK21680	msrB	-2,662	1,124	0,093	567	1,155
BKK03460	hxlA	-2,666	1,100	0,096	72	1,130
BKK06800	yezA	-2,671	1,147	0,112	331	1,179
BKK14840	ylaN	-2,671	1,148	0,089	337	1,179
BKK10570	yhjN	-2,674	1,152	0,096	86	1,183
BKK15130	mraZ	-2,674	1,108	0,100	130	1,138
BKK04000	ycsA	-2,676	1,134	0,109	110	1,165
BKK26748	yrdD	-2,68	1,152	0,098	379	1,177
BKK28210	lonB	-2,683	1,129	0,093	223	1,160
BKK22460	ypjH	-2,687	1,120	0,100	398	1,151
BKK36890	upp	-2,692	1,084	0,127	283	1,114
BKK26360	yqaD	-2,695	0,949	0,088	33	0,975
BKK33290	fhuC	-2,696	1,135	0,092	100	1,166
BKK13370	ykoQ	-2,697	1,135	0,109	154	1,166
BKK02080	ybxH	-2,698	1,099	0,097	41	1,130
BKK13960	ykwC	-2,7	1,135	0,118	328	1,166
BKK11280	yjaU	-2,7	1,125	0,095	139	1,156
BKK21950	ypzA	-2,71	1,124	0,107	354	1,155
BKK40740	yyaR	-2,71	1,152	0,101	266	1,184
BKK36610	ywnC	-2,717	1,125	0,105	107	1,157
BKK14980	ylbE	-2,718	1,147	0,130	424	1,179
BKK23420	spoVAC	-2,718	1,120	0,110	328	1,151
BKK37200	ywjD	-2,721	0,948	0,072	47	0,975
BKK24880	yqgO	-2,728	1,132	0,098	228	1,164
BKK27970	spoIVFB	-2,733	1,128	0,102	269	1,160
BKK03470	hxlR	-2,74	1,099	0,087	84	1,130
BKK12750	xkdV	-2,743	1,134	0,127	291	1,166
BKK30860	ytcA	-2,746	1,188	0,074	183	1,222
BKK13570	mtnU	-2,747	1,134	0,099	260	1,166
BKK19790	cgeB	-2,751	1,098	0,091	379	1,129
BKK30260	msmR	-2,757	1,124	0,102	89	1,156
BKK34630	mdxR	-2,757	1,129	0,091	497	1,161
BKK17800	yndJ	-2,759	1,131	0,125	81	1,164
BKK03569	sfp	-2,765	1,099	0,093	46	1,130
BKK15210	spoVE	-2,766	1,173	0,125	72	1,206
BKK22280	yppD	-2,774	1,070	0,114	281	1,100
BKK29770	ytxH	-2,777	1,082	0,121	83	1,113
BKK34770	yvcJ	-2,778	1,104	0,114	215	1,135
BKK35730	tagE	-2,778	1,134	0,104	114	1,166
BKK15750	yloN	-2,779	1,103	0,097	89	1,134
BKK23660	yqkB	-2,783	1,154	0,103	246	1,187
BKK25240	yqfL	-2,784	1,131	0,102	140	1,164
BKK28380	gerM	-2,788	1,127	0,117	165	1,160
BKK00980	sigH	-2,792	1,154	0,109	586	1,187
BKK00960	rlmB	-2,8	1,154	0,093	297	1,187
BKK24910	yqgM	-2,8	1,119	0,105	522	1,151
BKK29830	ytpQ	-2,8	1,125	0,105	157	1,158
BKK05109	ydzN	-2,801	1,127	0,101	522	1,160
BKK13330	ykoL	-2,802	1,108	0,114	39	1,139
BKK23870	polYA	-2,803	1,154	0,100	293	1,187
BKK27640	yrvC	-2,807	1,127	0,093	225	1,160
BKK00830	ctsR	-2,808	1,069	0,121	202	1,100
BKK09890	natA	-2,808	1,174	0,151	90	1,208
BKK12020	manA	-2,808	1,156	0,138	122	1,189
BKK39370	hutI	-2,808	0,948	0,072	30	0,975
BKK36180	sacC	-2,81	1,134	0,098	114	1,166
BKK02600	cwlJ	-2,812	1,168	0,061	37	1,201
BKK38370	ywbC	-2,813	1,087	0,222	31	1,119
BKK32280	yutG	-2,816	1,116	0,105	62	1,148
BKK15640	yloA	-2,818	1,116	0,165	31	1,148
BKK27850	nadA	-2,823	1,127	0,098	241	1,160
BKK35609	tuaA	-2,823	1,117	0,109	518	1,149
BKK14360	yknY	-2,829	1,106	0,096	114	1,138
BKK27530	yrvN	-2,83	1,127	0,093	414	1,160
BKK32719	yuzK	-2,83	1,102	0,097	169	1,134
BKK20490	yoqW	-2,843	1,097	0,126	55	1,129
BKK09430	citR	-2,849	1,143	0,109	184	1,177
BKK27030	sacV	-2,849	1,143	0,095	115	1,177
BKK28180	ysxD	-2,85	1,127	0,099	313	1,160
BKK02560	ycbM	-2,851	1,098	0,099	170	1,130
BKK07920	yfkE	-2,852	1,126	0,101	420	1,159
BKK08630	ydzN	-2,855	1,126	0,105	319	1,159
BKK11290	yjaV	-2,86	1,149	0,123	133	1,183
BKK08580	yfhL	-2,862	1,126	0,095	507	1,159
BKK40320	argI	-2,862	1,118	0,105	351	1,151
BKK26640	yqqG	-2,866	1,129	0,102	411	1,162
BKK06510	purN	-2,871	1,145	0,103	417	1,179
BKK08090	acoL	-2,873	1,129	0,106	386	1,163
BKK25400	yqeY	-2,873	1,130	0,103	178	1,164
BKK16845	ymfF	-2,877	1,130	0,114	127	1,164
BKK36000	alsD	-2,881	1,124	0,100	167	1,157
BKK20340	yorL	-2,89	1,097	0,144	56	1,129
BKK31760	pncA	-2,894	1,106	0,096	129	1,139
BKK20100	yosI	-2,898	1,097	0,133	51	1,129
BKK01060	yaaH	-2,903	1,125	0,122	113	1,159
BKK13590	mtnW	-2,908	1,132	0,083	62	1,166
BKK28860	rpmI	-2,908	1,118	0,090	208	1,151
BKK36200	ywqI	-2,909	1,082	0,114	70	1,114
BKK00660	yabT	-2,912	1,096	0,161	81	1,129
BKK37940	ywdI	-2,916	1,086	0,115	64	1,119
BKK40430	yycE	-2,918	1,132	0,099	185	1,166
BKK01860	ybcF	-2,921	1,126	0,100	422	1,160
BKK07690	yfmP	-2,928	1,125	0,111	181	1,159
BKK11060	ymzD	-2,929	1,129	0,099	74	1,164
BKK21670	yqgP	-2,929	1,125	0,104	256	1,159
BKK15170	spoVD	-2,931	1,129	0,117	221	1,163
BKK31130	yonJ	-2,931	1,186	0,079	95	1,222
BKK21100	yonG	-2,932	1,117	0,114	347	1,151
BKK27468	yrzC	-2,932	1,126	0,093	407	1,160
BKK21570	yokJ	-2,94	1,121	0,112	417	1,155
BKK36020	alsR	-2,95	1,081	0,116	336	1,114
BKK26410	psiE	-2,953	1,142	0,103	251	1,177
BKK10070	yhaM	-2,954	1,142	0,100	150	1,177
BKK30370	bceB	-2,959	1,117	0,122	241	1,151
BKK39040	yxiS	-2,961	1,130	0,109	60	1,165
BKK21330	yomK	-2,963	1,119	0,098	639	1,153
BKK21310	yorL	-2,965	1,121	0,112	299	1,155
BKK21910	metA	-2,965	1,123	0,107	137	1,158
BKK12610	xkdG	-2,968	1,132	0,119	584	1,166
BKK18080	yneT	-2,968	1,129	0,103	100	1,164
BKK32469	yuzJ	-2,968	1,114	0,108	58	1,148
BKK06083	ydzW	-2,972	1,133	0,102	143	1,167
BKK32330	lipA	-2,976	1,127	0,102	309	1,161
BKK20270	yorS	-2,98	1,149	0,106	101	1,184
BKK37360	sboX	-2,982	1,085	0,096	95	1,119
BKK28270	leuB	-2,983	1,125	0,107	318	1,160
BKK40770	tetB	-2,985	1,149	0,108	84	1,184
BKK00620	yfhP	-2,99	1,124	0,103	526	1,159
BKK40220	yydB	-2,99	1,131	0,098	113	1,166
BKK31480	yuxJ	-2,992	1,114	0,129	184	1,148
BKK31490	pbpD	-2,993	1,185	0,074	103	1,222
BKK37530	ywhC	-2,999	1,085	0,097	81	1,119
BKK08670	ygaB	-3	1,124	0,115	301	1,159
BKK13420	ykoW	-3,003	1,172	0,147	169	1,208
BKK34750	whiA	-3,003	1,101	0,144	238	1,135
BKK23329	ypuC	-3,016	1,116	0,097	316	1,151
BKK33600	smpB	-3,02	1,100	0,093	70	1,134
BKK20060	nrdEB	-3,021	1,133	0,108	417	1,168
BKK02820	rapJ	-3,026	1,096	0,099	54	1,130
BKK01570	ybaN	-3,027	1,141	0,098	283	1,177
BKK20950	yopB	-3,03	1,120	0,111	297	1,155
BKK15510	pyrAA	-3,032	1,121	0,094	126	1,156
BKK29130	icd	-3,037	1,079	0,108	216	1,113
BKK03560	ycxD	-3,039	1,096	0,099	61	1,130
BKK40920	engD	-3,041	1,148	0,097	110	1,184
BKK09680	nhaC	-3,044	1,141	0,116	113	1,177
BKK29020	gapB	-3,044	1,079	0,093	142	1,113
BKK12880	ykcB	-3,053	1,131	0,111	385	1,166
BKK07200	yetJ	-3,058	1,143	0,100	309	1,179
BKK06550	yecA	-3,059	1,143	0,108	290	1,179
BKK17070	ymcC	-3,059	1,128	0,120	77	1,164
BKK14630	speA	-3,064	1,143	0,111	578	1,179
BKK11780	cotV	-3,066	1,153	0,125	942	1,183
BKK24590	yqhG	-3,067	0,945	0,091	78	0,975
BKK19190	desK	-3,071	1,169	0,108	71	1,206
BKK23700	yqjX	-3,071	1,151	0,099	168	1,187
BKK27890	nadR	-3,074	1,124	0,100	193	1,160
BKK25720	yqeD	-3,075	1,116	0,107	389	1,151
BKK09690	nhaX	-3,08	1,141	0,099	150	1,177
BKK26090	yqbJ	-3,086	1,116	0,102	571	1,151
BKK06900	cotJB	-3,088	1,142	0,116	414	1,179
BKK28820	ysdC	-3,09	1,104	0,103	125	1,139
BKK17400	ymaB	-3,092	1,126	0,106	441	1,162
BKK15670	ylzA	-3,11	1,099	0,073	125	1,134
BKK27250	mccB	-3,11	1,140	0,114	181	1,177
BKK20880	yopI	-3,118	0,945	0,070	69	0,975
BKK31690	comP	-3,118	1,130	0,098	91	1,166
BKK39790	yxcE	-3,119	1,128	0,075	63	1,165
BKK27520	cymR	-3,12	1,123	0,099	388	1,160
BKK24370	spoIIIAG	-3,121	1,115	0,095	412	1,151
BKK40520	yybS	-3,126	1,130	0,104	157	1,166
BKK11690	thiG	-3,13	1,131	0,106	235	1,168
BKK09040	yhcD	-3,132	1,140	0,100	215	1,177
BKK22110	kdgK	-3,133	1,066	0,117	155	1,100
BKK08710	gsaB	-3,15	1,122	0,108	543	1,159
BKK35640	lytA	-3,153	1,125	0,107	340	1,161
BKK05343	ydzS	-3,154	1,104	0,100	137	1,139
BKK36730	ywmD	-3,154	1,079	0,104	233	1,114
BKK10540	ntdB	-3,157	1,170	0,147	103	1,208
BKK19170	yocD	-3,157	1,135	0,086	143	1,172
BKK34360	epsB	-3,157	1,113	0,096	410	1,149
BKK01010	nusG	-3,16	1,150	0,133	98	1,187
BKK08720	ygaF	-3,163	1,122	0,108	460	1,159
BKK24480	yghR	-3,164	1,150	0,108	225	1,187
BKK30850	ytdA	-3,165	1,183	0,059	48	1,222
BKK27750	bofC	-3,168	1,123	0,096	254	1,160
BKK23500	drm	-3,182	1,117	0,098	469	1,153
BKK21140	yopX	-3,183	1,118	0,103	283	1,155
BKK40573	yyzI	-3,185	1,129	0,123	86	1,166
BKK23310	yutD	-3,19	1,124	0,097	399	1,161
BKK25020	sodA	-3,192	1,126	0,100	159	1,164
BKK19970	yosX	-3,193	1,093	0,142	297	1,129
BKK40910	rpsF	-3,199	1,147	0,108	212	1,184
BKK14540	ykzG	-3,2	0,944	0,093	45	0,975
BKK22420	panC	-3,206	1,121	0,112	139	1,158
BKK25280	recO	-3,206	1,126	0,110	82	1,164
BKK06930	yesK	-3,208	1,163	0,082	103	1,201
BKK09610	yhdV	-3,211	1,139	0,098	139	1,177
BKK11580	xtmB	-3,213	1,151	0,158	645	1,189
BKK21350	yomI	-3,215	1,118	0,106	363	1,155

Sup. Table 3: Cell width of mutants of the BKK collection (continued)

BKK name [1]	gene	screening delta[2] (%)	average width (µ	+/-	nb	ADP
BKK09790	yheB	-3,22	1,139	0,102	160	1,177
BKK26100	yabI	-3,22	1,139	0,092	273	1,177
BKK31790	yueG	-3,222	1,124	0,102	275	1,161
BKK29880	malS	-3,223	1,077	0,100	138	1,113
BKK27770	yrbE	-3,227	1,122	0,092	155	1,160
BKK14040	ykuD	-3,239	1,129	0,144	63	1,166
BKK37660	pta	-3,241	1,082	0,103	43	1,119
BKK00300	yaaR	-3,253	1,110	0,098	509	1,147
BKK06559	yezF	-3,254	1,140	0,111	323	1,179
BKK07010	yesS	-3,259	1,140	0,121	322	1,179
BKK14569	ykzV	-3,26	1,141	0,108	517	1,179
BKK17340	ymaH	-3,269	1,125	0,100	83	1,164
BKK17780	yndG	-3,27	1,125	0,105	111	1,164
BKK30610	ytlC	-3,287	1,182	0,097	538	1,222
BKK14870	ctaA	-3,294	1,140	0,100	700	1,179
BKK05970	rex	-3,303	1,119	0,082	66	1,158
BKK10660	yisB	-3,316	1,144	0,111	81	1,183
BKK32040	yuiF	-3,321	1,110	0,106	107	1,148
BKK11320	yjzB	-3,327	1,144	0,115	105	1,183
BKK06650	sapB	-3,332	1,140	0,101	446	1,179
BKK26970	adhB	-3,333	1,113	0,113	266	1,151
BKK33560	yvaD	-3,335	1,127	0,112	111	1,166
BKK32580	frlM	-3,337	1,110	0,135	34	1,148
BKK09590	yhdT	-3,344	1,138	0,080	117	1,177
BKK05020	phrI	-3,351	1,109	0,099	471	1,147
BKK34300	epsH	-3,353	1,111	0,099	194	1,149
BKK06870	yesE	-3,354	1,139	0,095	217	1,179
BKK21470	sunT	-3,362	1,116	0,106	467	1,155
BKK10200	yhfE	-3,37	1,119	0,098	274	1,158
BKK20190	yosA	-3,372	1,091	0,129	93	1,129
BKK33280	yvrP	-3,372	1,127	0,100	87	1,166
BKK13710	ykvI	-3,373	1,127	0,095	412	1,166
BKK18979	yoyA	-3,38	1,123	0,105	426	1,162
BKK22980	ypbG	-3,381	1,114	0,101	575	1,153
BKK39400	pdp	-3,384	1,125	0,092	89	1,165
BKK09150	yhcN	-3,386	1,137	0,103	110	1,177
BKK10930	yitB	-3,389	1,143	0,117	93	1,183
BKK25790	arsB	-3,389	0,942	0,067	55	0,975
BKK33330	lysP	-3,395	1,096	0,103	76	1,134
BKK17267	ymzE	-3,396	1,124	0,105	209	1,164
BKK26500	yrkI	-3,397	1,114	0,105	511	1,153
BKK03250	ycgR	-3,398	1,092	0,106	46	1,130
BKK21700	ypoP	-3,398	1,114	0,113	383	1,153
BKK30900	cotS	-3,401	1,117	0,109	124	1,156
BKK28890	yscB	-3,405	1,180	0,075	73	1,222
BKK03600	tcyB	-3,418	1,160	0,074	478	1,201
BKK11730	cotO	-3,419	1,144	0,087	86	1,184
BKK11460	oppD	-3,422	1,167	0,138	139	1,208
BKK09450	yhdF	-3,424	1,137	0,090	185	1,177
BKK04250	lrpC	-3,436	1,125	0,106	439	1,165
BKK11140	yitU	-3,455	1,142	0,113	73	1,183
BKK23780	yqjQ	-3,467	1,146	0,097	190	1,187
BKK05050	lrpA	-3,473	1,127	0,144	150	1,167
BKK36960	ywlB	-3,477	1,076	0,112	120	1,114
BKK05290	ycbP	-3,478	1,091	0,094	50	1,130
BKK07650	yflK	-3,479	0,941	0,063	173	0,975
BKK11960	yjcR	-3,489	1,148	0,130	929	1,189
BKK38470	ywaD	-3,499	1,080	0,109	176	1,119
BKK32610	frlB	-3,5	1,108	0,106	42	1,148
BKK41010	trmF	-3,505	1,143	0,084	169	1,184
BKK28350	ysnB	-3,509	1,119	0,091	379	1,160
BKK19760	cgeD	-3,515	1,090	0,119	447	1,129
BKK25550	rpsT	-3,519	1,123	0,107	179	1,164
BKK26030	yqbO	-3,521	1,135	0,098	271	1,177
BKK20320	yorN	-3,526	1,089	0,138	92	1,129
BKK04670	rsbRA	-3,527	1,124	0,113	335	1,165
BKK39800	yxcD	-3,532	1,124	0,094	35	1,165
BKK06660	opuE	-3,537	1,159	0,081	125	1,201
BKK03110	ycgH	-3,54	1,159	0,079	59	1,201
BKK21710	dinF	-3,54	1,113	0,091	278	1,153
BKK39430	deoR	-3,547	1,123	0,097	111	1,165
BKK17678	ynzJ	-3,551	1,122	0,114	97	1,164
BKK35010	nagA	-3,553	1,095	0,121	463	1,135
BKK03780	phrC	-3,557	1,107	0,095	545	1,147
BKK17970	cotM	-3,572	1,163	0,126	121	1,206
BKK26240	yqaO	-3,575	1,135	0,121	168	1,177
BKK30300	bceR	-3,578	1,115	0,092	136	1,156
BKK00800	yozB	-3,586	1,061	0,121	391	1,100
BKK08370	yfiR	-3,589	0,940	0,072	60	0,975
BKK36080	ywrF	-3,595	1,074	0,089	41	1,114
BKK10410	yhzC	-3,599	1,141	0,102	84	1,183
BKK09830	yhaX	-3,6	1,135	0,110	118	1,177
BKK07150	yetG	-3,602	1,136	0,119	289	1,179
BKK27180	yrhH	-3,605	1,112	0,102	274	1,153
BKK00710	hslO	-3,607	1,113	0,099	356	1,155
BKK27020	yraA	-3,608	1,134	0,098	379	1,177
BKK14480	abh	-3,61	1,121	0,102	535	1,163
BKK17260	aprX	-3,611	1,122	0,098	158	1,164
BKK07540	yfmA	-3,613	1,118	0,096	627	1,160
BKK28840	ysdA	-3,614	1,073	0,116	106	1,113
BKK39950	yxaJ	-3,614	1,123	0,101	114	1,165
BKK02080	yosL	-3,619	0,940	0,080	38	0,975
BKK29249	ytzI	-3,62	1,073	0,117	108	1,113
BKK05280	ydeO	-3,645	1,125	0,088	99	1,167
BKK23880	yqzI	-3,651	1,144	0,098	324	1,187
BKK02930	yceG	-3,654	1,106	0,100	616	1,147
BKK19749	yoyG	-3,667	1,088	0,152	196	1,129
BKK36250	ptkA	-3,668	1,107	0,105	358	1,149
BKK19720	yodR	-3,67	1,088	0,093	404	1,129
BKK32150	paiA	-3,672	1,106	0,100	272	1,148
BKK13850	zosA	-3,676	1,162	0,141	155	1,206
BKK37370	albA	-3,678	1,078	0,095	136	1,119
BKK33760	sdpB	-3,694	1,123	0,098	97	1,166
BKK37930	ywdK	-3,709	1,107	0,090	210	1,149
BKK14460	ykpC	-3,721	1,135	0,136	765	1,179
BKK24410	spoVAD	-3,724	1,143	0,096	457	1,187
BKK34310	epsG	-3,724	1,093	0,119	56	1,135
BKK21580	yokI	-3,73	1,112	0,091	566	1,155
BKK24320	nusB	-3,732	0,939	0,070	33	0,975
BKK37870	spsE	-3,735	1,107	0,098	536	1,149
BKK36590	clsA	-3,737	1,073	0,122	124	1,114
BKK22270	yutH	-3,738	1,105	0,106	90	1,148
BKK23590	ansR	-3,751	1,143	0,095	227	1,187
BKK12660	xkdM	-3,759	1,144	0,132	612	1,189
BKK38280	efeU	-3,772	1,077	0,104	58	1,119
BKK04750	ydcF	-3,778	1,121	0,136	564	1,165
BKK32880	yusP	-3,779	1,176	0,100	100	1,222
BKK09940	yhaL	-3,779	1,132	0,097	159	1,177
BKK00860	clpC	-3,799	1,058	0,117	107	1,100
BKK18700	yoaQ	-3,803	1,124	0,106	271	1,168
BKK26960	yraF	-3,807	1,132	0,108	202	1,177
BKK11250	argF	-3,819	1,138	0,104	86	1,183
BKK40450	yycD	-3,82	1,122	0,096	163	1,166
BKK02810	cwlK	-3,825	0,938	0,082	36	0,975
BKK36520	glnK	-3,83	1,072	0,132	124	1,114
BKK10650	sbcC	-3,832	1,138	0,096	97	1,183
BKK15310	spoIIGA	-3,832	1,160	0,124	93	1,206
BKK11480	sunA	-3,836	1,111	0,108	268	1,155
BKK18899	yozX	-3,839	1,123	0,110	308	1,168
BKK27170	dacA	-3,839	1,112	0,102	83	1,179
BKK15550	pyrR	-3,84	1,096	0,098	60	1,139
BKK14660	ykzI	-3,854	1,134	0,129	292	1,179
BKK00090	guaB	-3,863	1,133	0,135	122	1,179
BKK35180	csbA	-3,866	1,091	0,106	540	1,135
BKK37540	ywhB	-3,866	1,075	0,114	68	1,119
BKK21600	yokB	-3,871	1,110	0,108	369	1,155
BKK28830	ysdB	-3,876	1,070	0,094	74	1,113
BKK19840	yotL	-3,883	1,085	0,134	281	1,129
BKK02030	ydbM	-3,888	1,086	0,090	31	1,130
BKK03420	nin	-3,897	1,086	0,104	116	1,130
BKK23300	ypuD	-3,899	1,141	0,100	84	1,187
BKK15740	rsmB	-3,92	1,090	0,099	114	1,134
BKK14250	yknT	-3,924	1,133	0,114	285	1,179
BKK11055	yitO	-3,936	1,137	0,110	90	1,183
BKK35550	tuaG	-3,947	1,104	0,093	553	1,149
BKK13440	ykoY	-3,951	1,117	0,107	228	1,163
BKK37630	ywfM	-3,953	1,104	0,104	290	1,149
BKK21830	ypjQ	-3,954	1,112	0,097	125	1,158
BKK26200	yqaS	-3,957	1,130	0,107	234	1,177
BKK13320	ykzB	-3,975	1,120	0,092	92	1,166
BKK13789	ykzQ	-3,975	1,159	0,133	121	1,206
BKK24990	pstS	-3,978	1,158	0,156	108	1,206
BKK24360	spoIIIAH	-3,997	1,158	0,136	37	1,206
BKK16340	fliZ	-4,004	1,115	0,095	163	1,162
BKK16710	fadG	-4,011	1,116	0,099	391	1,163
BKK40760	yyaP	-4,018	1,137	0,098	178	1,184
BKK04790	ydcK	-4,026	1,118	0,146	53	1,165
BKK03010	amhX	-4,043	1,084	0,074	32	1,130
BKK17559	ynzI	-4,045	1,115	0,093	526	1,162
BKK35620	lytC	-4,049	1,114	0,103	339	1,161
BKK13540	ogt	-4,073	1,119	0,105	204	1,166
BKK04760	ydcG	-4,077	1,117	0,152	168	1,165
BKK04950	yddF	-4,078	1,120	0,120	192	1,167
BKK25170	yqfO	-4,091	1,116	0,097	313	1,164
BKK16610	ylxR	-4,097	1,120	0,116	327	1,168
BKK34490	yvdS	-4,098	1,089	0,113	297	1,135
BKK37120	fbaA	-4,113	1,073	0,088	41	1,119
BKK10400	yhxC	-4,116	1,135	0,095	95	1,183
BKK26850	yrpG	-4,125	1,128	0,100	173	1,177
BKK15360	ylmC	-4,127	1,130	0,156	58	1,179
BKK36360	mscL	-4,129	1,113	0,099	373	1,161
BKK04430	ydbD	-4,132	1,112	0,098	537	1,160
BKK09030	yhcC	-4,143	1,128	0,101	105	1,177
BKK20560	yoqO	-4,143	1,083	0,119	63	1,129
BKK00360	yaaT	-4,151	1,130	0,121	206	1,179
BKK27360	yrrM	-4,152	1,111	0,095	162	1,160
BKK11310	comZ	-4,157	1,134	0,126	120	1,183
BKK20240	yorV	-4,158	1,082	0,152	32	1,129
BKK14020	ykyB	-4,169	1,118	0,115	263	1,166
BKK36970	spoIIR	-4,174	1,101	0,097	368	1,149
BKK28420	ysmA	-4,186	0,934	0,072	45	0,975
BKK10220	gltT	-4,205	1,127	0,108	206	1,177
BKK19260	yocM	-4,211	1,122	0,115	39	1,172
BKK10290	yhfN	-4,215	1,133	0,099	135	1,183
BKK24390	spoVAD	-4,219	1,156	0,152	36	1,206
BKK04330	mutT	-4,22	1,116	0,137	405	1,165
BKK14400	yknW	-4,227	1,129	0,124	701	1,179
BKK19910	yotE	-4,23	1,082	0,126	111	1,129
BKK15010	ylbH	-4,237	1,129	0,093	295	1,179
BKK37050	maeA	-4,238	1,067	0,145	60	1,114
BKK25410	rpsU	-4,24	1,114	0,102	277	1,164
BKK37600	ywfO	-4,243	1,071	0,116	355	1,119
BKK34810	yvcA	-4,246	1,087	0,120	38	1,135
BKK05830	gmuC	-4,253	1,118	0,098	150	1,167
BKK35319	yvcK	-4,258	1,117	0,104	141	1,166
BKK16710	mlpA	-4,265	1,114	0,105	137	1,164
BKK09300	yqjN	-4,267	1,127	0,096	306	1,177
BKK25710	cwlH	-4,267	1,114	0,108	127	1,164
BKK18210	yngE	-4,273	1,122	0,083	93	1,172
BKK01560	kbaA	-4,282	1,126	0,100	278	1,177
BKK09290	yabC	-4,282	1,127	0,095	239	1,177
BKK20430	yorC	-4,284	1,081	0,134	168	1,129
BKK16560	rseP	-4,286	1,112	0,095	344	1,161
BKK20550	yoqP	-4,289	1,081	0,113	131	1,129
BKK23920	yqjD	-4,3	1,129	0,094	539	1,187
BKK18819	yozU	-4,31	1,121	0,121	242	1,172
BKK31720	degQ	-4,329	1,111	0,098	262	1,161
BKK16320	fliY	-4,343	1,117	0,102	233	1,168
BKK05960	moaC	-4,346	1,116	0,143	429	1,167
BKK22370	aspB	-4,347	1,052	0,113	246	1,100
BKK09410	phoA	-4,354	1,126	0,096	124	1,177
BKK38750	cydB	-4,356	1,115	0,093	133	1,166
BKK17560	ynaI	-4,358	1,113	0,098	106	1,164
BKK20660	yogE	-4,36	1,111	0,108	325	1,162
BKK34960	yvoF	-4,365	1,086	0,132	229	1,135
BKK05099	yvoG	-4,366	1,116	0,108	265	1,167
BKK38700	sigY	-4,369	1,114	0,126	108	1,165
BKK12430	rapA	-4,37	1,137	0,122	337	1,189
BKK37880	spsD	-4,37	1,070	0,088	48	1,119
BKK16460	ytkL	-4,375	1,117	0,097	396	1,168
BKK24130	prpD	-4,381	1,135	0,104	362	1,187
BKK23690	yqjY	-4,394	1,133	0,091	466	1,187
BKK14050	ykuE	-4,402	1,115	0,087	38	1,166
BKK11920	yjcN	-4,404	1,137	0,112	461	1,189
BKK11990	yjdB	-4,405	0,932	0,069	59	0,975
BKK22240	ypqA	-4,438	1,100	0,108	384	1,151
BKK14800	ylaJ	-4,439	1,153	0,128	77	1,206
BKK30290	ymzC	-4,441	1,168	0,100	39	1,222
BKK29030	ytcD	-4,444	1,089	0,087	120	1,139
BKK32430	ucaI	-4,444	1,097	0,125	117	1,148
BKK04300	ydaM	-4,447	1,113	0,116	664	1,165
BKK18370	yoeA	-4,451	1,153	0,128	99	1,206
BKK22470	ypjG	-4,455	1,051	0,124	189	1,100
BKK36710	fadH	-4,455	1,065	0,100	67	1,114
BKK36260	ywqC	-4,467	1,098	0,099	537	1,149
BKK05010	rapI	-4,478	1,096	0,098	520	1,147
BKK25310	dgkA	-4,491	1,111	0,099	473	1,164
BKK11430	oppA	-4,499	1,131	0,091	184	1,184
BKK36030	ywrK	-4,499	1,109	0,094	476	1,161
BKK11030	yitL	-4,506	1,130	0,105	101	1,183
BKK01910	skfA	-4,507	1,063	0,112	125	1,113
BKK21500	uvrX	-4,507	1,099	0,087	250	1,151
BKK36190	ywqJ	-4,507	1,064	0,100	195	1,114
BKK34540	clpP	-4,517	0,931	0,054	36	0,975
BKK26470	yrkL	-4,523	1,123	0,100	311	1,177
BKK38210	ywcD	-4,527	1,113	0,104	138	1,166
BKK36470	pucI	-4,534	1,064	0,108	62	1,114
BKK33790	sdpR	-4,55	1,113	0,091	226	1,166
BKK05860	gmuE	-4,552	1,114	0,088	305	1,167
BKK10850	yisS	-4,558	1,129	0,109	73	1,183
BKK25970	yqcA	-4,558	0,931	0,068	64	0,975
BKK34700	yvcR	-4,564	1,097	0,098	406	1,149
BKK09270	glpP	-4,568	1,123	0,107	161	1,177
BKK01030	rplA	-4,57	1,110	0,095	223	1,164
BKK23450	sigF	-4,571	1,133	0,090	301	1,187
BKK27630	yrvD	-4,583	1,106	0,096	377	1,160
BKK06950	yesM	-4,585	1,146	0,096	151	1,201
BKK29470	ackA	-4,595	1,062	0,133	35	1,113
BKK02100	cypC	-4,597	1,078	0,117	37	1,130
BKK03370	yckA	-4,601	1,078	0,094	106	1,130
BKK35380	fliW	-4,612	1,083	0,113	531	1,135
BKK23770	dsdA	-4,62	1,132	0,095	216	1,187
BKK19310	dhaS	-4,642	1,117	0,092	136	1,172
BKK40670	yybE	-4,648	1,129	0,094	248	1,184
BKK16150	clpQ	-4,649	1,106	0,101	470	1,160
BKK05220	ydeJ	-4,679	1,113	0,145	91	1,167
BKK38190	galT	-4,685	1,066	0,107	79	1,119
BKK36330	ywpF	-4,692	1,107	0,101	437	1,161
BKK19130	yocA	-4,707	1,116	0,105	99	1,172
BKK23480	dacF	-4,71	1,150	0,165	58	1,206
BKK29120	mdh	-4,727	1,060	0,088	105	1,113
BKK12410	yjoA	-4,729	1,133	0,143	638	1,189
BKK32260	hom	-4,73	1,086	0,079	69	1,139
BKK11150	yitV	-4,739	1,127	0,105	113	1,183
BKK34670	yvdA	-4,739	1,081	0,111	207	1,135
BKK04530	ydbN	-4,74	1,110	0,132	519	1,164
BKK34640	yvdD	-4,745	1,081	0,141	94	1,135
BKK24120	prpB	-4,757	1,131	0,098	252	1,187
BKK34890	hisH	-4,766	1,081	0,123	256	1,135
BKK00650	yabS	-4,774	1,075	0,096	189	1,129
BKK09180	yhcQ	-4,784	1,121	0,096	245	1,177
BKK41000	rsmG	-4,785	1,128	0,101	132	1,184
BKK37820	spsK	-4,787	1,094	0,095	637	1,149
BKK37950	ywdI	-4,791	1,065	0,104	135	1,119
BKK12690	xkdP	-4,799	1,112	0,109	284	1,168
BKK14620	slp	-4,824	1,148	0,097	30	1,206
BKK23490	pupG	-4,83	1,130	0,095	606	1,187
BKK00200	yaaK	-4,842	1,120	0,098	112	1,167
BKK37890	spsC	-4,844	1,122	0,106	420	1,149
BKK07240	yetN	-4,848	1,122	0,106	358	1,179
BKK12749	ykzA	-4,85	1,110	0,111	304	1,166
BKK10050	ecsB	-4,852	1,120	0,097	133	1,177
BKK35610	tuaA	-4,853	1,094	0,109	551	1,149
BKK00810	senS	-4,855	1,103	0,093	320	1,159
BKK09920	yhaN	-4,864	1,120	0,104	87	1,177
BKK13910	ptsI	-4,878	1,109	0,141	37	1,166
BKK37100	murAB	-4,883	1,093	0,107	138	1,149
BKK10820	yisQ	-4,889	1,082	0,099	56	1,138
BKK28000	minC	-4,891	1,103	0,092	148	1,160
BKK32850	fadM	-4,908	1,079	0,092	71	1,134
BKK03990	mtlD	-4,91	1,108	0,104	181	1,165
BKK36668	ywzE	-4,915	1,060	0,103	164	1,114
BKK12160	yjgC	-4,947	1,130	0,130	416	1,189
BKK33540	yvaB	-4,95	0,927	0,077	67	0,975
BKK20140	yosF	-4,958	1,073	0,145	34	1,129
BKK19660	yozD	-4,959	1,073	0,138	455	1,129
BKK32730	metN	-4,964	1,161	0,124	157	1,222
BKK11839	yjzE	-4,985	1,130	0,103	859	1,189
BKK04240	pucI	-4,989	1,107	0,105	284	1,165
BKK09800	yheA	-4,993	1,118	0,103	137	1,177
BKK18910	rapK	-5,013	1,113	0,103	123	1,172
BKK19210	yocH	-5,019	1,146	0,139	59	1,206
BKK22200	cotD	-5,019	1,045	0,142	141	1,100
BKK12860	ykbA	-5,021	1,108	0,126	173	1,166
BKK33020	cssS	-5,025	1,108	0,098	126	1,166

Sup. Table 3: Cell width of mutants of the BKK collection (continued)

BKK name[1]	gene	screening delta[2] (%)	erage width (μ	+/-	nb	ADP
BKK35490	degU	-5,028	1,103	0,094	353	1,161
BKK14220	ykuU	-5,047	1,120	0,137	196	1,179
BKK19770	cgeC	-5,049	1,072	0,116	409	1,129
BKK00640	spoIIE	-5,063	1,072	0,165	177	1,129
BKK32230	yuxL	-5,063	1,082	0,082	98	1,139
BKK05210	ydeI	-5,069	1,108	0,096	109	1,167
BKK28370	rph	-5,069	1,099	0,101	155	1,158
BKK39720	iolE	-5,075	1,106	0,118	77	1,165
BKK21850	ypiP	-5,086	1,096	0,100	509	1,155
BKK13270	ykoI	-5,104	1,107	0,112	287	1,166
BKK04080	kipI	-5,122	1,089	0,094	445	1,147
BKK07110	lplB	-5,123	1,118	0,096	317	1,179
BKK04360	mntH	-5,141	1,100	0,112	447	1,160
BKK29090	polA	-5,16	1,056	0,129	60	1,113
BKK37250	narI	-5,167	1,061	0,096	52	1,119
BKK34265	epsK	-5,179	1,076	0,097	424	1,135
BKK23760	coaA	-5,185	1,126	0,092	230	1,187
BKK05900	thiL	-5,19	1,107	0,121	370	1,167
BKK05980	tatAY	-5,201	1,088	0,096	277	1,147
BKK36210	ywqH	-5,201	1,056	0,105	299	1,114
BKK36290	ywpJ	-5,219	1,056	0,121	169	1,114
BKK38290	thiE	-5,222	1,060	0,102	69	1,119
BKK13350	ykoN	-5,223	1,105	0,103	305	1,166
BKK06100	ydiS	-5,237	1,106	0,112	395	1,167
BKK00740	pabB	-5,238	1,043	0,145	55	1,100
BKK14320	yknU	-5,243	1,143	0,143	136	1,206
BKK28130	hemB	-5,281	1,098	0,126	56	1,160
BKK37080	rho	-5,283	1,060	0,114	150	1,119
BKK24040	bkdAB	-5,286	1,124	0,096	57	1,187
BKK16970	ymdB	-5,293	1,102	0,099	95	1,164
BKK36780	ywzB	-5,303	1,055	0,123	105	1,114
BKK38690	yxlC	-5,32	1,138	0,098	184	1,201
BKK32130	guaC	-5,331	1,087	0,117	176	1,148
BKK18600	yozQ	-5,334	1,077	0,106	38	1,138
BKK31945	yukJ	-5,337	1,079	0,103	69	1,139
BKK37470	phrF	-5,346	1,088	0,098	284	1,149
BKK28730	araQ	-5,352	1,078	0,106	33	1,139
BKK00850	mcsB	-5,355	1,041	0,117	384	1,100
BKK03080	ycgE	-5,36	1,069	0,115	43	1,130
BKK13890	ptsG	-5,372	1,125	0,113	244	1,189
BKK36500	ywoB	-5,372	1,054	0,112	54	1,114
BKK22690	aroH	-5,383	1,041	0,110	175	1,100
BKK36750	spoIID	-5,389	1,054	0,112	82	1,114
BKK12120	yjfB	-5,402	1,125	0,150	170	1,189
BKK35850	ywtE	-5,418	1,054	0,109	232	1,114
BKK32750	metQ	-5,42	1,156	0,076	124	1,222
BKK34500	yvdR	-5,445	1,103	0,089	116	1,166
BKK31770	yueI	-5,453	1,085	0,111	190	1,148
BKK24810	yqgV	-5,456	1,120	0,085	254	1,184
BKK38460	tyrZ	-5,457	1,058	0,120	209	1,119
BKK13360	ykoP	-5,467	1,103	0,103	133	1,166
BKK06090	ydiR	-5,478	1,103	0,116	63	1,167
BKK16900	ymfK	-5,503	1,104	0,095	484	1,168
BKK31120	lytG	-5,51	1,155	0,114	35	1,222
BKK24770	mgsR	-5,511	1,099	0,120	148	1,164
BKK14940	ylbA	-5,527	1,114	0,085	267	1,179
BKK20370	yorI	-5,528	1,067	0,130	64	1,129
BKK11520	mecA	-5,533	1,118	0,103	81	1,183
BKK19950	sspC	-5,539	1,067	0,150	182	1,129
BKK03760	yclK	-5,543	1,135	0,108	308	1,201
BKK21730	ypmS	-5,56	1,091	0,100	321	1,155
BKK18790	yoaZ	-5,564	1,106	0,126	34	1,172
BKK23040	fer	-5,575	1,039	0,127	159	1,100
BKK12540	xkdD	-5,596	1,122	0,146	699	1,189
BKK36480	ywoD	-5,598	1,052	0,104	113	1,114
BKK35000	hprK	-5,65	1,071	0,123	123	1,135
BKK35870	pgsE	-5,661	1,051	0,132	83	1,114
BKK40510	yybT	-5,677	1,100	0,103	128	1,166
BKK11040	yitM	-5,698	1,116	0,103	82	1,183
BKK19579	yoyD	-5,717	1,065	0,104	462	1,129
BKK13080	ykkB	-5,729	1,100	0,122	341	1,166
BKK19730	yodS	-5,735	1,065	0,085	361	1,129
BKK13740	queE	-5,743	1,099	0,117	417	1,166
BKK09930	yhaM	-5,753	1,109	0,088	144	1,177
BKK05930	riml	-5,775	1,100	0,114	469	1,167
BKK15340	ylmA	-5,795	1,137	0,138	72	1,206
BKK00150	dgk	-5,807	1,110	0,110	188	1,179
BKK18650	pelB	-5,827	1,103	0,123	37	1,172
BKK37860	spsF	-5,842	1,053	0,133	51	1,119

BKK name[1]	gene	screening delta[2] (%)	erage width (μ	+/-	nb	ADP
BKK37570	mmr	-5,904	1,082	0,096	313	1,149
BKK12370	exuR	-5,917	1,119	0,124	591	1,189
BKK36770	ywmB	-5,935	1,048	0,115	118	1,114
BKK36680	ywmF	-6,051	1,047	0,104	38	1,114
BKK09050	yhcE	-6,061	1,106	0,098	147	1,177
BKK15420	divIVA	-6,061	0,916	0,082	59	0,975
BKK14610	pdhD	-6,065	1,108	0,135	50	1,179
BKK37690	bacF	-6,105	1,050	0,091	51	1,119
BKK15580	cysP	-6,125	1,078	0,113	180	1,148
BKK21840	ypjP	-6,132	1,084	0,092	301	1,155
BKK22230	ypqE	-6,15	1,033	0,094	68	1,100
BKK28440	sdhA	-6,151	0,915	0,055	47	0,975
BKK35830	ywtG	-6,225	1,078	0,098	291	1,149
BKK40620	yybJ	-6,234	1,094	0,142	88	1,167
BKK28770	araL	-6,238	1,044	0,104	85	1,113
BKK22320	ponA	-6,259	1,081	0,128	394	1,153
BKK14629	ykzW	-6,274	1,105	0,123	561	1,179
BKK18709	yozT	-6,295	1,095	0,097	377	1,168
BKK20510	yoqU	-6,361	1,057	0,110	64	1,129
BKK18909	yozZ	-6,405	1,097	0,128	169	1,172
BKK19820	yotN	-6,465	1,056	0,100	822	1,129
BKK14950	ylbB	-6,466	1,103	0,092	122	1,179
BKK34980	yvoD	-6,499	1,061	0,109	241	1,135
BKK00470	purR	-6,679	1,086	0,103	97	1,164
BKK34900	hisB	-6,697	1,059	0,132	224	1,135
BKK12330	yjmD	-6,79	1,108	0,132	661	1,189
BKK18480	proH	-6,817	1,092	0,113	162	1,172
BKK38780	yxkl	-6,825	1,085	0,104	133	1,165
BKK19280	yocN	-6,839	1,091	0,131	344	1,172
BKK36300	glcR	-6,844	1,038	0,122	159	1,114
BKK29550	ytcJ	-6,86	1,037	0,127	55	1,113
BKK22510	ypjC	-6,867	1,025	0,117	53	1,100
BKK36900	glyA	-6,868	0,908	0,093	50	0,975
BKK20520	yoqT	-6,872	1,052	0,130	198	1,129
BKK22520	ypjB	-6,873	1,025	0,127	101	1,100
BKK13900	ptsH	-6,914	1,107	0,116	41	1,189
BKK04710	rsbV	-6,939	1,084	0,114	273	1,165
BKK23130	resC	-6,943	1,024	0,116	207	1,100
BKK16070	ylqG	-6,983	1,023	0,148	134	1,100
BKK13210	thiX	-7,014	1,085	0,142	40	1,166
BKK22820	yphE	-7,032	1,023	0,103	46	1,100
BKK04510	ydbL	-7,047	1,083	0,126	663	1,165
BKK20290	yorQ	-7,086	1,049	0,132	74	1,129
BKK35370	csrA	-7,091	1,055	0,099	132	1,135
BKK06110	ydjA	-7,15	1,084	0,102	369	1,167
BKK33360	yvgJ	-7,2	1,134	0,104	215	1,222
BKK29640	ytsP	-7,208	1,033	0,094	123	1,113
BKK28760	araM	-7,221	1,033	0,100	80	1,113
BKK36460	ywoF	-7,232	1,034	0,090	30	1,114
BKK20270	yorS	-7,257	1,047	0,136	40	1,129
BKK28120	hemL	-7,261	1,075	0,093	185	1,160
BKK22390	ypmA	-7,278	1,020	0,133	164	1,100
BKK37440	ywhL	-7,291	1,037	0,104	101	1,119
BKK01920	skfB	-7,317	1,032	0,098	110	1,13
BKK29350	tcyM	-7,331	1,055	0,089	48	1,138
BKK10980	yitG	-7,353	1,119	0,116	43	1,208
BKK05680	ydgK	-7,423	1,112	0,082	220	1,201
BKK20260	yorT	-7,443	1,045	0,130	38	1,129
BKK29440	argH	-7,459	1,030	0,111	30	1,113
BKK36430	usd	-7,477	1,031	0,092	84	1,114
BKK11770	cotW	-7,497	1,100	0,099	279	1,189
BKK37510	pbpG	-7,549	1,063	0,096	228	1,149
BKK18730	yoaS	-7,576	1,083	0,120	71	1,172
BKK14230	ykuV	-7,642	1,089	0,123	156	1,179
BKK00870	radA	-7,706	1,015	0,122	45	1,100
BKK14920	ctaF	-7,782	1,087	0,085	361	1,179
BKK37640	ywfL	-7,8	0,899	0,073	40	0,975
BKK40490	yycA	-7,876	1,073	0,118	255	1,165
BKK15600	cysC	-7,935	1,057	0,118	187	1,148
BKK32740	metP	-7,952	1,074	0,096	103	1,166
BKK35270	cccB	-8,013	1,044	0,129	273	1,135
BKK19290	yozO	-8,249	1,075	0,139	549	1,172
BKK18760	yoaU	-8,275	1,075	0,115	81	1,172
BKK11720	fabI	-8,301	1,090	0,127	90	1,189
BKK26710	azlC	-8,397	1,054	0,122	257	1,151
BKK05850	gmuR	-8,408	1,069	0,090	154	1,167
BKK22480	mgsA	-8,445	1,007	0,132	95	1,100
BKK36690	rapB	-8,446	1,020	0,110	88	1,114
BKK10140	hemY	-8,489	1,077	0,097	162	1,177

BKK name[1]	gene	screening delta[2] (%)	erage width (μ	+/-	nb	ADP
BKK38430	gspA	-8,529	1,023	0,108	128	1,119
BKK36910	ywlG	-8,643	1,018	0,105	199	1,114
BKK00630	yabR	-8,798	1,056	0,101	68	1,158
BKK28170	hemA	-8,839	1,057	0,124	323	1,160
BKK29180	**pyk**	**-8,872**	1,062	0,119	123	1,156
BKK15790	**rpe**	**-8,905**	1,034	0,115	35	1,135
BKK22210	**yprB**	**-8,945**	1,002	0,134	110	1,100
BKK13900	**ptsH**	**-8,969**	1,062	0,126	58	1,166
BKK19320	**sqhC**	**-9,157**	1,064	0,134	55	1,172
BKK34620	**mdxD**	**-9,39**	1,028	0,120	61	1,135
BKK03480	**srfAA**	**-9,413**	1,024	0,103	38	1,130
BKK20300	**yorP**	**-9,42**	1,023	0,145	52	1,129
BKK01889	**ybzH**	**-9,47**	0,883	0,054	42	0,975
BKK22340	**nth**	**-9,547**	0,995	0,104	85	1,100
BKK19020	**yobN**	**-9,585**	1,059	0,107	36	1,172
BKK38660	**yxlF**	**-9,702**	1,028	0,068	45	1,138
BKK06360	**guaA**	**-9,94**	0,878	0,055	50	0,975
BKK22220	**yprA**	**-10,181**	0,988	0,116	38	1,100
BKK22410	**panD**	**-10,738**	0,982	0,183	31	1,100
BKK19180	**des**	**-10,806**	1,045	0,088	87	1,172
BKK04310	**ydaN**	**-10,896**	1,038	0,123	218	1,165
BKK08680	**ygaC**	**-10,965**	1,032	0,117	185	1,159
BKK22360	**asnS**	**-12,67**	0,961	0,148	37	1,100
BKK40390	**walH**	**-13,873**	0,973	0,069	43	1,130
BKK00110	pdxS	n/a	n/a	n/a	n/a	n/a
BKK00690	ftsH	n/a	n/a	n/a	n/a	n/a
BKK01670	ybbE	n/a	n/a	n/a	n/a	n/a
BKK27260	mccA	n/a	n/a	n/a	n/a	n/a
BKK31290	yugT	n/a	n/a	n/a	n/a	n/a
BKK33920	tpiA	n/a	n/a	n/a	n/a	n/a
BKK33930	pgk	n/a	n/a	n/a	n/a	n/a
BKK38150	qoxC	n/a	n/a	n/a	n/a	n/a
BKK41030	jag	n/a	n/a	n/a	n/a	n/a

1: underscored and bold names indicate the top 10 % and 1% delta, respectively

2: n/a: 9 mutants have no values because they did not grow after several attempts

Sup. Table 4: Width of the 0.5% largest and thinnest selected strains

gene		screening step					checking step				post-backcross step					backcross strains without Mg[2+ 5]				
name	reference	width (µm)	+/-	delta[1] (%)	nb	ADP (µm)	width (µm)	+/-	delta[2] (%)	nb	width (µm)	+/-	delta[3] (%)	nb	P-value[4]	width (µm)	+/-	delta (%)	nb	P value[4]
cwlO	BKK34800	1,42	0,14	23,36	301	1,15	1,25	0,13	28,30	265	1,09	0,09	13,07	438	***	1,10	0,08	7,68	411	****
rodZ	BKK16910	1,29	0,11	11,40	287	1,16	1,12	0,10	14,77	199	1,08	0,08	12,55	331	****	1,07	0,09	5,52	370	**
rpe	BKK15790	1,03	0,12	-8,91	35	1,14	0,86	0,07	-11,39	136	1,07	0,18	11,02	207	****	1,09	0,11	6,96	357	****
ftsX	BKK35250	1,33	0,18	15,59	286	1,15	1,18	0,12	20,97	167	1,06	0,08	9,95	342	****	1,10	0,08	7,66	341	****
yaaA	BKK00030	1,28	0,12	9,94	189	1,17	1,02	0,07	4,87	188	1,06	0,09	9,72	349	***	1,13	0,09	11,38	294	****
ftsE	BKK35260	1,38	0,15	19,51	109	1,16	1,21	0,11	24,20	90	1,05	0,08	9,43	431	****	1,07	0,08	5,03	335	**
dacA	BKK00100	1,34	0,15	15,94	107	1,16	1,03	0,10	5,94	260	1,05	0,08	9,33	409	****	1,08	0,09	6,06	370	****
yuaC	BKK31070	1,29	0,13	11,33	238	1,16	0,99	0,08	2,04	150	1,02	0,08	5,48	405		1,01	0,07	-0,88	281	
natA	BKK02750	1,31	0,12	9,10	127	1,20	0,96	0,06	-0,86	129	1,00	0,07	4,03	331		1,03	0,09	1,63	90	
ymfD	BKK16825	1,31	0,17	12,67	82	1,16	0,93	0,07	-4,42	53	1,00	0,07	3,92	383		1,04	0,07	1,79	102	
comFC	BKK35450	1,24	0,10	11,47	90	1,11	1,03	0,06	6,07	238	1,00	0,07	3,64	351		1,06	0,07	3,60	97	
yvyF	BKK35440	1,22	0,11	9,58	161	1,11	0,99	0,07	2,07	131	1,00	0,07	3,30	367		1,02	0,08	-0,10	324	
greA	BKK27320	1,10	0,10	12,65	40	0,98	n/a	n/a	n/a	n/a	0,99	0,06	3,20	154		0,99	0,06	-2,32	154	
mdxD	BKK34620	1,03	0,12	-9,39	61	1,14	0,95	0,07	-2,70	174	0,99	0,07	3,08	405		0,99	0,07	-2,56	295	
yprB	BKK22210	1,00	0,13	-8,95	110	1,10	0,97	0,09	-0,22	127	0,99	0,08	3,06	180		0,99	0,07	-3,06	246	
srfAA	BKK03480	1,02	0,10	-9,41	38	1,13	0,92	0,08	-4,89	123	0,99	0,07	3,06	441		1,00	0,08	-1,46	287	
yobN	BKK19020	1,06	0,11	-9,59	36	1,17	0,98	0,09	1,31	124	0,99	0,08	3,03	324		0,97	0,07	-4,43	305	
yxlF	BKK38660	1,03	0,07	-9,70	45	1,14	0,97	0,06	-0,12	127	0,99	0,07	2,83	394		1,00	0,08	-1,91	324	
ypzH	BKK22849	1,28	0,13	15,99	401	1,10	1,02	0,08	4,54	78	0,99	0,08	2,73	356		0,99	0,07	-2,57	116	
ykhA	BKK13030	1,27	0,18	9,17	109	1,17	0,99	0,06	1,31	155	0,99	0,09	2,71	355		1,04	0,10	1,93	138	
yoqC [6]	BKK20680	1,32	0,18	13,23	411	1,17	0,94	0,07	-3,46	101	n/a	n/a	n/a	n/a		n/a	n/a	n/a	n/a	
yorP [6]	BKK20300	1,02	0,15	-9,42	52	1,13	1,01	0,07	3,90	189	n/a	n/a	n/a	n/a		n/a	n/a	n/a	n/a	
des	BKK19180	1,04	0,09	-10,81	87	1,17	0,98	0,07	0,72	165	0,99	0,09	2,47	424		1,02	0,07	0,29	204	
yprA	BKK22220	0,99	0,12	-10,18	38	1,10	0,99	0,08	2,14	107	0,99	0,07	2,46	276		0,99	0,07	-3,10	183	
xkdX	BKK12770	1,29	0,12	10,49	135	1,17	0,96	0,07	-0,86	224	0,98	0,07	2,04	283		1,02	0,08	-0,30	155	
yqiW	BKK23990	1,26	0,14	9,50	142	1,15	0,97	0,07	-0,31	237	0,98	0,07	1,61	324		0,97	0,08	-4,31	131	
nth	BKK22340	1,00	0,10	-9,55	85	1,10	1,00	0,08	2,44	160	0,98	0,07	1,33	267		0,99	0,08	-2,52	277	
ygaC	BKK08680	1,03	0,12	-10,97	185	1,16	0,94	0,07	-3,16	121	0,98	0,07	1,24	366		1,01	0,08	-0,55	236	
xkdW	BKK12760	1,42	0,12	21,84	150	1,17	0,99	0,08	2,14	79	0,97	0,07	1,14	316		0,98	0,06	-4,25	64	
sqhC	BKK19320	1,06	0,13	-9,16	55	1,17	0,97	0,07	-0,06	163	0,97	0,07	1,07	322		0,98	0,07	-3,88	270	
ydaN	BKK04310	1,04	0,12	-10,90	218	1,16	0,97	0,07	-0,04	119	0,97	0,06	0,74	376		1,01	0,08	-0,87	173	
kbl	BKK17000	1,28	0,16	9,83	152	1,16	0,96	0,07	-1,70	161	0,97	0,07	0,62	333		0,99	0,06	-2,71	75	
asnS	BKK22360	0,96	0,15	-12,67	37	1,10	0,97	0,07	-0,24	103	0,97	0,07	0,17	322		0,98	0,08	-3,34	100	
minJ	BKK35220	1,24	0,13	9,38	104	1,14	0,98	0,08	0,49	211	0,95	0,09	-1,02	677		0,97	0,11	-4,60	225	
walH	BKK40390	0,97	0,07	-13,87	43	1,13	0,89	0,06	-8,04	126	0,92	0,07	-4,54	366		0,92	0,07	-9,55	361	
pyk	BKK29180	1,06	0,12	-8,87	123	1,16	0,82	0,04	-15,67	105	0,88	0,06	-8,32	314	***	0,94	0,08	-7,47	289	****
ybzH	BKK01889	0,88	0,05	-9,47	42	0,98	n/a	n/a	n/a	n/a	0,87	0,06	-9,30	369	****	0,86	0,06	-15,19	115	****
panD	BKK22410	0,98	0,18	-10,74	31	1,10	0,82	0,05	-15,65	135	0,85	0,06	-11,39	301	****	0,87	0,06	-15,04	256	****
guaA	BKK06360	0,88	0,05	-9,94	50	0,98	n/a	n/a	n/a	n/a	0,84	0,05	-12,73	351	****	0,81	0,05	-20,58	182	****
ptsH	BKK13900	1,06	0,13	-8,97	58	1,17	0,83	0,06	-14,44	163	0,84	0,07	-13,30	392	****	0,91	0,06	-10,27	390	****

Text on right margin: confirmed positively affected · confirmed negatively

1: δ relative to ADP
2: δ relative to wild type width
3: δ relative to wild type strain, average of 3 independant replicates
4: *P*-value (summary) of nested t-tests, comparing the widths of mutants with that of the wild type cells   (**** = P<0.0001; *** = 0.0001<P<0.001; ** = 0.001<P<0.01; * = 0.01<P<0.05; ns = P>0.05)
5: width values for confirmed affected mutants are the average of 3 independant replicates
6: PCR checking revealed that the BKK strains are wt for the tested loci. Deletions were therefore not backcrossed into the 168 strain

Sup. Table 5. *B. subtilis* strains used in this study

Strain (*B. subtilis*)	Relevant genotype	Source or reference[1]
168	(wt)	Laboratory stock
PY79	(wt)	Laboratory stock
RCL413	Ω*neo3427* - Δ*mreB*	Billaudeau, 2019
PS2062	Δ*ponA*::*spc*	Popham, 1995
CcBs351	Δ*rodZ*::*cat*	Gibson assembly → 168
CcBs628	Δ*rodZ*::*cat*	CcBs351→ PY79
BKK00030	Δ*yaaA*::*km*	Koo, 2017
BKK00100	Δ*dacA*::*km*	Koo, 2017
BKK01889	Δ*ybzH*::*km*	Koo, 2017
BKK02750	Δ*natA*::*km*	Koo, 2017
BKK03480	Δ*srfAA*::*km*	Koo, 2017
BKK04310	Δ*ydaN*::*km*	Koo, 2017
BKK06360	Δ*guaA*::*km*	Koo, 2017
BKK08680	Δ*ygaC*::*km*	Koo, 2017
BKK12760	Δ*xkdW*::*km*	Koo, 2017
BKK12770	Δ*xkdX*::*km*	Koo, 2017
BKK13030	Δ*ykhA*::*km*	Koo, 2017
BKK13900	Δ*ptsH*::*km*	Koo, 2017
BKK15790	Δ*rpe*::*km*	Koo, 2017
BKK16825	Δ*ymfD*::*km*	Koo, 2017
BKK16910	Δ*rodZ*::*km*	Koo, 2017
BKK17000	Δ*kbl*::*km*	Koo, 2017
BKK19020	Δ*yobN*::*km*	Koo, 2017
BKK19180	Δ*des*::*km*	Koo, 2017
BKK19320	Δ*sqhC*::*km*	Koo, 2017
BKK20300	Δ*yorP*::*km*	Koo, 2017
BKK20680	Δ*yoqC*::*km*	Koo, 2017
BKK22210	Δ*yprB*::*km*	Koo, 2017
BKK22220	Δ*yprA*::*km*	Koo, 2017
BKK22340	Δ*nth*::*km*	Koo, 2017
BKK22360	Δ*asnS*::*km*	Koo, 2017
BKK22410	Δ*panD*::*km*	Koo, 2017
BKK22849	Δ*ypzH*::*km*	Koo, 2017
BKK23990	Δ*yqiw*::*km*	Koo, 2017
BKK27320	Δ*greA*::*km*	Koo, 2017
BKK29180	Δ*pyk*::*km*	Koo, 2017
BKK31070	Δ*yuaC*::*km*	Koo, 2017
BKK34620	Δ*mdxD*::*km*	Koo, 2017
BKK34800	Δ*cwlo*::*km*	Koo, 2017
BKK35220	Δ*minJ*::*km*	Koo, 2017
BKK35250	Δ*ftsX*::*km*	Koo, 2017
BKK35260	Δ*ftsE*::*km*	Koo, 2017
BKK35440	Δ*yvyF*::*km*	Koo, 2017
BKK35450	Δ*comFC*::*km*	Koo, 2017
BKK38660	Δ*yxlF*::*km*	Koo, 2017
BKK40390	Δ*walH*::*km*	Koo, 2017
RCL0820	Δ*cwlo*::*km*	BKK34800 DNA → 168
RCL0821	Δ*xkdW*::*km*	BKK12760 DNA → 168
RCL0822	Δ*ftsE*::*km*	BKK35260 DNA → 168
RCL0823	Δ*ypzH*::*km*	BKK22849 DNA → 168
RCL0824	Δ*dacA*::*km*	BKK00100 DNA → 168
RCL0825	Δ*ftsX*::*km*	BKK35250 DNA → 168
RCL0826	Δ*ymfD*::*km*	BKK16825 DNA → 168
RCL0827	Δ*comFC*::*km*	BKK35450 DNA → 168
RCL0828	Δ*rodZ*::*km*	BKK16910 DNA → 168
RCL0829	Δ*yuaC*::*km*	BKK31070 DNA → 168
RCL0830	Δ*xkdX*::*km*	BKK12770 DNA → 168
RCL0831	Δ*yaaA*::*km*	BKK00030 DNA → 168
RCL0832	Δ*kbl*::*km*	BKK17000 DNA → 168
RCL0833	Δ*yvyF*::*km*	BKK35440 DNA → 168

RCL0834	Δ*minJ* ::*km*	BKK35220 DNA → 168
RCL0835	Δ*yqiw* ::*km*	BKK23990 DNA → 168
RCL0837	Δ*ykhA* ::*km*	BKK13030 DNA → 168
RCL0838	Δ*natA* ::*km*	BKK02750 DNA → 168
RCL0839	Δ*ywqA* ::*km*	BKK36280 DNA → 168
RCL0840	Δ*walH* ::*km*	BKK40390 DNA → 168
RCL0841	Δ*asnS* ::*km*	BKK22360 DNA → 168
RCL0842	Δ*ygaC* ::*km*	BKK08680 DNA → 168
RCL0843	Δ*ydaN* ::*km*	BKK04310 DNA → 168
RCL0844	Δ*des* ::*km*	BKK19180 DNA → 168
RCL0845	Δ*panD* ::*km*	BKK22410 DNA → 168
RCL0846	Δ*yprA* ::*km*	BKK22220 DNA → 168
RCL0848	Δ*yobN* ::*km*	BKK19020 DNA → 168
RCL0849	Δ*nth* ::*km*	BKK22340 DNA → 168
RCL0850	Δ*srfAA* ::*km*	BKK03480 DNA → 168
RCL0851	Δ*mdxD* ::*km*	BKK34620 DNA → 168
RCL0852	Δ*sqhC* ::*km*	BKK19320 DNA → 168
RCL0853	Δ*pyk* ::*km*	BKK29180 DNA → 168
RCL0854	Δ*ptsH* ::*km*	BKK13900 DNA → 168
RCL0855	Δ*yprB* ::*km*	BKK22210 DNA → 168
RCL0856	Δ*rpe* ::*km*	BKK15790 DNA → 168
RCL0858	Δ*greA* ::*km*	BKK27320 DNA → 168
RCL0859	Δ*ybzH* ::*km*	BKK01889 DNA → 168
RCL0860	Δ*guaA* ::*km*	BKK06360 DNA → 168

1: Arrows indicate construction by transformation with chromosomal DNA

Sup. Table 6. Genes reported to affect cell width in *B. subtilis*

gene	fonction	essentiality	effect[1]	references [2]	this study[3]
CW homeostasis					
mreB	rod complex regulation	no	+	Jones *et al.,* 01	na
mbl	rod complex regulation	no	+	Abhayawardhane *et al.,* 95, Jones *et al.,* 01	na
mreBH	rod complex regulation	no	-	Carballido-López *et al.* , 06; Sassine *et al.* , 20	no
mreB+mbl+mreBH	rod complex regulation	yes[4]	R	Schirmer *et al.* , 09	na
mreC	rod complex transpeptidase regulator	no	R	Lee *et al.,* 03; Leaver *et al.,* 05	na
mreD	rod complex, unknown function	yes	R	Leaver *et al.,* 05	na
pbpA+pbpH	rod complex transpeptidases (bPBPs)	yes[4]	R	Wei *et al.,* 03	na
rodA	rod complex transglycosylase	yes	R	Henriques *et al.,* 98	na
rodZ	rod complex, unknown function	?[5]	+	Muchova *et al.,* 13; Van Beilen *et al.* , 16	yes
murB	PG precursor synthetic pathway	yes	+	Peters *et al.,* 16	na
ponA	transglycosylase/transpeptidase (aPBP)	no	-	Popham *et al.,* 96; Claessen *et al.,* 08	yes
lytE	autolysin	no	?[6]	Dominguez-Cuevas *et al.,* 13; Carballido-López *et al.* , 06; Sassine *et al.* 20	no
cwlO	autolysin	no	+	Dominguez-Cuevas *et al.,* 13; Meisner *et al.,* 13	yes
ftsE	ABC transporter, activator of CwlO	no	+	Dominguez-Cuevas *et al.,* 13; Meisner *et al.,* 13	yes
ftsX	ABC transporter, activator of CwlO	no	+	Dominguez-Cuevas *et al.,* 13; Meisner *et al.,* 13	yes
glmR (yvcK)	stimulates PG precursor synthetic pathway	yes[7]	+[7]	Foulquier *et al.,* 11	na
tagT+TagV	Teichoic acid synthesis	no	+	Kawai *et al* ., 11	na
tagO	TA synthesis (first step)	no[8]	R	D'Elia *et al.,* 06	na
tagA	TA synthesis (first commited step)	no[8]	R	D'Elia *et al.,* 09	na
Other cell envelop					
ltaS (yflE)	lipoteichoic acid synthase	no	-	Sassine, 20	no
yqgS	lipoteichoic acid synthase	no	-	Sassine, 20	no
yfnI (ltaSA)	lipoteichoic acid synthase	no	-	Sassine, 20	no
Division:					
ezrA+gpsB	FtsZ inhibitor and late divisome protein; *ponA* localization	no	-	Claessen *et al.,* 08	na

1: + or - impact on cell width upon gene deletion or depletion; R stands for round cell, a consequence of the absence of elongation

2: references for measurement of width

3: genes selected for width defect in the present screen; na (not available) indicates that the gene could not be be find here du to its absence in BKK library, its essentiality or because the phenotype is synthetic

4: synthetically lethal

5: essential in Muchova *et al.,* 13 but inactivated in Van Beilen *et al.,* 16, Koo *et al.* , 17, Kobayashi *et al.,* 03

6: thinner in Dominguez *et al., 20* 13, Carballido *et al., 20* 06, and Sassine, 2020 but wild type width in Meisner *et al.* , 13

7: only during neoglucogenic condition

8: non lethal but absent from the BKK library (wrongly annotated into the subtiwiky database)

Sup. Table 7. Average length differences (%) across replicates[1]

	WT 1	WT 2	WT 3	WT 4	WT 5	WT 6
WT 1	-	+2.9 %	+1.2 %	+4.8 %	+3.0 %	-0.5 %
WT 2	-	-	-1.7 %	+2.0 %	+0.1 %	-3.5 %
WT 3	-	-	-	+3.7 %	+1.8 %	-1,7 %
WT 4	-	-	-	-	-2.0 %	-5.6 %
WT 5	-	-	-	-	-	-3.6 %
length (µm)	3,384	3,484	3,424	3,556	3,487	3,367
+/-	0,708	0,760	0,805	0,822	0,705	0,777

1: ordinary one-way ANOVA test between replicates conclude to no significant differences