Micro-colony observation: A rapid and simple approach to count bacterial colony forming units

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ABSTRACT

For enumerating viable bacteria, traditional dilution plating to count colony forming units (CFU) has always been the preferred method in microbiology owing to its simplicity, albeit laborious and time-consuming. Similar CFU counts can be obtained by quantifying growing microcolonies in conjunction with the perks of a microscope. Here, we employed a simple method of five microliter spotting of differently diluted bacterial culture multiples times on a single petri plate followed by finding out CFU by counting microcolonies using phase contrast microscope. In this method within four-hour period CFU of an *Escherichia coli* culture can be found out. Further, within ten-hour period, CFU in a culture of *Ralstonia solanacearum*, a bacterium with generation time around 3 h, can be estimated. The CFU number determined by microcolonies observed is comparable with that obtained by the dilution plating method. Microcolonies number observed in the early hours of growth (2 h in case of *E. coli* and 8 h in case of *R. solanacearum*) were found to remain consistent at later h, though there was noticeable increase in size of the colonies. It suggested that microcolonies observed in the early hours indeed represent the bacterial number in the culture. Though manual, the method is less labor intensive apart from being simple, easy, economical and rapid.

Keywords: Colony forming unit; Microcolony; Microspotting; *Escherichia coli*; *Ralstonia solanacearum*; Phase contrast microscopy.

Introduction

Quantifying bacterial number by colony forming units (CFU) by employing serial dilution and spread plate techniques is an age-old fundamental microbiology technique used in research and diagnostics [1-3]. Though popular among the microbiologists because of its ease, the method demands large volumes of growth media, time and labor. Besides, working with bacteria having high doubling time such as *Ralstonia solanacearum* that takes ~48 h to form a full-grown colony on solid agar nutrient medium only doubles the misery [4,5]. Several other methods for enumerating the number of bacteria in a culture have come to light over the years. These include the measurement of absorbance using spectrophotometer or plate reader; flow cytometry, microscopy in conjunction with staining, qPCR and several microarray-based methods with each method having its own advantages over others [6-9]. Despite all advancements

and fast approaches to enumerate bacteria in a culture [1,2,6], CFU counting has been the most dependable and popular method among microbiologists because of its viable nature and visual impact.

Therefore, it will be a big boon for the microbiologists to count CFU in a bacterial culture with reduced labor, time and cost.

In this study we are describing a subtle method of enumerating bacteria in a culture by observing microcolonies under the microscope. The basic principle of this method is to spot a small amount (~ 5.0 µl) of a serially diluted bacterial culture followed by imaging and quantification of the micro-colonies obtained using the microscope [10]. In an *E. coli* culture, CFU can be found out within four hours after observing the microcolonies by this method. Similarly, in case of *R. solanacearum*, a bacterium with generation time more than 3.0 h, CFU can be found out within ten hours after observing the microcolonies. The method is simple to use, scalable and costeffective that maintains the benefits of a microscopic platform. Further, validation of this method was done by comparison with other standard quantification approaches. It is pertinent to note that the microcolonies count method is done by spotting a few microliters, which enables us to analyse different dilutions in a single petridish. Therefore, the number of CFU observed under the microscope can be confirmed by observing the CFU through naked eyes in later days depending upon the growth rate of the bacteria.

Materials and methods

Strains and growth conditions

Escherichia coli DH5α cells were grown in Luria–Bertani (LB) medium at 37° C. *R. solanacearum* F1C1 (1) cells were grown in BG medium (contains 1% peptone, 0.1% yeast extract and 0.1% casamino acids) supplemented with 5 g/L glucose at 28°C. For solid medium agar 1.5% was used.

Colony-forming unit (CFU) count

The bacterial strains were inoculated in their respective liquid medium and incubated at their optimal growth conditions for 6 h in *E. coli* and overnight in *R. solanacearum* with 150 rpm shaking condition. For counting CFU, the overnight grown culture of *E. coli* cells was serially diluted and spotted on LB agar plates as illustrated in Fig. 1. First, 100 µL overnight grown culture of *E. coli* was diluted in 900 µL of sterile distilled water. Since the dilution was one part sample to nine parts diluents, it was referred to as a 10¹ dilution. Further, we took 100 µL of the 10¹ dilutions and added it to 900 µL of the sterile distilled water. This dilution was a 10² dilution. Lastly, we made a 10¹⁰ dilution by taking 100 µL of the 10⁹ dilutions and adding 900 µL of the sterile distilled water. For each dilution, we now had 1 mL of diluted bacterial culture. After two hours of incubation at 37° C, micro-colonies were counted by using Life technologies EVOS FL inverted microscope with 40X magnification using the 4/10 PH condenser annulus. Under the microscope, the area covered by the cells was calculated by extrapolating the length of the 1000 µm scale provided. Thus, we calculated the total CFU in the area covered by 5 µL spot and subsequently in 1 mL of *E. coli* culture.

This micro-colony counting was further validated by on-plate CFU counting of overnight incubation at 37°C. The mean value of ten independent spots was used to show the growth parameters. Similar approach was used to count CFU in case of *R. solanacearum* culture. However, the culture was incubated at 28°C. As *R. solanacearum* is a slow growing bacterium with a doubling time of more than ~ 3.0 h, its microcolonies at different dilution could be observed distinctly at 40X magnification in 10.0 h of incubation after spotting. The bacterial CFU was also counted on Life technologies EVOS FL inverted microscope with 40X magnification using the 4/10 PH condenser annulus.

CFU from microcolonies method

Microcolonies observed under the microscope of different dilutions such as 10^4 , 10^5 , 10^6 fold etc. were counted at different time intervals manually. In case of *E. coli* observation of microcolonies was done from 2.0 h onwards up to 8.0 h at every 2.0 h intervals in the EVOS FL microscope with an objective 4X magnification. The area covered under the microscope was calculated from the scale bar 1000 μm provided in the microscope. Total area covered under the microscope was calculated as 5.67 mm^2 . A 5 μl culture covers an average area of 19.62 mm^2 (a circle of average diameter of 5.0 mm). From a single spot of bacterial culture, a minimum of three sections (each 5.67 mm^2) were counted and the average number of colonies per section was found out, which was then multiplied with 3.46 to find out number of bacterial colonies in 5.0 μl of the culture at that dilution. The same approach was used to find CFU at different dilution used for spotting the culture. In case of *R. solanacearum*, observation of microcolonies was done from 8.0 h onwards up to 14.0 h at every 2.0 h intervals.

CFU from microliter spotting

After observing the microcolonies under the microscope in their early hours of growth, the petridishes were kept back in the incubator. After overnight growth (in case of *E. coli*) colonies could be visible by naked eyes. In case of 10^6 dilution, average 15 colonies per spot can be observed, while in case of 10^7 dilution average 2.0 colonies per spot were observed. This method proved the observations made earlier through microcolonies counting.

CFU from spread plate method

In the usual spread plate method to count CFU, 100 μl of overnight grown serially diluted bacterial cultures were evenly distributed on the solid growth medium by using a glass spreader. The petridishes were kept in the incubator and after overnight incubation, we observed the colonies by naked eyes. All experiments were performed in triplicates.

Results and Discussion

We calculated the micro-colonies in an *E. coli* culture as described in Materials and Methods section. We observed the micro-colonies of *E. coli* by the microscope from ten different spots of different dilutions such as 10^4 , 10^5 , 10^6 from 2.0 h onwards after incubation till 8.0 h after every 2 h interval [SFig1a-1c]. Number of microcolonies observed and CFU calculated are listed in Table 1. After 2.0 h of incubation, the micro-colonies counts were 2.22×10^9 and 3.62×10^9 for the 10^4 and 10^5 dilution spots, respectively. Subsequently, in 4.0 h of incubation period at 37°C, the micro-colonies counts were calculated as 1.99×10^9 , 2.40×10^9 and 4.98×10^9 for the 10^4 , 10^5 and 10^6 dilution spots, respectively [Fig. 2a, Table 1]. After 6.0 h of incubation period, the micro-colonies counts were found out

1.8×10^9 , 2.35×10^9 and 3.04×10^9 for the 10^4 , 10^5 and 10^6 dilution spots, respectively. While, in 8.0 h of incubation, the micro-colonies counts were calculated as 1.26×10^9 , 2.10×10^9 and 3.46×10^9 for the 10^4 , 10^5 and 10^6 dilution spots, respectively. Further, these micro-colonies counts were confirmed by spot dilution [SFig 2] and plate spreading [SFig 3] methods and results showed that almost similar CFU counts were observed with different dilutions at different time periods as listed in Table 3. It supported that microcolonies count can represent the CFU in a culture in case of *E. coli*.

Likewise, the CFU count using microliter spotting and microcolony counting was done in *R. solanacearum*, a bacterium with generation time more than ~ 3.0 h, as well. Microcolonies could be observed from 8 h onwards at 10^4 dilution spotting. However, it could be counted properly after 10 h of spotting. Fig 2b demonstrate microcolonies of *R. solanacearum* after 10 h of spotting of dilutions such 10^4 , 10^5 , and 10^6 . It is pertinent to note that at this hour of growth it is not easy to find colonies in case of 10^7 dilutions as the number of colonies are very low. We observed that the micro-colonies of *R. solanacearum* by the microscope from six different spots of 10^4 dilution and ten different spots of 10^5 and 10^6 each from 8 h of incubation till 14 h after every 2 h interval [SFig4a4c]. Number of microcolonies observed and CFU calculated are listed in Table 2. After 8.0 h of incubation, the micro-colonies counts were 2.52×10^9 , 3.66×10^9 and 4.84×10^9 for the 10^4 , 10^5 and 10^6 dilution spots, respectively. Subsequently, in 10.0 h of incubation period at 28°C , the micro-colonies counts were calculated as 2.60×10^9 , 3.46×10^9 and 5.07×10^9 for the 10^4 , 10^5 and 10^6 dilution spots, respectively [Table 2]. After 12.0 h of incubation period, the micro-colonies counts were found out 3.92×10^9 , 4.57×10^9 and 4.61×10^9 for the 10^4 , 10^5 and 10^6 dilution spots respectively. Similar results were obtained with plate spreading method [Table 3, SFig 5]. We further counted 10 spots for 10^6 and 10^7 dilutions each, as well as 10 plates each of dilutions 10^6 and 10^7 of *E. coli* and the observation revealed that the co-efficient of variance was comparable by both spotting method and the dilution plating method, respectively [Table 4]. The size of micro-colonies has increased at different time periods, however, these results show that there is a similarity of calculated number of micro-colonies at different time periods [Fig. 3]. It supported that microcolonies count could represent the CFU in a culture in case of *R. solanacearum*. In *R. solanacearum*, the microcolonies grow in size, exhibit twitching motility and aggregate together. Twitching motility is, however, absent in *E. coli*.

Serial dilution is an easy and convenient approach to measure viable bacterial concentration in a culture. In early days of microbiology, enumerating bacterial number in a culture was made in test tubes by finding out the minimum dilution at which no bacterial growth was observed [11]. Usually, turbidity development in the medium used to be measured as bacterial growth [SFig 6]. Later with the use of agar, colony formation by bacteria could be observed on solid agar medium [12]. With the availability of micropipettes, which can pipette out liquids of volume 1.0 or 2.0 μl with accuracy, it became an aid for us to think of counting bacteria in 5 μl volume.

The findings that microcolonies number remain similar at various point of time at a specific dilution supported that the method can be used to count bacteria in a culture by observing micro-colonies. In contrast to the conventional spread plate method, calculating CFU by dilution spotting is both time-saving and economical. While we could calculate *R. solanacearum* and *E. coli* CFU using dilution spotting at 10hrs and 4hrs respectively, we could obtain

similar results of *R. solanacearum* and *E. coli* using spread plate method at only 48hrs and 16hrs of incubation respectively.

Dilution spotting and subsequent back calculations are a fast and easy technique to enumerate bacterial population in a culture. Besides, CFU can be estimated from different dilutions, such as, 10^4 , 10^5 , 10^6 and 10^7 . Counting CFU from 10^4 and 10^5 dilution by spread plate method is difficult because the colonies merged with each other due to their high number. Also less variation of CFU numbers across different spots of same dilution is observed. A dilution spot of 10^5 was the most convenient in counting microcolonies. Phase-contrast micrographs not only eliminates the use of stain to distinguish between live and dead cells but also help us identify contaminants to some extent. The advantages to this technique are many, but it too has some cons. Phase-contrast microscope itself comes with a hefty price tag but this method of CFU counting can be performed using any microscope that comes with a scale to extrapolate. Also, this technique demands manual labor; albeit lesser labor-intensive than the traditional counting ways.

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Conflict of interest

We declare that there is no conflict of interest among the authors of this manuscript.

References

1. Jett BD, Hatter KL, Huycke MM, Gilmore MS (1997) Simplified agar plate method for quantifying viable bacteria. *BioTechniques* 23(4):648–650. <https://doi.org/10.2144/97234BM22>
2. McNulty JJ, Dunn JJ (1999) High-throughput transformation and plating using petristrips. *BioTechniques* 26(3):390–392. <https://doi.org/10.2144/99263BM04>

3. Yadav M, Rathore JS (2022) Functional and transcriptional analysis of chromosomal encoded hipBAXn2 type II toxin-antitoxin (TA) module from *Xenorhabdus nematophila*. Microbial Pathogenesis 162:105309. <https://doi.org/10.1016/J.MICPATH.2021.105309>
4. Kumar R, Barman A, Jha G, Ray SK (2013) Identification and establishment of genomic identity of *Ralstonia solanacearum* isolated from a wilted chilli plant at Tezpur, North East India. Curr Sc 105(11):1571–1578.
5. Xu P, Iwata M, Leong S, Sequeira L (1990) Highly virulent strains of *Pseudomonas solanacearum* that are defective in extracellular-polysaccharide production. J Bacteriol 172(7):3946. <https://doi.org/10.1128/JB.172.7.3946-3951.1990>
6. Hsieh K, Zec HC, Chen L, Kaushik AM, MacH KE, Liao JC, Wang TH (2018) Simple and Precise Counting of Viable Bacteria by Resazurin-Amplified Picoarray Detection. In Analytical Chem 90(15):9449-9456. <https://doi.org/10.1021/acs.analchem.8b02096>
7. Koch AL (2014) Growth Measurement. Methods for Gen & Mol Microbiol 172–199. <https://doi.org/10.1128/9781555817497.CH9>
8. Pan H, Zhang Y, He GX, Katagori N, Chen H (2014) A comparison of conventional methods for the quantification of bacterial cells after exposure to metal oxide nanoparticles. BMC Microbiol 14(1):222. <https://doi.org/10.1186/S12866-014-0222-6>
9. Stevenson K, McVey AF, Clark IBN, Swain PS, Pilizota T (2016) General calibration of microbial growth in microplate readers. Sc Reports 6(1):1–7. <https://doi.org/10.1038/srep38828>
10. Ray SK, Kumar R, Peeters N, Boucher C, Genin S (2015) rpoN1, but not rpoN2, is required for twitching motility, natural competence, growth on nitrate, and virulence of *Ralstonia solanacearum*. Frontiers in Microbiol 6:229. <https://doi.org/10.3389/FMICB.2015.00229/ABSTRACT>
11. Koch R. (1883) Über die neuen Untersuchungsmethoden zum Nachweis der Mikrokosmen in Boden, Luft und Wasser. 1.
12. Breed RS, Dotterrer WD (1916) The number of colonies allowable on satisfactory agar plates. J Bacteriol 1(3):321. <https://doi.org/10.1128/jb.1.3.321-331.1916>

Figures

Fig. 1: Schematic representation of the microcolonies counting method

In a given bacterial culture considered for counting CFU, dilution of the culture was performed as shown in the figure. 100 µl of the culture was added to 900 µl of water to make it 10-fold (10^1) dilution, from the diluted culture 100 µl was taken and added to 900 µl of water to dilute it another 10 times. The culture was diluted 10^{10} fold. Only 5 µl from different dilutions were spotted on a LB-Agar plate in case of *E. coli*. Approximately, ten such spots can be made on a Petri-plates (90 mm diameter). After 2.0 h of incubation, micro-colonies were observed at 40x magnification. Three different sections of 10^4 dilution spotted have been shown here. Distinct colonies could be observed.

Fig 2a. *E. coli* microcolonies number observed at different time points

At 2.0 h, 4.0 h, 6.0 h and 8.0 h duration microcolonies were observed for different dilutions such as 10^4 , 10^5 and 10^6 of *E. coli* culture. Counted microcolonies found to be similar at different time points though there is growth in colony size. Further, number of CFU estimated by counting the microcolonies at different dilutions is observed to be similar.

Fig 2b. *R. solanacearum* microcolonies number observed at different time points

At 8.0 h, 10.0 h, 12.0 h and 14.0 h duration microcolonies were observed for different dilutions such as 10^4 , 10^5 and 10^6 of *R. solanacearum* culture. Counted microcolonies found to be similar at different time points though there is growth in colony size. Further, number of CFU estimated by counting the microcolonies at different dilutions is observed to be similar.

Fig. 3. *R. solanacearum* microcolonies at different dilutions

R. solanacearum microcolonies observed at different dilutions such as 10^4 , 10^5 , 10^6 and 10^7 after at 10 h incubation. Microcolony formed are distinctly observed at different dilutions. The number are decreasing with increase in dilution. This suggest that CFU count is possible after 10 h of incubation in case of *R. solanacearum*.

Table 1: CFU calculated in *E. coli* culture by micro-colony counting method

Dilution	Time (h)	CFU/mL
10^4	2	2.22×10^9
	4	1.99×10^9
	6	1.80×10^9
	8	1.26×10^9
10^5	2	3.62×10^9
	4	2.40×10^9
	6	2.35×10^9
	8	2.10×10^9
10^6	4	4.98×10^9
	6	3.04×10^9
	8	3.46×10^9

Table 2: CFU calculate in *R. solanacearum* by micro-colony counting method

Dilution	Time (h)	CFU/ml
10^4	8	2.52×10^9
	10	2.60×10^9
	12	3.92×10^9
	14	1.26×10^9
10^5	8	3.66×10^9
	10	3.46×10^9
	12	4.57×10^9
	14	3.83×10^9
10^6	8	4.84×10^9
	10	5.07×10^9
	12	4.61×10^9
	14	4.28×10^9

Table 3: Comparison of CFU calculation by 5 µl spotting and 100 µl spreading method

Dilutions	<i>R. solanacearum</i>		<i>E. coli</i>	
	Dilution spotting	Spread plate	Dilution spotting	Spread plate
10 ⁴	2.1 x 10 ⁹ cells		Colonies merge with each other	
10 ⁵	2.3 x 10 ⁹ cells		6.04 x 10 ⁸	4.5 x 10 ⁸
10 ⁶	3.5 x 10 ⁹	10 ⁹	6.2 x 10 ⁸	7.1 x 10 ⁸
10 ⁷	7 x 10 ⁹	1.1 x 10 ⁹	1.6 x 10 ⁹	7 x 10 ⁸
10 ⁸	No colonies found	6.7 x 10 ⁸	4 x 10 ⁹	6.7 x 10 ⁸

Table 4: Comparison of visible colonies observed by both the methods in *E. coli*

	10 ⁶ dilution	10 ⁷ dilution		10 ⁶ dilution	10 ⁷ dilution
	5μL	5μL		100μL	100μL
S	20	2	S	121	13
P	21	1	P	233	13
O	11	0	R	106	30
T	15	1	E	102	13
T	19	1	A	156	14
I	16	0	D	80	19
N	9	5	I	199	21
G	14	1	N	242	29
	11	3	G	210	23
	17	1		205	27
MEAN	15.3	1.5	MEAN	165.4	20.2
SD	4.083844	1.509231	SD	59.65121	6.860515
CV	0.266918	1.006154	CV	0.360648	0.339629