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An improved vector for baculovirus-mediated protein production in mammalian cells

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Abstract

BacMam system utilizes baculovirus to deliver exogenous gene into mammalian cells and are

extensively used for recombinant production of eukaryotic proteins. Here, we described an

improved BacMam vector (pBMCL1) which allows convenient tracing of virus production,

provides higher infection efficiency towards mammalian cells, minimizes unwanted transcription

of toxic gene in insect cells, and provides the capability for co-expression of multiple proteins

via a single virus. We demonstrate the successful application of pBMCL1 vector for the

expression of not only human TRPC3 channel but also heteromeric K_{ATP} channel.

1. Introduction

Baculoviruses are double-stranded DNA viruses that can infect and replicate in insect

populations. *Autographa californica* multiple nuclear polyhedrosis virus (AcMNPV) is one of

the most common isolates of baculovirus. A modified version of AcMNPV, of which the

polyhedrin gene is replaced by a target gene, is widely used for over-expression of recombinant

protein in insect cells. Since it was found that the modified AcMNPV could mediate gene-

transfer in mammalian cells [1-3], baculovirus is more and more extensively used as a vehicle for

recombinant protein production in mammalian cells and this procedure is called BacMam

system. BacMam system has several advantages over other eukaryotic expression systems, such

as insect cells or yeast expression systems. BacMam system allows exogenous gene transcription

from several available strong promoters such as CMV in mammalian cells. Moreover, the

BacMam system provides genuine mammalian cell environment for the posttranslational

modifications of target proteins, and the glycosylation pattern of membrane protein or secreted

protein can be modified using available cell line such as GnT1[-4]. Furthermore, BacMam system

often provides high level of membrane protein production compared to other expression systems,

allowing the access to milligram-level purified sample of previously intractable membrane

protein targeted for structural or biochemical studies [5]. Moreover, the large physical size of

baculovirus provides high capacity for packing exogenous DNA, allowing delivering multiple

genes in one single virus. However, the current available BacMam vectors have not fully

exploited these advantages of the system and suffer from low virus titer due to leak expression of

toxic target gene, inconvenient tracing of virus production, and lack of the capability of co-

expression of multiple proteins in a single virus. Here, we describe the design and application of

an improved vector for BacMam system.

2. Materials and methods

2.1. Cell lines and reagents

FreeStyle 293F cells (Thermo Fisher Scientific, R79007) are cultured in either FreeStyle 293

Expression Medium (Thermo Fisher Scientific, 12338018) or SMM 293-TI (Sino Biological

Inc., M293TI) with 1% Fetal bovine serum (Vistech). Sf9 cells (Thermo Fisher Scientific,

12659017) were cultured in Sf-900 III SFM medium (Thermo Fisher Scientific, 12658019) or

SIM SF Expression Medium (Sino Biological Inc, MSF1).

2.2. Construction of pBMCL1 vector

We synthesized two DNA sequences that contain the open reading frame of (GP64 signal

peptide)-(VSV-GED)-(CreiLOV) fusion protein and (LINK1)-(CMV promoter)-(multiple

cloning site, MCS)-(Woodchuck Hepatitis Virus Posttranscriptional Regulatory Element,

WPRE)-(SV40 polyA signal)-(LINK2) expression cassette. We inserted (GP64 signal peptide)-

(VSV-GED)-(CreiLOV) into pFastBac1 vector and replaced its SV40 polyA signal with

HSVtkPolyA signal to generate a vector that displays VSV-GED protein on insect cell

membrane and baculovirus envelope. Then we inserted the expression cassette of (LINK1)-

(CMV promoter)-(MCS)-(WPRE)-(SV40 polyA signal)-(LINK2) into the vector to the opposite

direction of the polyhedrin promoter to minimize mammalian gene expression in insect cells. We

name this vector pBMCL1 (Figure 1). The vector has been deposited onto Addgene (#80316).

2.3. Generation of expression constructs

The cDNA of human TRPC3 (hTRPC3) [6] with N-terminal GFP was cloned into the MCS of

pBMCL1 vector for protein expression using BacMam system. The cDNA of SUR1 of

Mesocricetus auratus (XP_005084737) [7] and the cDNA of Kir6.2 with C-terminal GFP tag from

Mus musculus [8] were cloned into the MCS of pBMCL1 BacMam expression vector to generate

pBMCL1-Kir6.2-CGFP and pBMCL1-SUR1 vector. Then these two vectors were merged into

one vector by the LINK sequences using ligation-independent cloning method [9] to generate the

pBMCL1-K_{ATP} vector.

2.4. Generation of baculoviruses and infection of mammalian cells

pBMCL1 vectors containing target proteins were transformed into DH10Bac *Escherichia coli*

cells. Recombinant bacmid was identified by the blue-white screening. Baculoviruses were

generated using the Bac-to-Bac Baculovirus Expression system (Invitrogen). Sf9 cells were

cultured in Sf-900 III SFM medium (Gibco) at 27°C and seeded in 12-well plate. After

transfection of bacmid for 7 days, the fluorescence of CreiLOV was detected and the supernatant

was collected. Higher titer P2 virus was amplified using sf9 suspension cells cultured in SIM SF

medium. At the time of cell mortality over 20% determined by trypan blue staining, the

supernatant that containing baculovirus was collected by centrifugation and storage at 4 degree.

P2 virus were added into FreeStyle 293F suspension cell culture directly for infection.

2.5. Fluorescence-detection size-exclusion (FSEC)

Cells that expressed hTRPC3 channel were solubilized in TBS (20 mM Tris pH 8.0 at 4°C, 150

mM NaCl) with 10 mM Lauryl Maltose Neopentyl Glycol (MNG, Antrance, NG310), 0.1%

Cholesteryl Hemisuccinate Tris Salt, (CHS, Antrance, CH210) and protease inhibitors including

1 μg/ml aprotinin, 1 μg/ml leupeptin, and 1 μg/ml pepstatin on ice for 30 min. Cell lysates were

clarified by centrifugation at 40,000 rpm for 30 min and supernatants were loaded onto Superose

6 Increase (GE Healthcare) for FSEC analysis.

Cells that expressed K_{ATP} channels were harvested and solubilized in TBS with 1% digitonin

(Biosynth) and protease inhibitors including 1 μg/ml aprotinin, 1 μg/ml leupeptin, and 1 μg/ml

pepstatin on ice for 30 min. Cell lysates were processed by centrifugation at 40,000 rpm for 30

min and supernatant was loaded onto Superose 6 Increase for FSEC analysis.

2.6. Microscopy imaging

Successful generation of virus was indicated by green fluorescence of sf9 cells 7 days post

transfection of bacmid. Images were taken using EVOS FL imaging system. For visualization of

protein expression in mammalian cells, FreeStyle 293F cells (cultured in Freestyle 293 medium)

were plated in 35 mm dish without shaking for 30 min, then baculoviruses were added into the

medium. 48 h post infection, GFP signal was detected in live cells and images were obtained

using Axio Image M2 microscope.

2.7. Electrophysiology

For patch clamp recording, pBMCL1-K_{ATP} plasmids were transfected into FreeStyle 293F cells

(cultured in FreeStyle 293 medium) at density of 1×10^6 cells/ml using PEI (Polyscience, 687600)

and cells were cultured for 32 h before recording. Patch electrodes were pulled from a horizontal

microelectrode puller (P-1000, Sutter Instrument Co, USA) to a tip resistance of 2-3 MΩ.

Currents were recorded using inside-out mode at +60 mV using an Axopatch 200B amplifier

(Axon Instruments, USA). We used KINT solution which contained (mM):140 KCl, 1EGTA, 10

HEPES (pH 7.4, KOH) in both pipette and bath. Signals were acquired at 20 kHz and low-pass

filtered at 5 kHz. Data were further analyzed with pClampFit 9.0 software and filtered at 1 kHz.

3. Results and discussion

3.1 Design of the pBMCL1 vector

We adopted the popular Bac-to-Bac system which utilizes the site-specific Tn7-mediated

transposition for bacmid generation. We modified the established BacMam expression cassette [5]

by minimizing the lengths of linkers between each elements and removing commonly used

restriction enzyme cutting sites outside of the MCS. To confer the capability to express multiple

genes simultaneously, we appended LINK1 and LINK2 sequences at 5' and 3' of the expression

cassette respectively. LINK sequences allow the unidirectional fusion of unlimited number of

expression cassettes using a ligation-independent and PCR-independent cloning method [9]. LINK

sequences are found to be functional in BacMam system based on our previous successful

application on the expression of DUOX1-DUOXA1 complex [10]. The resulting expression

cassette contains a 5' LINK1 sequence, a CMV promoter, an MCS, a WPRE, a SV40 polyA

signal and a 3' LINK2 sequence.

Figure 1. The design of pBMCL1 vector. **a**, Map of pBMCL1 vector. Elements that are essential
for function are shown on the vector. **b**, The sequence of MCS and restriction enzyme
recognition sites. **c**, LINK1 and LINK2 sequences and their restriction enzyme recognition sites
by SwaI and PacI. These sequences are used for merging multiple expression cassettes into one
vector by ligation-independent cloning.

The successful application of BacMam system highly relies on the titer of baculovirus produced

from insect cells. The strong expression of some proteins, such as certain ion channels, in insect

cells might be toxic and lead to low virus-titer. However, in the commonly used BacMam vector,

pEG BacMam [5], the polyhedrin promoter is in the same direction as target protein, resulting in

leak expression in insect cells. Therefore, we intentionally inserted the aforementioned

expression cassette in reverse direction to the polyhedrin promoter of pFastbac1 vector to

prevent the high level transcription of target gene in insect cells and avoid target protein toxicity

to the virus-producing cell. On the membrane envelope of baculovirus, there are several copies

of GP64 surface protein which might bind to the heparin sulfate proteoglycans on the plasma

membrane of mammalian cells [11]. The virus then enters the cell through clathrin-mediated

endocytosis process [12]. Many exogenous membrane proteins can be functionally displayed on the

baculovirus envelope during virus assembly and budding. It is reported that the vesicular

stomatitis virus glycoprotein G (VSV-G) protein displayed on baculovirus envelope can enhance

the infectivity of baculovirus towards mammalian cells [13] and a short fragment of VSV-G (VSV-

GED) is sufficient to facilitate baculovirus-mediated gene delivery in vertebrate cells [14,15]. In

order to trace the successful transfection of bacmid and production of baculovirus, we utilize the

polyhedrin promoter to drive the expression of a fluorescent reporter, VSV-GED-LOV fusion

protein in insect cells. A GP64 signal peptide directs the fusion protein to the plasma membrane

of insect cell and virus envelope. VSV-GED fragment is used to enhance the infectivity of

baculovirus [14,15]. CreiLOV is a small, thermostable, photostable, and fast-maturing monomeric

flavin-based green fluorescent protein [16]. The commonly available green channel on fluorescent

microscope for GFP detection can be used to visualize the bright CreiLOV signals. More

importantly, the CreiLOV has no sequence homology to GFP, which is commonly used as a

versatile fusion tag for the tracing of target protein expression and purification. Because the

presence of multiple copies of similar sequences on the same plasmid might lead to DNA

instability during molecular biology manipulations such as QucikChange mutagenesis, the

presence CreiLOV would avoid the instability of plasmid when target protein has GFP tags. If

the insect cells are successfully transfected with bacmids and are producing baculovirus, they

would show green signals generated from CreiLOV, which is under the control of polyhedrin

promoter (Figure 2).

Figure 2. Successful baculovirus generation indicated by green fluorescence signal in sf9 cells.
a, CreiLOV signal in sf9 cells after transfection for 7 days detected by a fluorescent microscope
using green channel. **b**, Bright filed image of transfected sf9 cells. **c**, The merged image of green
signal and bright field signal.

3.2 Using pBMCL1 vector to express human TRPC3 channel in mammalian cells

To test the function of pBMCL1 vector, we used it to express human TRPC3 (hTRPC3) channel

in FreeStyle 293F cells. Full length hTRPC3 is a canonical TRP channel with molecular weight

around 90 kDa. Functional hTRPC3 is a homo-tetrameric channel. We cloned the full length

hTRPC3 with N-terminal GFP tag into the pBMCL1 vector. After transfection of the bacmid into

sf9 insect cells for 7 days, we observed green fluorescence signal using GFP channel (Figure 3a)

and P1 viruses were harvested and further amplified to obtain the P2 virus. P2 viruses were used

for overexpression of hTRPC3 in FreeStyle 293F cells. 60 h post infection, we observed GFP

signal from the majority of FreeStyle 293F cells (Figure 3b). We also found the band of NGFP-

hTRPC3 protein at correct molecular weight on SDS-PAGE using in-gel fluorescence. (Figure

3c).

To optimize the protein expression condition, we screened different virus-cell ratio, expression

duration and expression medium. We found that at the volume ratio of 1:10 (virus: cell), the

expression level was highest (Figure 3d). Moreover, the expression level of hTRPC3 peaked at

48 h post-infection but decreased thereafter (Figure 3e). Furthermore, we found 293-TI medium

supported higher level of hTRPC3 expression than the FreeStyle 293 medium under conditions

tested (Figure 3f).

Figure 3. Expression of hTRPC3 channel via BacMam system using pBMCL1 vector. **a**, CreiLOV signal of sf9 cell 7 days post transfection of bacmid. **b**, GFP signal of FreeStyle 293F cells 48 h post infection by BacMam virus with virus-cell ratio 1:10 (v/v). **c**, In-gel fluorescence of NGFP-hTRPC3 protein. Band of NGFP-hTRPC3 protein is boxed in red. **d**, FSEC profiles of NGFP-hTRPC3 tetramer produced from FreeStyle 293F cells infected with different virus to cell ratio (v/v). **e**, FSEC profiles of NGFP-hTRPC3 tetramer produced from FreeStyle 293F cells that were cultured in different medium. **f**, FSEC profiles of NGFP-hTRPC3 tetramer produced from FreeStyle 293F cells expressed for various time.

3.3 Using pBMCL1 vector to express pancreatic K$_{ATP}$ channel in mammalian cells

To explore the capability of co-expressing more than one protein using a single pBMCL1 vector,

we expressed the hetero-octameric pancreatic K$_{ATP}$ channel in FreeStyle 293F cells. K$_{ATP}$

channel is a potassium channel, the activity of which is sensitive to intracellular ATP/ADP ratio.

Pancreatic K$_{ATP}$ channel in β cells couples blood glucose level to insulin secretion and is a drug

target for diabetes. Pancreatic K$_{ATP}$ channel is assembled from four Kir6.2 subunits and four

SUR1 subunits. Kir6.2 subunit is the ion channel pore with molecular weight 43.5 kDa. SUR1

subunit is the regulatory subunit with molecular weight 177 kDa. To trace the expression of K_{ATP}

channel, we fused mouse Kir6.2 with a C-terminal GFP tag and cloned it into pBMCL1 vector to

generate pBMCL1-Kir6.2-CGFP. We cloned non-tagged SUR1 subunit into pBMCL1 to

generate pBMCL1-SUR1 vector. The expression cassette of pBMCL1-SUR1 was inserted into

pBMCL1-Kir6.2-CGFP to obtain the pBMCL1-K_{ATP} vector. Transient transfection of pBMCL1-

K_{ATP} into FreeStyle 293F cells allowed the recording of robust ATP-sensitive potassium currents

in inside-out mode (Figure 4a). Moreover, the currents were sensitive to the insulin secretagogue

repaglinide, suggesting functional K_{ATP} channels were correctly assembled and trafficked onto

the plasma membrane. When solubilized, the K_{ATP} complex ran faster than Kir6.2 channel

expressed alone on FSEC, indicating the hetero-octamer assembly (Figure 4b). We generated

BacMam virus using pBMCL1-K_{ATP} vector for infection of FreeStyle 293F cells. We found K_{ATP}

complex showed robust expression 48 h post-infection compared to non-infected cells (Figure

4c). We also showed the expression of both Kir6.2 and SUR1 subunits by western blot (Figure

4d). These data collectively demonstrate pBMCL1 vector is capable for the expression of

eukaryotic membrane protein complex.

Figure 4. Expression of K_{ATP} channel via BacMam system using pBMCL1 vector. **a**, Recording of robust ATP-sensitive potassium currents in inside-out mode in FreeStyle 293F cells which were transfected by pBMCL1-K_{ATP} plasmid, indicating functional co-expression of both Kir6.2 and SUR1 subunits. **b**, The FSEC profile of K_{ATP} complex (using pBMCL1-K_{ATP} plasmid) and Kir6.2 channel in transfected cells. The earlier elution of K_{ATP} complex on size-exclusion chromatography suggests the successful hetero-complex formation between Kir6.2 and SUR1 subunits. **c**, The FSEC profile of K_{ATP} complex in FreeStyle 293F cells infected by pBMCL1-K_{ATP} BacMam virus. **d**, In virus-infected cells, the expression of Kir6.2 was detected by in-gel GFP fluorescence and the expression of SUR1 subunit was detected by western blot using anti-SUR1 antibody.

4. Conclusion

In this paper, we have described the design of pBMCL1 vector with several new features. First,

this vector facilitates the tracing of successful virus generation by green fluorescence signal from

LOV. Second, this vector might enhance the infectivity of BacMam virus towards mammalian

cells by displaying VSV-GED fragment on baculovirus surface. Third, the expression of target

protein in insect cells is minimized by placing CMV promoter in reverse direction to the

polyhedrin promoter. Fourth, the incorporation of LINK sequences permits co-expression of

multiple genes using BacMam system at the same time. Using this vector, we successfully

expressed homo-tetrameric ion channel hTRPC3. Furthermore, we demonstrated the functional

expression of hetero-octameric pancreatic K_{ATP} channel using this vector. We believe the

aforementioned new features of the pBMCL1 vector would help researchers to fully unleash the

capability of BacMam system for recombinant protein expression. Notably, a recently developed

MultiBacMam system uses a modified Bacmid that contains VSV-G and mCherry to achieve the

tracing of virus production [17]. However, the mCherry tracer is away from target gene on virus

genome. Therefore, the mCherry might show expression in insect cells even for viruses that

contains no target gene. In contrast, in pBMCL1 vector, the CreiLOV is closely associated with

the expression cassette, allowing more faithful tracing of the production of virus containing

target gene.

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Credit authorship contribution statement

Wenjun Guo: Methodology, Validation, Formal analysis, Investigation, Writing - review &

editing; **Mengmeng Wang:** Methodology, Validation, Formal analysis, Investigation, Writing -

review & editing; **Lei Chen:** Conceptualization, Methodology, Validation, Formal analysis,

Investigation, Writing - original draft, Writing - review & editing, Visualization, Supervision,

Project administration, Funding acquisition.

Competing interests

The authors declare no competing interests

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