

Optimization of Electroporation Conditions for *Bacillus pumilus* 3–19 Strain

I. V. Danilova¹ · N. L. Rudakova¹ · Y. A. Vasilyeva¹ · A. I. Gilmutdinova¹ · I. V. Diadkina¹ · D. I. Khasanov¹ · M. R. Sharipova¹

Accepted: 19 April 2022

© The Author(s), under exclusive licence to Springer Science+Business Media, LLC, part of Springer Nature 2022

Abstract

Electroporation is the process of using electrical impulses to create temporary pores in the plasma membrane which, in turn, enables the penetration of nucleic acids into the cytoplasm of a cell. This method is widely used for the rapid and efficient introduction of foreign DNA into a wide range of cells. Cell viability and electrotransfection efficiency depend on various experimental factors, including the impulse form, vector concentration, cell type (density), properties of the electroporation buffer, and the growth phase of the bacterial culture. In this work, we investigated the optimal conditions for the transformation of the *Bacillus pumilus* 3–19 strain using electroporation. Competent *B. pumilus* cells were obtained on the 4th hour of culture growth using PEB1 electroporation buffer and SOC1 (Super Optimal broth with Catabolic repressor) medium, while the electric field strength was 12 kV/cm. With these parameters, the transformation efficiency of bacillus cells was 56.3 transformants/µg DNA. Thus, the rational choice of pulsation conditions and buffering compositions is critical for the design of electroporation protocols to maximize the viability and efficiency of electrotransfection.

Keywords *Bacillus pumilus* · Transformation · Competent cells · Bacterial growth phase · Electric field strength · Electroporation

1 Introduction

Spore-forming gram-positive bacteria of the *Bacillus* genus belong to one of the most diverse and widely represented groups of microorganisms in nature. Besides their characteristics such as storage stability, manufacturability in the production process, and environmental safety, the natural ability of this group of bacteria to synthesize and secrete several important metabolites of industrial, agricultural, and medicinal significance makes bacilli an important object of industrial biotechnology. Due to the rapid development of post-genomic research, effective genetic tools are now necessary for targeted modification of the *Bacillus* genomes.

B. pumilus are gram-positive bacteria, known for their ability to produce industrially significant hydrolytic enzymes with the prospect of being used in producing gluten-degrading detergents, hydrolyzing bird feathers, and breaking down biofilms [1–5]. *B. pumilus* bacteria are also potential probiotic agents that positively affect the growth and development of farm animals and plants, as well as humans [6–9]. *B. pumilus* can produce biologically active substances with antibacterial [10], insecticidal (toxins acting on invertebrates) [11], and antifungal [12] activities, as well as inhibitors of metastasis [13] and substances for the biodegradation of industrial waste [14].

The high potential of the practical application of exoproteins and metabolites of *B. pumilus* actualizes the search for methods of genetic editing of the genome of these bacteria to increase the yield of useful products. On the path of genetic research, an important stage is the delivery of heterogeneous DNA into bacilli cells. Methods of transformation, electroporation, and conjugation are most often used for this purpose. These methods are used with high efficiency for laboratory strains; however, strains isolated from their natural habitat have natural defense mechanisms against the introduction of foreign DNA (DNA methylation, nucleases, antibiotic secretion). In this connection, transformation (and

[⊠] I. V. Danilova Danilova146@mail.ru

¹ Department of Microbiology, Institute of Fundamental Medicine and Biology, Kazan (Volga Region) Federal University, Kazan 420008, Russia

other methods of transfer of exogenous DNA) into *B. pumilus* cells could have some setbacks [15].

For gram-positive bacteria, chemical treatment of cells to increase competence is an ineffective method compared to other protocols, since the composition of the gram-positive cell wall prevents the absorption of foreign DNA, despite its possibility of use [16–18]. The electroporation method, originally developed for the transfection of eukaryotic cells, was subsequently adapted to transform a wide variety of bacteria, including gram-negative *Pseudomonas, Salmonellae, Vibrios, Serratiae*, and *Shigellae* and gram-positive *Clostridia, Bacilli, Lactobacilli*, and *Enterococci* [19]. The efficiency of transformation by electroporation is reportedly 10–20-fold higher than chemical competence or heat shock [20].

In the process of selecting the transformation conditions, we faced the problem of stable resistance of *B. pumilus* 3–19 to the acceptance of foreign DNA. The chemical transformation methods described for other strains of bacilli are inapplicable to *B. pumilus* 3–19. In using the electroporation method, few studies have been conducted on the transformation of *B. pumilus*. However, results showed a lower efficiency as compared to *B. subtilis* cells [15, 21]. Shen et al. developed an efficient transformation method for the *B. pumilus* DX01 strain by optimizing the electroporation conditions [22]. In this study, we designed transformation conditions for *B. pumilus* 3–19 by electroporation based on the method described in [22].

The development of an effective transformation system for *B. pumilus* will pave the way for the genome editing of this microorganism, creating recombinant strains with altered (enhanced and suppressed) functions of significant proteins.

2 Materials and Methods

We used the strains *B. pumilus* 3-19 (strR) and *E. coli* DH5 α for the isolation of plasmids. Strains were obtained from the collection of the Agrobioengineering Research Laboratory (IFMaB, KFU). The plasmid vector pJOE9282.1 (9059 bps) used was acquired from Prof. Altenbuchner of the Institute of Industrial Genetics, Stuttgart University, Stuttgart, Germany.

LB agar and LB broth were used as nutrient media. The antibiotic kanamycin was added to the medium at a final concentration of 15 μ g/ml. For electroporation, a SOC1 recovery medium (2% tryptone, 0.5% yeast extract, 0.05% NaCl, 2.5 mmol/l KCl and 10% sucrose, pH 7.0) and electroporation buffer PEB1 (0.5 mol/l sucrose, 0.1 mmol/l KH₂ ²PO₄⁴/K₂HPO₄, pH 7.4) were used [Shen et al., 2013].

Isolation of plasmid DNA from *E. coli* cells was performed using the GeneJET Plasmid Miniprep Kit (Thermo Scientific). To isolate plasmids, 1-5 ml of an overnight culture of *E. coli* DH5 α was used according to protocol A, for a high copy number of plasmids. Centrifugation was carried out at 13,000 rpm.

Before plasmid DNA isolation from *B. pumilus* 3-19 strain, we harvested up to 2×10^9 bacterial cells in a 1.5- or 2-ml microcentrifuge tube by centrifugation for 10 min at 5000 g. Discarded the supernatant. Resuspended the pellet in 180 µl of Gram-positive bacteria lysis buffer (20 mM Tris–HCl, pH 8.0, 2 mM EDTA, 1.2% Triton X-100, add lysozyme to 20 mg/ml immediately before use). Incubated for 30 min at 37 °C. After these manipulations, the GeneJET Plasmid Miniprep Kit (Thermo Scientific) was used for isolation of plasmid DNA from *B. pumilus* 3–19 cells.

To obtain competent cells, an overnight culture of *B*. *pumilus* 3–19 strain was introduced into LB medium (at $OD_{600}=0.1$) and cultured at 37 °C with aeration (180 rpm) for 4 h (until $OD_{600}=0.6$).

B. pumilus 3–19 competent cells were cooled on ice for 15 min, and then the pellet was collected by centrifugation at $4000 \times g$ for 10 min and washed twice with an equal volume of electroporation buffer (PEB1). Finally, the bacterial cells were suspended in 800 µl of electroporation buffer; 200 µl of competent cells and 30 µl of plasmid were carefully mixed in a 1.5-ml Eppendorf centrifuge tube. The mixture was added to a precooled 2-mm-gap electroporation cuvette (Bio-Rad, USA) and kept on ice for 40 min before electroporation.

Electroporation was performed using MicroPulser Electroporation equipment (Bio-Rad). Aliquots were electroporated at a field strength of 12 kV/cm.

The transformed *B. pumilus* cells were immediately mixed with 800 μ l of SOC1 recovery medium and incubated for 4 h at 200 rpm and 37 °C for regenerative growth, and then plated on LB agar medium containing 15 μ g/ml kanamycin [22].

DNA electrophoresis was performed in 2% agarose gel in tris-acetate buffer. A 1-Kb kit from SibEnzyme (M11, Russia) was used as markers.

3 Results and Discussion

Competence in many bacterial species occurs in a short period at a certain stage of growth. For example, *Streptococcus* cells are electrocompetent in the early exponential growth phase, *B. subtilis* cells — at later stages of the logarithmic growth phase, *Haemophilus* cells — at a stationary growth stage [23–25]. Shen et al. showed that the maximum transformation frequency was achieved at an OD_{600} of 1.0 (after incubation for 4 h) in the exponential phase of *B. pumilus* DX01 [22]. When the *B. pumius* 3–19 strain was cultured in 1000-ml flasks, we obtained competent cells after 4 h (until $OD_{600} = 0.6$) (Fig. 1). In some





experiments, the *B. pumius* 3-19 strain culture reached a density of $OD_{600} = 1.0$ only after 8 h. At the same time, the efficiency of cell transformation under these growth conditions did not change.

A recovery medium with high osmolarity and optimized composition increases the frequency of electrotransformation of recipient cells, presumably because bacterial cells were osmotically or electrochemically sensitive after electroporation [26]. By using the PEB1 electroporation buffer and SOC1 recovery medium as recommended by Shen et al.,



Fig. 2 Colonies of *B. pumius* 3-19 transformants, grown on a selective medium with kanamycin. The eight indicated clones were chosen for further validation (by pDNA isolation)

we achieved a transformation efficiency of 56.3 transformants/ μ g DNA (Fig. 2). A mixture of competent cells and DNA resuspended in PEB1 was incubated on ice for 40 min before electroporation. The 10 min recommended by Shen et al. was insufficient for *B. pumilus* 3–19 cells. Subjected to Shen's protocol, the transformation of *B. pumius* 3–19 was unsuccessful.

High-voltage pulsed electric fields are widely used in the genetic transformation of bacteria [27]. Field strength affects the transformation rate, and for most prokaryotic cells, the corresponding electrical voltage range is from 6 to 14 kV [28, 29]. Shen et al. obtained the maximum transformation frequency at 12 kV/cm, which we adopted for the *B. pumilus* 3–19 strain. We also carried out the transformation of bacilli cells at an electric field voltage of 9 kV/cm. However, the transformation frequency was slightly lower. Confirmation of transformation was



Fig. 3 Electrophoresis of plasmids isolated from colonies of transformants. M, marker; K+, positive control (plasmid pJOE9282.1, isolated from *E. coli* DH5 α); 1–8, transformant colonies. Indicated by the corresponding numbers in Fig. 2

performed by the electrophoresis of pJOE9282.1 plasmids isolated from transformants (Fig. 3). All transformants obtained were observed to carry the vector pJOE9282.1. Thus, the transformation frequency for the *B. pumilus* 3–19 strain was increased by optimizing the electroporation conditions. Extension of the co-incubation time of competent cells and donor DNA before electroporation (for 40 min) was a critical parameter for the successful transformation of *B. pumilus* 3–19 cells.

Funding This work was supported by the RFBR grant (Russian Foundation for Basic Research) №19–08-00853 and Kazan Federal University Strategic Academic Leadership Program.

Declarations

Conflict of Interest The authors declare no competing interests.

Research Involving Humans and Animals Statement The experiments were carried out without involving humans or animals.

Informed Consent All authors consent to the publication of the article.

References

- Zhao, H.Y., & Feng, H. (2018). Engineering *Bacillus pumilus* alkaline serine protease to increase its low-temperature proteolytic activity by directed evolution. *BioMed Central Biotechnology*, *18*, 34. https://doi.org/10.1186/s12896-018-0451-0
- Baweja, M., Tiwari, R., Singh, P. K., Nain, L. & Shukla, P. (2016). An alkaline protease from *Bacillus pumilus* MP 27: Functional analysis of its binding model toward its applications as detergent additive. *Frontiers in microbiology*, *7*, 1195. https://doi.org/10. 3389/fmicb.2016.01195
- Huang Y., Lezyk M., Herbst F. A., Busk P. K., & Lange L. (2020). Novel keratinolytic enzymes, discovered from a talented and efficient bacterial keratin degrader. *Scientific reports*, 10, 1. https:// doi.org/10.1038/s41598-020-66792-2
- Koiv, V., Adamberg, K., Adamberg, S., Sumeri, I., Kasvandik, S., Kisand, V., Maivali, U., & Tenson, T. (2020). Microbiome of root vegetables-A source of gluten-degrading bacteria. *Applied microbiology and biotechnology*, *104*(20), 8871–8885. https://doi. org/10.1007/s00253-020-10852-0
- Mitrofanova, O., Mardanova, A., Evtugyn, V., Bogomolnaya, L., & Sharipova, M. (2017). Effects of *Bacillus* serine proteases on the bacterial biofilms. *BioMed research international*. https://doi. org/10.1155/2017/8525912
- Lee, N. K., Kim, W. S., & Paik, H. D. (2019). *Bacillus* strains as human probiotics: Characterization, safety, microbiome, and probiotic carrier. *Food science and biotechnology*, 28(5), 1297–1305. https://doi.org/10.1007/s10068-019-00691-9
- Masood, S., Zhao, X. Q., & Shen, R. F. (2019). *Bacillus pumilus* increases boron uptake and inhibits rapeseed growth under boron supply irrespective of phosphorus fertilization. *AoB Plants*, *11*, 4. https://doi.org/10.1093/aobpla/plz036
- Zhang, N., Wang, L., & Wei, Y. (2020). Effects of *Bacillus amyloliquefaciens* and *Bacillus pumilus* on rumen and intestine morphology and microbiota in weanling Jintang black goat.

🖉 Springer

Animals (Basel), 10(9), 1604. https://doi.org/10.3390/ani10 091604

- Opriessnig, T., Karuppannan, A. K., Beckler, D., Ait-Ali, T., Cubas-Atienzar, A., & Halbur, P. G. (2019). *Bacillus pumilus* probiotic feed supplementation mitigates *Lawsonia intracellularis* shedding and lesions. *Veterinary research*, 50(1), 85. https://doi. org/10.1186/s13567-019-0696-1
- Zhou, S. Y., Hu, Y. J., Meng, F. C., Qu, S. Y., Wang, R., Andersen. R. J., Liao, Z. H., & Chen, M. (2018). Bacillamidins A-G from a marine-derived *Bacillus pumilus*. *Marine drugs*, 16, 9. https:// doi.org/10.3390/md16090326
- Garcia-Ramon, D. C., Berry, C., Tse, C., Fernández-Fernández, A., Osuna, A., & Vílchez, S. (2018). The parasporal crystals of Bacillus pumilus strain 15.1: A potential virulence factor? *Microbial biotechnology*, *11*(2), 302–316. https://doi.org/10.1111/1751-7915.12771
- Gu, X., Sun, J., Cui, Y., Wang, X., & Sang, Y. (2019). Biological degradation of aflatoxin M₁ by *Bacillus pumilus* E-1–1–1. *Microbiology Open.* 8, 3. https://doi.org/10.1002/mbo3.663
- Zelenikhin, P. V., Ead, Mohamed I. S., Nadyrova, A. I., Sirotkina, A. A., Ulyanova, V. V., Mironova, N. L., Mitkevich, V. A., Makarov, A. A., Zenkova, M. A., Ilinskaya, O. N. (2020). Bacillus pumilus ribonuclease inhibits migration of human duodenum adenocarcinoma HuTu 80 cells. *Molekuliarnaia biologiia (Mosk)*, 54(1), pp.146–152. Russian. https://doi.org/10.31857/S002689842 0010176
- Bonifer, K. S., Wen, X., Hasim, S., Phillips, E. K., Dunlap, R. N., Gann, E. R., DeBruyn, J. M., & Reynolds, T. B. (2019). *Bacillus pumilus* B12 degrades polylactic acid and degradation is affected by changing nutrient conditions. *Frontiers in microbiology*, 10, 2548. https://doi.org/10.3389/fmicb.2019.02548
- Toimentseva, A. A., & Altenbuchner, D. (2018). Obtaining Bacillus pumilus deletion mutants for proteinase genes based on CRISPR-CAS9 technology. International research journal, 11(77), 136–142. Russian.
- Sirajuddin, S. A., & Sundram, S. (2020). Evaluation of two transformation protocols and screening of positive plasmid introduction into Bacillus cereus EB2, a gram-positive bacterium using qualitative analyses. *Brazilian journal of microbiology*, 51(3), 919–929. https://doi.org/10.1007/s42770-020-00241-0
- Rahmer, R., Morabbi Heravi, K., & Altenbuchner, J. (2015). Construction of a super-competent *Bacillus subtilis* 168 using the PmtlA-comKS inducible cassette. *Frontiers in microbiology.*, 6, 1431. https://doi.org/10.3389/fmicb.2015.01431
- Vojcic, L., Despotovic, D., Martinez, R., Maurer, K.-H., & Schwaneberg, U. (2012). An efficient transformation method for *Bacillus subtilis* DB104. *Applied microbiology and biotechnol*ogy., 94(2), 487–493. https://doi.org/10.1007/s00253-012-3987-2
- Gonzales, M. F., Brooks, T., Pukatzki, S. U. & Provenzano, D. (2013). Rapid protocol for preparation of electrocompetent *Escherichia coli* and *Vibrio cholerae. Journal of visualized experiments: JoVE.* 80, 50684. https://doi.org/10.3791/50684
- Dower, W. J., Miller, J. F., & Ragsdale, C. W. (1988). High efficiency transformation of E. coli by high voltage electroporation. *Nucleic Acids Research.*, 16(13), 6127–6145.
- Shao, H., Cao, Q., Zhao, H., Tan, X., & Feng, H. (2015). Construction of novel shuttle expression vectors for gene expression in *Bacillus subtilis* and *Bacillus pumilus*. *The Journal of general and applied microbiology.*, 61(4), 124–131. https://doi.org/10.2323/jgam.61.124
- Shen, X., Chen, Y., Liu, T., Hu, X., & Gu, Z. (2013). Development of a high-efficient transformation system of *Bacillus pumilus* strain DX01 to facilitate gene isolation via gfp-tagged insertional mutagenesis and visualize bacterial colonization of rice roots. *Folia microbiologica.*, 58, 409–417. https://doi.org/10.1007/s12223-013-0223-0

- Black, H. F., Mastromatteo, S., Sinha, S., Ehrlich, R. L., Nislow, C., Mell, J. C. & Redfield, R. J. (2020). A competence-regulated toxin-antitoxin system in *Haemophilus influenza*. *PLoS One*, *15*, 1. https://doi.org/10.1371/journal.pone.0217255
- Blokesch, M. (2016). Natural competence for transformation. *Current biology: CB*, 26(23), 3255. https://doi.org/10.1016/j.cub. 2016.11.023
- Perry, D., & Kuramitsu, H. K. (1981). Genetic transformation of Streptococcus mutans. Infection and immunity, 32, 1295–1297. https://doi.org/10.1128/iai.32.3.1295-1297.1981
- Haynes, J. A., & Britz, M. L. (1990). The effect of growth conditions of *Corynebacterium glutamicum* on the transformation frequency obtained by electroporation. *Journal of general microbiology*, 136, 255–263.
- Mcintyre, D. A., & Harlander, S. K. (1989). Genetic transformation of intact *Lactococcus lactis subsp.* lactis by high-voltage electroporation. *Applied and environmental microbiology*, 55, 604–610.

- Aune, T. E. V., & Aachmann, F. L. (2010). Methodologies to increase the transformation efficiencies and the range of bacteria that can be transformed. *Applied microbiology and biotechnology*, 85, 1301–1313. https://doi.org/10.1007/s00253-009-2349-1
- Yang, M. M., Zhang, W. W., Bai, X. T., Li, H. X., & Cen, P. L. (2010). Electroporation is a feasible method to introduce circularized or linearized DNA into *B. subtilis* chromosome. *Molecular biology reports*, 37, 2207–2213. https://doi.org/10.1007/ s11033-009-9704-2

Publisher's Note Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.