# A method for screening the suppressor genes of siRNA and piRNA pathways using cultured silkworm cells

Haruka Sugiyama<sup>1\*</sup>, Susumu Katsuma<sup>1§\*</sup>

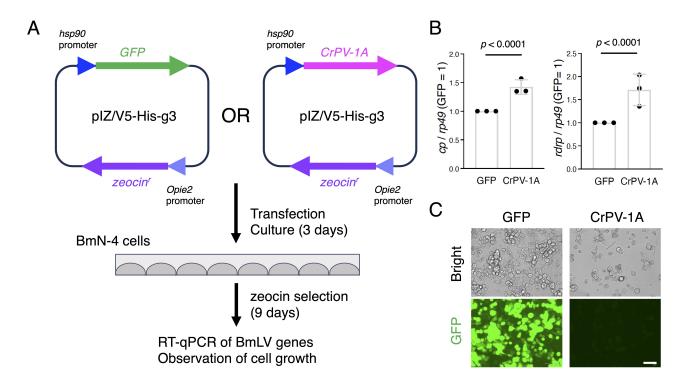
<sup>1</sup>Department of Agricultural and Environmental Biology, Graduate School of Agricultural and Life Sciences, The University of Tokyo, Tokyo, Japan

<sup>§</sup>To whom correspondence should be addressed: skatsuma@g.ecc.u-tokyo.ac.jp

<sup>\*</sup>These authors contributed equally.

# Abstract

The BmN-4 cell line originates from the ovaries of silkworm, *Bombyx mori*, and possesses endogenous small interfering RNA (siRNA) and PIWI-interacting RNA (piRNA) pathways. BmN-4 cells are latently infected with *Bombyx mori latent virus* (BmLV), an RNA virus whose replication is strictly controlled by both siRNA and piRNA pathways. Knockdown or knockout of the core factors of these two small RNA pathways increases BmLV RNA amount, which in turn inhibits cell growth. Here, we used the known RNAi suppressor CrPV-1A to assess whether the BmN-4 cell line can be used for screening the suppressors of siRNA and piRNA pathways.



#### Figure 1. A screening method of the suppressors of small RNA pathways using cultured silkworm cells:

(A) An experimental flow.

(B) RT-qPCR results for the BmLV *cp* and *rdrp* genes. BmN-4 cells were transfected with a plasmid expressing GFP- or CrPV-1A, cultured in zeocin-containing medium, and then subjected to RT-qPCR of two BmLV genes. The *cp* and *rdrp* 



#### 9/15/2023 - Open Access

mRNA levels were normalized to that of *B. mori rp49*. The data are shown as means ± standard deviation of three independent experiments. *p*-values were calculated via one sample *t* test (two-tailed).

(C) Fluorescence microscopy of BmN-4 cells stably expressing GFP or CrPV-1A. The scale bar represents 100 µm.

## Description

Cultured cell lines are an important biological resource for producing recombinant proteins and evaluating gene function. The silkworm *Bombyx mori* cell line BmN-4 is a well-known ovary-derived cell line (Grace, 1967) that has been routinely used for protein production by a baculovirus expression system (Maeda et al. 1985). In 2009, our group identified PIWI proteins and PIWI-interacting RNAs (piRNAs) that were endogenously expressed in BmN-4 cells. This was the first discovery of a cultured cell line that expressed PIWI/piRNA complexes (Kawaoka et al. 2009). Many groundbreaking studies involving piRNA factors used this piRNA-producing cell line (Kawaoka et al. 2011; Izumi et al. 2020; Matsumoto et al. 2016), and it has become a valuable resource for small RNA research (Tsukioka et al. 2006; Kawaoka et al. 2009).

*Bombyx mori latent virus* (formerly known as *Bombyx mori macula-like virus*, BmMLV) is a positive, single-stranded insect RNA virus that is closely related to plant maculaviruses. BmLV was first discovered in BmN-4 cells, and has been found to infect almost all *B. mori*-derived cultured cell lines (Katsuma et al. 2005). Surprisingly, BmLV accumulates to extremely high levels (approximately 15% of total mRNA) in BmN-4 cells (Katsuma et al. 2018). Knockdown or knockout of the core biogenesis genes for either small interfering RNA (siRNA) and piRNA revealed that disruption of these small RNA pathways results in increased BmLV accumulation and inhibition of BmN-4 cell growth (Katsuma et al. 2018; Katsuma et al. 2021). These findings show that the siRNA and piRNA pathways function cooperatively to silence BmLV RNA and that both pathways are required for the normal growth of BmLV-infected silkworm cells.

In this study, we tested whether BmN-4 cells can be used to assess the suppressor activity of foreign genes by measuring the expression levels of BmLV genes and observing the degree of inhibition of cell growth. We selected the cricket paralysis virus 1A protein (CrPV-1A) gene as the suppressor gene. CrPV-1A has been found to inhibit Ago2-dependent RNAi via blocking the initial target searching by Ago2-RISC (Watanabe et al. 2017; Nayak et al. 2018).

First, we cloned *CrPV-1A* into the vector, pIZ/His-V5-g3, so that the cloned gene would be expressed under the control of the *B. mori hsp90* promoter (Hirota et al., 2021). Next, pIZ/His-V5-g3-CrPV-1A or pIZ/His-V5-g3-GFP (i.e., control vector) was transfected into BmN-4 cells and treated with zeocin from 3 days after transfection (Fig. 1A). Selection was conducted for 9 days, after which cells were photographed and then collected for RNA isolation (Fig. 1A). Reverse transcription-quantitative polymerase chain reaction (RT-qPCR) experiments revealed that CrPV-1A expression increased the mRNA levels of two BmLV genes, *coat protein* (*cp*) and *RNA-dependent RNA polymerase* (*rdrp*) (Fig. 1B), suggesting that CrPV-1A may block one or both of the siRNA and piRNA pathways. In addition, CrPV-1A expression also strongly inhibited the cell growth of BmN-4 cells (Fig. 1C). Taken together, these results demonstrate that BmN-4 cells can be used for a simple screening system to identify novel putative suppressors of the siRNA and/or piRNA pathways.

## Methods

#### Cell line

BmN-4 cells (provided by Chisa Yasunaga-Aoki, Kyushu University, and maintained in our laboratory) (Maeda et al. 1985) were cultured at 27°C in IPL-41 medium (Applichem) supplemented with 10% fetal bovine serum.

#### Plasmid construction

A CrPV-1A fragment was amplified from pCold II-CrPV-1A (provided by Yukihide Tomari) (Watanabe et al. 2017). This fragment was then cloned into the vector pIZ/V5-His-g3 (Hirota et al., 2021) using the In-Fusion HD Cloning Kit (Clontech). pIZ/V5-His-g3-GFP (Hirota et al., 2021) was used as a control.

#### Transfection and generation of stably transfected BmN-4 cells

BmN-4 cells (2 × 10<sup>5</sup> cells per 35 mm dish) were transfected with 1  $\mu$ g of pIZ/V5-His-g3-GFP or pIZ/V5-His-g3-GFP-CrPV-1A using FuGENE HD (Promega). Three days after transfection, zeocin (InvivoGen, final concentration of 500  $\mu$ g/mL) was added to the medium (Kawaoka et al. 2009). Nine days after drug selection, cells were observed using a FLoid<sup>TM</sup> cell imaging station (Life Technologies). Thereafter they were collected for RNA extraction.

#### RT-qPCR

Total RNA was isolated using TRI Reagent® (Sigma-Aldrich) and then subjected to reverse transcription with avian myeloblastosis virus reverse transcriptase and an oligo-dT primer (TaKaRa). RT-qPCR was performed using a KAPA SYBR



#### 9/15/2023 - Open Access

FAST qPCR kit (Kapa Biosystems) and the specific primers. The expression values were calculated using the  $2^{-\Delta\Delta Ct}$  method.

## Reagents

PCR primers for CrPV-1A CrPV1A-f: TACCGAGCTCGGATCatgtcttttcaacaaacaacaacaacacc CrPV1A-r: GCCACTGTGCTGGATctagaaggctctgcattcatcattac qPCR primers for BmLV *cp* coat-2F: TCCTCTCGCATTACTATTGG coat-2R: ATGGAGCCTCTGATGACAAC qPCR primers for BmLV *rdrp* rdrp-2F: TCTCTCATGAAATCAGCACC rdrp-2R: TCACGATATGGTTTGAGATG qPCR primers for *B. mori rp49* rp49-F: CCCAACATTGGTTACGGTTC

rp49-R: GCTCTTTCCACGATCAGCTT

**Acknowledgements:** We thank H. Hikida and N. Matsuda-Imai for technical assistance, C. Yasunaga-Aoki for providing BmN-4 cells, and Y. Tomari for providing pCold II-CrPV-1A.

### References

Grace TD. 1967. Establishment of a line of cells from the silkworm Bombyx mori. Nature 216(5115): 613. PubMed ID: <u>4868367</u>

Hirota K, Matsuda-Imai N, Kiuchi T, Katsuma S. 2021. Characterization of nuclear localization signal in Ostrinia furnacalis Masculinizer protein. Arch Insect Biochem Physiol 106(3): e21768. PubMed ID: <u>33644912</u>

Izumi N, Shoji K, Suzuki Y, Katsuma S, Tomari Y. 2020. Zucchini consensus motifs determine the mechanism of pre-piRNA production. Nature 578(7794): 311-316. PubMed ID: <u>31996847</u>

Katsuma S, Tanaka S, Omuro N, Takabuchi L, Daimon T, Imanishi S, et al., Shimada T. 2005. Novel macula-like virus identified in Bombyx mori cultured cells. J Virol 79(9): 5577-84. PubMed ID: <u>15827172</u>

Katsuma S, Kawamoto M, Shoji K, Aizawa T, Kiuchi T, Izumi N, et al., Iwanaga M. 2018. Transcriptome profiling reveals infection strategy of an insect maculavirus. DNA Res 25(3): 277-86. PubMed ID: <u>29360973</u>

Katsuma S, Shoji K, Suzuki Y, Kiuchi T. 2021. CRISPR/Cas9-mediated mutagenesis of Ago2 and Siwi in silkworm cultured cells. Gene 768: 145314. PubMed ID: <u>33220342</u>

Kawaoka S, Hayashi N, Suzuki Y, Abe H, Sugano S, Tomari Y, Shimada T, Katsuma S. 2009. The Bombyx ovary-derived cell line endogenously expresses PIWI/PIWI-interacting RNA complexes. RNA 15(7): 1258-64. PubMed ID: <u>19460866</u>

Kawaoka S, Izumi N, Katsuma S, Tomari Y. 2011. 3' end formation of PIWI-interacting RNAs in vitro. Mol Cell 43(6): 1015-22. PubMed ID: <u>21925389</u>

Maeda S, Kawai T, Obinata M, Fujiwara H, Horiuchi T, Saeki Y, Sato Y, Furusawa M. 1985. Production of human alphainterferon in silkworm using a baculovirus vector. Nature 315(6020): 592-4. PubMed ID: <u>2989694</u>

Matsumoto N, Nishimasu H, Sakakibara K, Nishida KM, Hirano T, Ishitani R, et al., Nureki O. 2016. Crystal Structure of Silkworm PIWI-Clade Argonaute Siwi Bound to piRNA. Cell 167(2): 484-497.e9. PubMed ID: <u>27693359</u>

Nayak A, Kim DY, Trnka MJ, Kerr CH, Lidsky PV, Stanley DJ, et al., Andino R. 2018. A Viral Protein Restricts Drosophila RNAi Immunity by Regulating Argonaute Activity and Stability. Cell Host Microbe 24(4): 542-557.e9. PubMed ID: <u>30308158</u>

Tsukioka H, Takahashi M, Mon H, Okano K, Mita K, Shimada T, et al., Kusakabe T. 2006. Role of the silkworm argonaute2 homolog gene in double-strand break repair of extrachromosomal DNA. Nucleic Acids Res 34(4): 1092-101. PubMed ID: <u>16478716</u>



#### 9/15/2023 - Open Access

Watanabe M, Iwakawa HO, Tadakuma H, Tomari Y. 2017. Biochemical and single-molecule analyses of the RNA silencing suppressing activity of CrPV-1A. Nucleic Acids Res 45(18): 10837-10844. PubMed ID: <u>28977639</u>

Funding: This work was supported by JSPS KAKENHI 17H06431.

**Author Contributions:** Haruka Sugiyama: investigation, writing - review editing, formal analysis. Susumu Katsuma: conceptualization, formal analysis, funding acquisition, investigation, methodology, supervision, writing - original draft.

#### Reviewed By: Anonymous

History: Received August 14, 2023 Revision Received August 30, 2023 Accepted August 29, 2023 Published Online September 15, 2023 Indexed September 29, 2023

**Copyright:** © 2023 by the authors. This is an open-access article distributed under the terms of the Creative Commons Attribution 4.0 International (CC BY 4.0) License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.

**Citation:** Sugiyama, H; Katsuma, S (2023). A method for screening the suppressor genes of siRNA and piRNA pathways using cultured silkworm cells. microPublication Biology. <u>10.17912/micropub.biology.000953</u>