

# **Virus-host protein-protein interactions of mycobacteriophage Giles**

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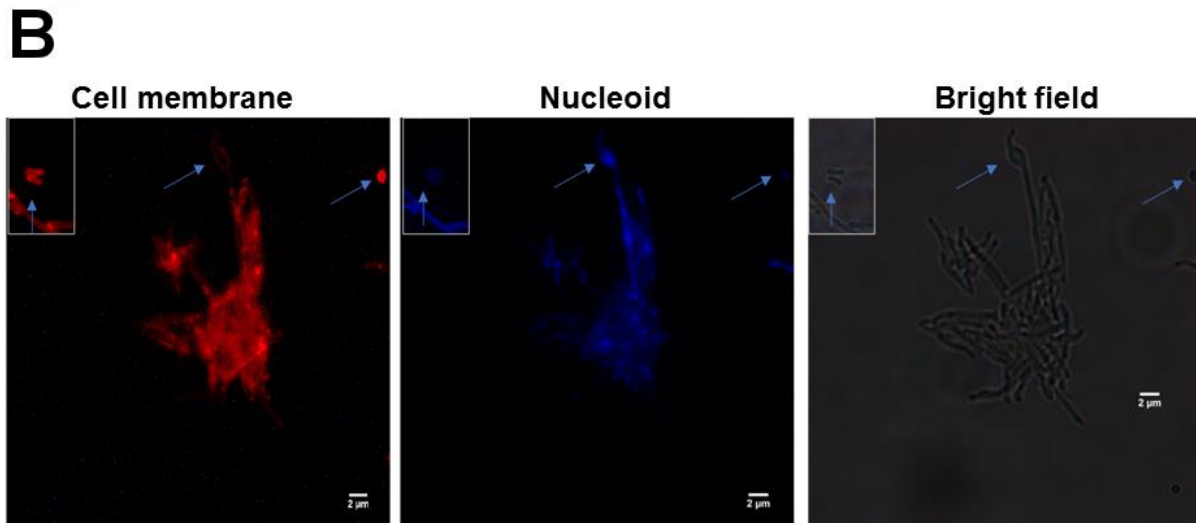
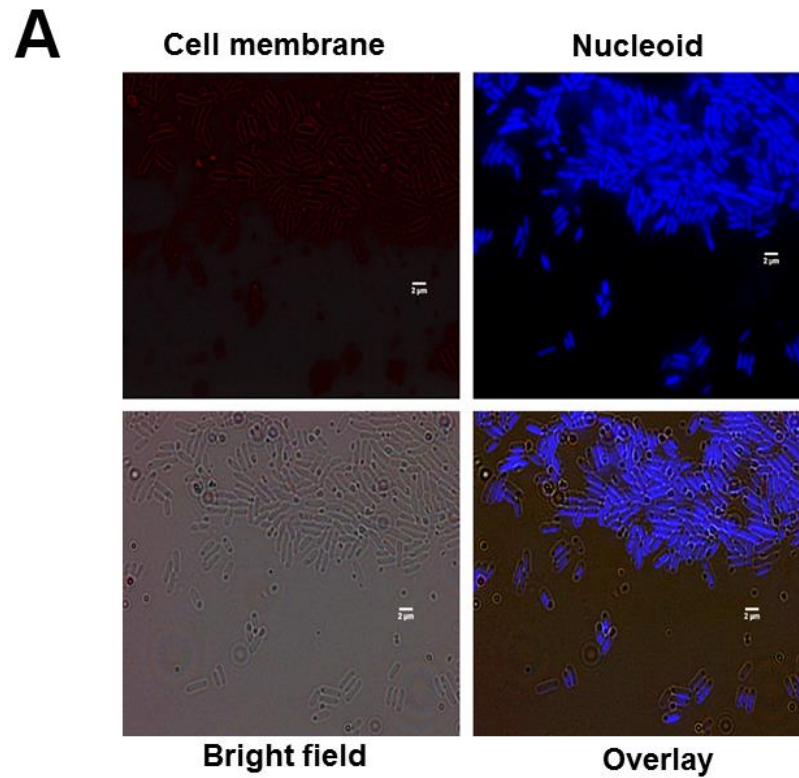
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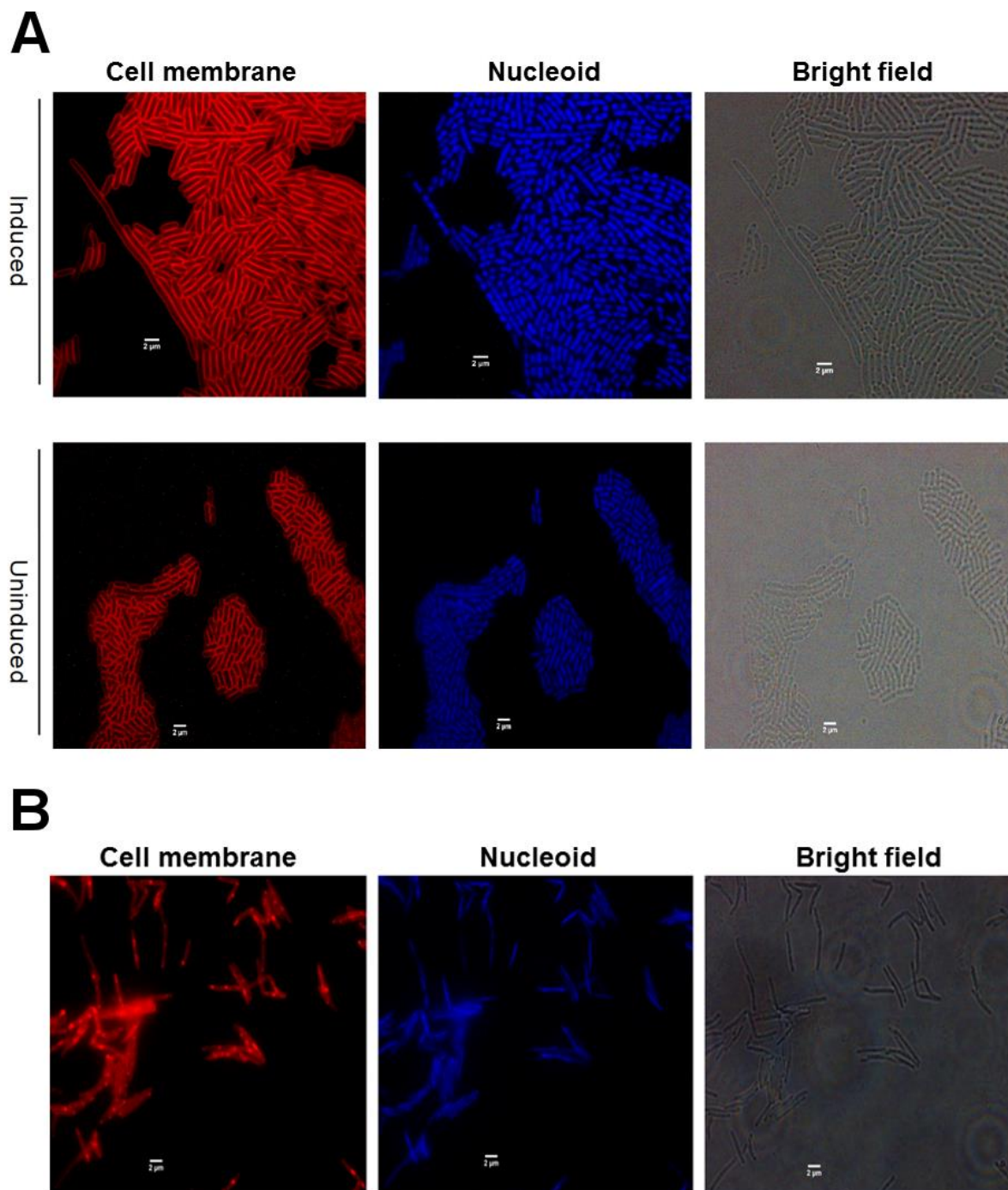
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## Supplementary Figures and Tables

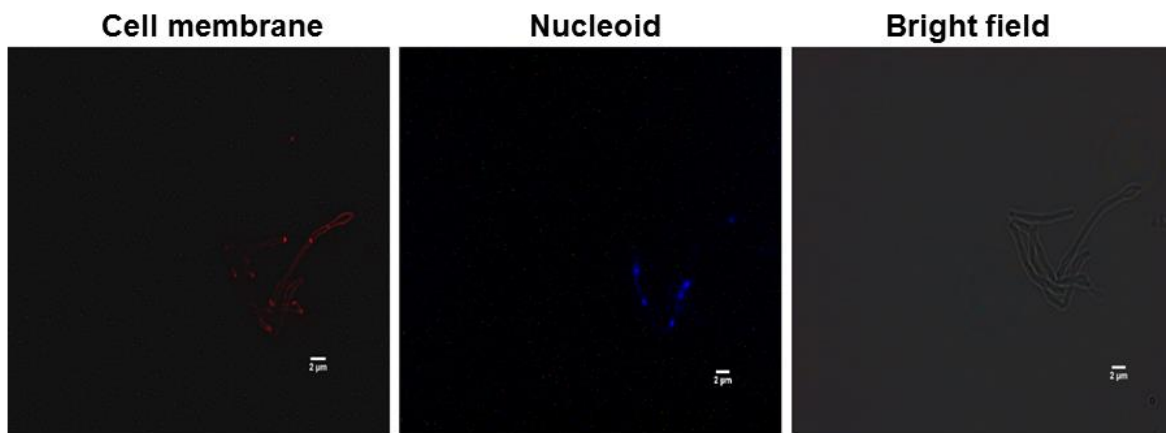
**Figures S1- S4** The phenotypic alterations of natural and heterologous hosts upon overexpression of Giles proteins. Giles Gp17 in *E. coli* (**Fig S1 A**) and *M. smegmatis* (**Fig S1 B**; **arrow indicates formation of small cells**); Gp47 in *E. coli* (**Fig S2 A**) and *M. smegmatis* (**Fig S2 B**); Gp54 in *M. smegmatis* (**Fig S3**) and Gp64 in *E. coli* (**Fig S4 top**) and *M. smegmatis* (**Fig S4 bottom**).



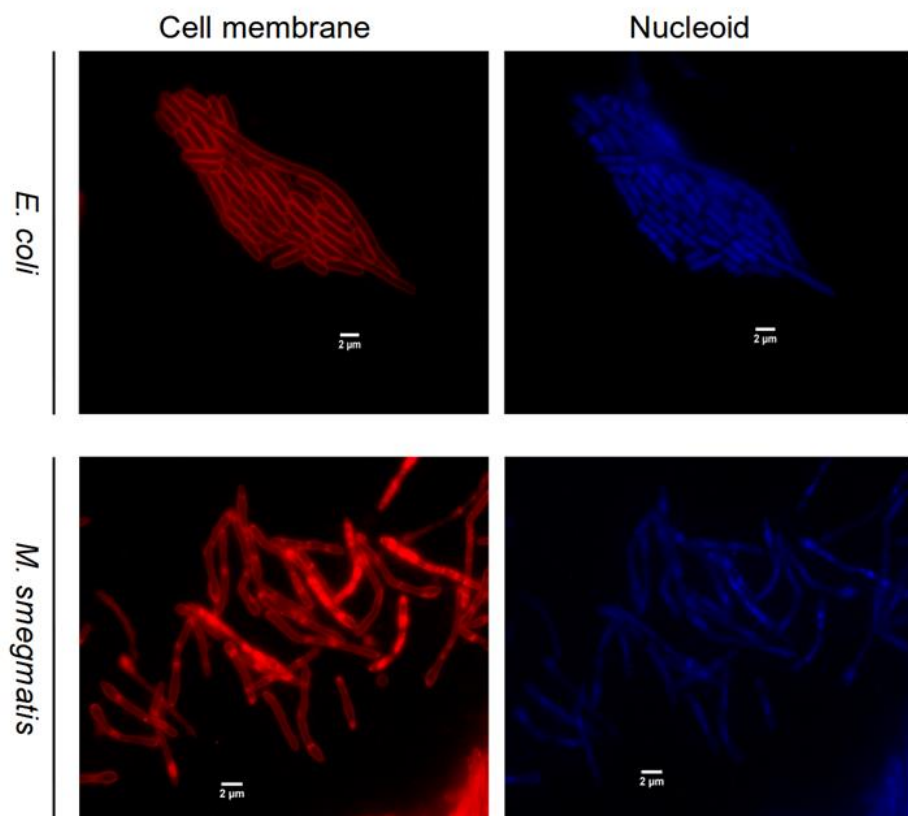
**Supplementary Figure S1.** The phenotypic alterations of natural and heterologous hosts upon overexpression of Giles proteins. Giles Gp17 in *E. coli* (**Fig S1 A**) and *M. smegmatis* (**Fig S1 B**; arrow indicates formation of small cells).



**Supplementary Figure S2.** The phenotypic alterations of natural and heterologous hosts upon overexpression of Giles proteins. Giles Gp47 in *E. coli* (Fig S2 A) and *M. smegmatis* (Fig S2 B).



**Supplementary Figure S3.** The phenotypic alterations of *M. smegmatis* upon overexpression of Giles protein Gp54.



**Supplementary Figure S4.** The phenotypic alterations of natural and heterologous hosts upon overexpression of Giles proteins. Giles Gp64 in *E. coli* (**top**) and *M. smegmatis* (**bottom**).

**Table S1.** The sequences of the host interactors for Giles proteins with other details, are shown here. Separate Excel spreadsheet.

**Table S2.** List of oligonucleotides used in this study.

<b>Oligo Name</b>	<b>Sequence</b>
MSMEG_3746A	GCACGGATCGGGAACCGTCC
MSMEG_3746B	TGGTGAGGGAGATGAGGTCTGAAGAAGCATCCTCCGTGACGAAGGG
MSMEG_3746C	GTTGAGGTGTGAGGTGTGCTGAAGGCTGGAGAAATCCGACGTCCGTG
MSMEG_3746D	TGGCGGTATTGGTACACCAGTACCAC
MSMEG_4430A	CCAGCGATACAAAAGGTTTGAGGAAGATG
MSMEG_4430B	TGGTGAGGGAGATGAGGTCTGAAGGTCCGAACTCCACGAATTGAATGAATCC
MSMEG_4430C	GTTGAGGTGTGAGGTGTGCTGAAGGGTCGTGAACTCGGCTTGCTC
MSMEG_4430D	GGAGGATGTCCGATCGGTTGC
MSMEG_5773A	GTCGGCGAGGAACTGGGATGG
MSMEG_5773B	TGGTGAGGGAGATGAGGTCTGAAGGGCTTCTCCTGGTAGCTTCGGC
MSMEG_5773C	GTTGAGGTGTGAGGTGTGCTGAAGTCAAATCATTCTCCGGTGACAGGAG
MSMEG_5773D	CGACGCCTACACCGTCACACC