

Regular paper

Chlamydial genomic MinD protein does not regulate plasmiddependent genes like Pgp5

Yina Sun^{1,2}, Jie Kong^{1,3}, Jingyue Ma^{1,3}, Manli Qi^{1,3}, Ying Zhang^{1,3}, Long Han^{1,3}, Quanzhong Liu^{1,3} and Yuanjun Liu^{1,3}

¹Department of Dermatovenereology, Tianjin Medical University General Hospital, Tianjin 300052, China; ²Key Laboratory of Hormones and Development (Ministry of Health), Tianjin Key Laboratory of Metabolic Diseases, Tianjin Metabolic Diseases Hospital and Tianjin Institute of Endocrinology, Tianjin Medical University, Tianjin 300070, China; ³Tianjin Neurological Institute, Key Laboratory of Post-neurotrauma Neuro-repair

and Regeneration in Central Nervous System, Ministry of Education and Tianjin City, Tianjin 300052, China

Chlamydia has a unique intracellular developmental cycle, which has hindered the single protein function study of Chlamydia. Recently developed transformation system of Chlamydia has greatly advanced the chlamydial protein's function research and was used to find that a chlamydial plasmid-encoded Pgp5 protein can down-regulate plasmid-dependent genes. It is assumed, that chlamydial genomic MinD protein has a similar function to Pgp5. However, it is unknown whether MinD protein regulates the same plasmid-dependent genes. We replaced pgp5 gene in the shuttle vector pGFP::CM with minD gene of C. trachomatis (CT0582) or C. muridarum (TC0871). The recombinant plasmid was transformed into plasmid-free organisms-CMUT3 and gRT-PCR was used to detect the transcription level of plasmid-encoded and -dependent genes in these pgp5 deficient organisms. As a readout, GlgA, one of the plasmid-regulated gene products was detected by immunofluorescence assay. After recombination, transformation and plaque purification, the stable transformants CT0582R and TC0871R were generated. In these transformants, the plasmid-dependent genes were upregulated, alike in the pgp5 premature stop mutant and pgp5 replacement with mCherry mutant. GlgA protein level was also increased in all pgp5 mutants, including CT0582R and TC0871R. Thus, our study showed that genomic MinD protein had different function than Pgp5, which was useful for further understanding the chlamydiae.

Key words: Chlamydia, MinD, Pgp5, Transformation system

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e-mail: liuyuanjun1980@126.com

Abbreviations: ÉB, Elementary body; RB, Reticular body; PCR, polymerase chain reaction; C. *trachomatis, Chlamydia trachomatis;* C. *muridarum, Chlamydia muridarum*; qRT-PCR, quantitative real-time polymerase chain reaction; GFP, green fluorescent protein; CPAF, chlamydial protease-like activity factor; WT, wild type; PBS, phosphate buffer saline; CT0582R, CMUT3-pGFP::CM CT0582Rpgp5; TC0871R, CMUT3-pGFP::CM TC0871Rpgp5; pgp5S, the pgp5 premature stop mutant organism; mCherryR, the pgp5 replacement with mCherry mutant organism

| Plasmids | PCR | Forward primer | Reverse primer | |
|----------|---------|----------------------------------|------------------------------|--|
| _ | product | | | |
| CT0582 | F1 | cttgtcgtcatcgtccttgtagtccatcgtta | | |
| Replace | | gagtaaaacatctattctgccttag | ttagagaatcgttctctttgag | |
| pgp5 | F2 | cacaaacaagacttttcccagaggacact | t ctcaaagagaacgattctctaa | |
| | F3(CT05 | cggggcggtggtgggtggtggcggc | tctgggaaaagtcttgtttgtgttagat | |
| | 82) | ggatctaaaacaatcgctgttaatagtttca | gttccttaacaaaaatagcagttc | |
| | | aag | | |
| | F4(CT05 | gactacaaggacgatgacgacaagtctg | tctgggaaaagtcttgtttgtgttagat | |
| | 82) | gtggcggtggctcgggcggtggtgggtc | gttccttaacaaaaatagcagttc | |
| | | ggg | | |
| TC0871 | F1 | cttgtcgtcatcgtccttgtagtccatcgtta | | |
| Replace | | gagtaaaacatctattctgccttag | ttagagaatcgttctctttgag | |
| pgp5 | F2 | cacaaacaagacttttcccagaggacact | ctcaaagagaacgattctctaa | |
| | F3(TC08 | tctgggaaaagtcttgtttgtgttagatgtc | tctgggaaaagtcttgtttgtgttagat | |
| | 71) | ccttaacaaaaatagcagttc | gtcccttaacaaaaatagcagttc | |
| | F4(TC08 | gactacaaggacgatgacgacaagTcct | tctgggaaaagtcttgtttgtgttagat | |
| | 71) | ggtggcggtggctcgggcggtggtgggt | gtcccttaacaaaaatagcagttc | |
| | | cgggt | | |

Supplementary Table 1: Primers for generation of the recombinant plasmids

Supplementary Table 2: Primers for PCR screen to detect the insertion fragment in the recombinant plasmids

| | Forward primer | Reverse primer |
|-------------------------------------|---------------------|-----------------------|
| Primers for detecting the insertion | aaaacaatcgctgttaata | Gagaaacggattcctgattt |
| fragment CT0582 | gtttcaaag | tatttaaaaaagacgctagct |
| Primers for detecting the insertion | aaaacaatcgctgttaaca | Gagaaacggattcctgattt |
| fragment TC0871 | gttttaaagg | tatttaaaaaagacgctagct |