

[S9-5]

The *Methanothermobacter thermoautotrophicus* MCM Helicase is Active as a Hexameric Ring

Jae-Ho Shin^{1*} and Zvi Kelman²

¹*School of Applied Bioscience, Kyungpook National University, Daegu 702-701*

²*Center for Advanced Research in Biotechnology, University of Maryland Biotechnology Institute,
9600 Gudelsky Drive, Rockville, MD 20850, USA*

Replicative DNA helicases play an essential role during chromosomal DNA replication by separated the two strands of the duplex DNA providing the single-stranded (ss) DNA for the polymerase. In bacteria, the replicative helicase is the DnaB protein which forms hexameric rings (1, 2). The eukaryotic minichromosome maintenance (MCM) complex is a family of six related polypeptides (Mcm2-7), each of which is essential for cell viability. *In vivo* and *in vitro* studies have shown that in addition to forming a heterohexamers revealed the existence of several additional MCM complexes (3). It was shown that the Mcm4,6,7 heterotrimer forms hexameric rings (4) that possess DNA helicase activity. The enzyme also forms double-hexamers in the presence of forked DNA substrate (5).

The structure of the archaeal MCM complex is unclear. The MCM homologues of *Sulfolobus solfataricus* (6-8), *Archaeoglobus fulgidus*, *A. pernix* (9) and *Thermoplasma acidophilum* (10) form hexamers in solution. The *Methanothermobacter thermoautotrophicus* (Mth) enzyme appears to form dodecamers (11-13), and a dodecameric structure was also suggested by the crystal structure of the N-terminal portion of the protein (14) and by an electron microscopic (EM) analysis of the full-length enzyme (15-17). However, other EM reconstruction studies of the *M. thermoautotrophicus* enzyme showed hexameric (18), heptameric (19), open rings (15) and filamentous structures (20). Those conflicted structures of Mth MCM are summarized in Fig. 1.

The conflicting data regarding the structure of the *M. thermoautotrophicus* MCM hinder progress in elucidating the mechanism of enzyme action during initiation and elongation at the replication fork. The knowledge of the active form of the enzyme is needed as to form a working and testable hypothesis regarding the helicase activity during the initiation and elongation phases. One would expect, for example, that a hexamer would work differently than a filament. Therefore, a study was initiated to determine the

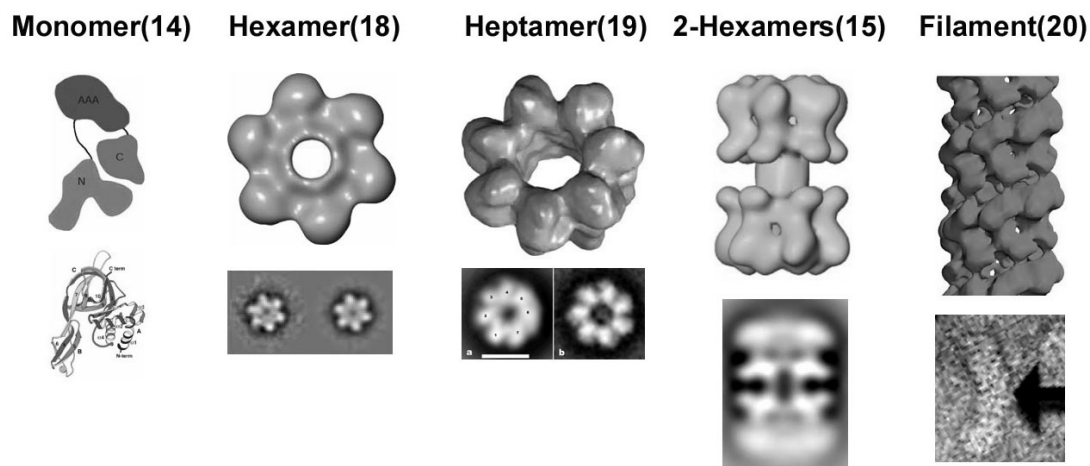


Figure 1. Structural polymorphism of MthMCM

active form of the enzyme using different substrates and experimental conditions. Using a combination of biochemical and structural analysis it is shown here that the *M. thermoautotrophicus* MCM helicase is active as a hexamer. The scientific impact of this result on DNA replication model is discussed.

References

1. Yu, X., Jezewska, M.J., Bujalowski, W. and Egelman, E.H. (1996) The hexameric *E. coli* DnaB helicase can exist in different Quaternary states. *J Mol Biol*, **259**, 7-14.
2. San Martin, M.C., Stamford, N.P., Dammerova, N., Dixon, N.E. and Carazo, J.M. (1995) A structural model for the *Escherichia coli* DnaB helicase based on electron microscopy data. *J Struct Biol*, **114**, 167-176.
3. Tye, B.K. and Sawyer, S.L. (2000) The Hexameric Eukaryotic MCM Helicase: Building Symmetry from Nonidentical Parts. *J Biol Chem*, **275**, 34833-34836.
4. Sato, M., Gotow, T., You, Z., Komamura-Kohno, Y., Uchiyama, Y., Yabuta, N., Nojima, H. and Ishimi, Y. (2000) Electron microscopic observation and single-stranded DNA binding activity of the Mcm4,6,7 complex. *J Mol Biol*, **300**, 421-431.
5. Lee, J.-K. and Hurwitz, J. (2001) Processive DNA helicase activity of the minichromosome maintenance proteins 4, 6, and 7 complex requires forked DNA structures. *Proc Natl Acad Sci U S A*, **98**, 54-59.
6. Carpentieri, F., De Felice, M., De Falco, M., Rossi, M. and Pisani, F.M. (2002) Physical and functional interaction between the MCM-like DNA helicase and the single-stranded DNA binding protein from the crenarchaeon *Sulfolobus solfataricus*. *J Biol Chem*, **277**, 12118-12127.
7. Pucci, B., De Felice, M., Rossi, M., Onesti, S. and Pisani, F.M. (2004) Amino Acids of the *Sulfolobus solfataricus* Mini-chromosome Maintenance-like DNA Helicase Involved in DNA Binding/Remodeling.

- J Biol Chem*, **279**, 49222-49228.
8. McGeoch, A.T., Trakselis, M.A., Laskey, R.A. and Bell, S.D. (2005) Organization of the archaeal MCM complex on DNA and implications for the helicase mechanism. *Nat Struct Mol Biol*, **12**, 756-762.
 9. Grainge, I., Scaife, S. and Wigley, D.B. (2003) Biochemical analysis of components of the pre-replication complex of *Archaeoglobus fulgidus*. *Nucleic Acids Res*, **31**, 4888-4898.
 10. Haugland, G.T., Shin, J.-H., Birkeland, N.-K. and Kelman, Z. (2006) Stimulation of MCM helicase activity by a Cdc6 protein in the archaeon *Thermoplasma acidophilum*. *Nucleic Acids Res.*, **34**, 6337-6344.
 11. Kelman, Z., Lee, J.K. and Hurwitz, J. (1999) The single minichromosome maintenance protein of *Methanobacterium thermoautotrophicum* Δ H contains DNA helicase activity. *Proc Natl Acad Sci U S A*, **96**, 14783-14788.
 12. Chong, J.P., Hayashi, M.K., Simon, M.N., Xu, R.M. and Stillman, B. (2000) A double-hexamer archaeal minichromosome maintenance protein is an ATP-dependent DNA helicase. *Proc Natl Acad Sci U S A*, **97**, 1530-1535.
 13. Shechter, D.F., Ying, C.Y. and Gautier, J. (2000) The intrinsic DNA helicase activity of *Methanobacterium thermoautotrophicum* Δ H minichromosome maintenance protein. *J Biol Chem*, **275**, 15049-15059.
 14. Fletcher, R.J., Bishop, B.E., Leon, R.P., Sclafani, R.A., Ogata, C.M. and Chen, X.S. (2003) The structure and Function of MCM from archaeal *M. thermoautotrophicum*. *Nature Struct Biol*, **10**, 160-167.
 15. Gomez-Llorente, Y., Fletcher, R.J., Chen, X.S., Carazo, J.M. and Martin, C.S. (2005) Polymorphism and Double Hexamer Structure in the Archaeal Minichromosome Maintenance (MCM) Helicase from *Methanobacterium thermoautotrophicum*. *J. Biol. Chem.*, **280**, 40909-40915.
 16. Costa, A., Pape, T., van Heel, M., Brick, P., Patwardhan, A. and Onesti, S. (2006) Structural basis of the Methanothermobacter thermotrophicus MCM helicase activity. *Nucl. Acids Res.*, **34**, 5829-5838.
 17. Costa, A., Pape, T., van Heel, M., Brick, P., Patwardhan, A. and Onesti, S. (2006) Structural studies of the archaeal MCM complex in different functional states. *Journal of Structural Biology*, **156**, 210-219.
 18. Pape, T., Meka, H., Chen, S., Vicentini, G., van Heel, M. and Onesti, S. (2003) Hexameric ring structure of the full-length archaeal MCM protein complex. *EMBO Rep*, **4**, 1079-1083.
 19. Yu, X., VanLoock, M.S., Poplawski, A., Kelman, Z., Xiang, T., Tye, B.K. and Egelman, E.H. (2002) The *Methanobacterium thermoautotrophicum* MCM protein can form heptameric rings. *EMBO Rep*, **3**, 792-797.
 20. Chen, Y.J., Yu, X., Kasiviswanathan, R., Shin, J.H., Kelman, Z. and Egelman, E.H. (2005) Structural Polymorphism of *Methanothermobacter thermotrophicus* MCM. *J Mol Biol*, **346**, 389-394.