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Stoichiometry and architecture of active DNA replication machinery in Escherichia coli.

Reyes-Lamothe R, Sherratt DJ, Leake MC Science 2010 Apr 23 328(5977):498-501 [abstract on PubMed] [citations on Google Scholar] [related articles] [full text] [order article]

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Gottfried Otting

Australian National University, Australia Structural Biology

🐵 New Finding <u>s</u> Tech Advance

Amazingly, the number of protein molecules in a large functional complex can now be counted in vivo.

The authors expressed the individual proteins of the Escherichia coli replisome in fusion with YPet (a bright version of green fluorescent protein [GFP]) and counted the molecules by the intensity and step-wise bleaching observed by single-molecule fluorescence microscopy in living cells. Unexpectedly, the count of the replisome subunits epsilon, theta, tau, delta, chi, psi, beta and Ssb revealed three subunits of the polymerase III core (alpha, epsilon and theta), rather than two that had traditionally been assumed to perform the replication of leading and lagging strand DNA.

Competing interests: None declared Evaluated 10 May 2010 NEW

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