

Replicon dynamics during the cell cycle in *Escherichia coli*

During bacterial cell cycle, the replication of the chromosome and plasmids is followed by the segregation of each copy in daughter cells. As replication and segregation must be finely coordinated with cell division many proteins are recruited. Notably the MatP protein, and its DNA binding site *matS*, which participates in the positioning of the *ter* regions of sister chromatids at midcell; and the recombinases XerC and XerD which separate dimers by site-specific recombination at the *dif* site. Nevertheless, the entire mechanism that drive this coordination is not fully understood. It is known that a lot of proteins involved can interact with the Topoisomerase IV (TopoIV), which has a key role in the regulation through its catenane and pre-catenane resolution activity. Catenanes are interlinked DNA molecules, and these links must be removed to allow their segregation. To better understand the decatenation activity in vivo, I use the model bacterium *Escherichia coli* with temperature sensitive mutants of TopoIV. The activity is investigated by analyzing catenated plasmids accumulation at non-permissive temperature, as well as their resolution by recovering TopoIV.

Experiments are carried out with a small (2.7kb) and high copy number plasmid, pUC18; its derivative where a *dif* site has been added, TopoIV activity is strong at this site [4]. To study the effect of subcellular localization on the activity of TopoIV, a *matS* site has been added on these plasmids. My results suggest that subcellular position at midcell by *matS*, and *dif* sites on the plasmid can improve decatenation, but also that the combination of *matS* and *dif* have a synergic effect on TopoIV activity.

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