

# A tripartite-chromosome E. coli strain allows the chromosome isolation and implantation

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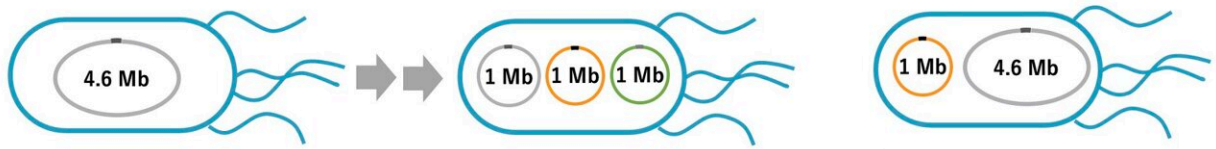


Figure 1: Chromosome splittingThe E. coli chromosome was split into three 1-Mb chromosomes, and the chromosome was used for genome implantation. Credit: Rikkyo University

The issue of concern was that the Escherichia coli (E. coli) genome, consisting of 4.6 million base pairs of a single circular DNA, is too large to manipulate following the extraction and transfer to other bacteria.

In the present study, a group of Rikkyo University researchers led by Assistant Professor Takahito Mukai and Professor Masayuki Su'etsugu has succeeded in splitting the E.coli [genome](#) into tripartite-genome of 1 million base pairs per genome (split-genome) using the smallest E. coli genome strain established so far. In addition, they successfully extracted the split-genome from bacteria and installed it in other E. coli.

It is a major breakthrough that E. coli could stably proliferate even after the bacterial genome was split into tripartite-genome. Going forward, it is imperative to clarify how the replication and distribution of the

tripartite-genome are controlled. Also, this research group has been developing the technology for synthesizing gigantic DNA without using cells (cell-free) and reported a cell-free technique for amplifying 1 million base pairs of circular DNA. In the future, the installation of cell-free synthesized split genomes in *E. coli* is expected to lead to the creation of artificial *E. coli* with designed valuable functions, such as material production.

This achievement is expected to lead to the clarification of the mechanism of genome replication/segregation and also to the application of tools in [synthetic biology](#) to convert the genome, the blueprint of life, so that we can create functionally designed life. The results in the present study have been published in the online version as a breakthrough paper in *Nucleic Acids Research* on April 28, 2021.

**More information:** Tatsuya Yoneji et al, Grand scale genome manipulation via chromosome swapping in *Escherichia coli* programmed by three one megabase chromosomes, *Nucleic Acids Research* (2021).  
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