

Spacer acquisition mechanism in type II-A CRISPR system.

Yibei Xiao¹, Jagat Budhathoki¹, Sherwin Ng^{1,#}, Ki Hyun Nam^{2,#}, Ailong Ke^{1,*}

¹ Department of Molecular Biology and Genetics, Cornell University, 253 Biotechnology Building, Ithaca, NY 14853, USA.

² Pohang Accelerator Laboratory, Pohang University of Science and Technology, Pohang, South Korea.

Molecular memory is created when a short foreign DNA-derived prespacer is integrated into the CRISPR array as a new spacer. Whereas the RNA-guided CRISPR interference mechanism varies widely among CRISPR-Cas systems, the spacer integration mechanism is essentially identical. The conserved Cas1 and Cas2 proteins form an integrase complex consisting of two distal Cas1 dimers bridged by a Cas2 dimer. The prespacer is bound by Cas1-Cas2 as a dual-forked DNA, and the terminal 3'-OH of each 3' overhang serves as an attacking nucleophile during integration. The prespacer is preferentially integrated into the leader-proximal region of the CRISPR array, guided by the leader sequence and a pair of inverted repeats inside the CRISPR repeat. Spacer integration in the well-studied *Escherichia coli* type I-E CRISPR system also relies on the bacterial integration host factor. In type II-A CRISPR, however, Cas1-Cas2 alone integrates spacers efficiently *in vitro*; other Cas proteins (such as Cas9 and Csn2) have accessory roles in the biogenesis phase of prespacers. I will present our structural and biochemical efforts in revealing the spacer acquisition mechanism in the *Enterococcus faecalis* type II-A CRISPR system.