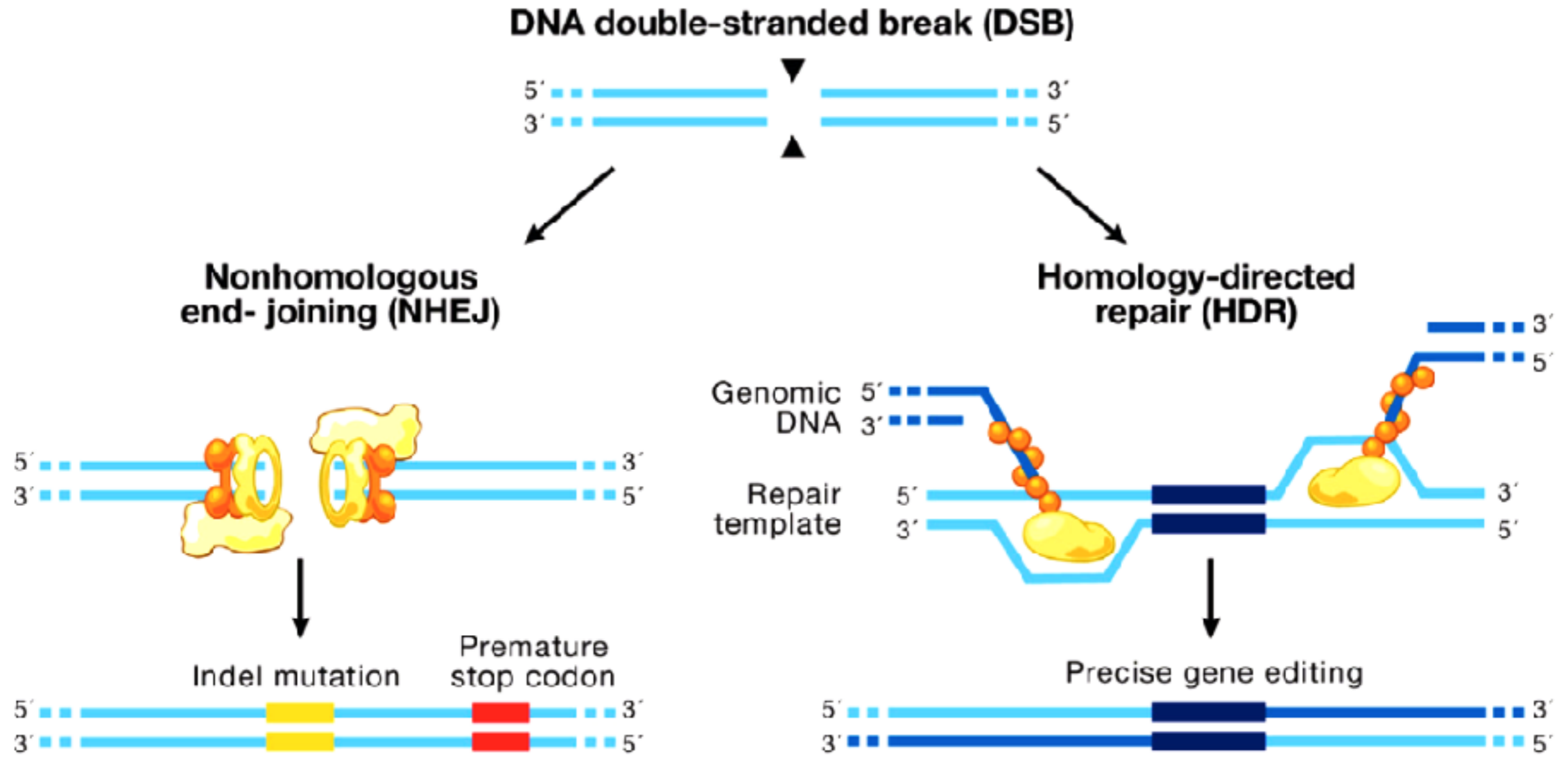
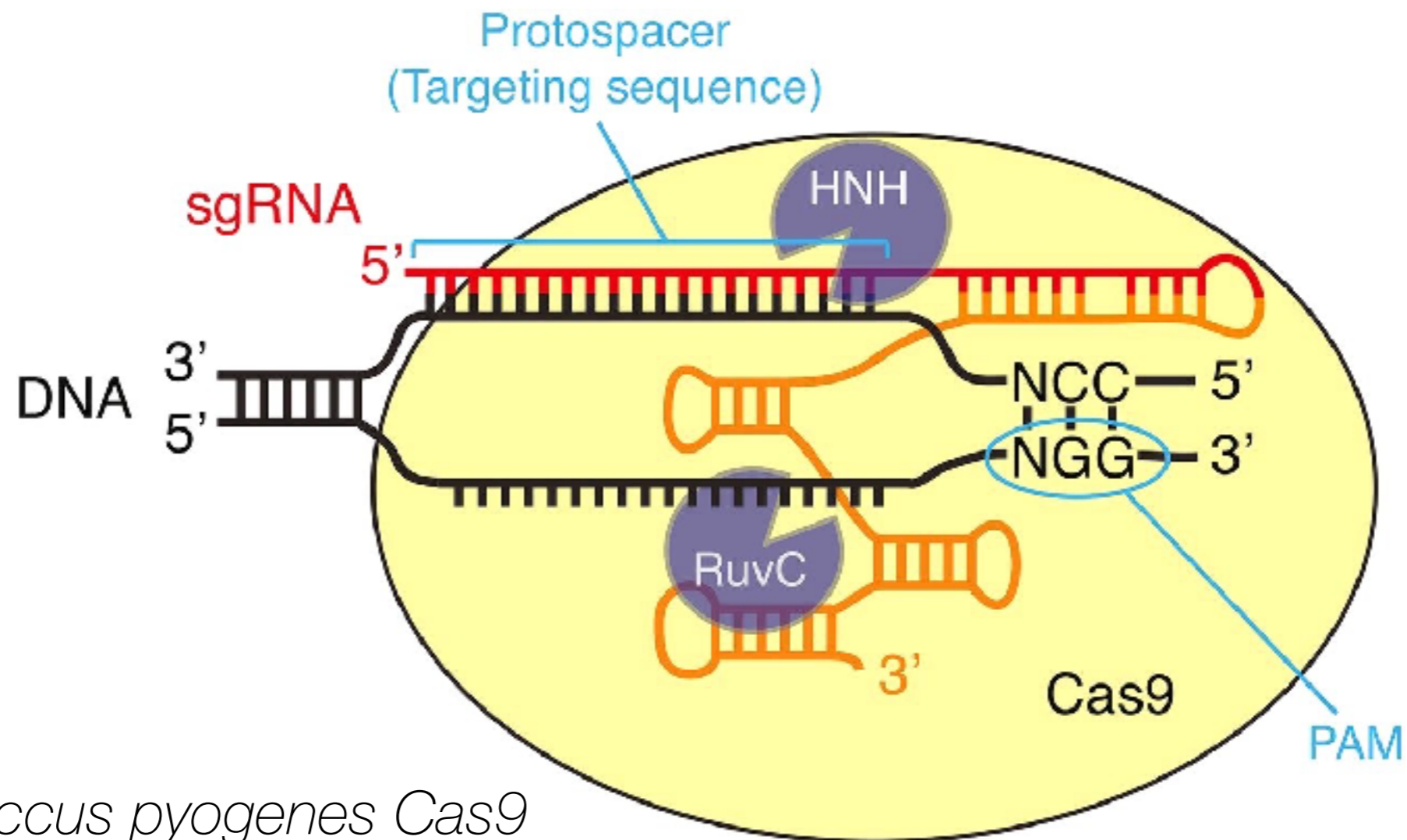


DNA damage repair pathways



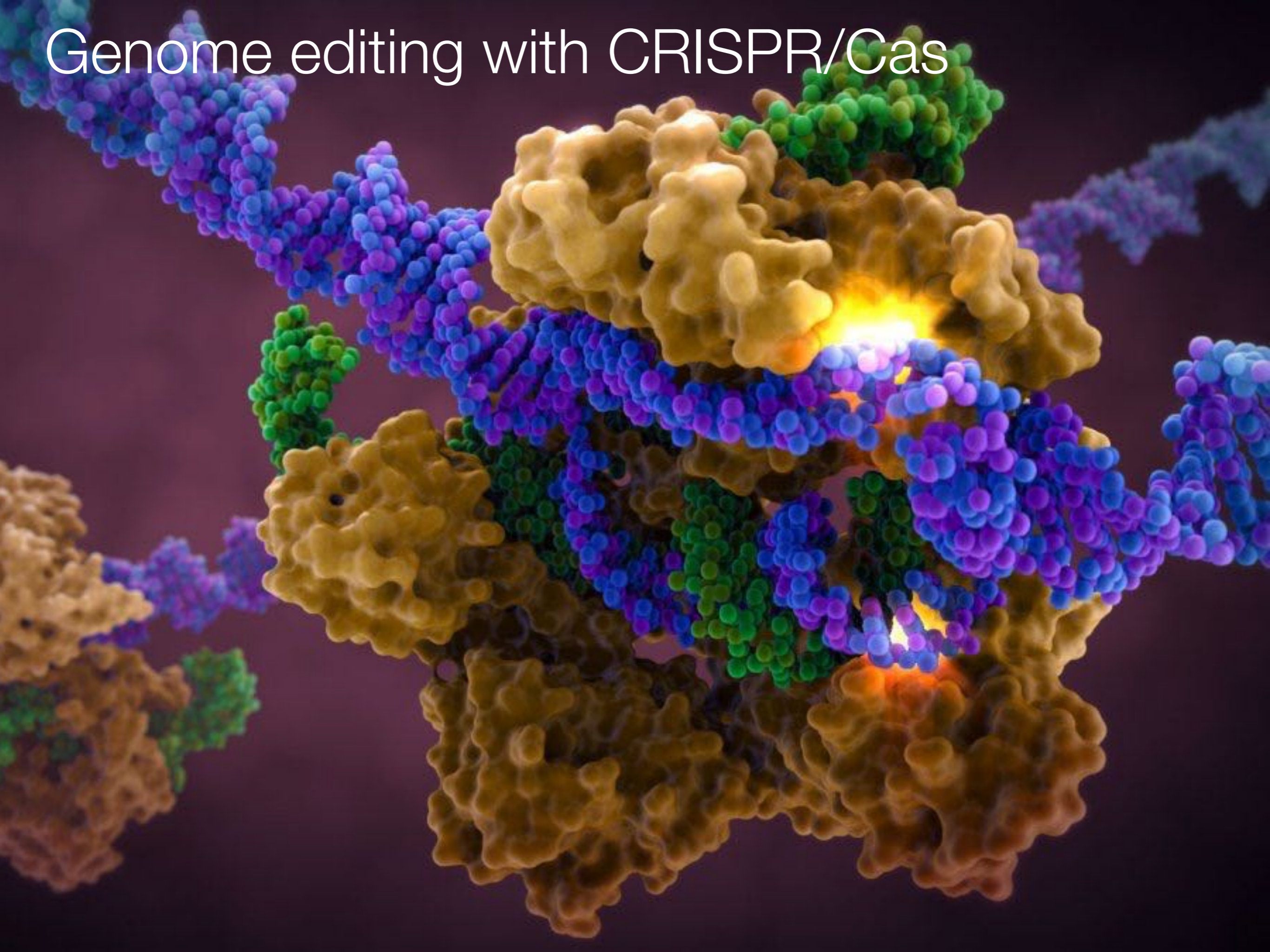
Genome editing with CRISPR/Cas



Streptococcus pyogenes Cas9

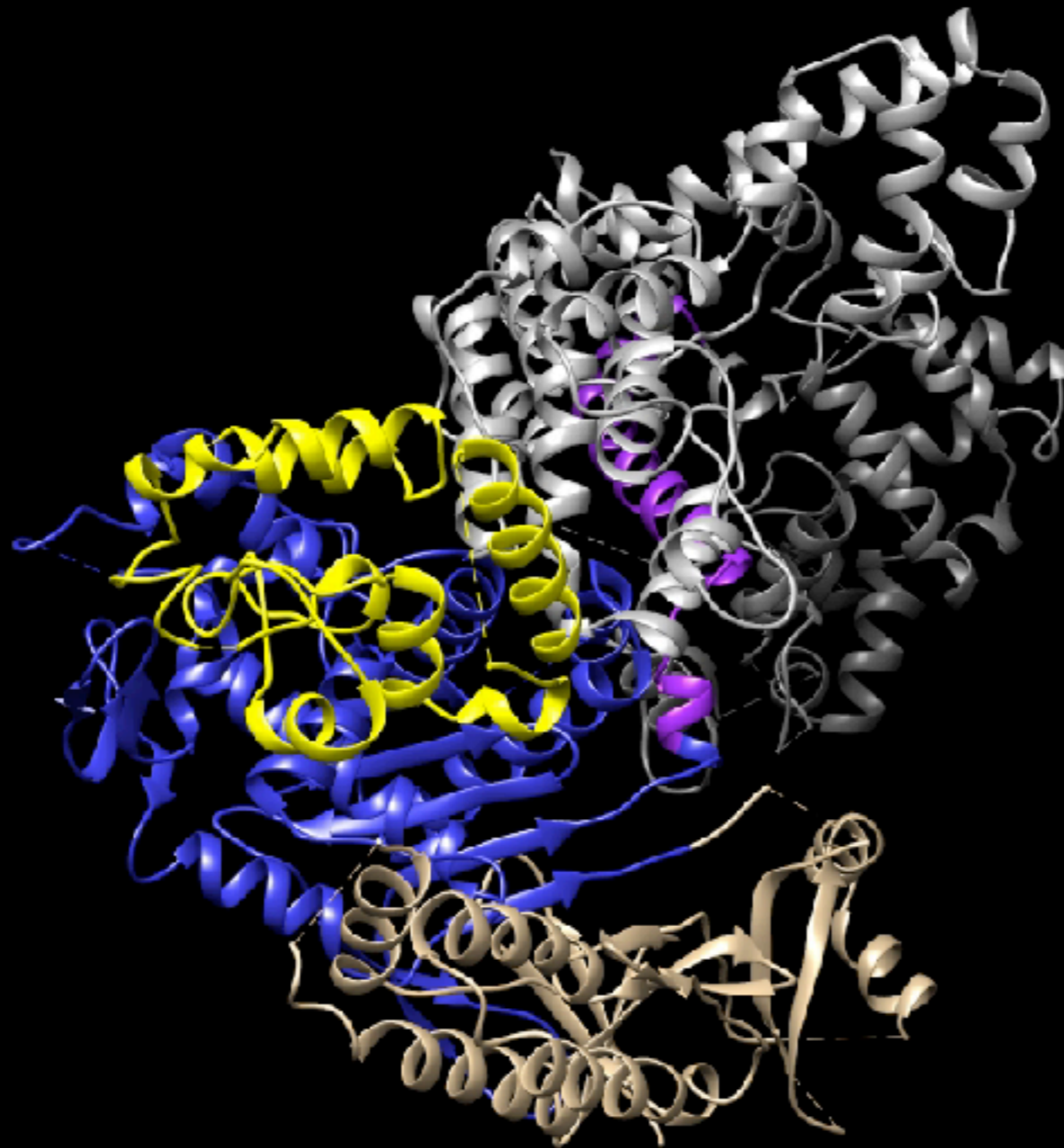
- ▶ Incredibly easy to program for any specific site
- ▶ HNH and RuvC can be mutated to create strand specific nickases or a non-cutting but still binding enzyme

Genome editing with CRISPR/Cas



Cas9 Conformational Dynamics

Apo (4CMP) → sgRNA-bound (4ZT0)



PI

Cas9 cutting has multiple checkpoints

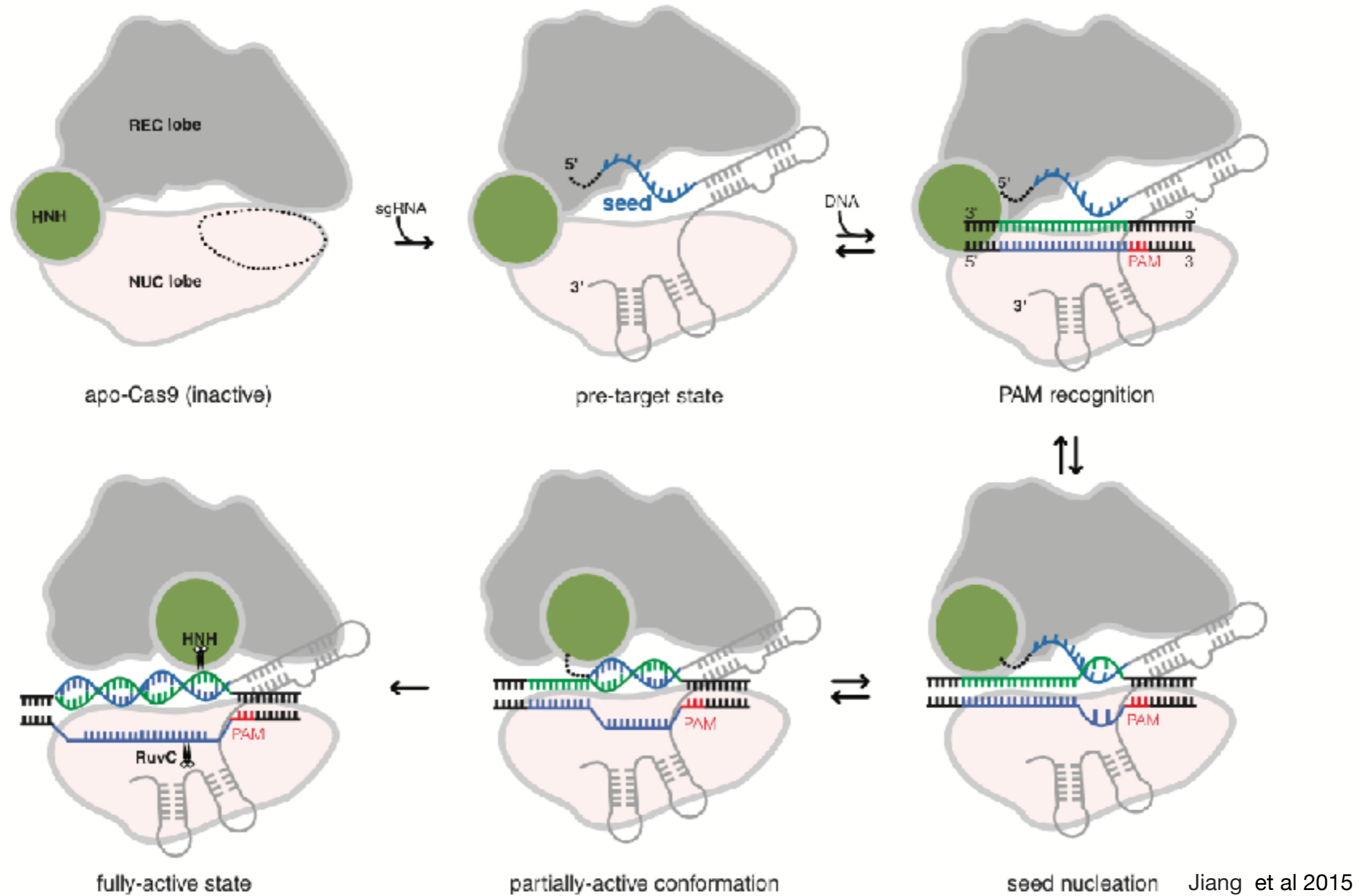


Fig. 4. Proposed mechanism for Cas9-mediated DNA targeting and cleavage. When Cas9 is in the apo state, its PAM-interacting cleft (dotted circle) is largely disordered. In the pretarget state, the PAM-interacting domain and seed sequence from guide RNA are preorganized for PAM recognition, followed by dsDNA melting next to PAM. The nonseed region is disordered and indicated as a dotted line. Base pairing between the seed sequence and the target DNA drives Cas9 into a near-active conformation; complete base pairing between the full guide segment and the target DNA strand enables Cas9 to reach a fully active state.

Classes of CRISPR

