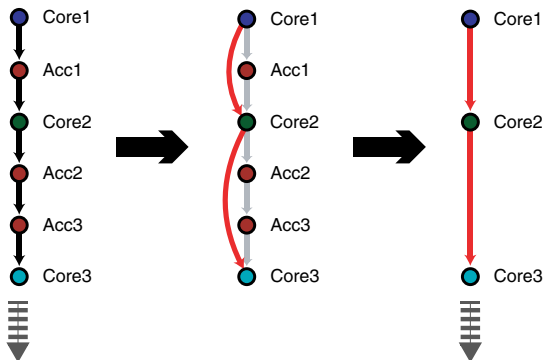


A

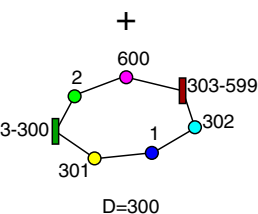
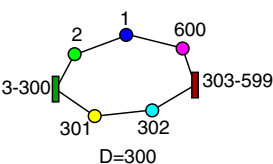


Genome organization
stability analysis

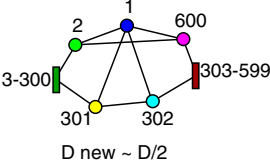
Backbone
stability analysis

B

Long-range movement



Take the sum of the corresponding
adjacency matrices



Nearby movement

