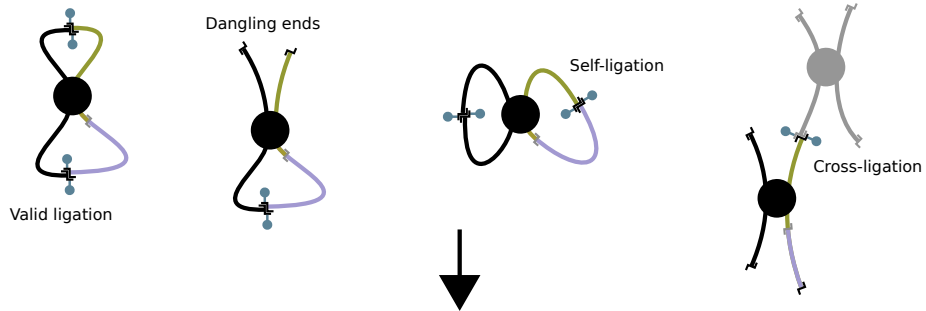


Digestion

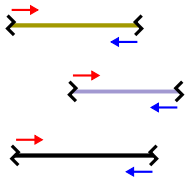


Ligation



Shearing, biotin pull-down, sequencing, mapping

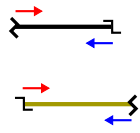
Same internal



Read pair orientations



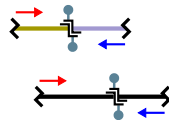
Dangling-end



Read pair orientations



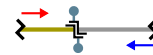
Self-ligation



Read pair orientations



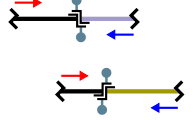
Cross-ligation fragments



Read pair orientations



Valid fragments



Read pair orientations

