

# De novo Genome Assembly Solutions

**bt1**  
technology\*

Assembly

Correction

Scaffolding

Gap-filling

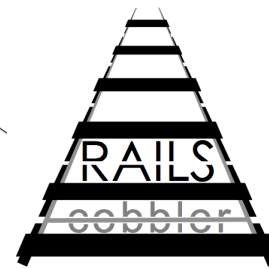
Polishing



ntjoin

long Stitch

mitLink



Scaler



Input

Sequencing reads

Short



Linked



Long



Genome

NGS

eg. Illumina/MGI

SMS

eg. ONT/PacBio

ALL DRAFT ASSEMBLIES SUPPORTED

\*<https://github.com/bcgsc>