#### **Read construction**

**Unique ind. barcode** 

**Sequence of interest** 

Illumina adapter

**Restriction site** 

#### **Read construction**

**Unique ind. barcode** 

**Sequence of interest** 

Illumina adapter

**Restriction site** 

## **Denovo pipeline**

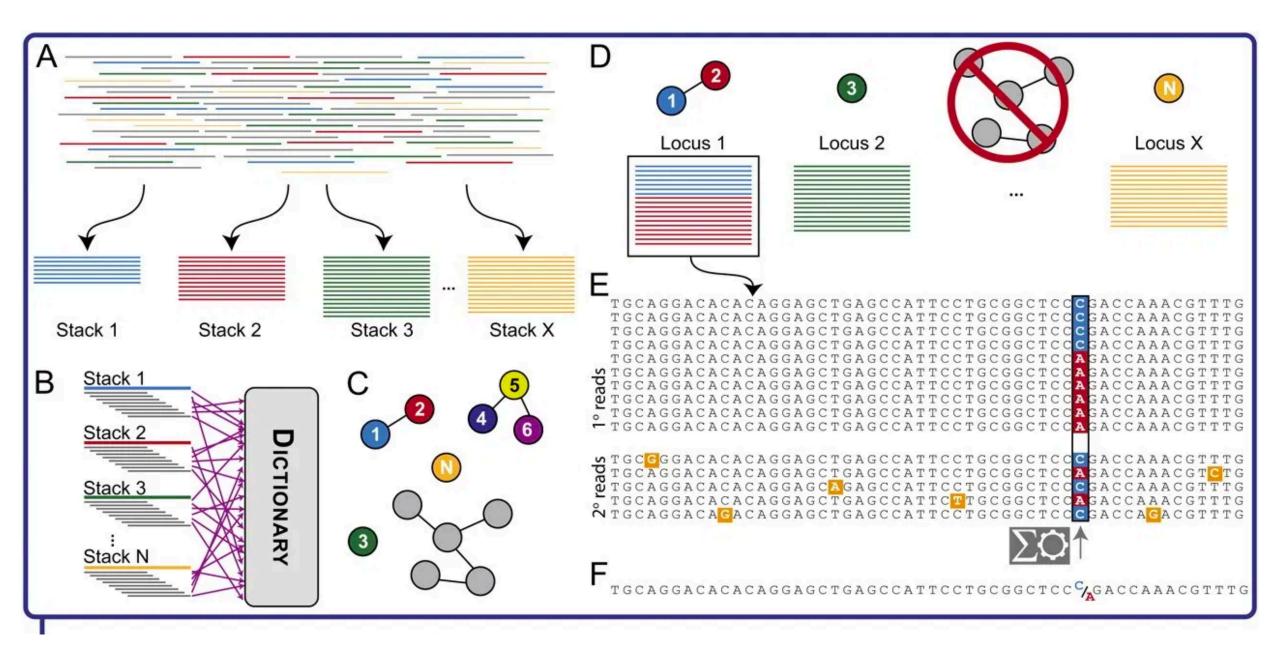
p

**USTACKS** Build loci de novo

**CSTACKS** Assemble catalog

**SSTACKS** Match to catalog

**Populations** Filter loci and compute Population statistics



Parameter Description	denovo_map.p  1 Parameter	Pipeline component	Component Parameter	Default Value
Minimum stack depth / minimum depth of coverage	-m	ustacks	-m	3
Distance allowed between stacks	-M	ustacks	-M	2
Distance allowed between catalog loci	-n	cstacks	-n	1

# reference pipeline

P
i
p
e
I
i

e

USTACKSBuild loci de novoCSTACKSAssemble catalogSSTACKSMatch to catalog

**Populations** 

Filter loci and compute Population statistics

### Reference pipeline

Alignment to a reference genome

**Populations** 

Filter loci and compute Population statistics