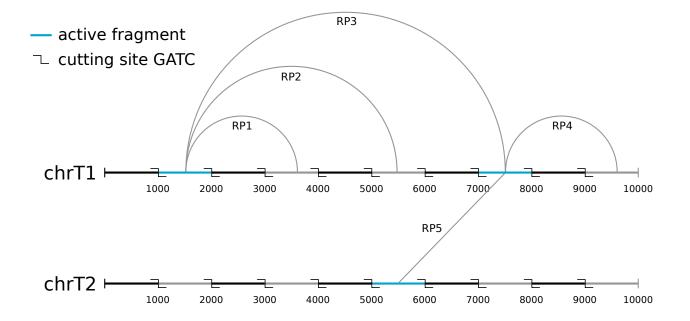
Test files for testing of class *InteractionCountsMap*



Prepare FASTQ file

Read length: 20 5 Read pairs

Filenames:

testInteractionCountsMap_R1.fq testInteractionCountsMap_R2.fq

- 1. Create $RP1_R1 = RP2_R1 = RP3_R1$
- 2. Create RP1 R2
- 3. Create RP2 R2
- 4. Create RP3 R2 = RP4 R1 = RP5 R1
- 5. Create RP4_R2
- 6. Create s

Prepare reference sequence file

Filename:

test Interaction Counts Map Reference. fa

One unit has 1000 bp and ends with GATC.

Fragments mainly consist of Ns.

Only the centers of interacting fragments contain read sequnces.

Create FASTA index for reference.

Perform digest with Diachromatic