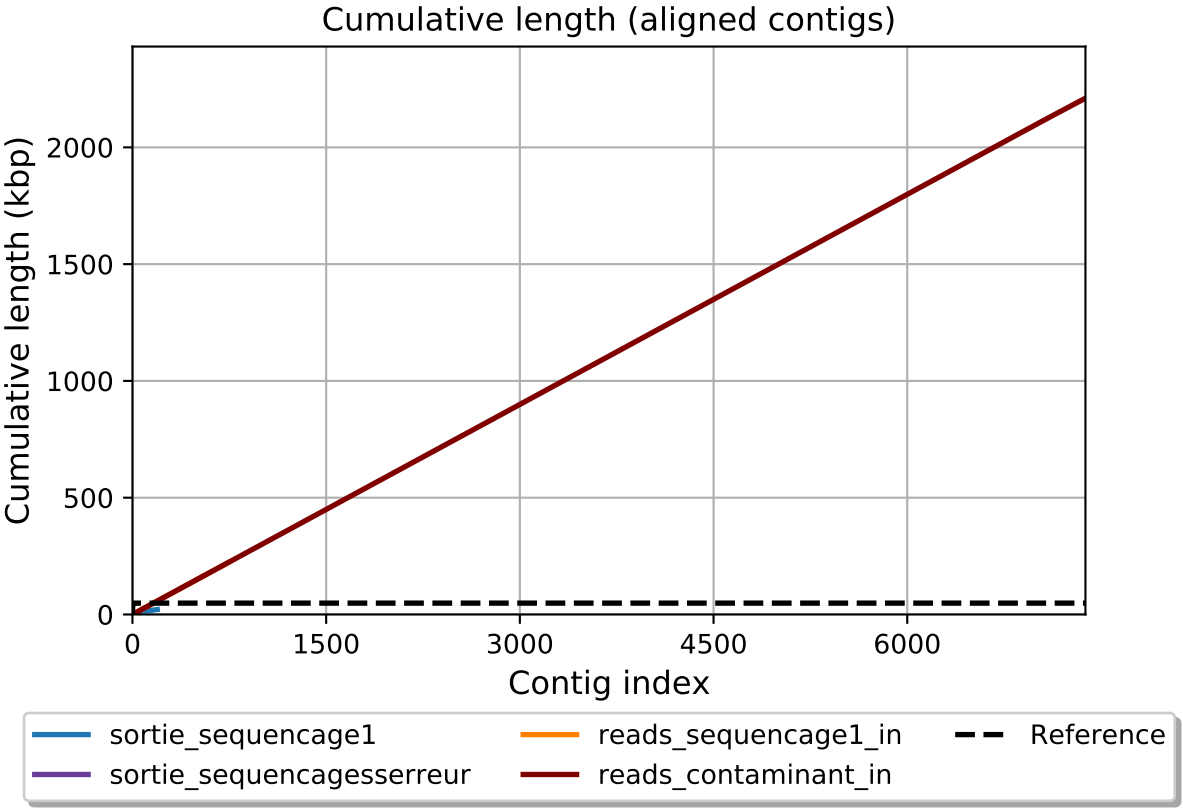
Cumulative length (aligned contigs)
Cumulative length (kbp)
2000
1500
1000
500
0
0
1500
3000
4500
6000
Contig index
sortie_sequencage1
sortie_sequencagesserreur
reads_sequencage1_in
reads_contaminant_in
Reference