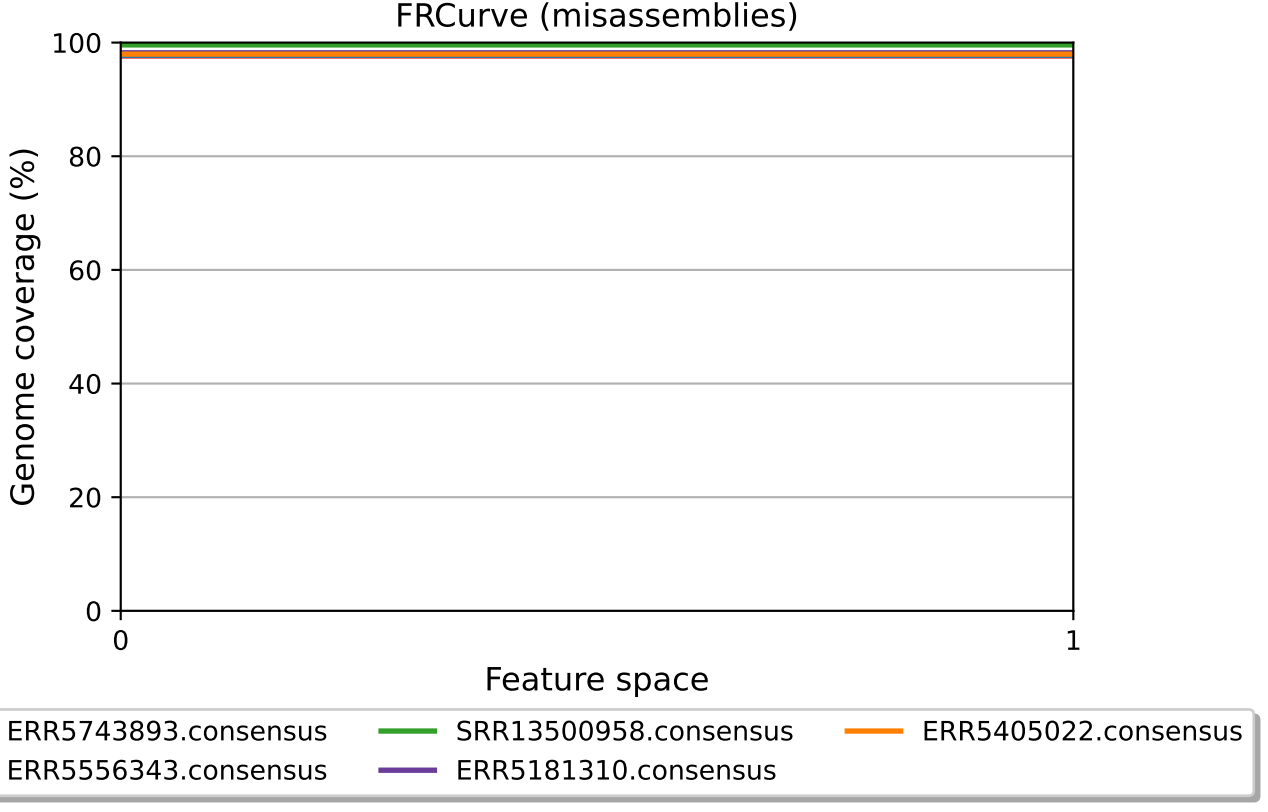
FRCurve (misassemblies)
Genome coverage (%)
100
80
60
40
20
0
0
1
Feature space
ERR5743893.consensus
ERR5556343.consensus
SRR13500958.consensus
ERR5181310.consensus
ERR5405022.consensus