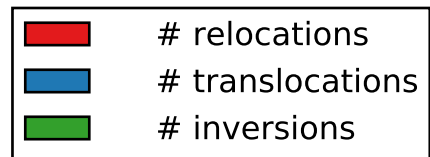


scaffolds

misassemblies

3.0
2.5
2.0
1.5
1.0
0.5
0.0



e_coli_k12

