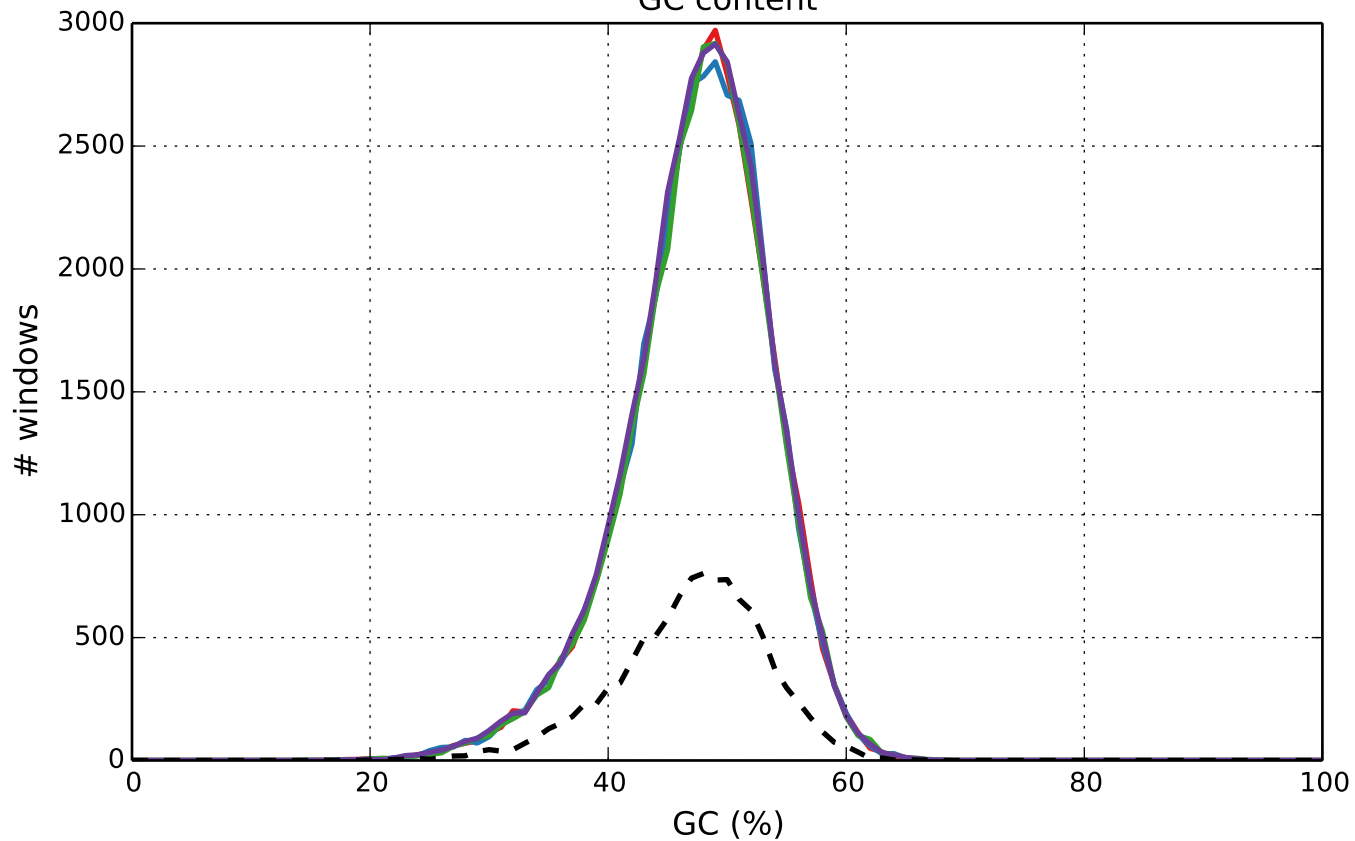


GC content



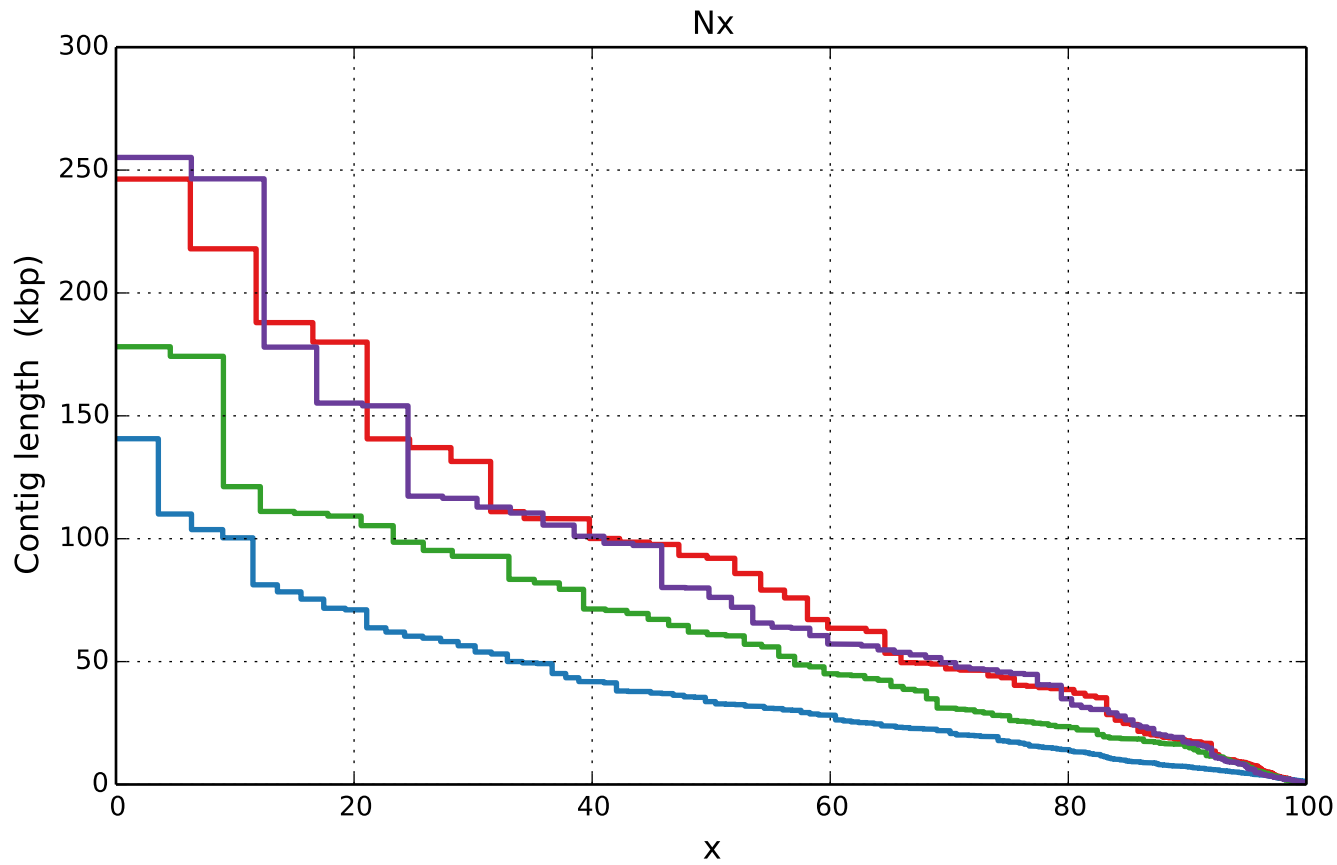
dataset\_390.dat

dataset\_392.dat

dataset\_393.dat

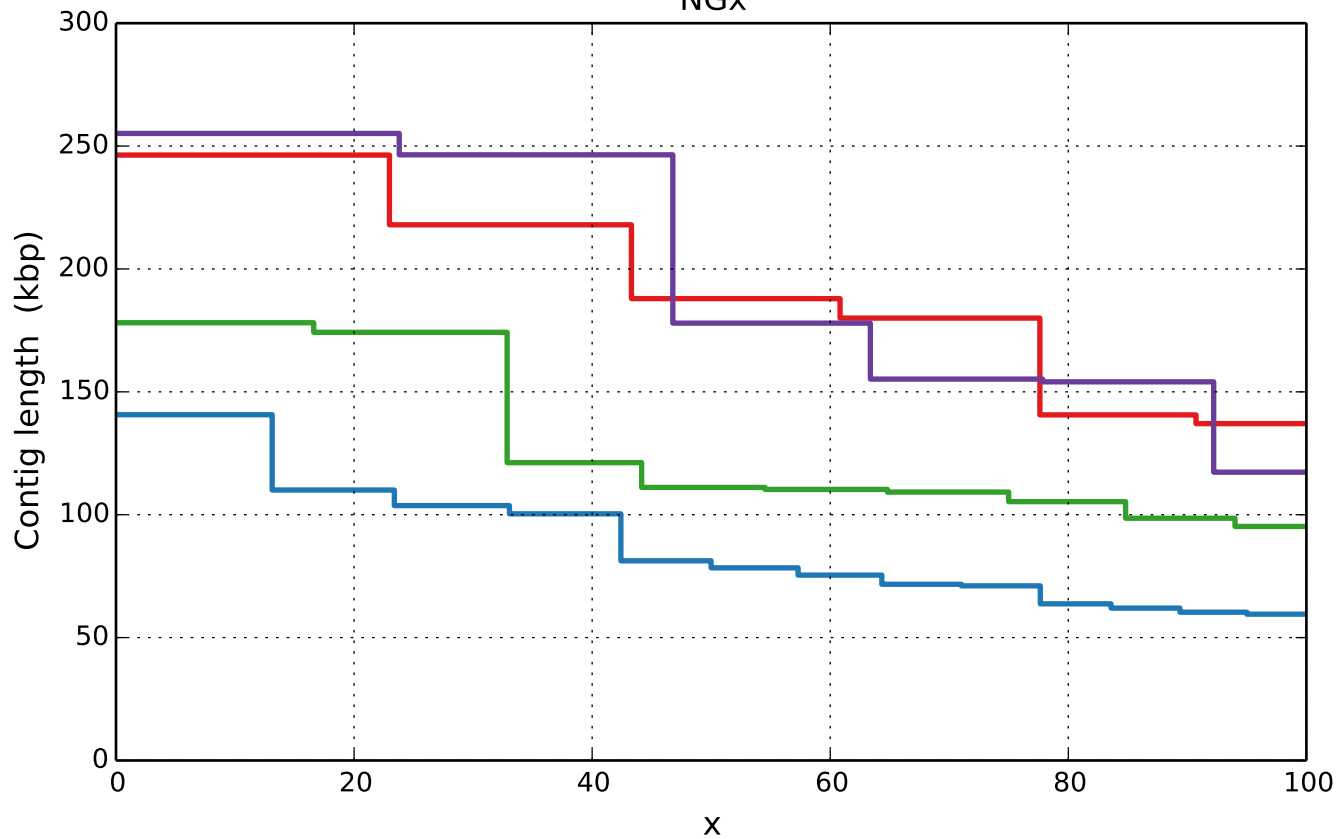
-- Reference

dataset\_391.dat



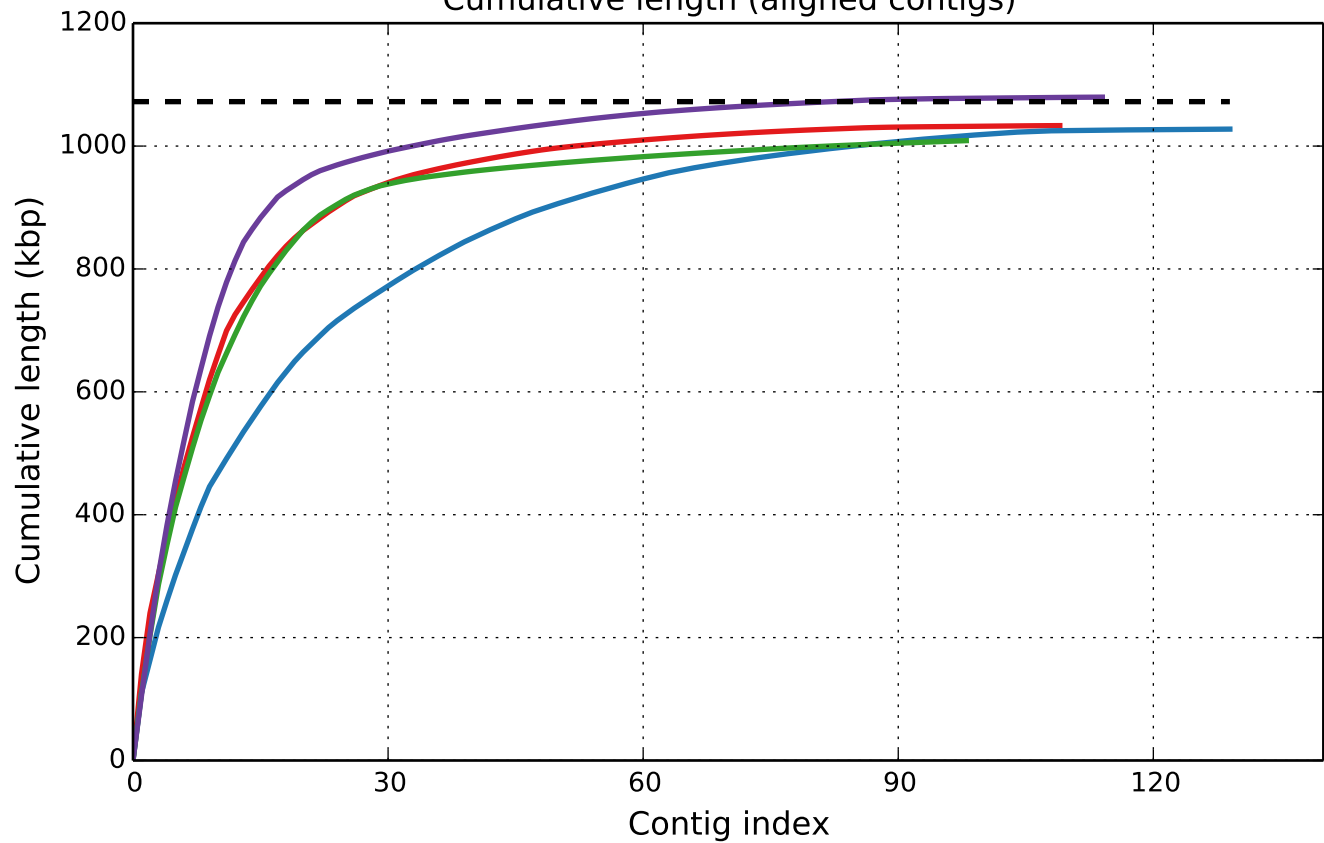
— dataset\_390.dat    — dataset\_391.dat    — dataset\_392.dat    — dataset\_393.dat

NGx



— dataset\_390.dat    — dataset\_391.dat    — dataset\_392.dat    — dataset\_393.dat

Cumulative length (aligned contigs)



dataset\_390.dat

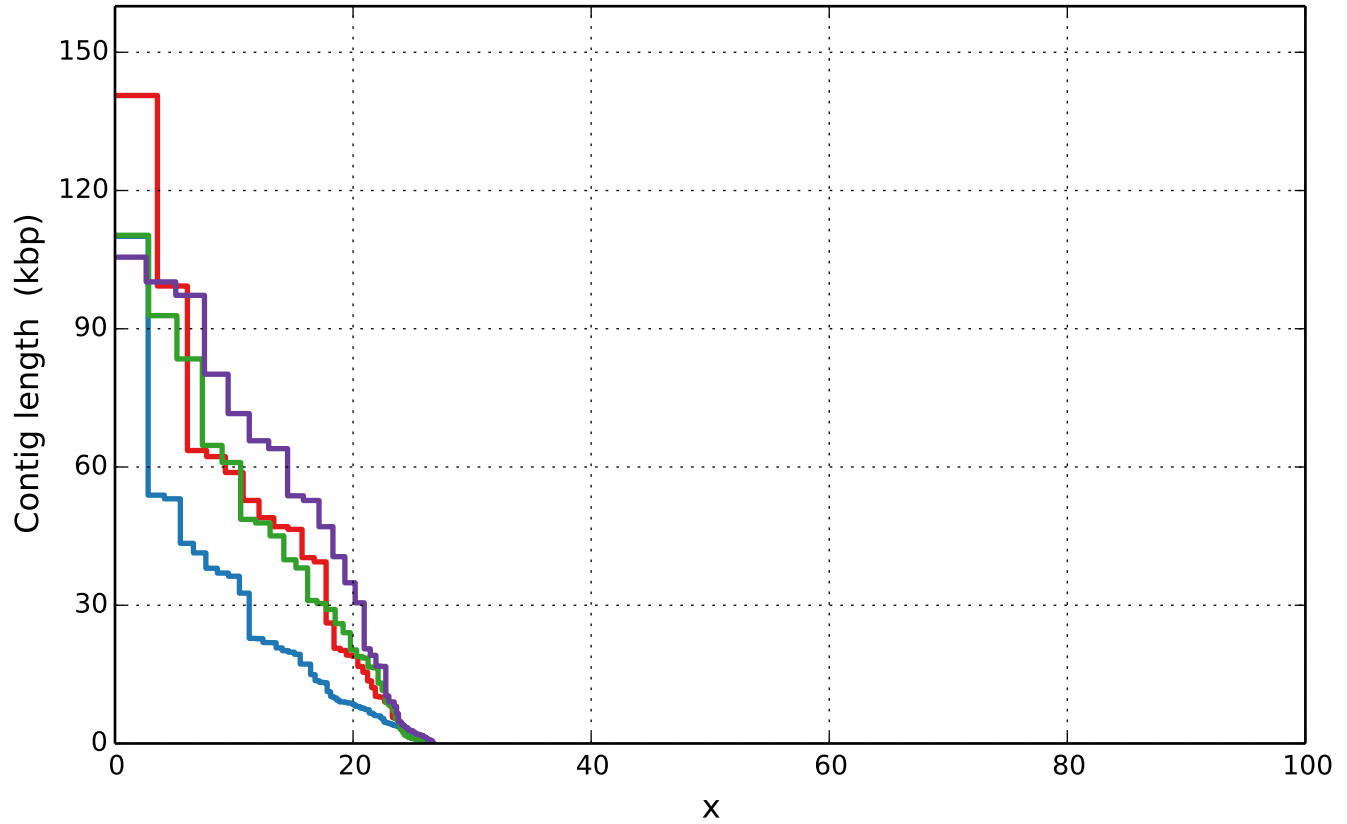
dataset\_392.dat

dataset\_393.dat

-- Reference

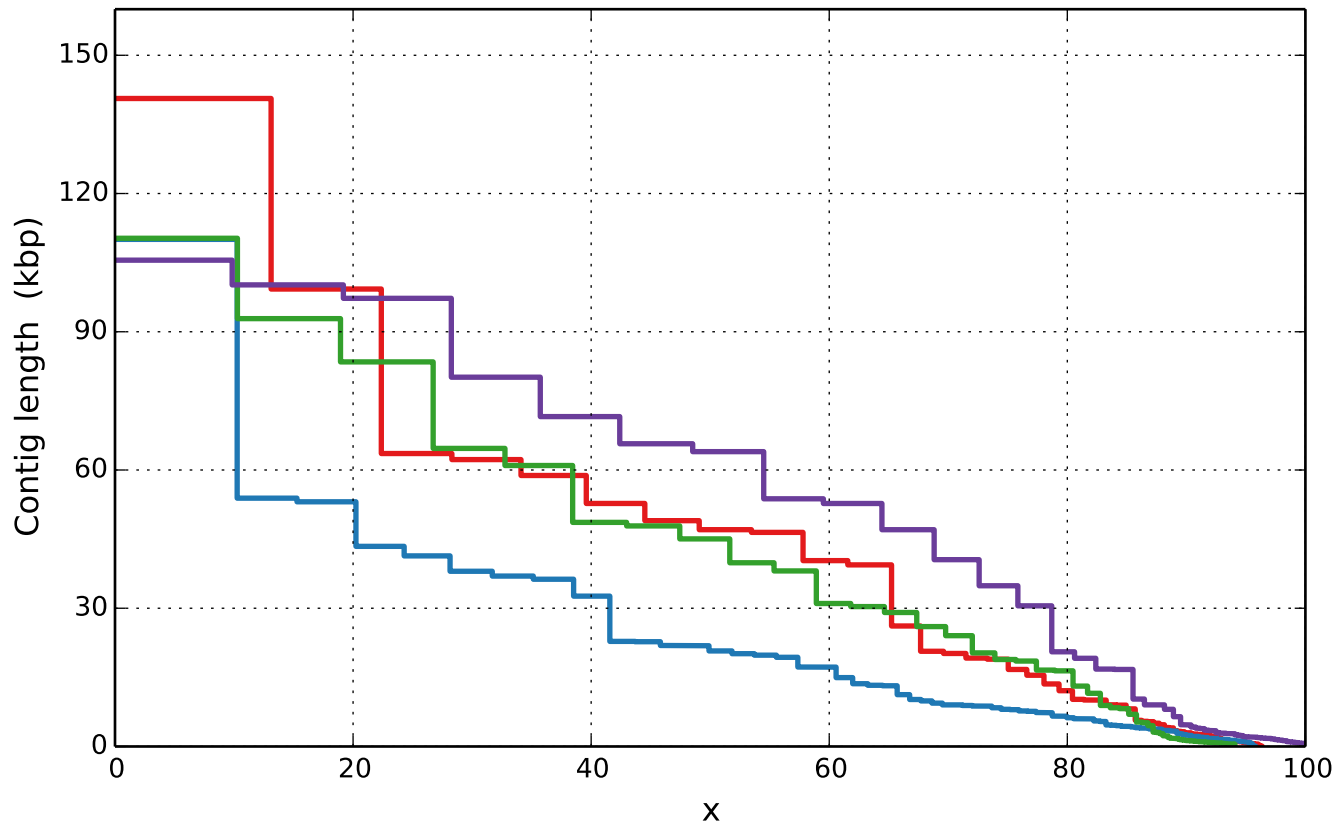
dataset\_391.dat

NAx



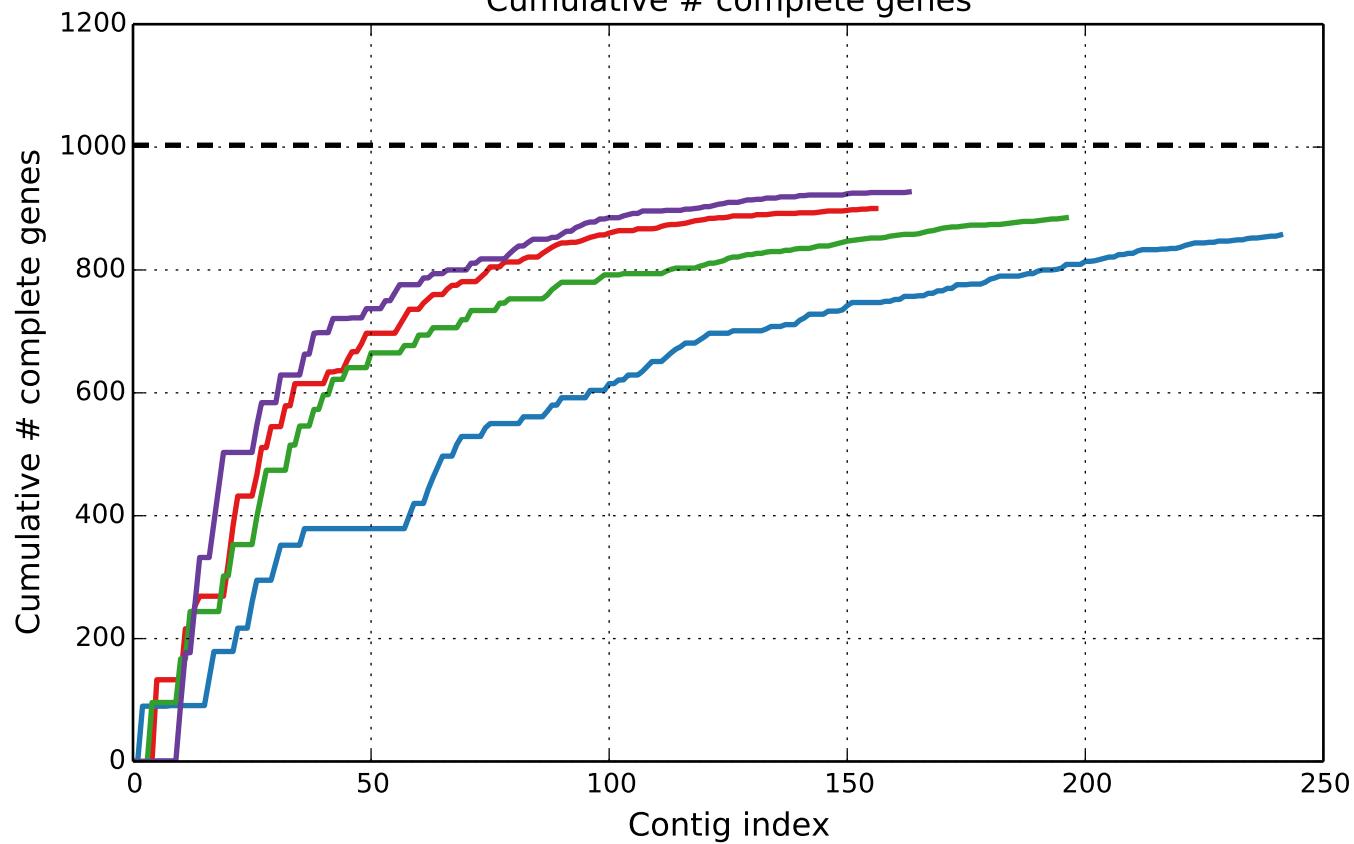
— dataset\_390.dat — dataset\_391.dat — dataset\_392.dat — dataset\_393.dat

# NGAx



— dataset\_390.dat    — dataset\_391.dat    — dataset\_392.dat    — dataset\_393.dat

Cumulative # complete genes



dataset\_390.dat

dataset\_392.dat

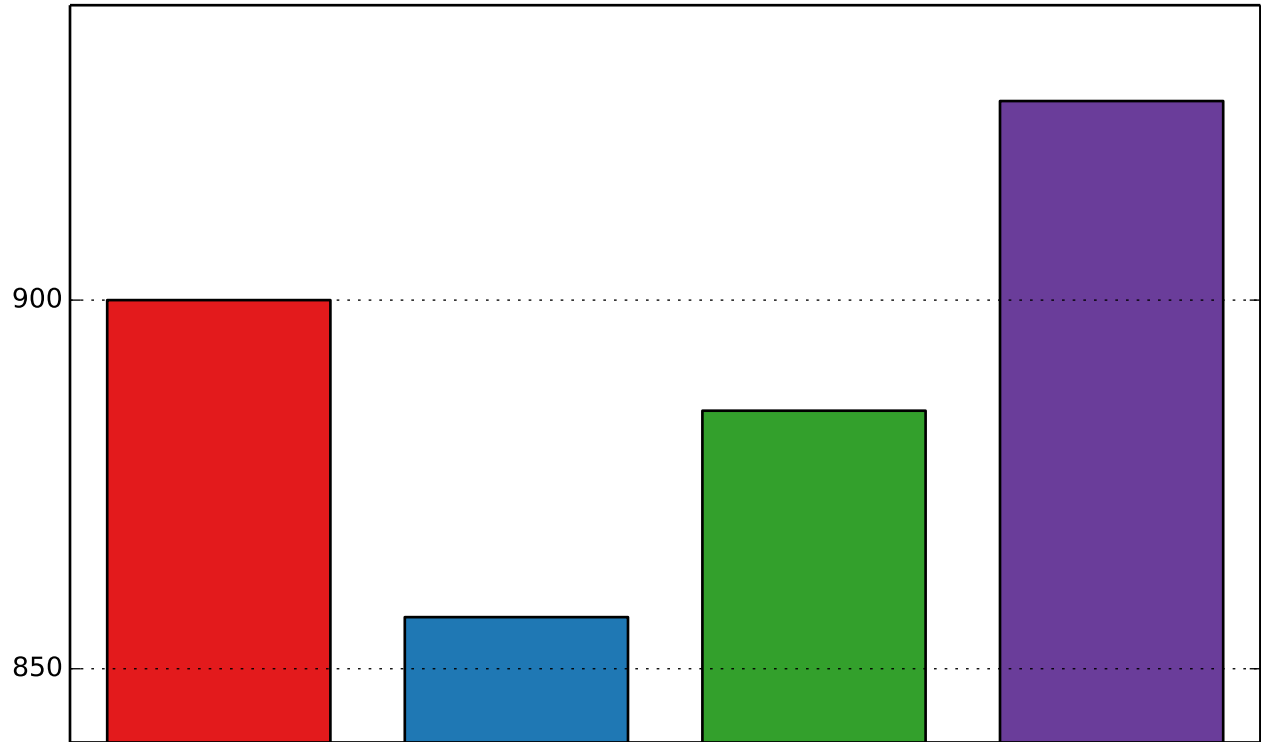
dataset\_393.dat

-- Reference

dataset\_391.dat



# complete genes



dataset\_390.dat   dataset\_391.dat   dataset\_392.dat   dataset\_393.dat

