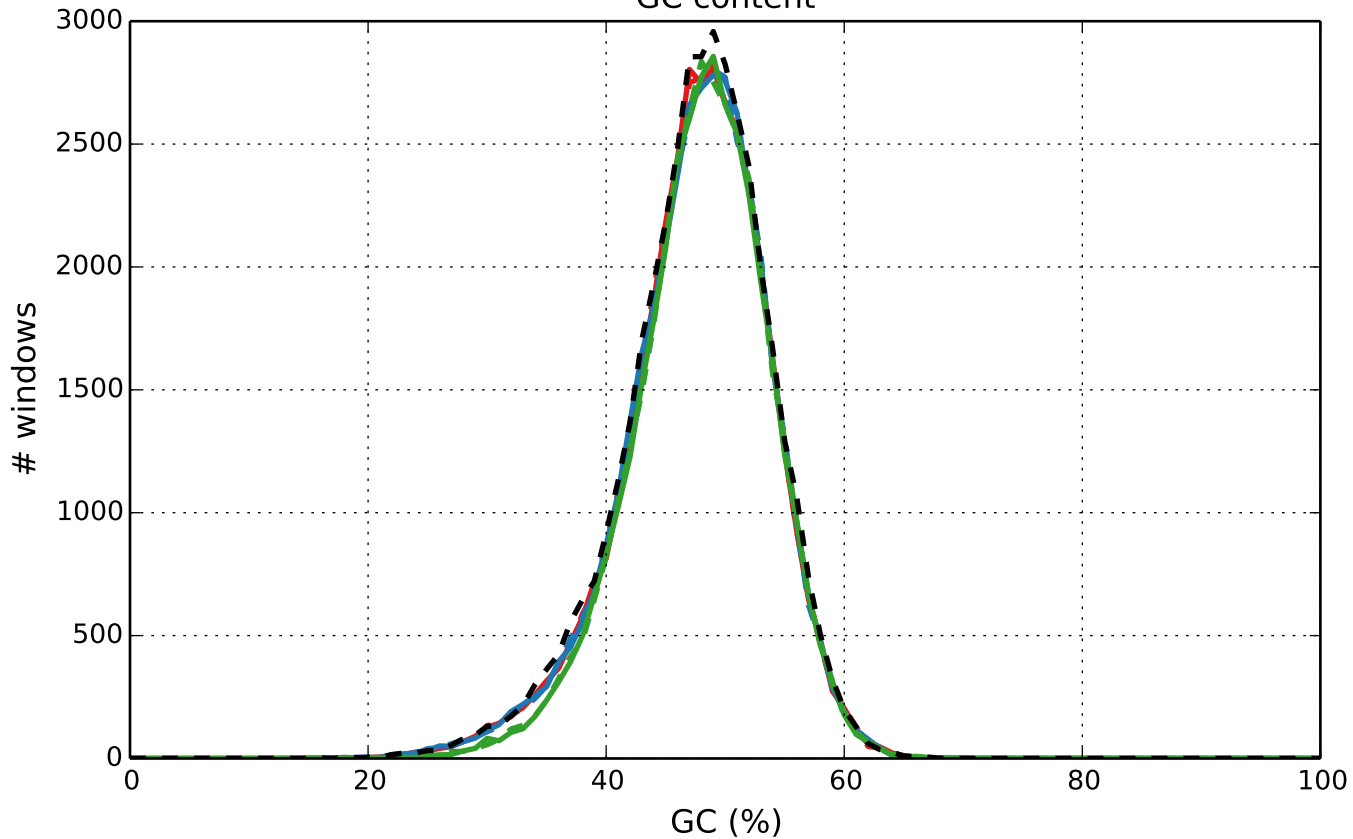
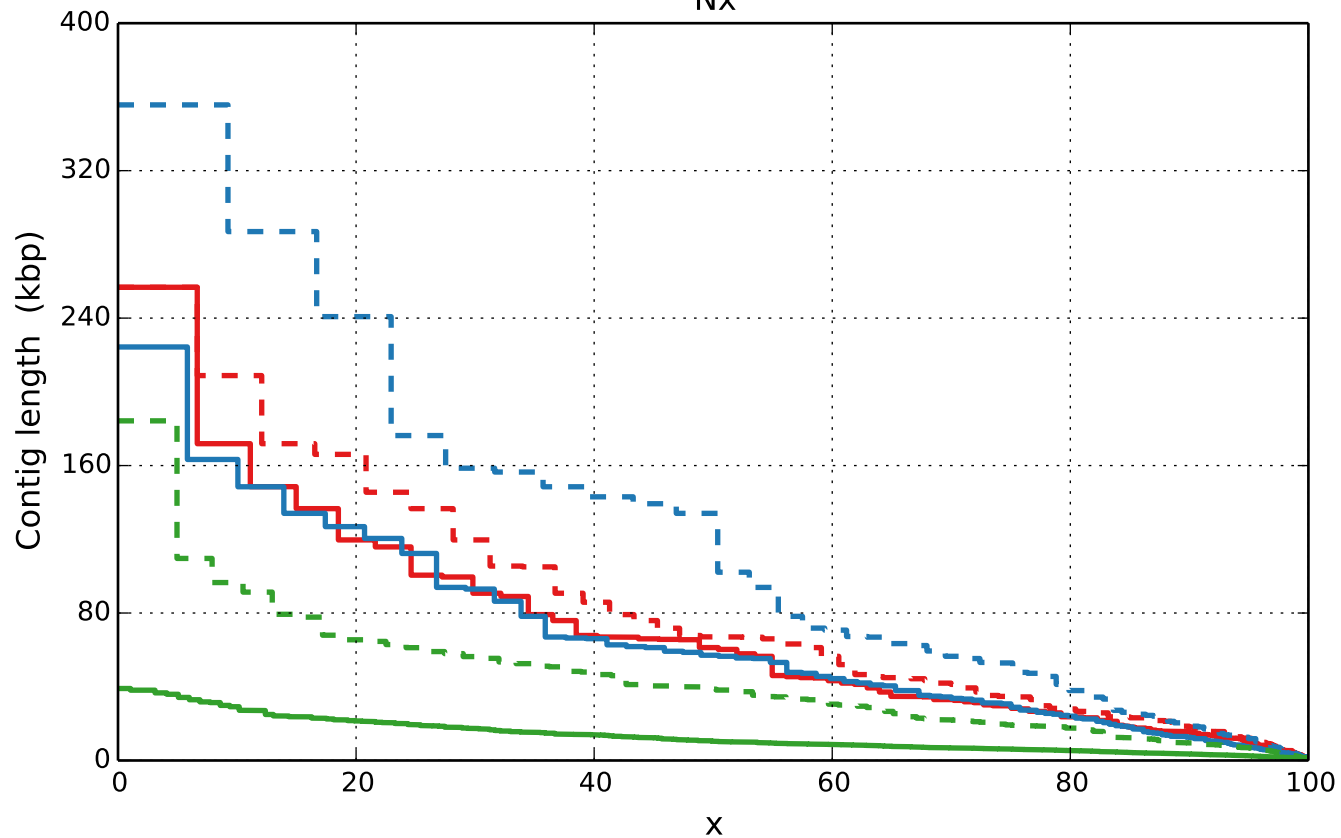


GC content



— GAGE-B broken
 — Cabog_7_0 broken
 — Cabog_8_1 broken
 - - Reference
- - GAGE-B
 - - Cabog_7_0
 - - Cabog_8_1

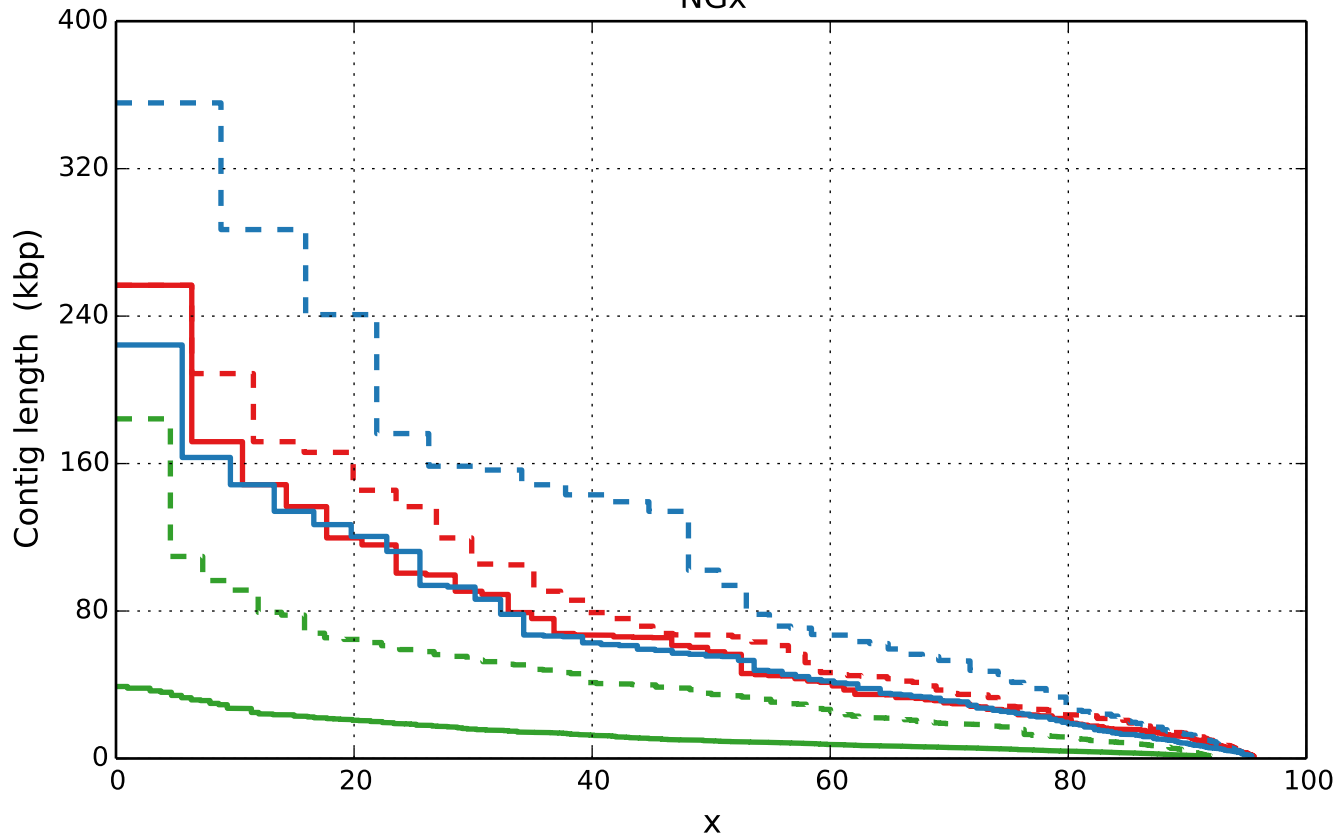
Nx



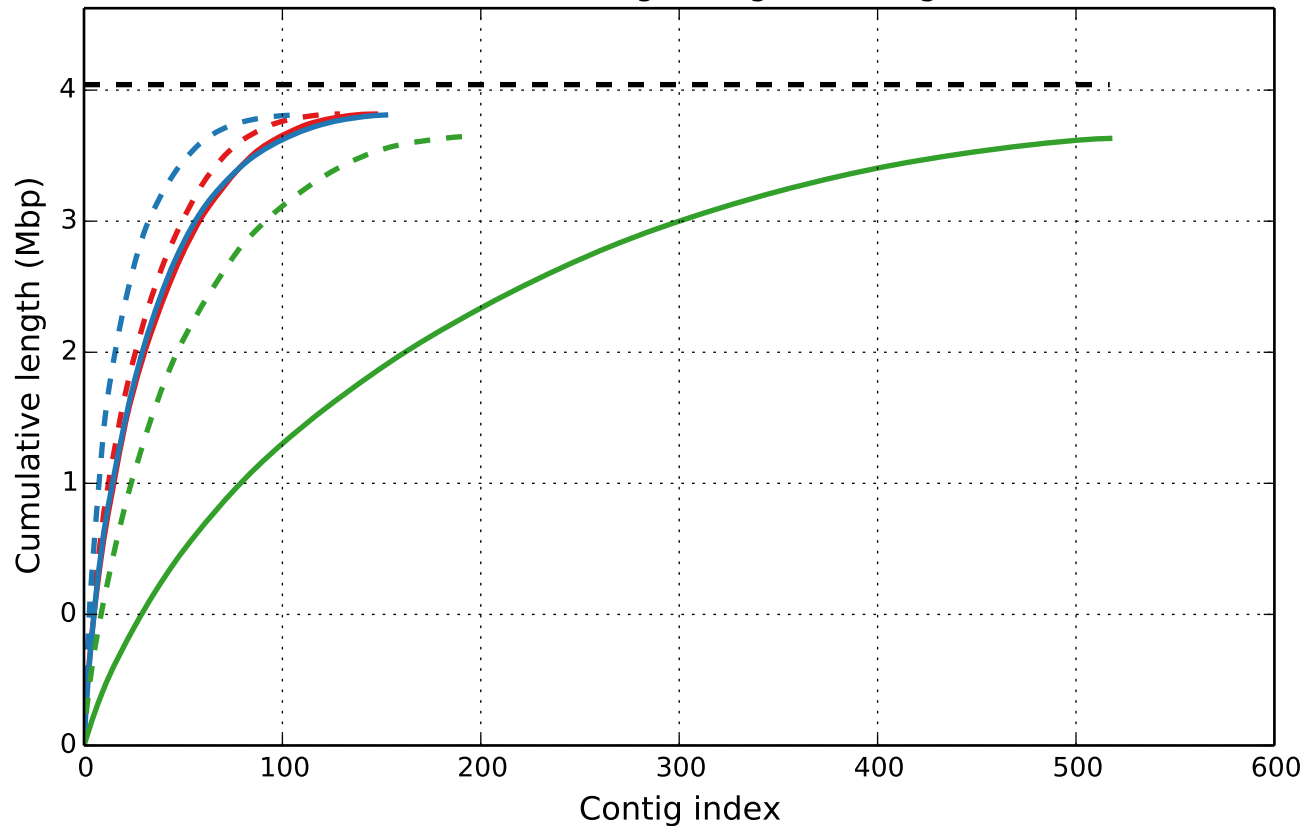
x

— GAGE-B broken — Cabog_7_0 broken — Cabog_8_1 broken - - - Cabog_8_1
- - - GAGE-B - - - Cabog_7_0

NGx

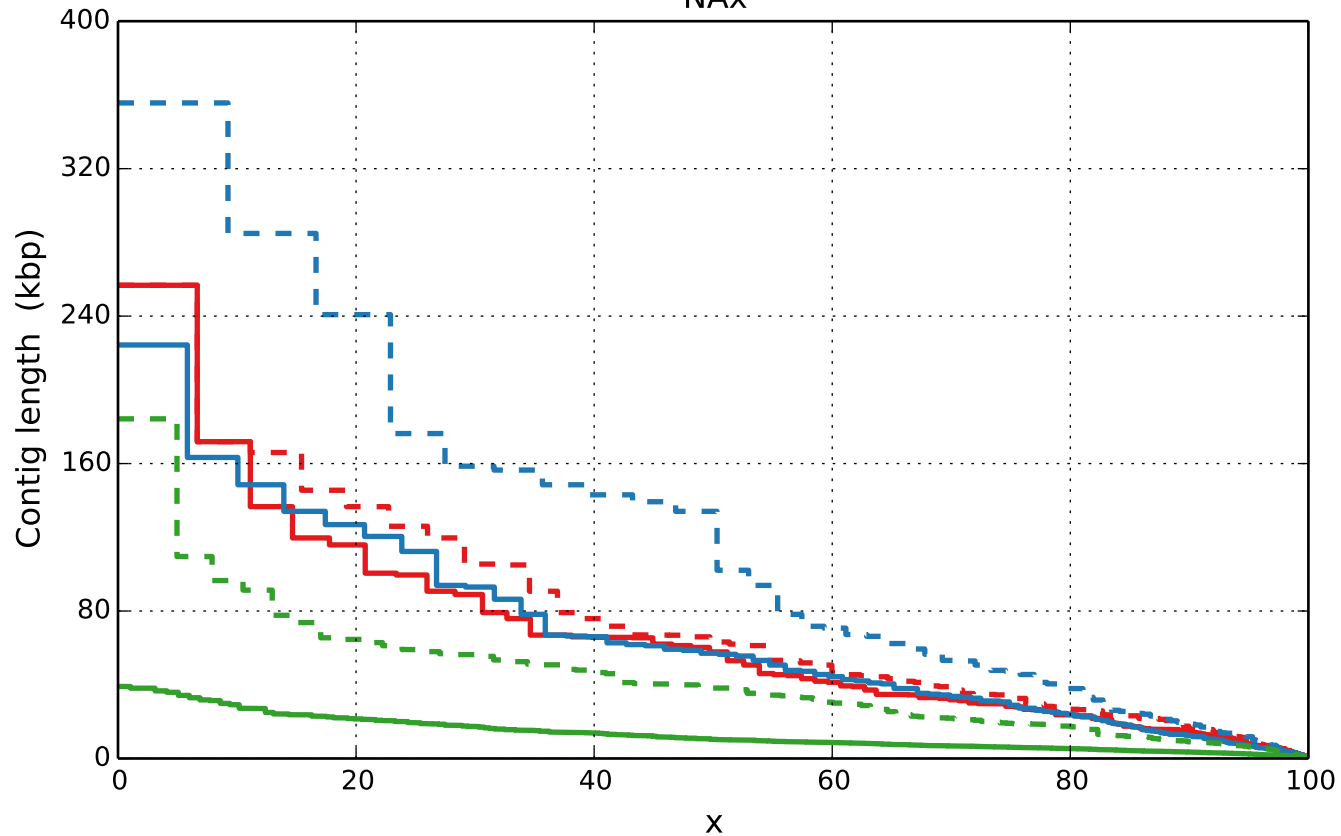


Cumulative length (aligned contigs)

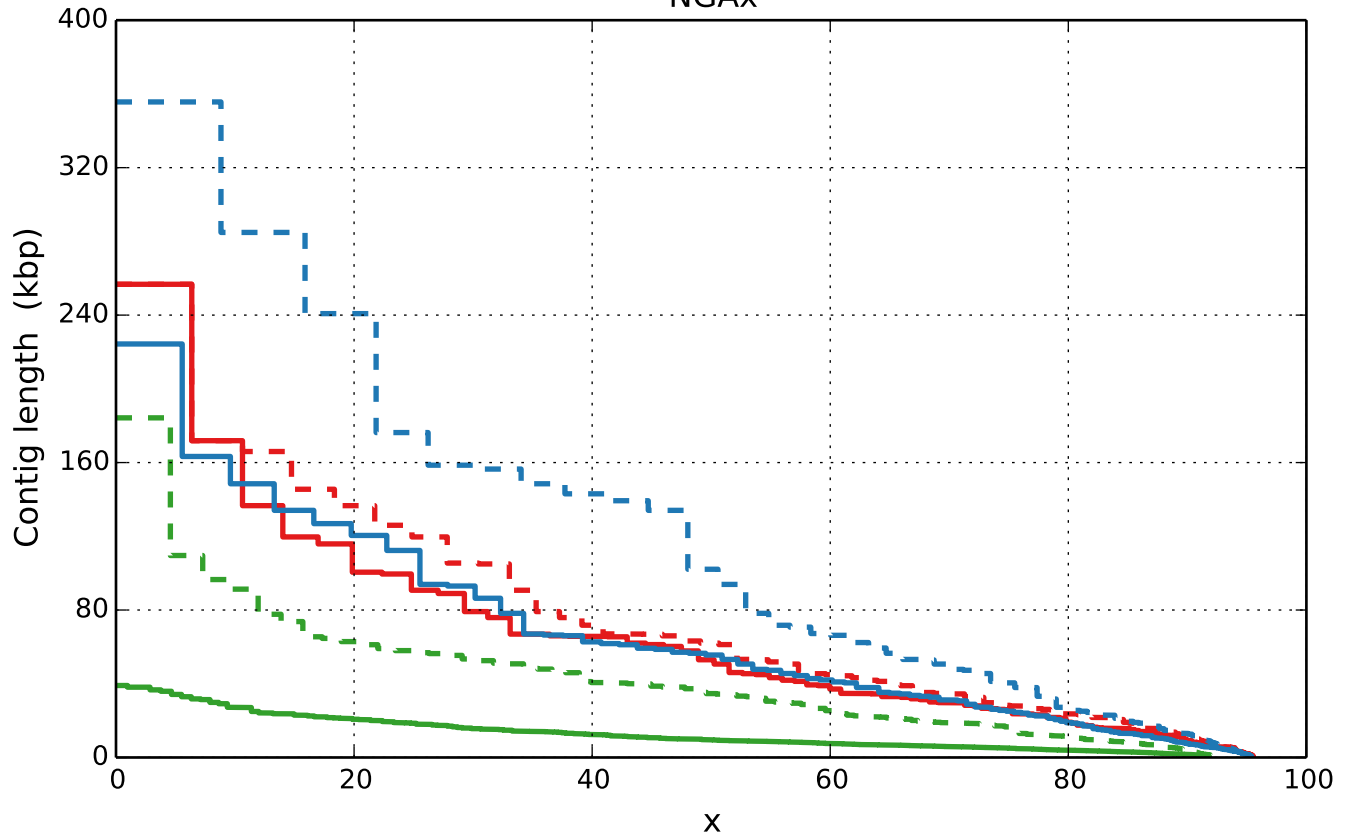


— GAGE-B broken	— Cabog_7_0 broken	— Cabog_8_1 broken	- - Reference
- - GAGE-B	- - Cabog_7_0	- - Cabog_8_1	

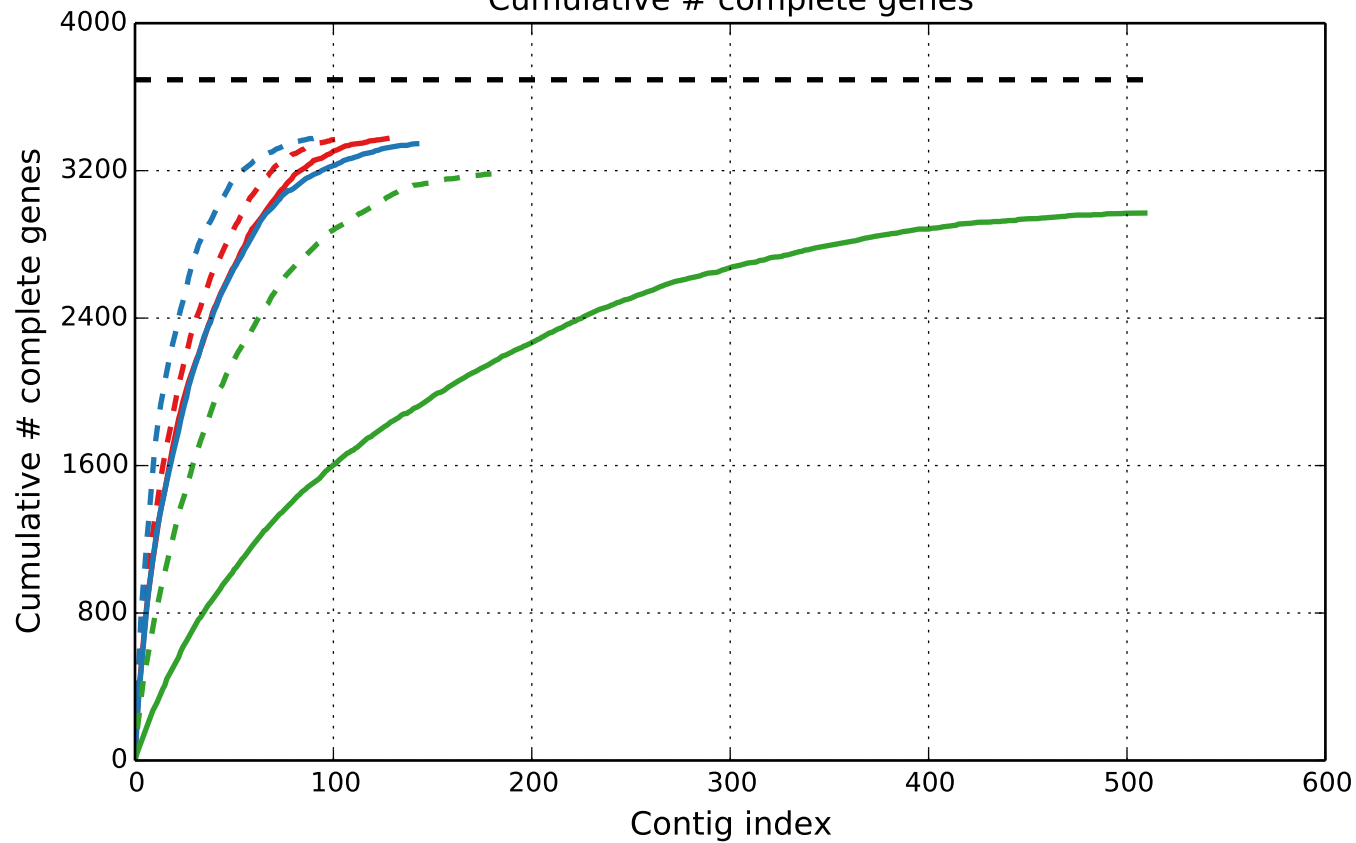
NAx



NGAx



Cumulative # complete genes



— GAGE-B broken — Cabog_7_0 broken — Cabog_8_1 broken - - Reference
- - GAGE-B - - Cabog_7_0 - - Cabog_8_1

complete genes

3500

3000



