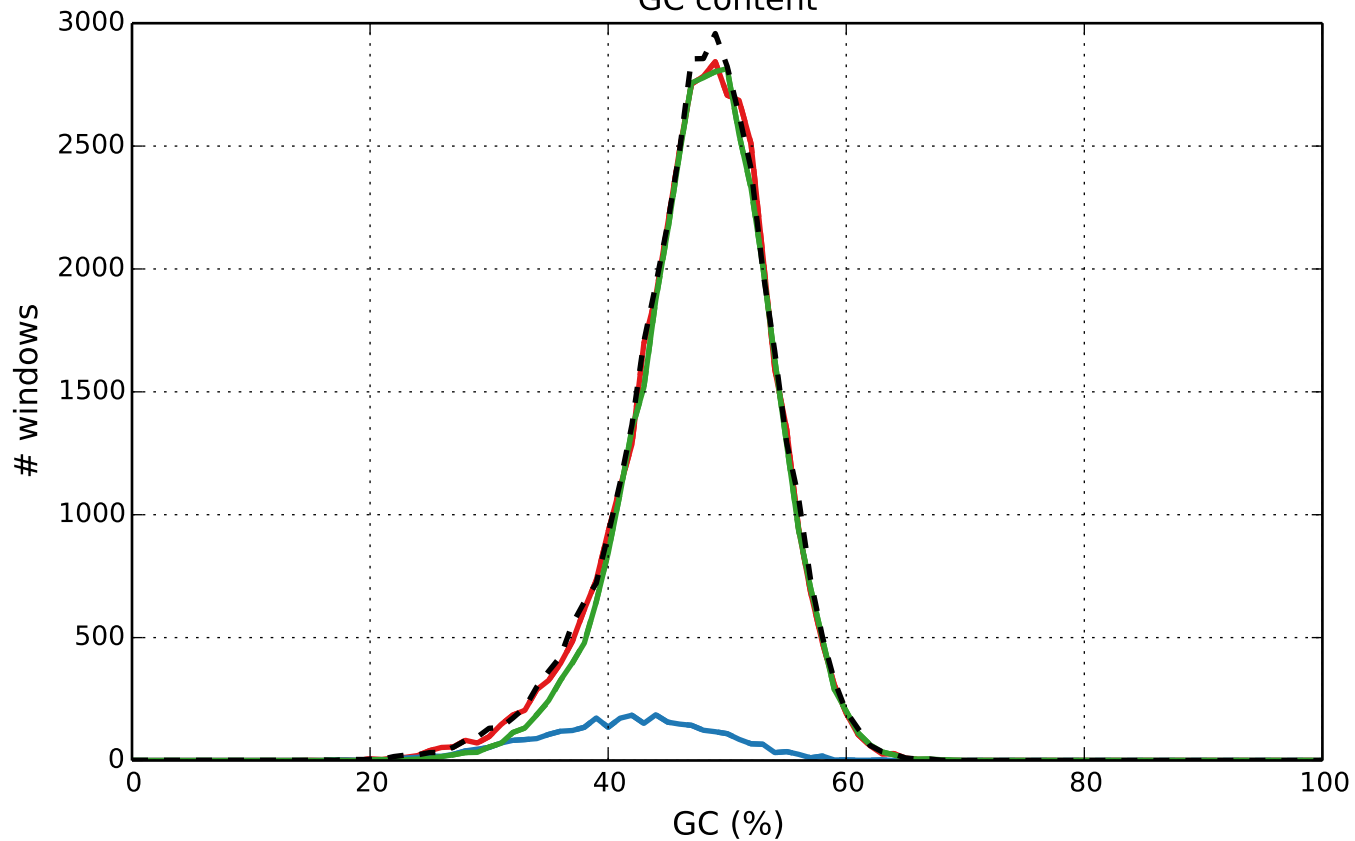
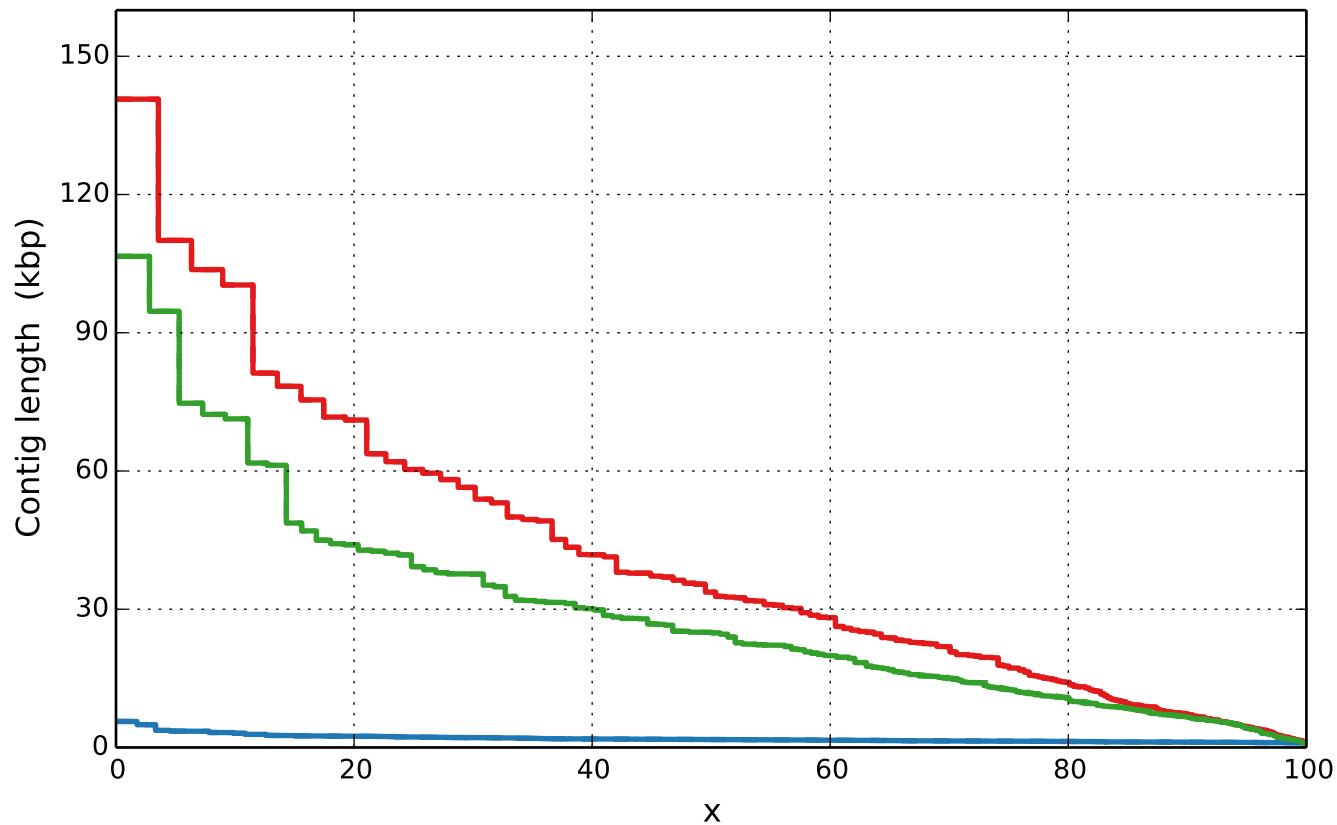


GC content



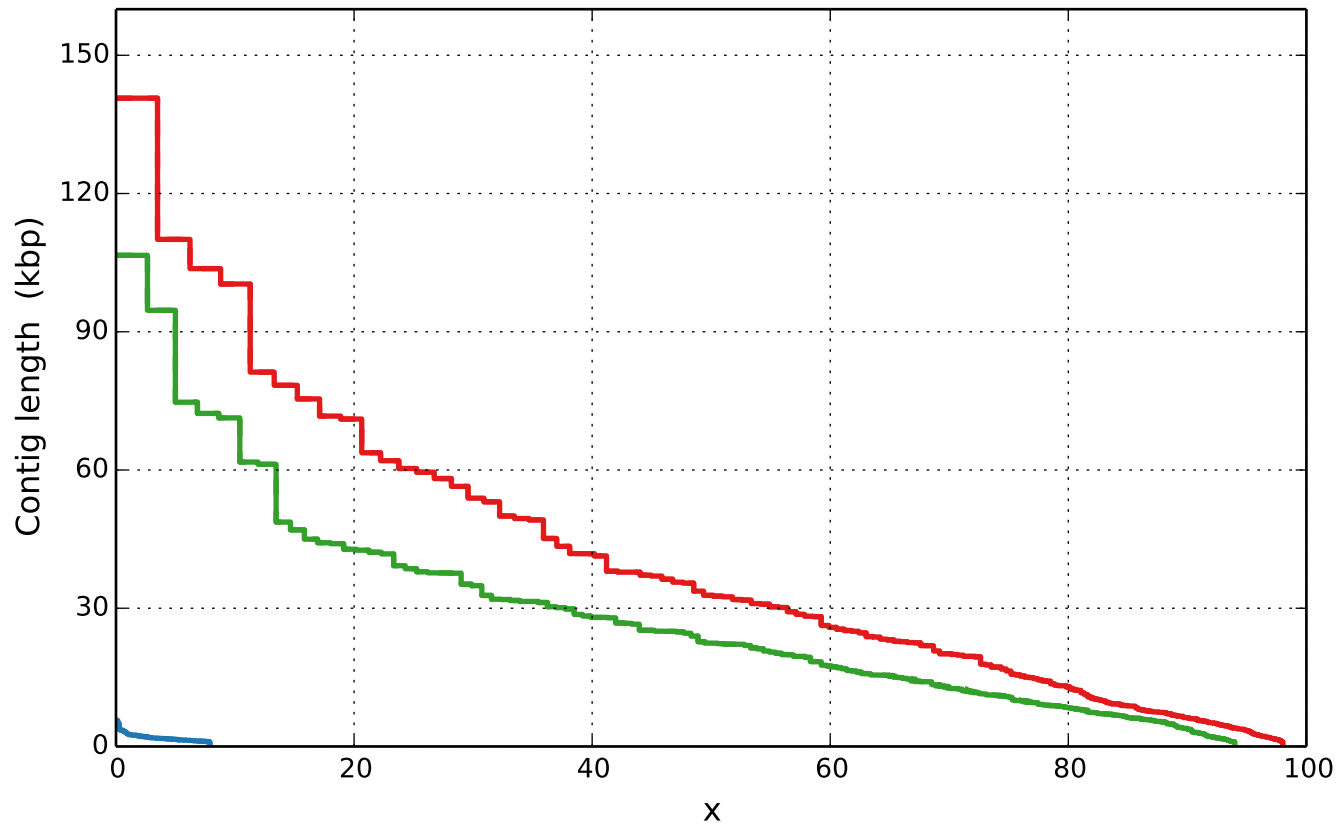
— GAGE-B broken — cabog_7.0 broken — cabog_8.1 broken - - Reference
- - GAGE-B - - cabog_7.0 - - cabog_8.1

Nx

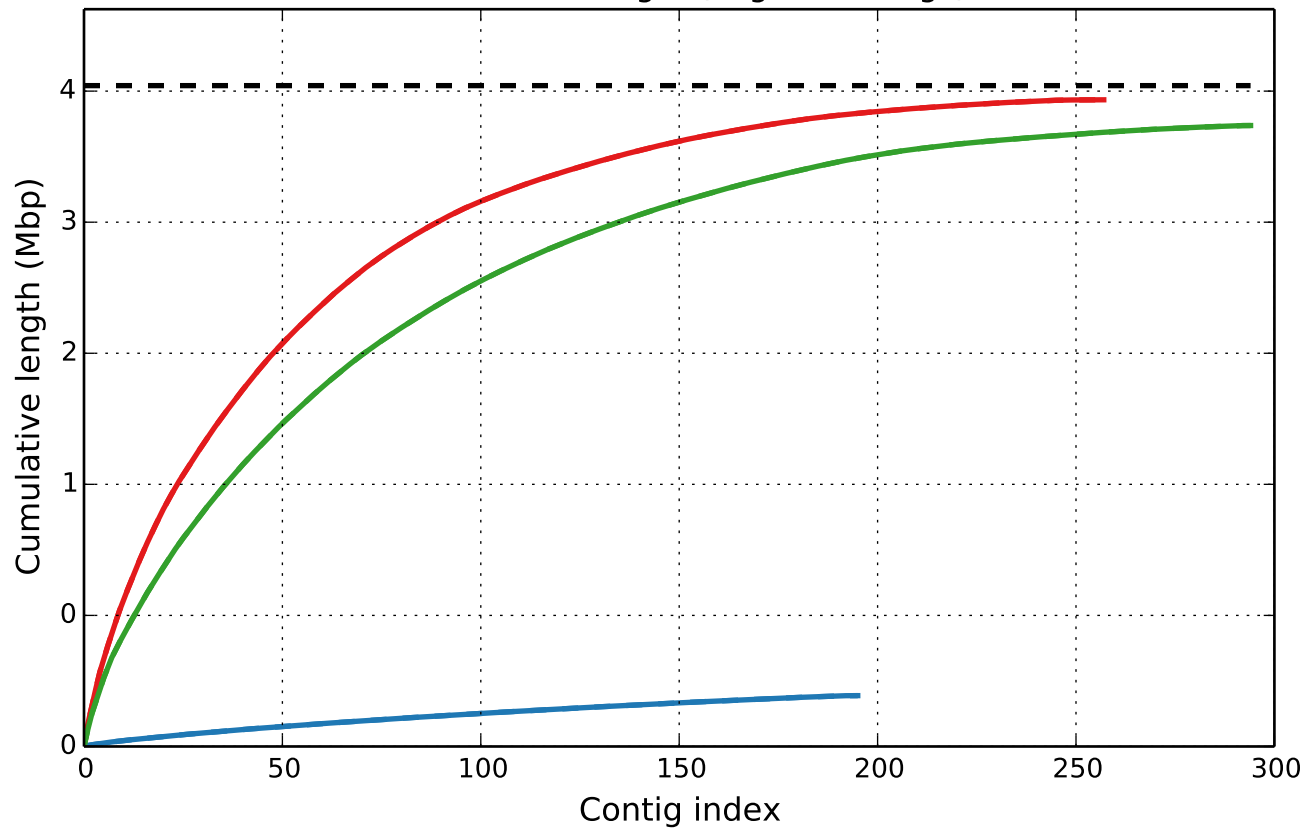


— GAGE-B broken — cabog_7.0 broken — cabog_8.1 broken - - cabog_8.1
- - GAGE-B - - cabog_7.0

NGx

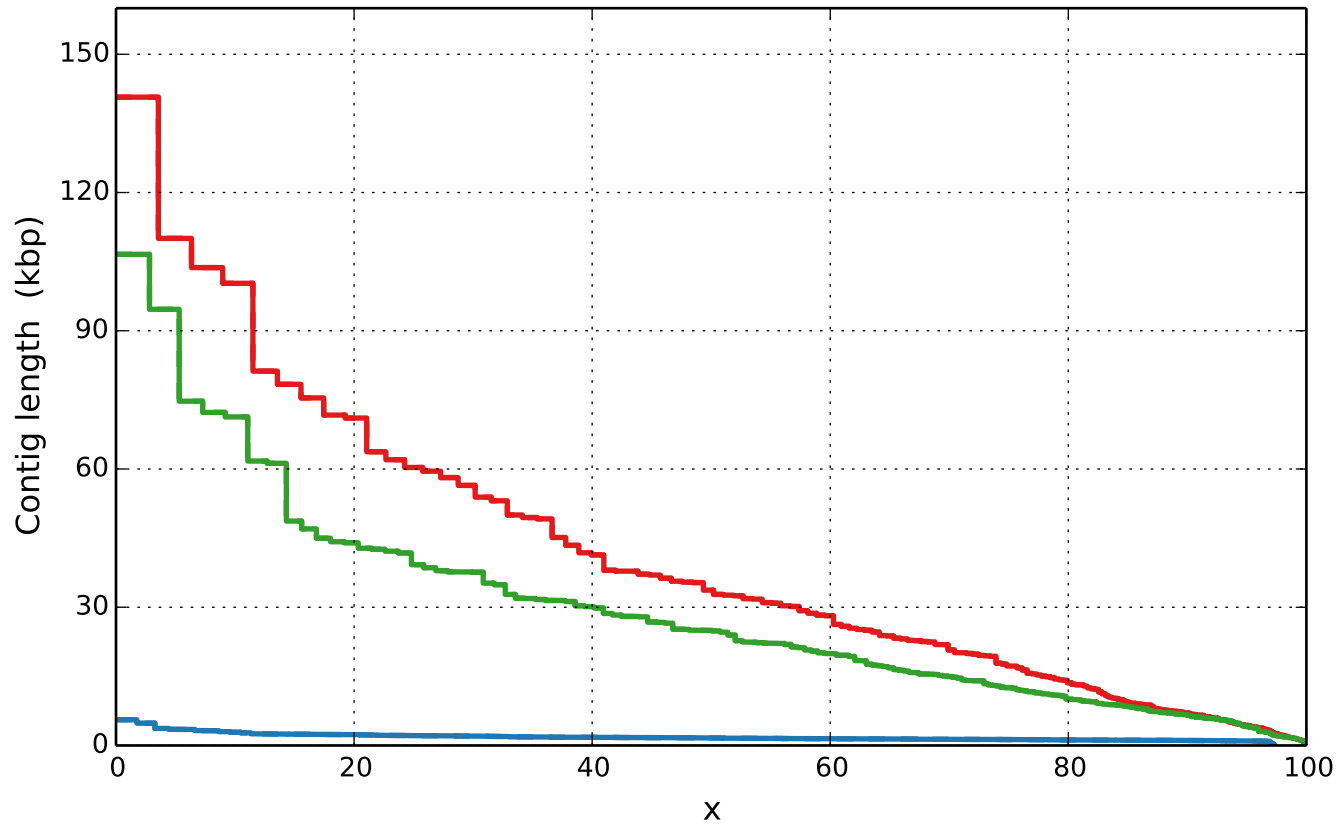


Cumulative length (aligned contigs)



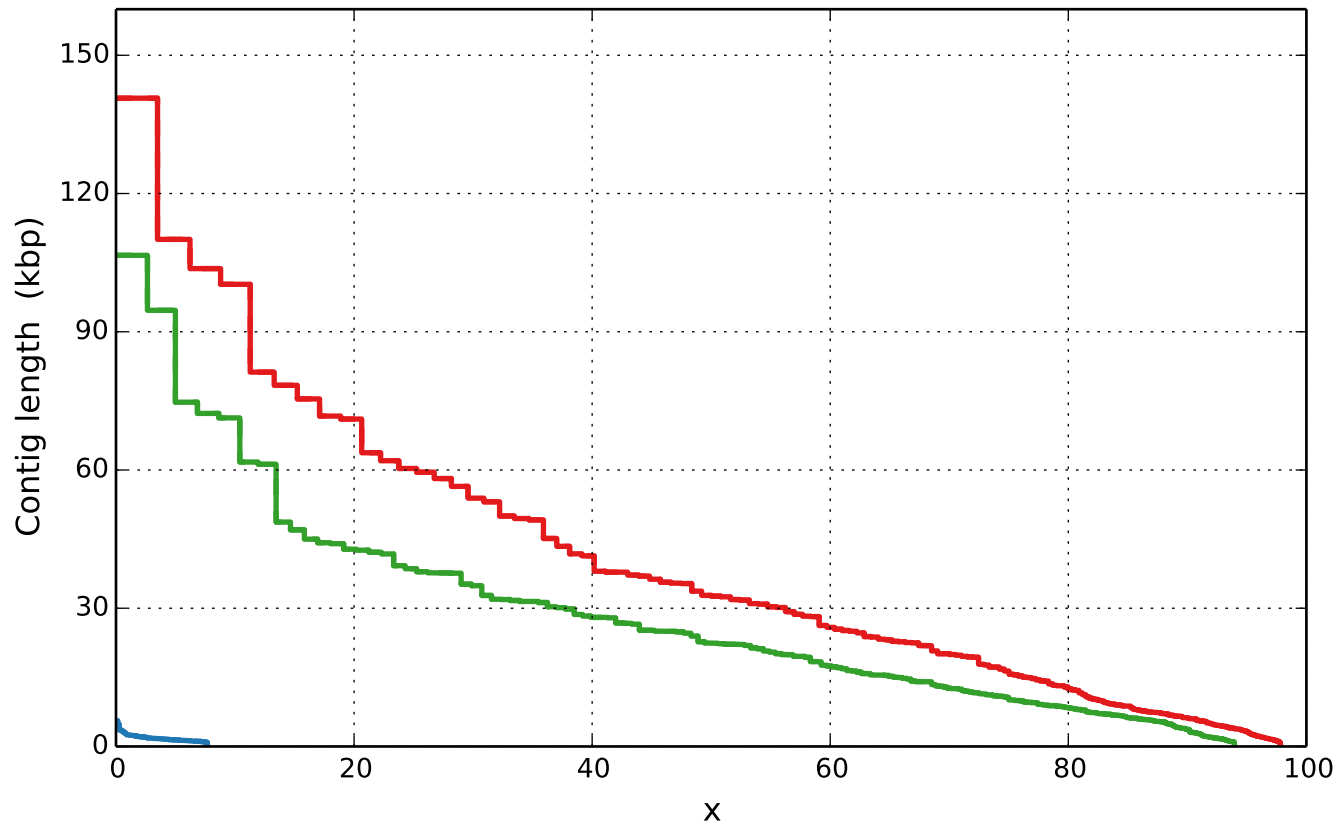
— GAGE-B broken	— cabog_7.0 broken	— cabog_8.1 broken	- - Reference
- - GAGE-B	- - cabog_7.0	- - cabog_8.1	

NAx

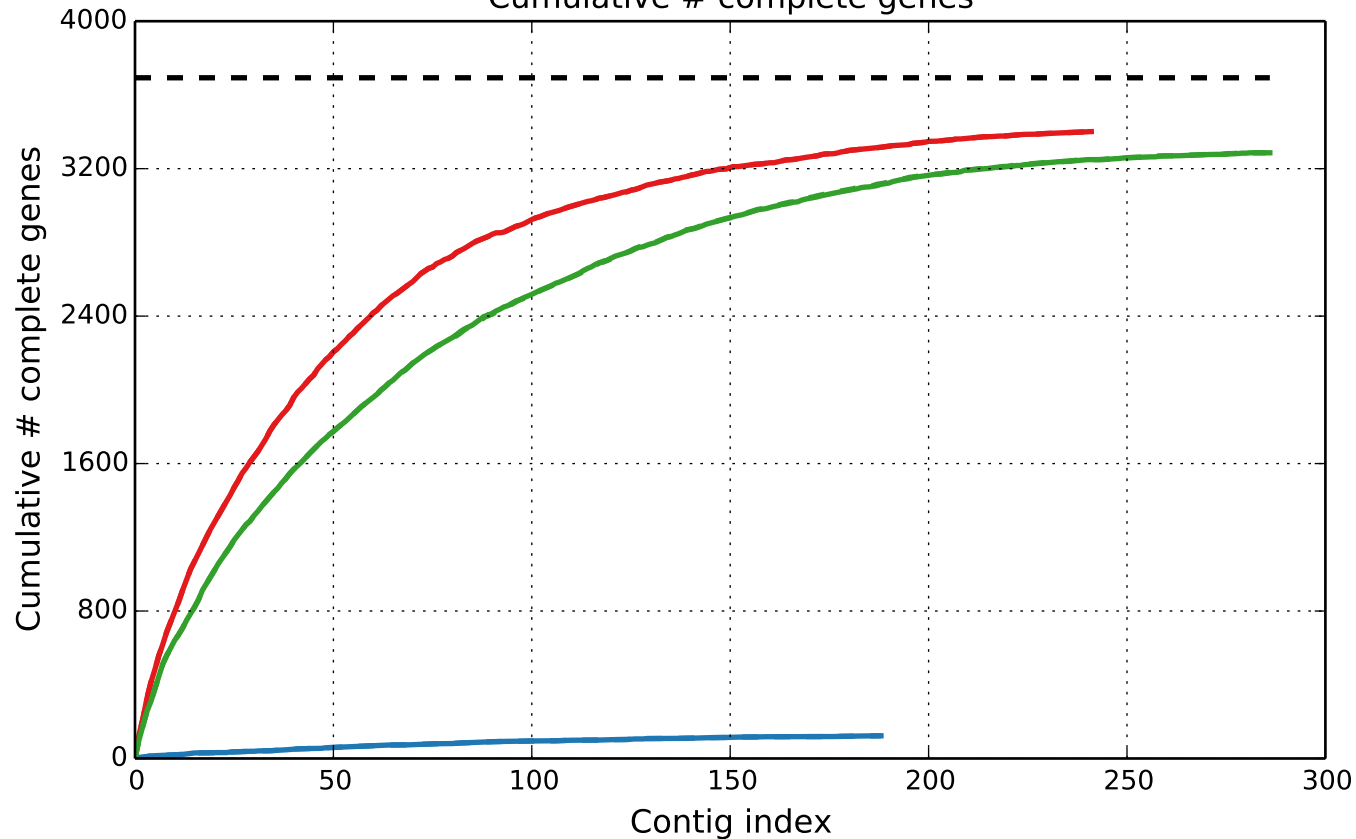


— GAGE-B broken — cabog_7.0 broken — cabog_8.1 broken - - cabog_8.1
- - GAGE-B - - cabog_7.0

NGAx

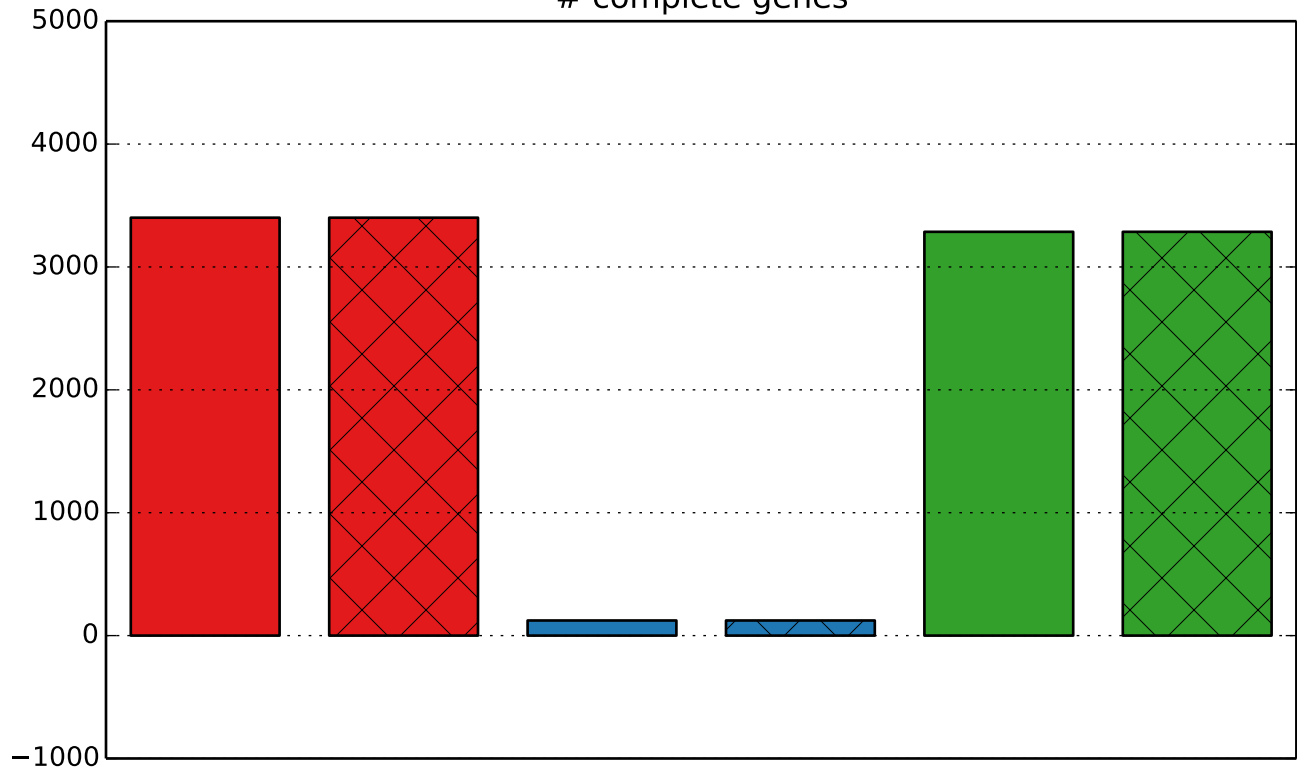


Cumulative # complete genes



— GAGE-B broken — cabog_7.0 broken — cabog_8.1 broken - - Reference
- - GAGE-B - - cabog_7.0 - - cabog_8.1

complete genes



GAGE-B broken

cabog_7.0 broken

cabog_8.1 broken

cabog_8.1

GAGE-B

cabog_7.0

