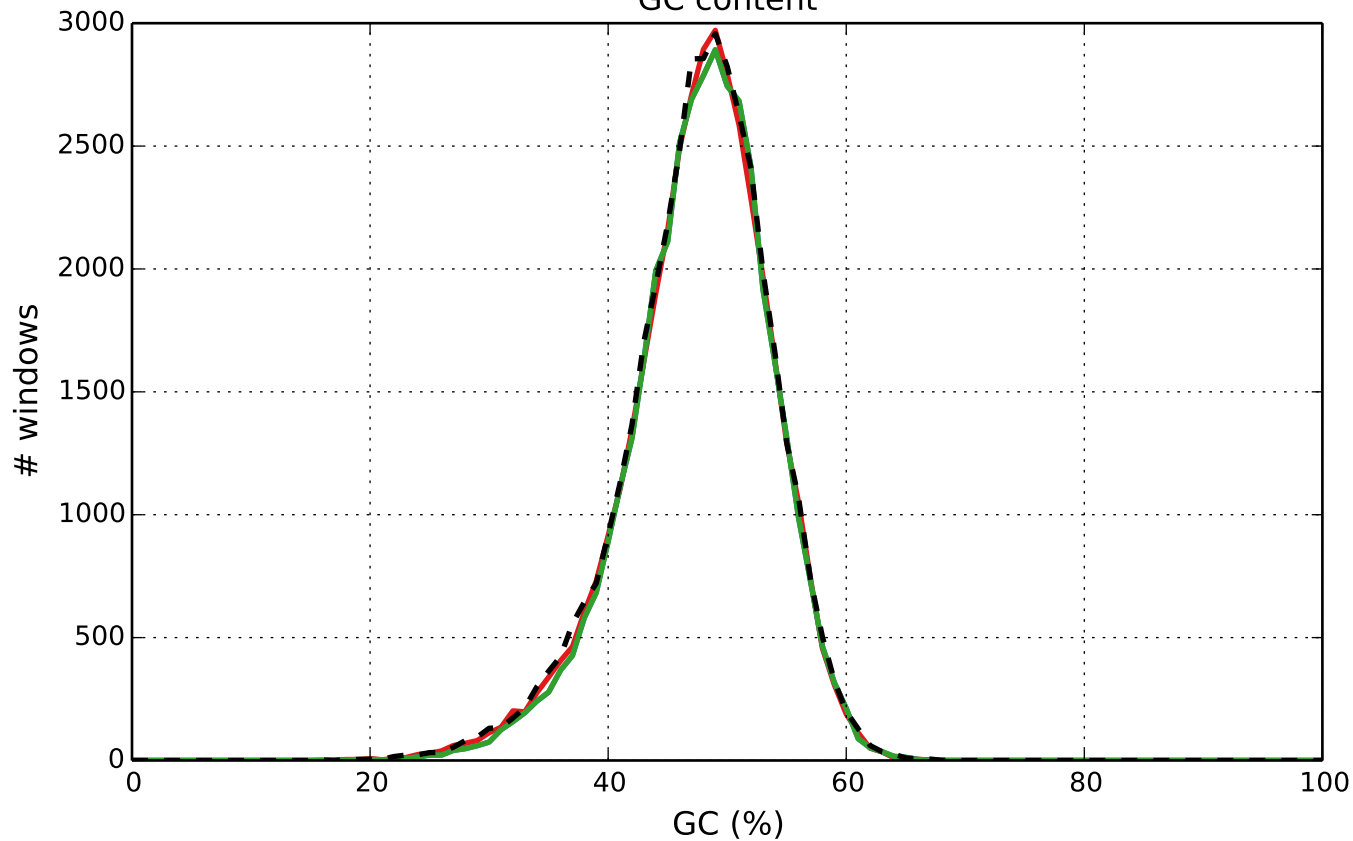


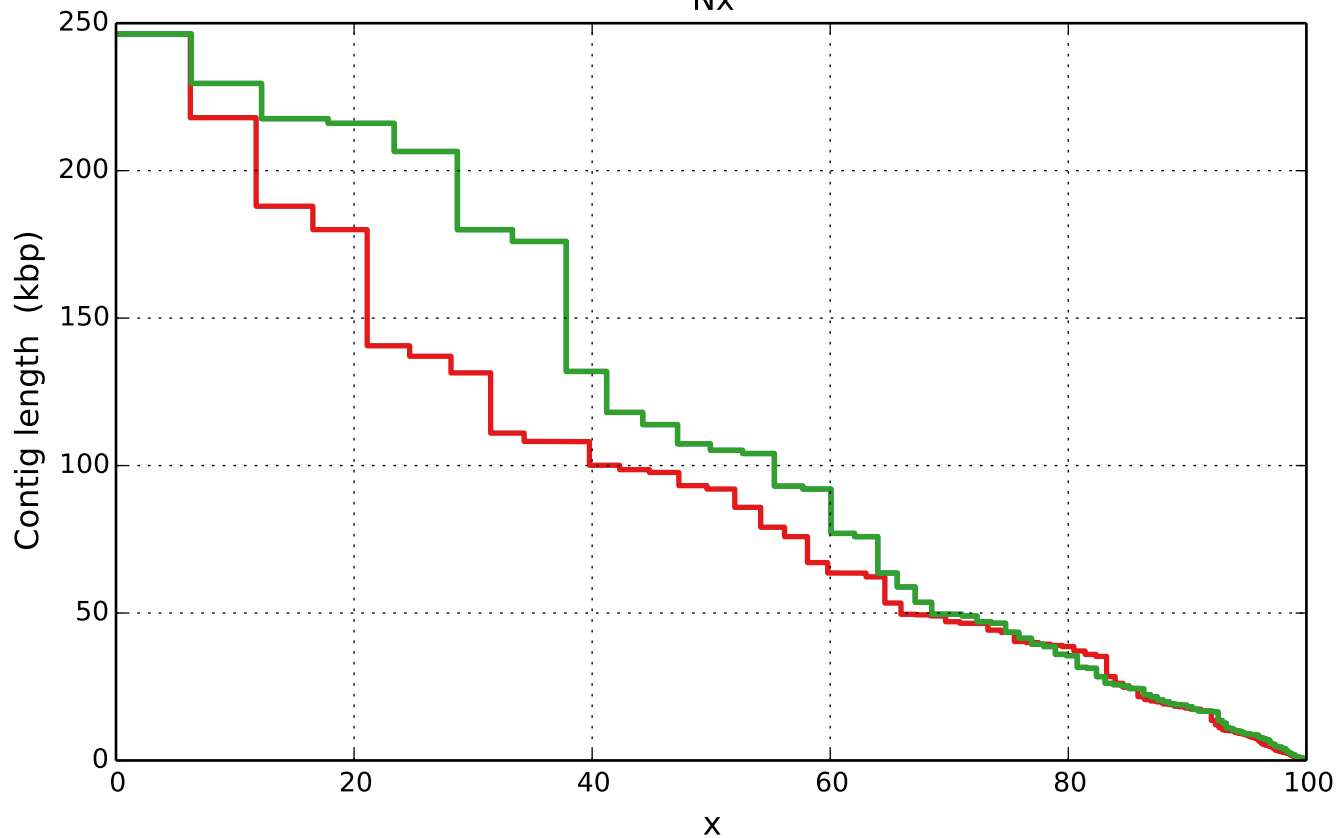
— GAGE-B-MiSeq — Velvet_1.2.08-MiSeq — Velvet_1.2.10-MiSeq - - Referen

GC content



— GAGE-B-MiSeq — Velvet_1.2.08-MiSeq — Velvet_1.2.10-MiSeq - - Referen

Nx

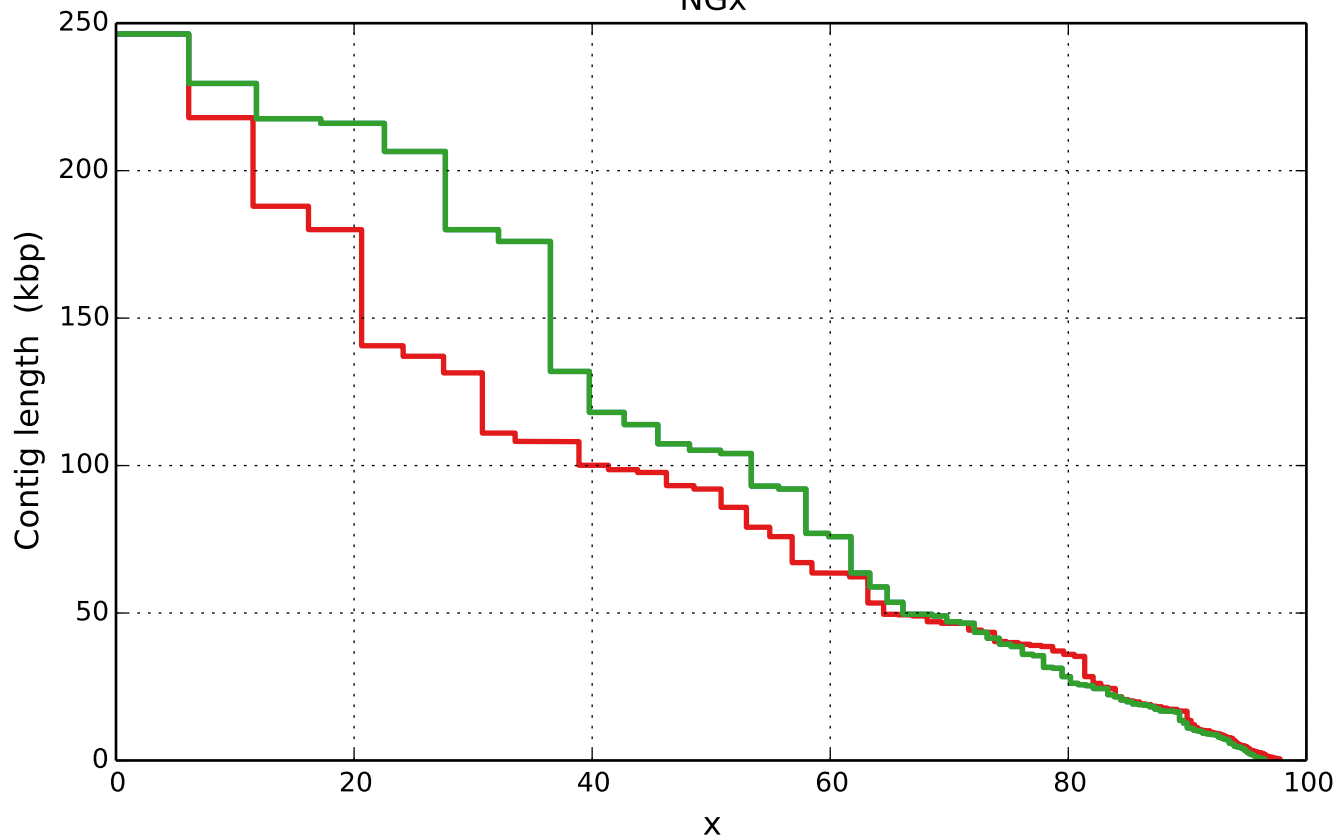


GAGE-B-MiSeq

Velvet_1.2.08-MiSeq

Velvet_1.2.10-MiSeq

NGx

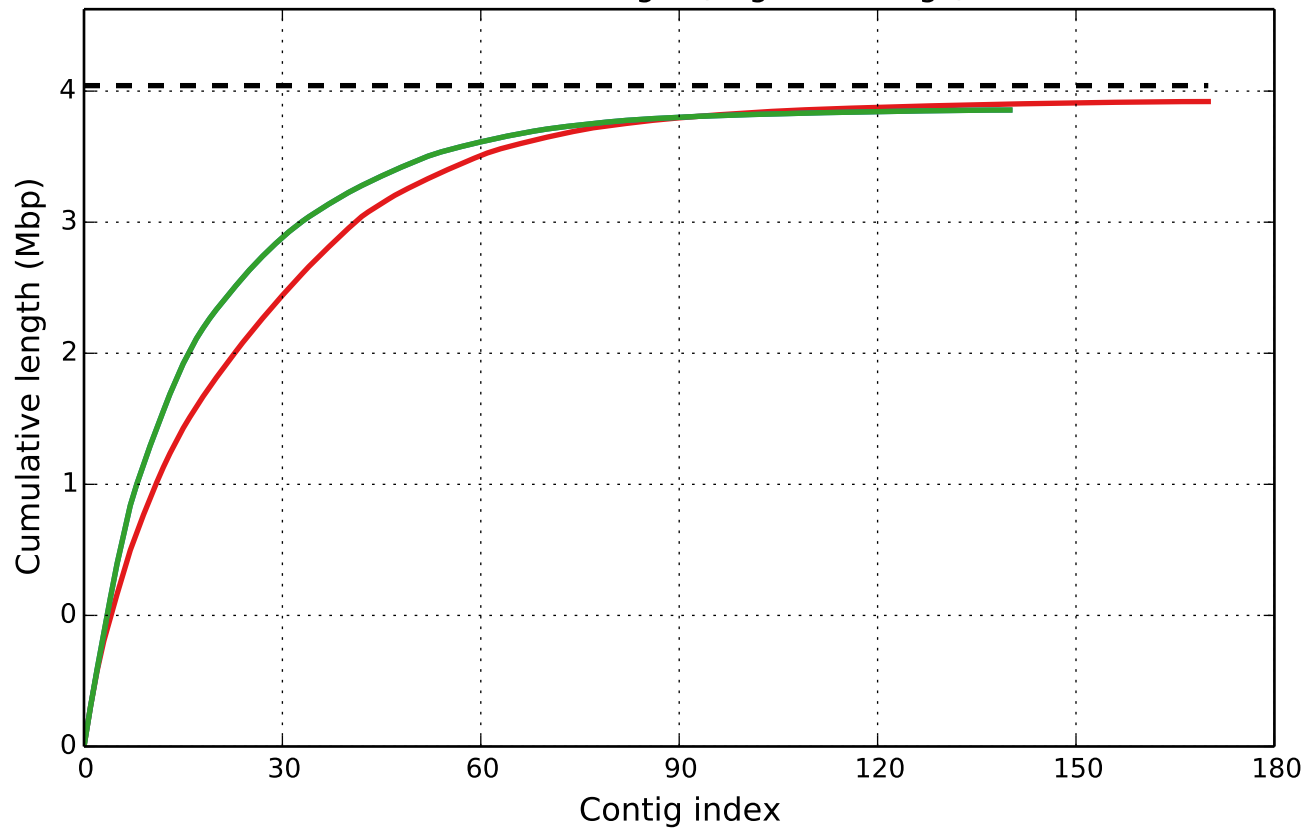


GAGE-B-MiSeq

Velvet_1.2.08-Miseq

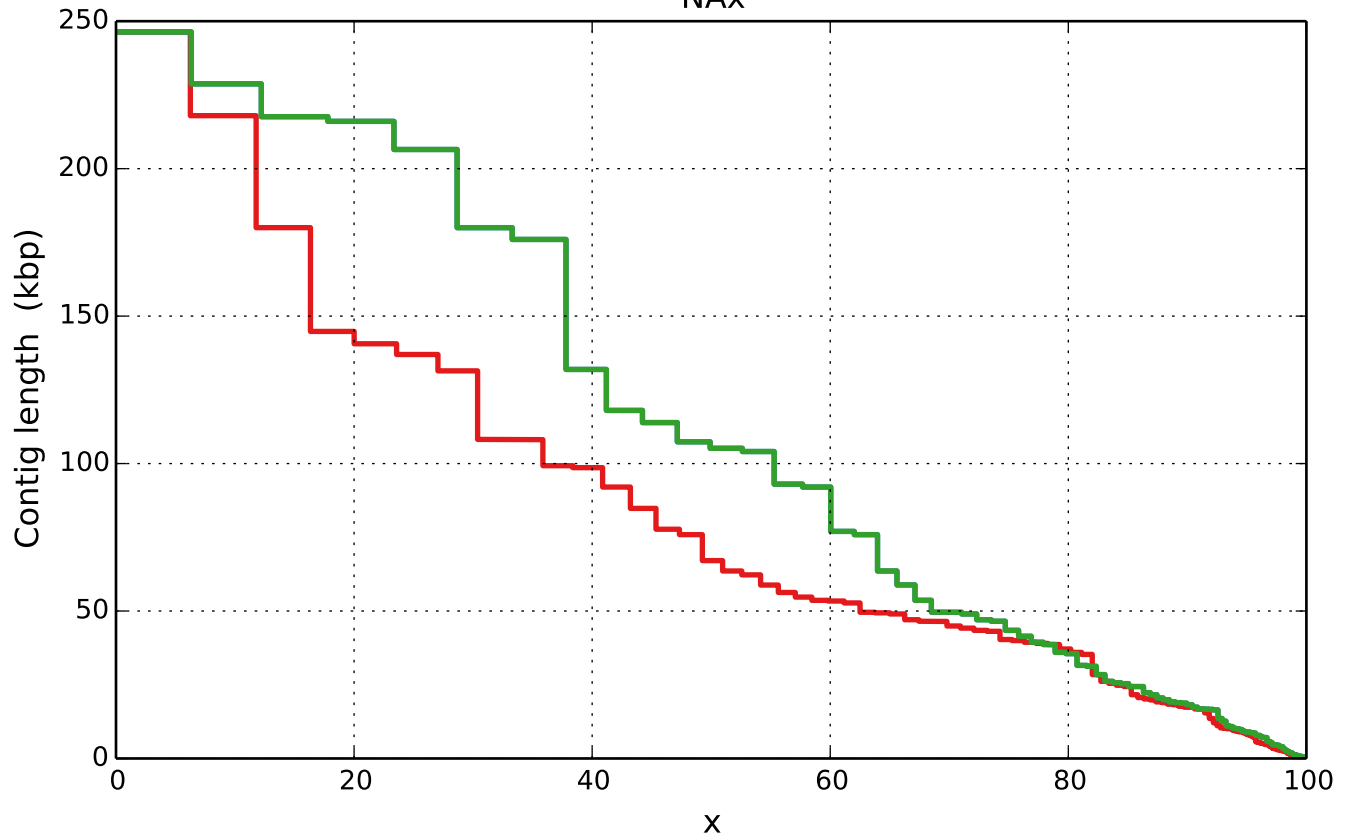
Velvet_1.2.10-Miseq

Cumulative length (aligned contigs)



— GAGE-B-MiSeq — Velvet_1.2.08-MiSeq — Velvet_1.2.10-MiSeq - - Referer

NAx

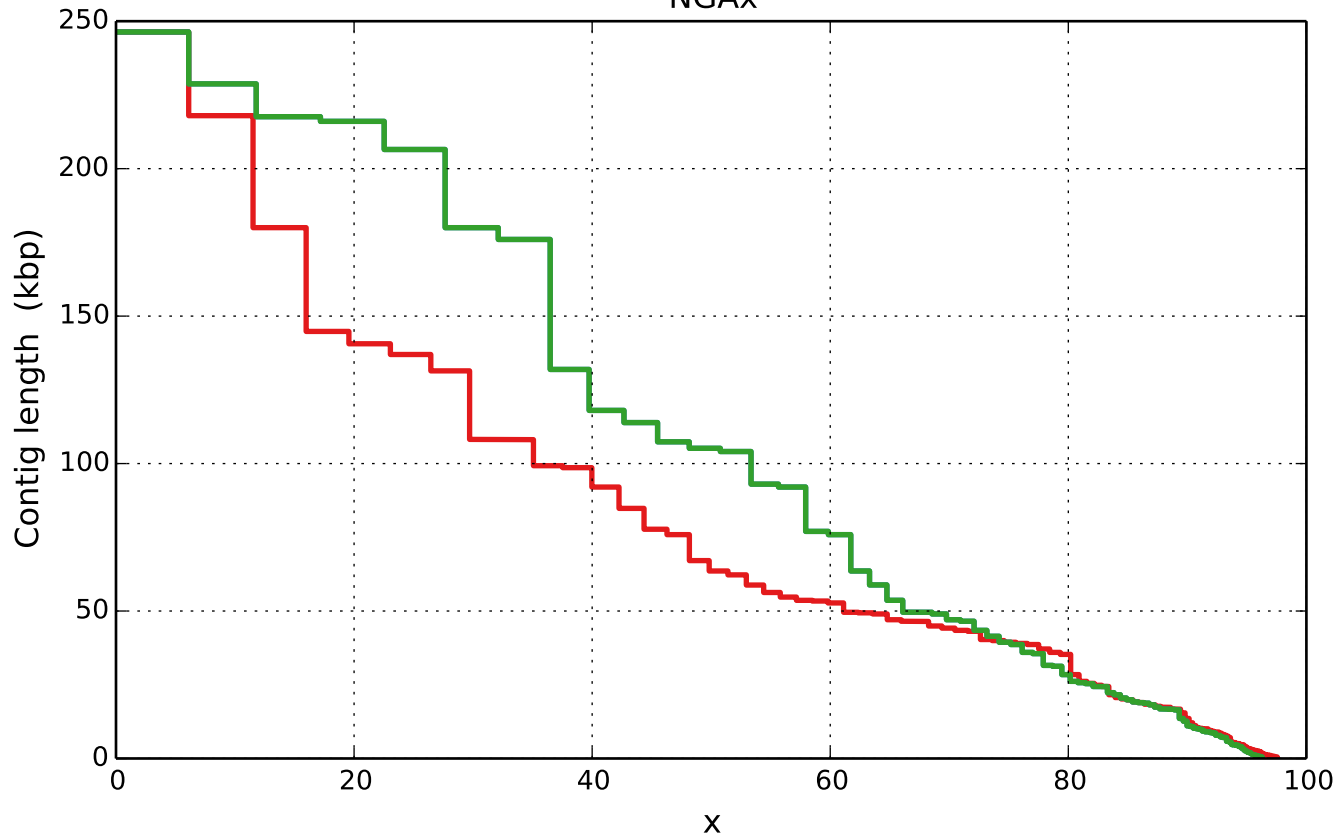


GAGE-B-MiSeq

Velvet_1.2.08-Miseq

Velvet_1.2.10-Miseq

NGAx

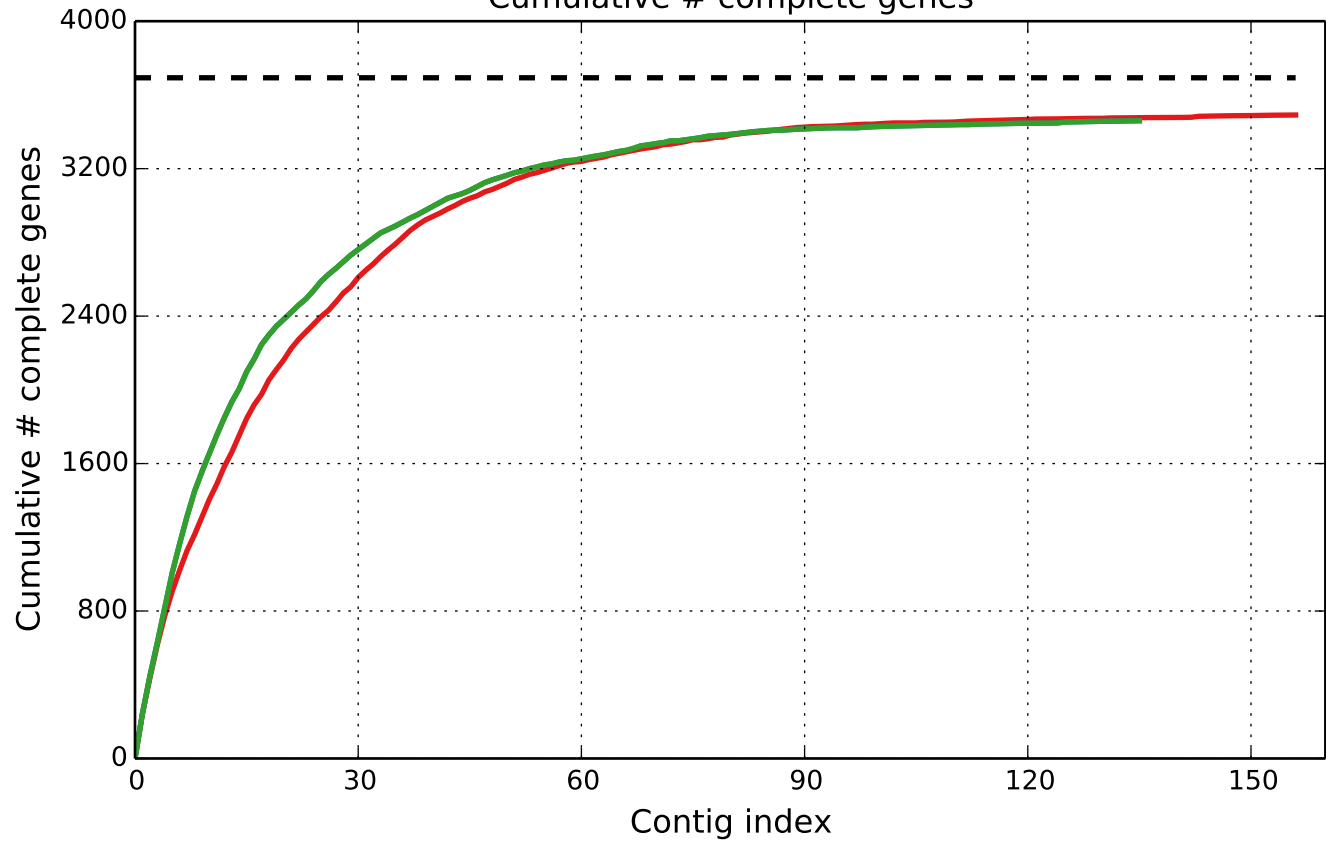


GAGE-B-MiSeq

Velvet_1.2.08-MiSeq

Velvet_1.2.10-MiSeq

Cumulative # complete genes



— GAGE-B-MiSeq — Velvet_1.2.08-MiSeq — Velvet_1.2.10-MiSeq - - Referer

complete genes

