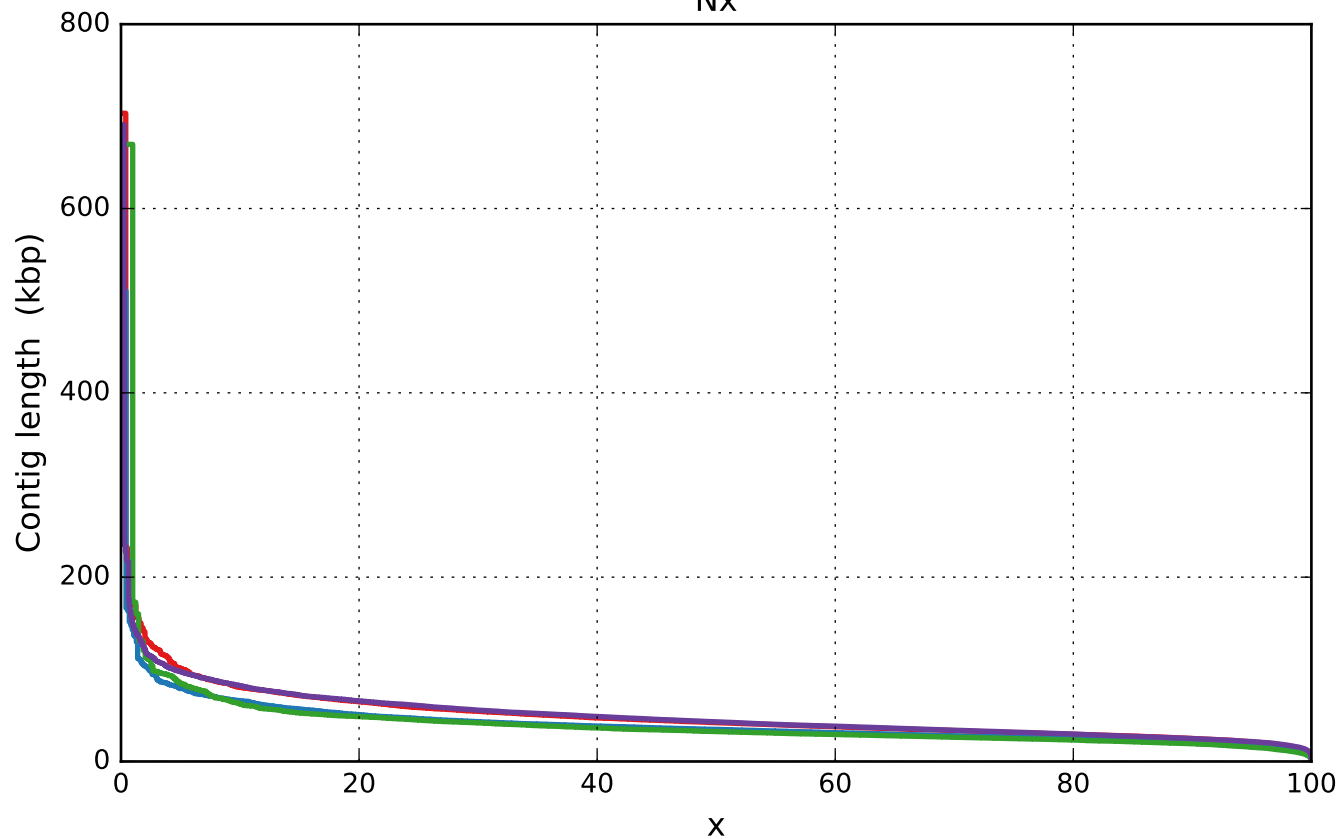


Report

	Q1133.091416.GS350M.contigs	Q1133.092116.GS170M.contigs	Q1171.091516.GS350M.contigs	Q1171.092116.GS170M.contigs
# contigs (≥ 0 bp)	4519	3856	2301	6350
# contigs (≥ 1000 bp)	4519	3856	2301	6350
# contigs (≥ 5000 bp)	4500	3825	2275	6338
# contigs (≥ 10000 bp)	4448	3715	2181	6306
# contigs (≥ 25000 bp)	3576	2406	1303	5035
# contigs (≥ 50000 bp)	909	369	171	1401
Total length (≥ 0 bp)	175616198	120328783	68041971	251289743
Total length (≥ 1000 bp)	175616198	120328783	68041971	251289743
Total length (≥ 5000 bp)	175558290	120219557	67947326	251244112
Total length (≥ 10000 bp)	175166554	119349548	67213729	250998195
Total length (≥ 25000 bp)	157882757	94781418	50479495	225400407
Total length (≥ 50000 bp)	63318102	24914965	12116294	96314572
# contigs	4519	3856	2301	6350
Largest contig	703356	510354	669621	690841
Total length	175616198	120328783	68041971	251289743
GC (%)	32.16	30.67	33.01	30.58
N50	42123	34787	32493	43021
N75	31292	26330	24823	31938
L50	1445	1237	726	2036
L75	2653	2230	1326	3736
# N's per 100 kbp	0.00	0.00	0.00	0.00
# predicted genes (unique)	14755	9835	7111	19289
# predicted genes (≥ 0 bp)	15011	9980	7227	19562
# predicted genes (≥ 300 bp)	10424	6881	5143	14022
# predicted genes (≥ 1500 bp)	2657	2091	1347	4259
# predicted genes (≥ 3000 bp)	800	753	468	1539

All statistics are based on contigs of size ≥ 500 bp, unless otherwise noted (e.g., "# contigs (≥ 0 bp)" and "Total length (≥ 0 bp)" include all contigs).

Nx



3.091416.GS350M.contigs
3.092116.GS170M.contigs

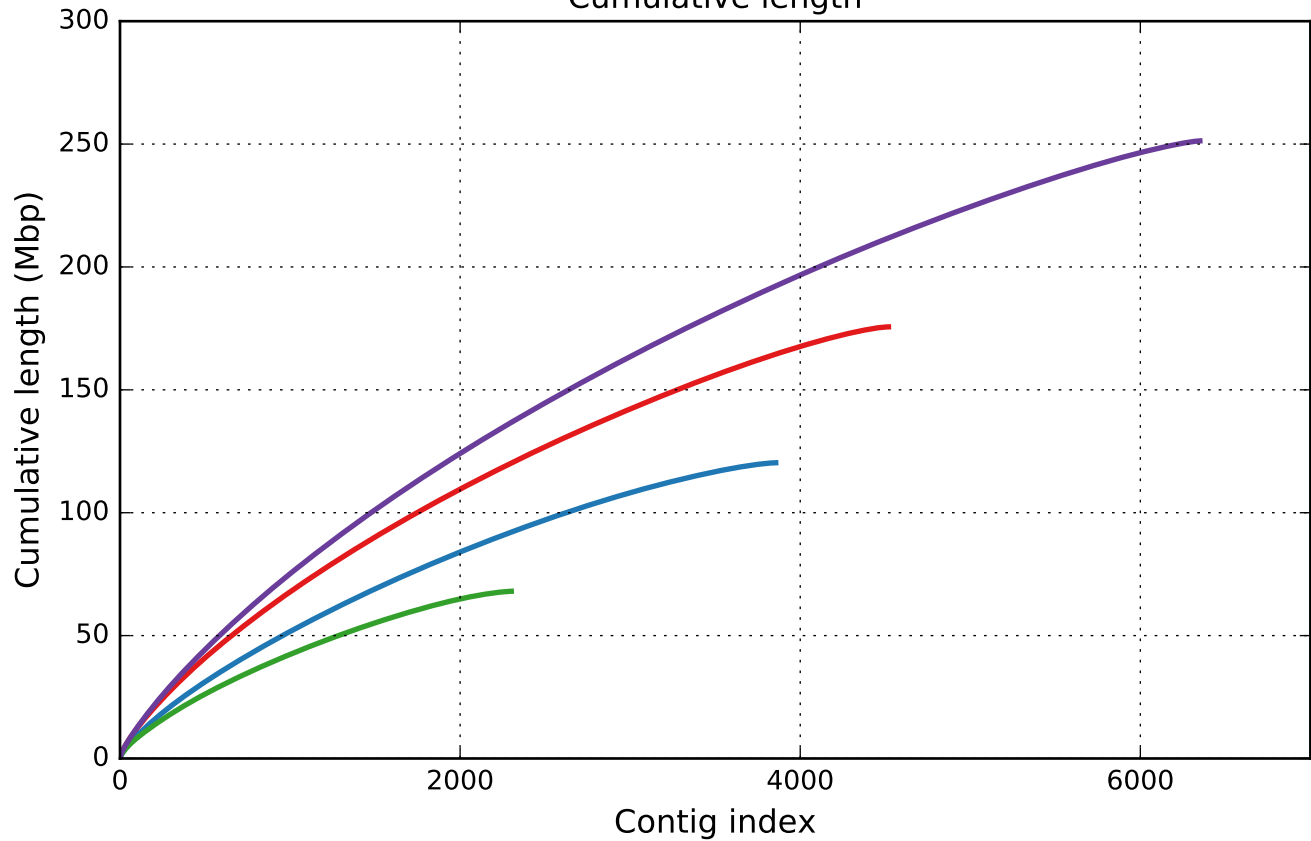


Q1171.091516.GS350M.contigs



Q1171.092116.GS

Cumulative length



3.091416.GS350M.contigs

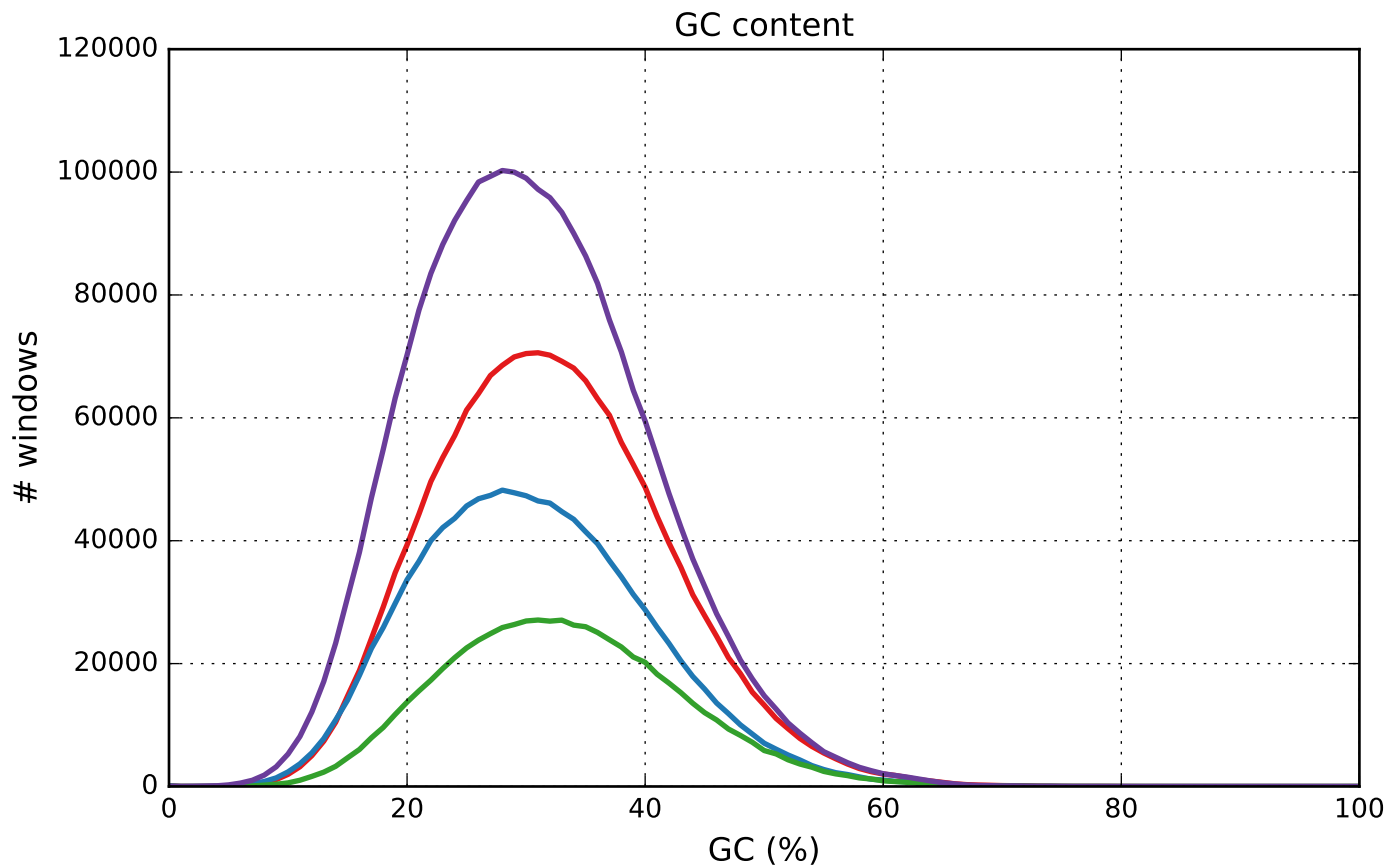


Q1171.091516.GS350M.contigs



Q1171.092116.GS350M.contigs

3.092116.GS170M.contigs



3.091416.GS350M.contigs



Q1171.091516.GS350M.contigs



Q1171.092116.GS

3.092116.GS170M.contigs