

MAPPING STATS

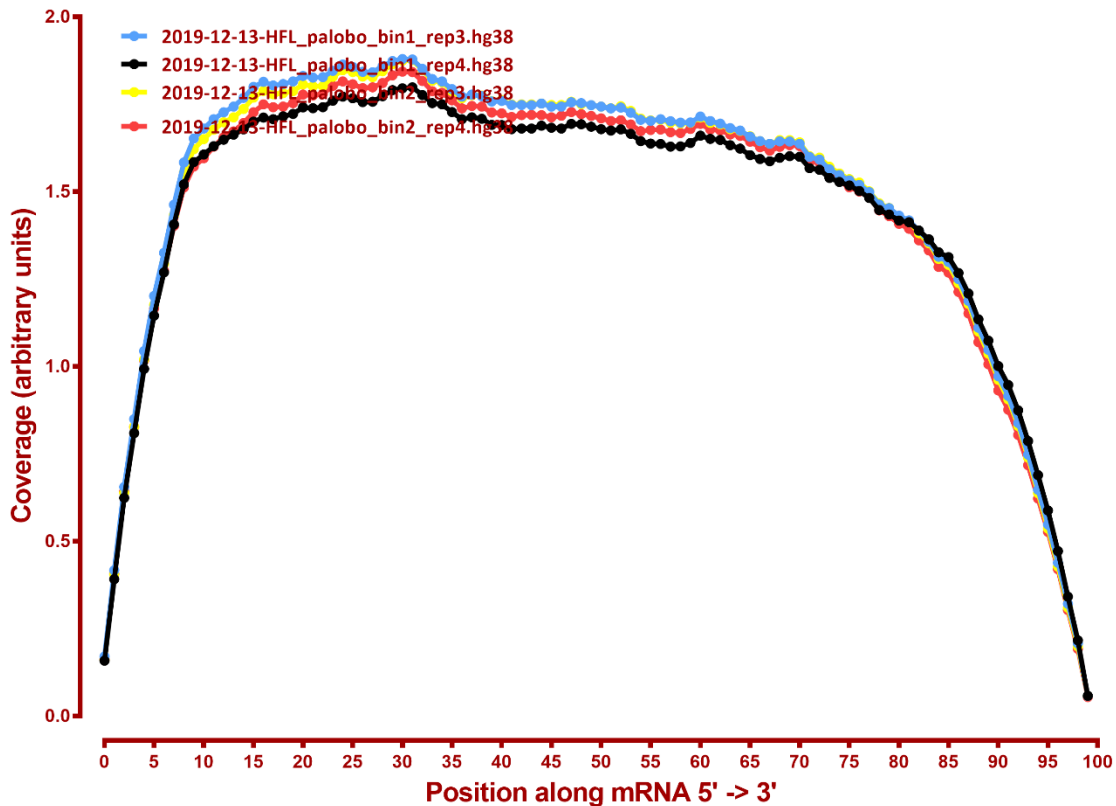
Library	Read Length	Mapping	Species	Raw fragments	Complexity	Unique	Unique Splices	Multi	Multi Splices	Fraction mapped
2019-12-13-HFL_palobo_bin1_rep3	2x150	STAR-2.5.3a	dm6	54,731,399	0.39	11,145,618	3,526,124	2,281,373	199,983	0.16
2019-12-13-HFL_palobo_bin1_rep3	2x150	STAR-2.5.3a	hg38	54,731,399	0.38	44,694,658	33,458,944	8,730,994	2,416,734	0.82
2019-12-13-HFL_palobo_bin1_rep4	2x150	STAR-2.5.3a	dm6	54,901,635	0.36	19,035,257	6,042,879	3,983,748	350,034	0.27
2019-12-13-HFL_palobo_bin1_rep4	2x150	STAR-2.5.3a	hg38	54,901,635	0.38	39,447,058	29,172,582	6,773,937	2,036,861	0.71
2019-12-13-HFL_palobo_bin2_rep3	2x150	STAR-2.5.3a	dm6	51,217,985	0.36	10,201,173	3,202,803	2,193,201	183,329	0.15
2019-12-13-HFL_palobo_bin2_rep3	2x150	STAR-2.5.3a	hg38	51,217,985	0.35	41,561,883	31,113,489	8,779,662	2,209,318	0.82
2019-12-13-HFL_palobo_bin2_rep4	2x150	STAR-2.5.3a	dm6	51,685,456	0.35	16,046,210	5,133,506	3,321,591	305,005	0.24
2019-12-13-HFL_palobo_bin2_rep4	2x150	STAR-2.5.3a	hg38	51,685,456	0.36	37,651,199	29,030,475	6,901,007	2,047,667	0.73
2019-12-13-HFL_palobo_bin3_rep3	2x150	STAR-2.5.3a	dm6	45,856,164	0.36	8,111,941	2,581,037	1,590,865	145,913	0.14
2019-12-13-HFL_palobo_bin3_rep3	2x150	STAR-2.5.3a	hg38	45,856,164	0.34	37,435,188	29,147,984	8,107,861	2,098,287	0.84
2019-12-13-HFL_palobo_bin3_rep4	2x150	STAR-2.5.3a	dm6	42,321,118	0.33	11,694,552	3,734,966	2,416,597	215,133	0.21
2019-12-13-HFL_palobo_bin3_rep4	2x150	STAR-2.5.3a	hg38	42,321,118	0.33	32,110,692	25,446,572	4,592,947	1,648,915	0.75
2019-12-13-HFL_palobo_bin4_rep3	2x150	STAR-2.5.3a	dm6	52,203,315	0.31	7,811,929	2,431,363	1,675,790	134,490	0.12
2019-12-13-HFL_palobo_bin4_rep3	2x150	STAR-2.5.3a	hg38	52,203,315	0.28	44,788,534	34,021,692	8,071,891	2,351,357	0.85
2019-12-13-HFL_palobo_bin4_rep4	2x150	STAR-2.5.3a	dm6	47,394,346	0.39	10,966,181	3,486,647	2,275,127	196,745	0.18
2019-12-13-HFL_palobo_bin4_rep4	2x150	STAR-2.5.3a	hg38	47,394,346	0.38	37,574,160	28,839,902	7,042,070	2,013,972	0.80
2019-12-13-HFL_palobo_pre-sort_rep3	2x150	STAR-2.5.3a	dm6	47,723,063	0.33	9,512,873	3,159,497	2,092,631	184,159	0.16
2019-12-13-HFL_palobo_pre-sort_rep3	2x150	STAR-2.5.3a	hg38	47,723,063	0.32	36,035,761	29,815,325	9,921,114	2,096,540	0.82
2019-12-13-HFL_palobo_pre-sort_rep4	2x150	STAR-2.5.3a	dm6	50,329,677	0.37	4,188,190	1,525,002	984,854	93,814	0.07
2019-12-13-HFL_palobo_pre-sort_rep4	2x150	STAR-2.5.3a	hg38	50,329,677	0.37	42,726,414	33,215,430	12,847,054	2,474,276	0.91
2019-12-13-HFL_palobo_sort-all_rep3	2x150	STAR-2.5.3a	dm6	48,823,436	0.36	9,854,674	3,050,136	2,061,742	171,320	0.16
2019-12-13-HFL_palobo_sort-all_rep3	2x150	STAR-2.5.3a	hg38	48,823,436	0.35	39,788,232	28,713,542	9,231,787	2,037,287	0.82
2019-12-13-HFL_palobo_sort-all_rep4	2x150	STAR-2.5.3a	dm6	51,158,724	0.35	10,455,707	3,393,799	2,107,526	186,770	0.16
2019-12-13-HFL_palobo_sort-all_rep4	2x150	STAR-2.5.3a	hg38	51,158,724	0.34	41,762,841	32,978,039	6,595,703	2,317,323	0.82
2019-12-13-HMEC_G1_bin1_rep1	2x150	STAR-2.5.3a	dm6	51,007,964	0.95	2,371	697	551	21	0.00
2019-12-13-HMEC_G1_bin1_rep1	2x150	STAR-2.5.3a	hg38	51,007,964	0.30	52,120,908	40,583,838	3,177,903	2,815,599	0.97
2019-12-13-HMEC_G1_bin1_rep2	2x150	STAR-2.5.3a	dm6	50,920,184	0.96	2,460	762	607	35	0.00
2019-12-13-HMEC_G1_bin1_rep2	2x150	STAR-2.5.3a	hg38	50,920,184	0.27	50,171,849	39,931,397	5,168,793	2,760,979	0.96
2019-12-13-HMEC_G1_bin4_rep1	2x150	STAR-2.5.3a	dm6	61,923,234	0.97	1,390	454	305	15	0.00
2019-12-13-HMEC_G1_bin4_rep1	2x150	STAR-2.5.3a	hg38	61,923,234	0.32	61,711,853	48,090,367	6,656,177	3,515,145	0.97
2019-12-13-HMEC_G1_bin4_rep2	2x150	STAR-2.5.3a	dm6	48,812,922	0.97	1,786	560	359	39	0.00
2019-12-13-HMEC_G1_bin4_rep2	2x150	STAR-2.5.3a	hg38	48,812,922	0.33	47,626,407	39,721,243	4,017,688	2,907,294	0.97

READ DISTRIBUTION

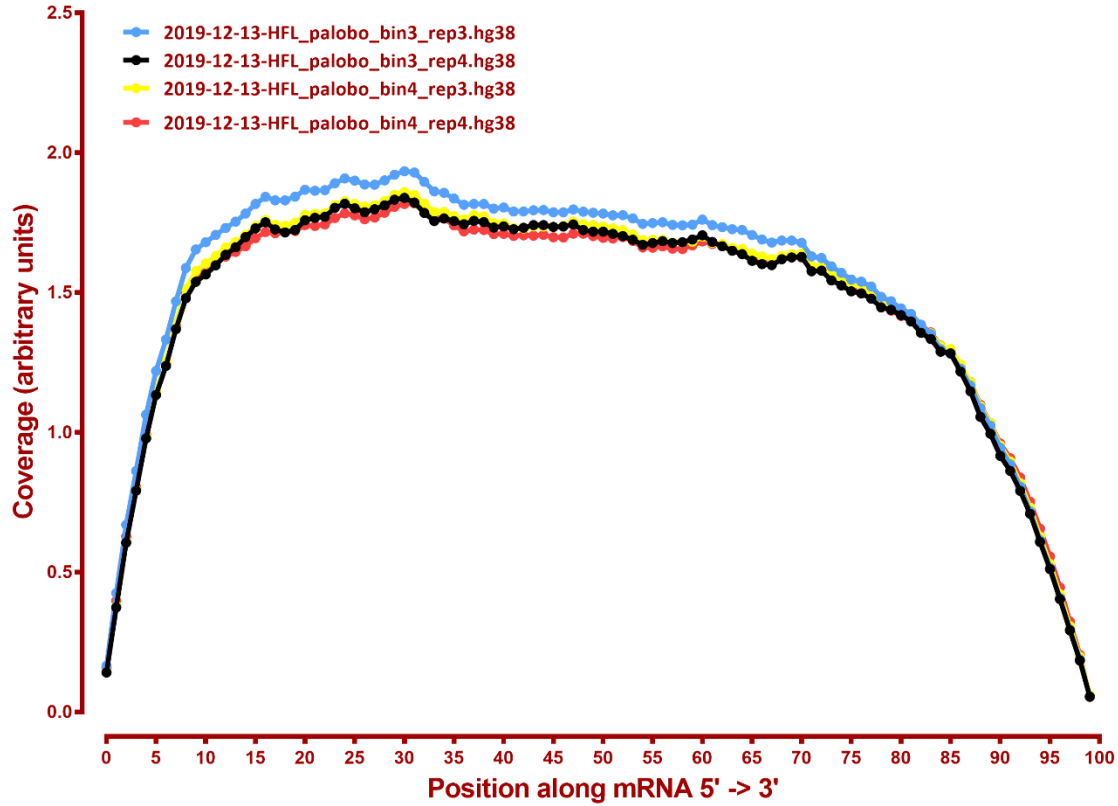
#	Exonic:	Intergenic:	Intronic:
2019-12-13-HFL_palobo_bin1_rep3.hg38	0.91	0.01	0.08
2019-12-13-HFL_palobo_bin1_rep4.hg38	0.91	0.01	0.08
2019-12-13-HFL_palobo_bin2_rep3.hg38	0.92	0.01	0.07
2019-12-13-HFL_palobo_bin2_rep4.hg38	0.92	0.01	0.07
2019-12-13-HFL_palobo_bin3_rep3.hg38	0.93	0.01	0.06
2019-12-13-HFL_palobo_bin3_rep4.hg38	0.92	0.01	0.07
2019-12-13-HFL_palobo_bin4_rep3.hg38	0.92	0.01	0.07
2019-12-13-HFL_palobo_bin4_rep4.hg38	0.92	0.01	0.07
2019-12-13-HFL_palobo_pre-sort_rep3.hg38	0.95	0.01	0.04
2019-12-13-HFL_palobo_pre-sort_rep4.hg38	0.92	0.01	0.07
2019-12-13-HFL_palobo_sort-all_rep3.hg38	0.91	0.01	0.08
2019-12-13-HFL_palobo_sort-all_rep4.hg38	0.92	0.01	0.06
2019-12-13-HMEC_G1_bin1_rep1.hg38	0.92	0.01	0.07
2019-12-13-HMEC_G1_bin1_rep2.hg38	0.92	0.01	0.07
2019-12-13-HMEC_G1_bin4_rep1.hg38	0.92	0.01	0.06
2019-12-13-HMEC_G1_bin4_rep2.hg38	0.94	0.01	0.06

COVERAGE QC

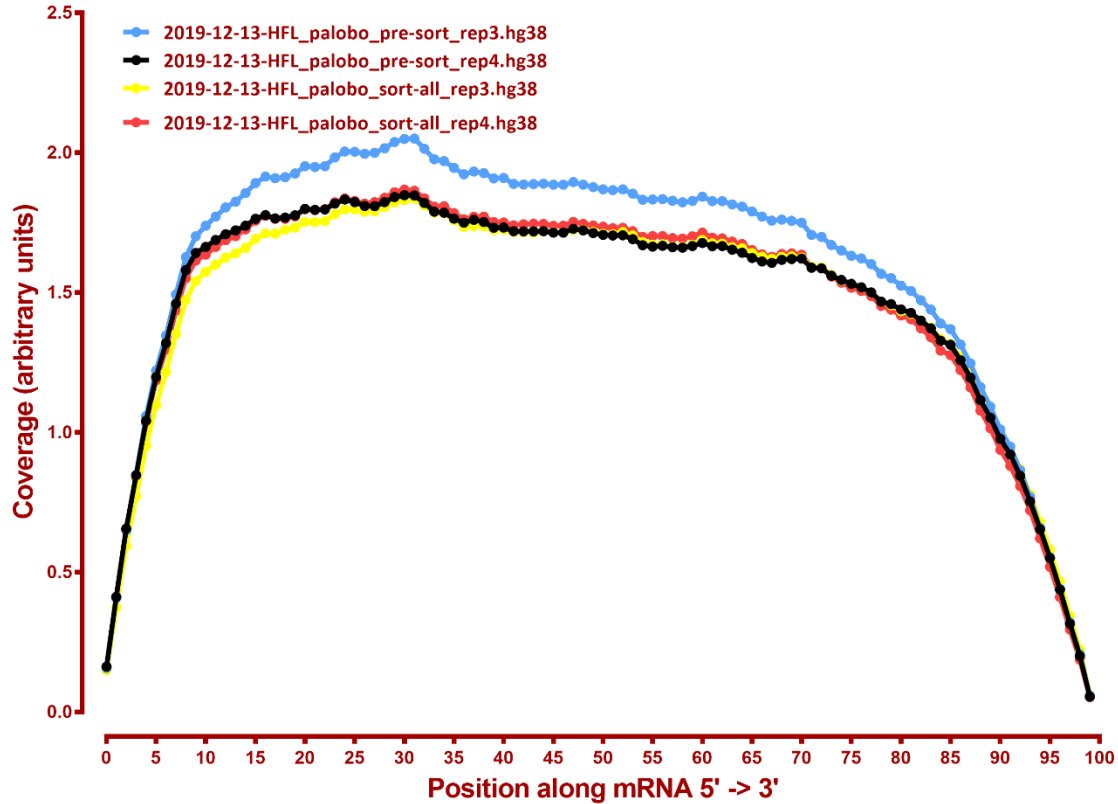
Coverage of genes; >1000bp, sacCer3



Coverage of genes; >1000bp, sacCer3



Coverage of genes; >1000bp, sacCer3



Coverage of genes; >1000bp, sacCer3

