

MAPPING STATS

Library	Read Length	Mapping	Species	Raw fragments	Complexity	Unique	Unique Splices	Multi	Multi Splices	Fraction mapped
2023-03-30-220328R03_exp211025_MS64_SCGE_F3_total-RNA-seq	2x36	STAR-2.5.3a	Candida	1,461,905	0.34	468,402	248	1,057,596	4,276	0.52
2023-03-30-220328R03_exp211025_MS64_SCGE_F3_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	1,461,905	0.86	118,525	1,773	2,065,088	3,508	0.75
2023-03-30-MS230306L01_exp211025_MS64_SCGE_F1_total-RNA-seq	2x36	STAR-2.5.3a	Candida	1,192,118	0.34	452,839	303	868,164	1,960	0.56
2023-03-30-MS230306L01_exp211025_MS64_SCGE_F1_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	1,192,118	0.87	97,659	1,553	1,610,476	1,628	0.72
2023-03-30-MS230306L02_exp211025_MS64_SCGE_F2_total-RNA-seq	2x36	STAR-2.5.3a	Candida	1,199,348	0.35	409,939	285	865,640	5,218	0.53
2023-03-30-MS230306L02_exp211025_MS64_SCGE_F2_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	1,199,348	0.87	100,401	1,401	1,655,488	3,596	0.73
2023-03-30-MS230306L04_exp211025_MS64_SCGE_F4_total-RNA-seq	2x36	STAR-2.5.3a	Candida	1,298,956	0.36	395,062	194	939,903	717	0.51
2023-03-30-MS230306L04_exp211025_MS64_SCGE_F4_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	1,298,956	0.86	104,483	1,669	1,819,933	14,243	0.75
2023-03-30-MS230306L05_exp211025_MS64_SCGE_F5_total-RNA-seq	2x36	STAR-2.5.3a	Candida	1,355,447	0.35	404,947	263	988,477	937	0.51
2023-03-30-MS230306L05_exp211025_MS64_SCGE_F5_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	1,355,447	0.86	105,780	1,702	1,948,675	2,485	0.76
2023-03-30-MS230306L06_exp211025_MS64_SCGE_F6_total-RNA-seq	2x36	STAR-2.5.3a	Candida	1,254,824	0.36	375,029	229	918,321	2,335	0.52
2023-03-30-MS230306L06_exp211025_MS64_SCGE_F6_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	1,254,824	0.86	91,070	1,420	1,820,473	3,111	0.76
2023-03-30-MS230306L07_exp211025_MS64_SCGE_F7_total-RNA-seq	2x36	STAR-2.5.3a	Candida	1,378,364	0.36	404,469	215	999,002	836	0.51
2023-03-30-MS230306L07_exp211025_MS64_SCGE_F7_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	1,378,364	0.84	97,742	1,776	1,967,867	6,029	0.75
2023-03-30-MS230306L08_exp211021_MS67_SCGE_F1_total-RNA-seq	2x36	STAR-2.5.3a	Candida	1,305,628	0.34	479,611	257	947,279	107	0.55
2023-03-30-MS230306L08_exp211021_MS67_SCGE_F1_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	1,305,628	0.84	122,313	1,657	1,772,900	2,472	0.73
2023-03-30-MS230306L09_exp211021_MS67_SCGE_F2_total-RNA-seq	2x36	STAR-2.5.3a	Candida	1,334,317	0.35	442,720	310	946,621	1,799	0.52
2023-03-30-MS230306L09_exp211021_MS67_SCGE_F2_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	1,334,317	0.86	116,534	1,896	1,798,001	6,467	0.72
2023-03-30-MS230306L10_exp211021_MS67_SCGE_F3_total-RNA-seq	2x36	STAR-2.5.3a	Candida	903,396	0.40	268,242	208	670,313	33	0.52
2023-03-30-MS230306L10_exp211021_MS67_SCGE_F3_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	903,396	0.87	87,896	1,158	1,279,096	3,294	0.76
2023-03-30-MS230306L11_exp211021_MS67_SCGE_F4_total-RNA-seq	2x36	STAR-2.5.3a	Candida	1,682,684	0.35	435,357	493	1,143,950	7,448	0.47
2023-03-30-MS230306L11_exp211021_MS67_SCGE_F4_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	1,682,684	0.84	139,444	2,008	2,276,939	6,151	0.72
2023-03-30-MS230306L12_exp211021_MS67_SCGE_F5_total-RNA-seq	2x36	STAR-2.5.3a	Candida	1,434,726	0.36	389,185	233	1,051,824	1,998	0.50
2023-03-30-MS230306L12_exp211021_MS67_SCGE_F5_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	1,434,726	0.85	119,152	1,608	2,100,845	1,417	0.77
2023-03-30-MS230306L13_exp211021_MS67_SCGE_F6_total-RNA-seq	2x36	STAR-2.5.3a	Candida	1,622,130	0.35	453,226	344	1,183,700	482	0.50
2023-03-30-MS230306L13_exp211021_MS67_SCGE_F6_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	1,622,130	0.84	124,068	1,940	2,353,399	4,905	0.77
2023-03-30-MS230306L14_exp211021_MS67_SCGE_F7_total-RNA-seq	2x36	STAR-2.5.3a	Candida	1,446,513	0.36	369,437	229	1,059,265	365	0.49
2023-03-30-MS230306L14_exp211021_MS67_SCGE_F7_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	1,446,513	0.84	113,536	1,708	2,131,131	1,507	0.78
2023-03-30-MS230306L15_exp211028_MS64_SCGE_F1_total-RNA-seq	2x36	STAR-2.5.3a	Candida	1,703,887	0.31	692,864	264	1,193,784	2,870	0.55
2023-03-30-MS230306L15_exp211028_MS64_SCGE_F1_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	1,703,887	0.87	117,702	1,954	2,259,461	2,887	0.70
2023-03-30-MS230306L16_exp211028_MS64_SCGE_F2_total-RNA-seq	2x36	STAR-2.5.3a	Candida	1,640,636	0.32	617,338	248	1,159,025	2,967	0.54
2023-03-30-MS230306L16_exp211028_MS64_SCGE_F2_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	1,640,636	0.87	115,002	1,830	2,244,418	4,660	0.72
2023-03-30-MS230306L17_exp211028_MS64_SCGE_F3_total-RNA-seq	2x36	STAR-2.5.3a	Candida	1,602,554	0.33	534,299	301	1,135,218	2,692	0.52
2023-03-30-MS230306L17_exp211028_MS64_SCGE_F3_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	1,602,554	0.87	110,570	1,916	2,262,502	1,890	0.74

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2023-03-30-MS230306L18_exp211028_MS64_SCGE_F4_total-RNA-seq	2x36	STAR-2.5.3a	Candida	1,567,448	0.33	528,354	264	1,092,688	1,472	0.52
2023-03-30-MS230306L18_exp211028_MS64_SCGE_F4_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	1,567,448	0.87	104,372	1,708	2,189,458	4,314	0.73
2023-03-30-MS230306L19_exp211028_MS64_SCGE_F5_total-RNA-seq	2x36	STAR-2.5.3a	Candida	1,626,161	0.34	476,281	219	1,153,391	1,453	0.50
2023-03-30-MS230306L19_exp211028_MS64_SCGE_F5_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	1,626,161	0.87	107,195	1,935	2,362,940	2,310	0.76
2023-03-30-MS230306L20_exp211028_MS64_SCGE_F6_total-RNA-seq	2x36	STAR-2.5.3a	Candida	1,912,546	0.32	563,592	210	1,355,991	2,597	0.50
2023-03-30-MS230306L20_exp211028_MS64_SCGE_F6_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	1,912,546	0.86	114,020	2,114	2,767,591	4,369	0.76
2023-03-30-MS230306L21_exp211028_MS64_SCGE_F7_total-RNA-seq	2x36	STAR-2.5.3a	Candida	1,585,556	0.34	480,343	201	1,127,566	3,646	0.51
2023-03-30-MS230306L21_exp211028_MS64_SCGE_F7_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	1,585,556	0.87	91,313	1,741	2,303,325	7,623	0.76
2023-03-30-MS230306L22_exp211101_MS67_SCGE_F1_total-RNA-seq	2x36	STAR-2.5.3a	Candida	1,608,890	0.33	537,962	284	1,145,345	8,561	0.53
2023-03-30-MS230306L22_exp211101_MS67_SCGE_F1_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	1,608,890	0.86	126,264	2,034	2,282,534	3,522	0.75
2023-03-30-MS230306L23_exp211101_MS67_SCGE_F2_total-RNA-seq	2x36	STAR-2.5.3a	Candida	1,792,977	0.33	547,535	343	1,280,828	2,118	0.51
2023-03-30-MS230306L23_exp211101_MS67_SCGE_F2_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	1,792,977	0.86	132,117	2,157	2,575,765	9,547	0.76
2023-03-30-MS230306L24_exp211101_MS67_SCGE_F3_total-RNA-seq	2x36	STAR-2.5.3a	Candida	1,669,158	0.34	474,193	203	1,175,902	3,214	0.50
2023-03-30-MS230306L24_exp211101_MS67_SCGE_F3_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	1,669,158	0.86	119,021	1,919	2,367,925	4,081	0.75
2023-03-30-MS230306L25_exp211101_MS67_SCGE_F4_total-RNA-seq	2x36	STAR-2.5.3a	Candida	1,784,264	0.33	513,283	191	1,277,203	403	0.50
2023-03-30-MS230306L25_exp211101_MS67_SCGE_F4_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	1,784,264	0.86	128,790	2,050	2,595,718	4,164	0.77
2023-03-30-MS230306L26_exp211101_MS67_SCGE_F5_total-RNA-seq	2x36	STAR-2.5.3a	Candida	1,887,644	0.33	525,534	262	1,353,098	2,268	0.50
2023-03-30-MS230306L26_exp211101_MS67_SCGE_F5_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	1,887,644	0.85	130,030	2,160	2,747,475	9,967	0.77
2023-03-30-MS230306L27_exp211101_MS67_SCGE_F6_total-RNA-seq	2x36	STAR-2.5.3a	Candida	1,835,475	0.34	475,915	171	1,307,251	1,319	0.49
2023-03-30-MS230306L27_exp211101_MS67_SCGE_F6_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	1,835,475	0.85	120,131	2,145	2,694,388	6,014	0.77
2023-03-30-MS230306L28_exp211101_MS67_SCGE_F7_total-RNA-seq	2x36	STAR-2.5.3a	Candida	1,793,374	0.33	510,222	252	1,304,251	1,941	0.51
2023-03-30-MS230306L28_exp211101_MS67_SCGE_F7_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	1,793,374	0.85	110,586	1,932	2,657,958	5,058	0.77
2023-03-30-MS230316L01_exp220519_MS01_SCGE_total-RNA-seq	2x36	STAR-2.5.3a	Candida	1,818,868	0.35	444,232	212	1,339,491	2,711	0.49
2023-03-30-MS230316L01_exp220519_MS01_SCGE_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	1,818,868	0.86	167,931	2,517	2,699,766	3,618	0.79
2023-03-30-MS230316L02_exp220519_MS80_SCGE_total-RNA-seq	2x36	STAR-2.5.3a	Candida	1,710,597	0.35	444,189	353	1,260,615	2,125	0.50
2023-03-30-MS230316L02_exp220519_MS80_SCGE_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	1,710,597	0.87	158,171	2,269	2,509,271	3,055	0.78
2023-03-30-MS230316L03_exp220519_MS81_SCGE_total-RNA-seq	2x36	STAR-2.5.3a	Candida	1,901,390	0.37	402,420	452	1,358,298	8,272	0.47
2023-03-30-MS230316L03_exp220519_MS81_SCGE_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	1,901,390	0.86	169,035	2,517	2,775,939	6,801	0.78
2023-03-30-MS230316L04_exp220519_MS63_SCGE-00nMBE_total-RNA-seq	2x36	STAR-2.5.3a	Candida	2,010,358	0.34	496,505	305	1,478,490	2,616	0.49
2023-03-30-MS230316L04_exp220519_MS63_SCGE-00nMBE_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	2,010,358	0.86	185,088	2,934	2,959,220	6,378	0.78
2023-03-30-MS230316L05_exp220519_MS63_SCGE-04nMBE_total-RNA-seq	2x36	STAR-2.5.3a	Candida	1,433,128	0.40	295,506	188	1,015,423	6,113	0.46
2023-03-30-MS230316L05_exp220519_MS63_SCGE-04nMBE_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	1,433,128	0.87	124,948	1,814	2,078,196	9,328	0.77
2023-03-30-MS230316L06_exp220519_MS63_SCGE-15nMBE_total-RNA-seq	2x36	STAR-2.5.3a	Candida	1,986,754	0.36	400,290	496	1,476,154	3,358	0.47
2023-03-30-MS230316L06_exp220519_MS63_SCGE-15nMBE_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	1,986,754	0.85	167,445	2,845	3,069,494	4,344	0.82

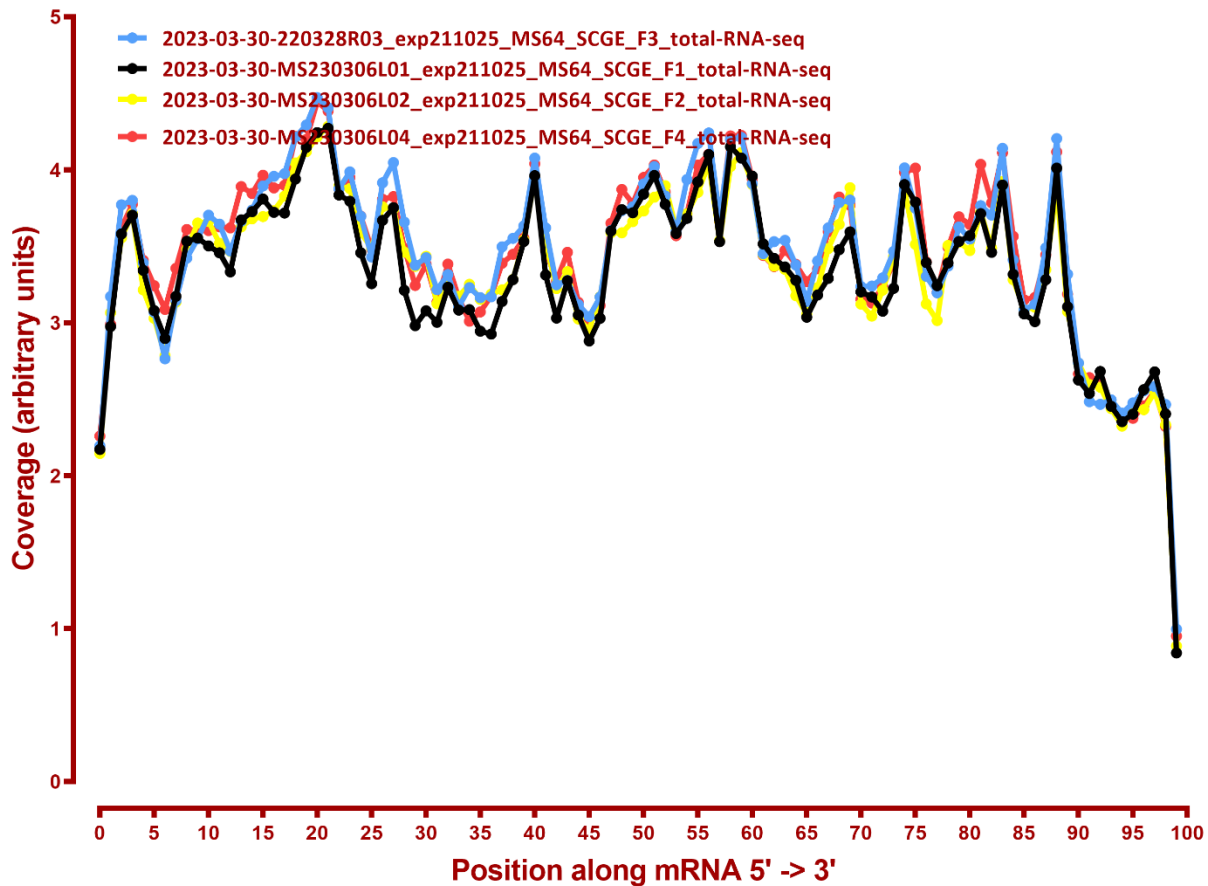
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2023-03-30-MS230316L07_exp220601_MS01_SCGE_total-RNA-seq	2x36	STAR-2.5.3a	Candida	2,063,490	0.32	699,313	443	1,526,748	9,010	0.54
2023-03-30-MS230316L07_exp220601_MS01_SCGE_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	2,063,490	0.86	165,014	2,724	2,892,900	9,742	0.74
2023-03-30-MS230316L08_exp220601_MS80_SCGE_total-RNA-seq	2x36	STAR-2.5.3a	Candida	1,878,812	0.32	672,139	443	1,384,368	7,606	0.55
2023-03-30-MS230316L08_exp220601_MS80_SCGE_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	1,878,812	0.87	148,373	2,229	2,578,764	7,072	0.73
2023-03-30-MS230316L09_exp220601_MS81_SCGE_total-RNA-seq	2x36	STAR-2.5.3a	Candida	1,966,144	0.33	659,613	447	1,458,228	1,676	0.54
2023-03-30-MS230316L09_exp220601_MS81_SCGE_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	1,966,144	0.87	147,617	2,409	2,742,306	4,590	0.74
2023-03-30-MS230316L10_exp220601_MS63_SCGE-00nMBE_total-RNA-seq	2x36	STAR-2.5.3a	Candida	2,065,584	0.32	716,260	402	1,524,488	7,464	0.54
2023-03-30-MS230316L10_exp220601_MS63_SCGE-00nMBE_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	2,065,584	0.87	164,014	2,658	2,881,239	8,165	0.74
2023-03-30-MS230316L11_exp220601_MS63_SCGE-04nMBE_total-RNA-seq	2x36	STAR-2.5.3a	Candida	2,122,419	0.33	634,338	710	1,561,349	13,005	0.52
2023-03-30-MS230316L11_exp220601_MS63_SCGE-04nMBE_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	2,122,419	0.86	162,687	2,781	3,032,565	16,839	0.76
2023-03-30-MS230316L12_exp220601_MS63_SCGE-15nMBE_total-RNA-seq	2x36	STAR-2.5.3a	Candida	2,039,843	0.34	520,585	459	1,507,381	5,529	0.50
2023-03-30-MS230316L12_exp220601_MS63_SCGE-15nMBE_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	2,039,843	0.86	154,703	2,655	3,021,152	5,900	0.78
2023-03-30-MS230316L13_exp220610_MS01_SCGE_total-RNA-seq	2x36	STAR-2.5.3a	Candida	2,137,700	0.33	638,244	452	1,584,147	9,189	0.52
2023-03-30-MS230316L13_exp220610_MS01_SCGE_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	2,137,700	0.86	177,932	2,758	3,078,831	9,395	0.76
2023-03-30-MS230316L14_exp220610_MS80_SCGE_total-RNA-seq	2x36	STAR-2.5.3a	Candida	2,073,602	0.33	621,602	316	1,523,044	2,306	0.52
2023-03-30-MS230316L14_exp220610_MS80_SCGE_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	2,073,602	0.86	183,329	2,607	2,950,223	5,279	0.76
2023-03-30-MS230316L15_exp220610_MS81_SCGE_total-RNA-seq	2x36	STAR-2.5.3a	Candida	2,088,139	0.35	495,294	486	1,507,548	2,142	0.48
2023-03-30-MS230316L15_exp220610_MS81_SCGE_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	2,088,139	0.86	177,804	2,730	3,052,638	6,768	0.78
2023-03-30-MS230316L16_exp220610_MS63_SCGE-00nMBE_total-RNA-seq	2x36	STAR-2.5.3a	Candida	2,779,529	0.30	853,845	753	2,051,608	4,032	0.52
2023-03-30-MS230316L16_exp220610_MS63_SCGE-00nMBE_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	2,779,529	0.85	235,638	4,098	3,976,146	7,748	0.76
2023-03-30-MS230316L17_exp220610_MS63_SCGE-3.5nMBE_total-RNA-seq	2x36	STAR-2.5.3a	Candida	1,818,509	0.35	485,239	287	1,335,241	6,311	0.50
2023-03-30-MS230316L17_exp220610_MS63_SCGE-3.5nMBE_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	1,818,509	0.86	152,482	2,280	2,661,940	6,904	0.78
2023-03-30-MS230316L18_exp220610_MS63_SCGE-16nMBE_total-RNA-seq	2x36	STAR-2.5.3a	Candida	1,766,545	0.36	411,517	335	1,299,471	1,887	0.48
2023-03-30-MS230316L18_exp220610_MS63_SCGE-16nMBE_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	1,766,545	0.87	141,741	2,461	2,662,101	6,395	0.80
2023-03-30-MS230316L19_exp220607_MS01_SCGE_total-RNA-seq	2x36	STAR-2.5.3a	Candida	1,997,027	0.34	569,188	506	1,466,479	5,757	0.51
2023-03-30-MS230316L19_exp220607_MS01_SCGE_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	1,997,027	0.87	167,510	2,534	2,849,109	10,081	0.76
2023-03-30-MS230316L20_exp220607_MS80_SCGE_total-RNA-seq	2x36	STAR-2.5.3a	Candida	2,015,369	0.34	568,609	449	1,435,806	3,542	0.50
2023-03-30-MS230316L20_exp220607_MS80_SCGE_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	2,015,369	0.86	178,483	2,711	2,796,865	5,355	0.74
2023-03-30-MS230316L21_exp220607_MS81_SCGE_total-RNA-seq	2x36	STAR-2.5.3a	Candida	1,768,406	0.36	453,967	429	1,322,811	1,667	0.50
2023-03-30-MS230316L21_exp220607_MS81_SCGE_total-RNA-seq	2x36	STAR-2.5.3a	sacCer3	1,768,406	0.85	164,178	2,484	2,643,888	5,986	0.80
2023-03-30-MS230316L22_exp220607_MS63_SCGE-00nMBE_total-RNA-seq	2x36	STAR-2.5.3a	Candida	1,782,044	0.35	509,306	288	1,316,136	2,572	0.51
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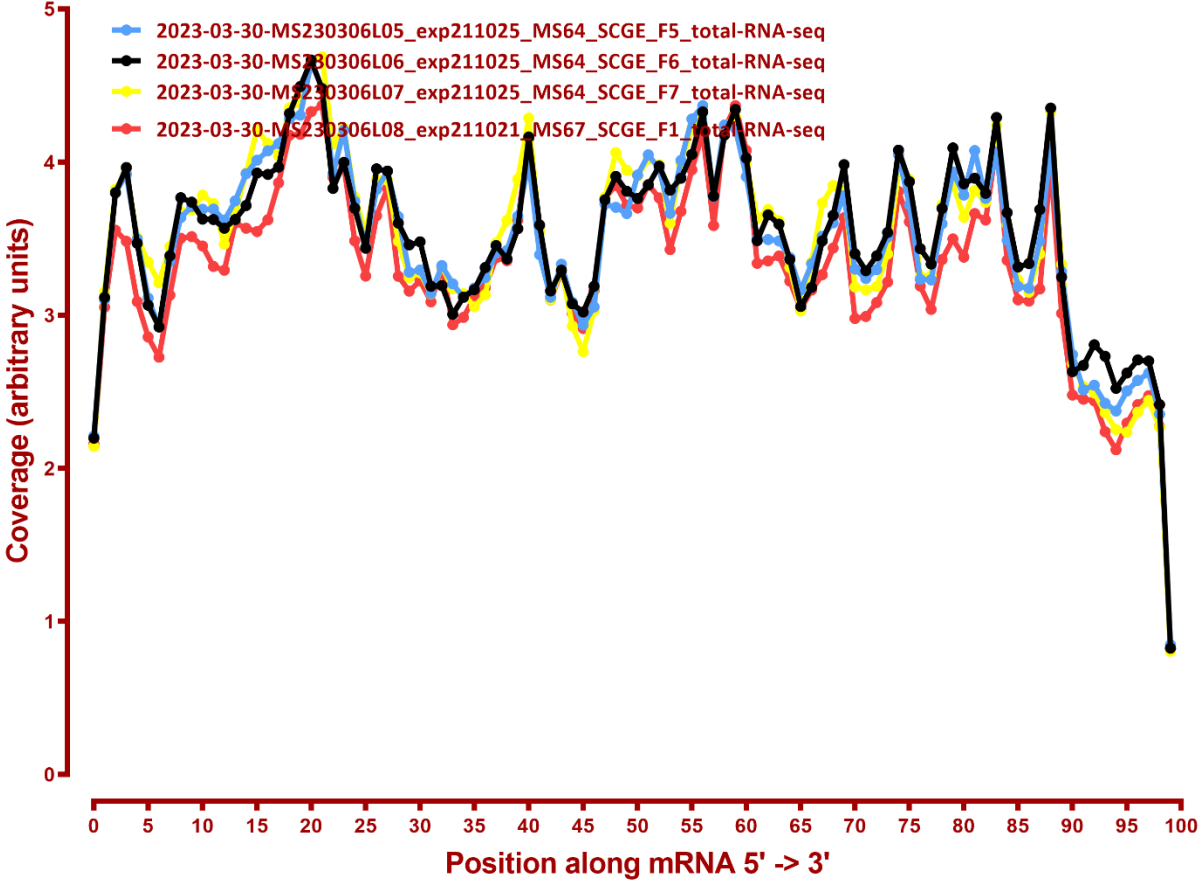
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2023-03-30-MS230316L23_exp220607_MS63_SCGE-3.5nMBE_total-RNA-seq	0.94	0.05	0.00
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COVERAGE

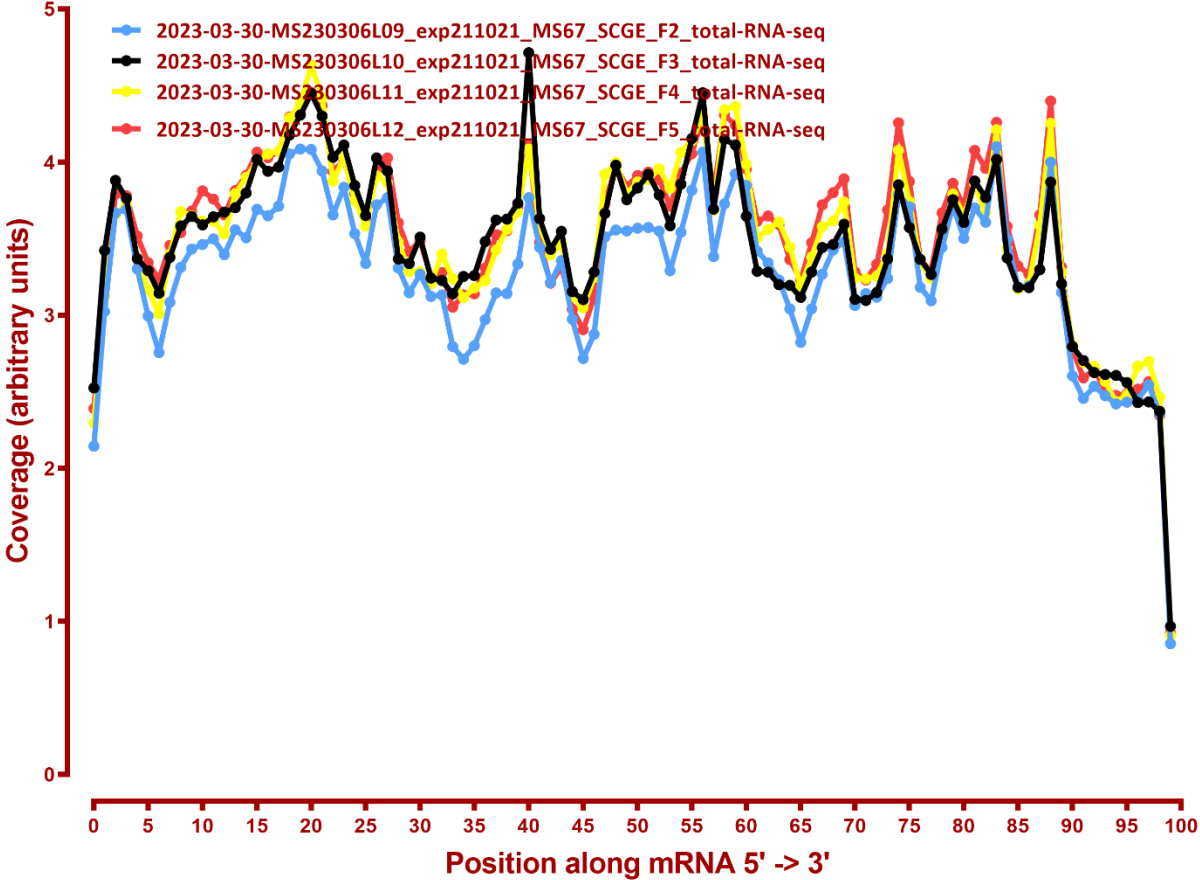
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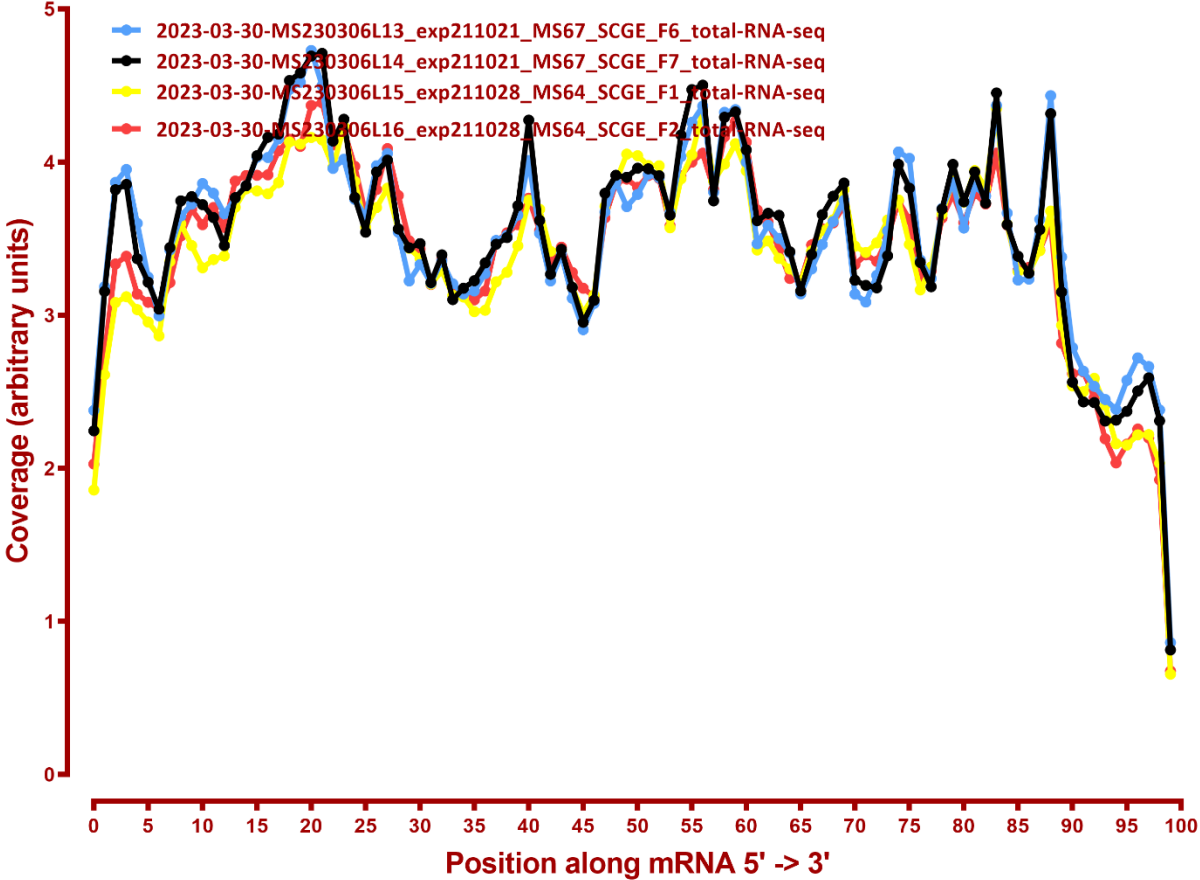
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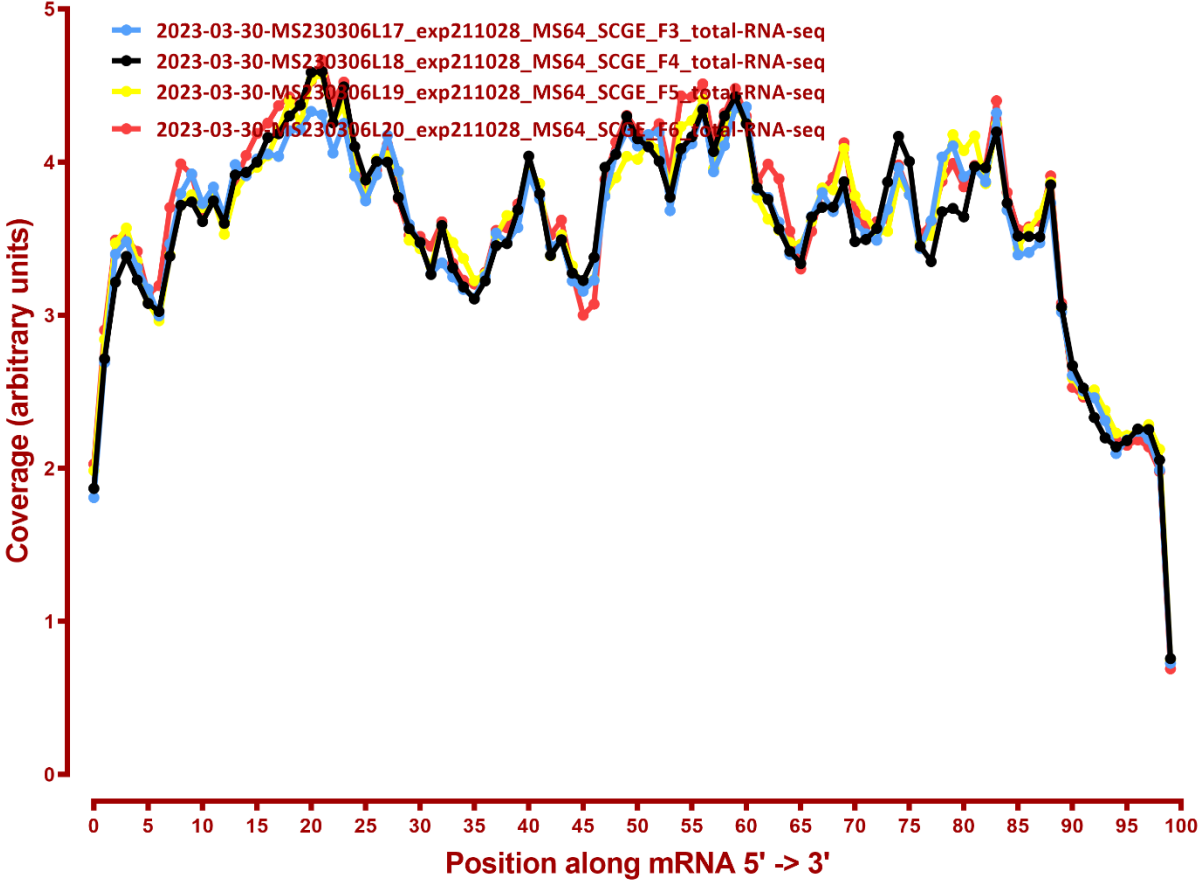
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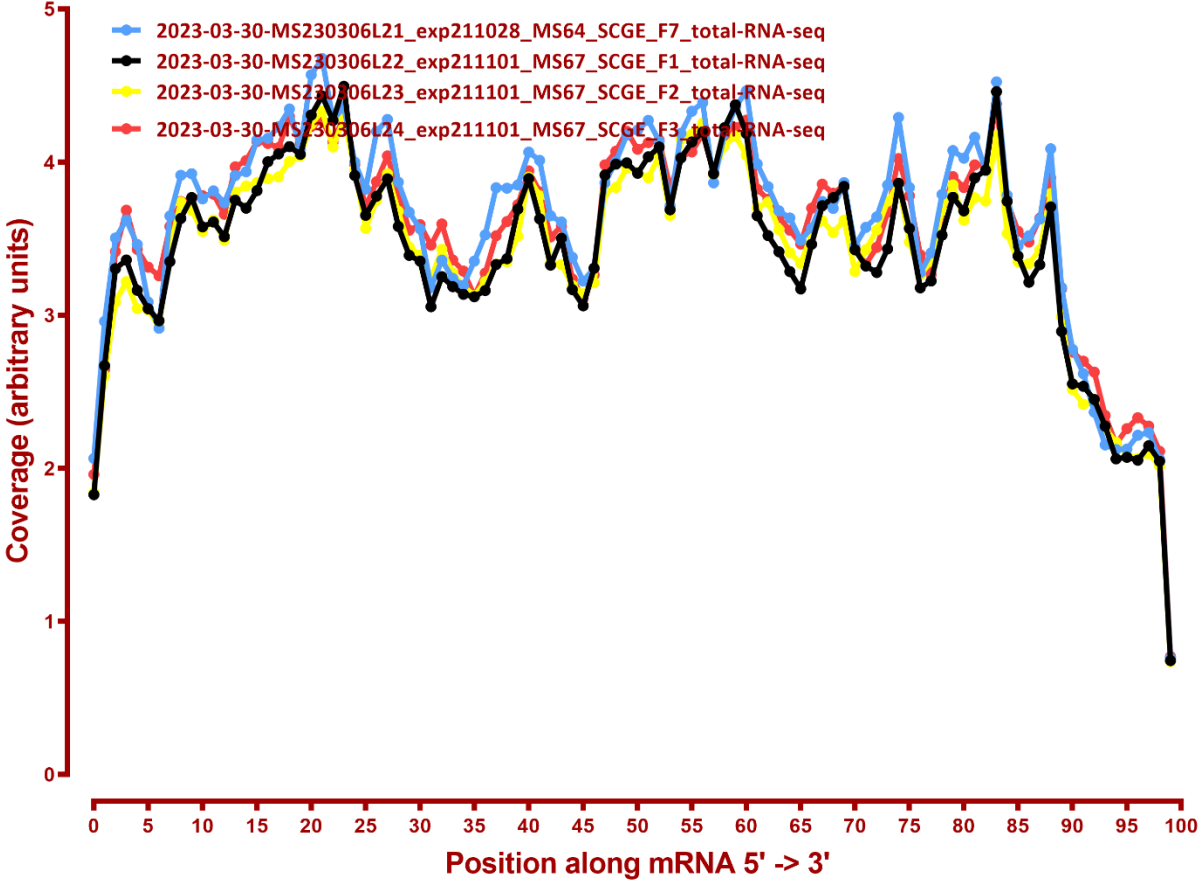
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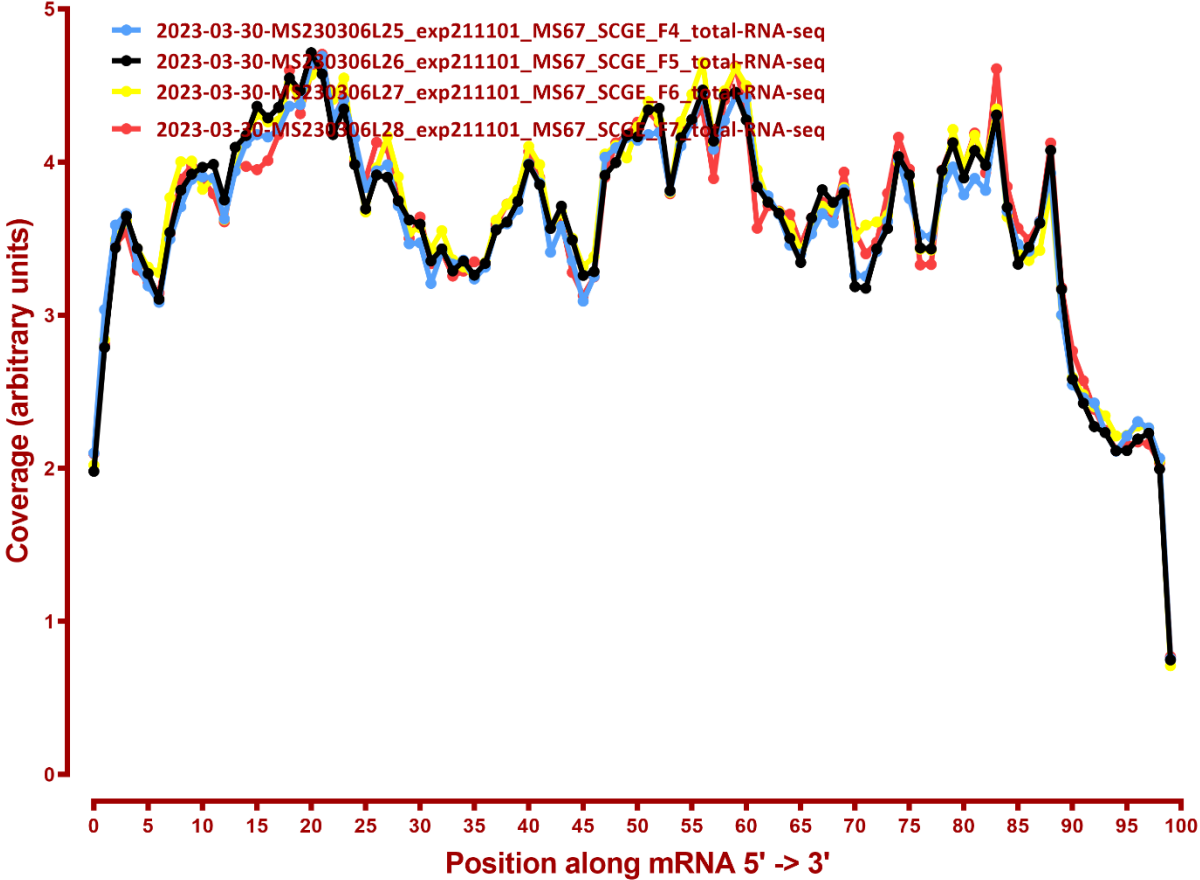
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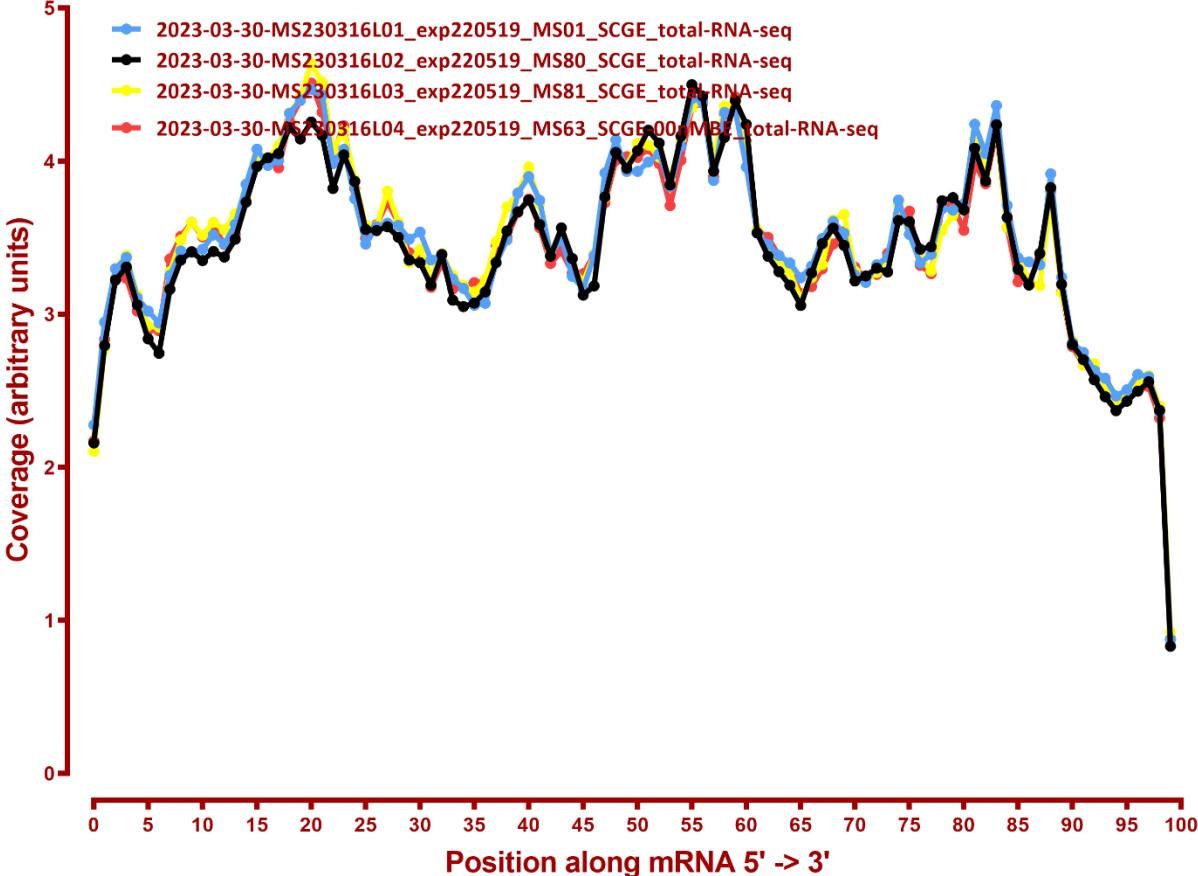
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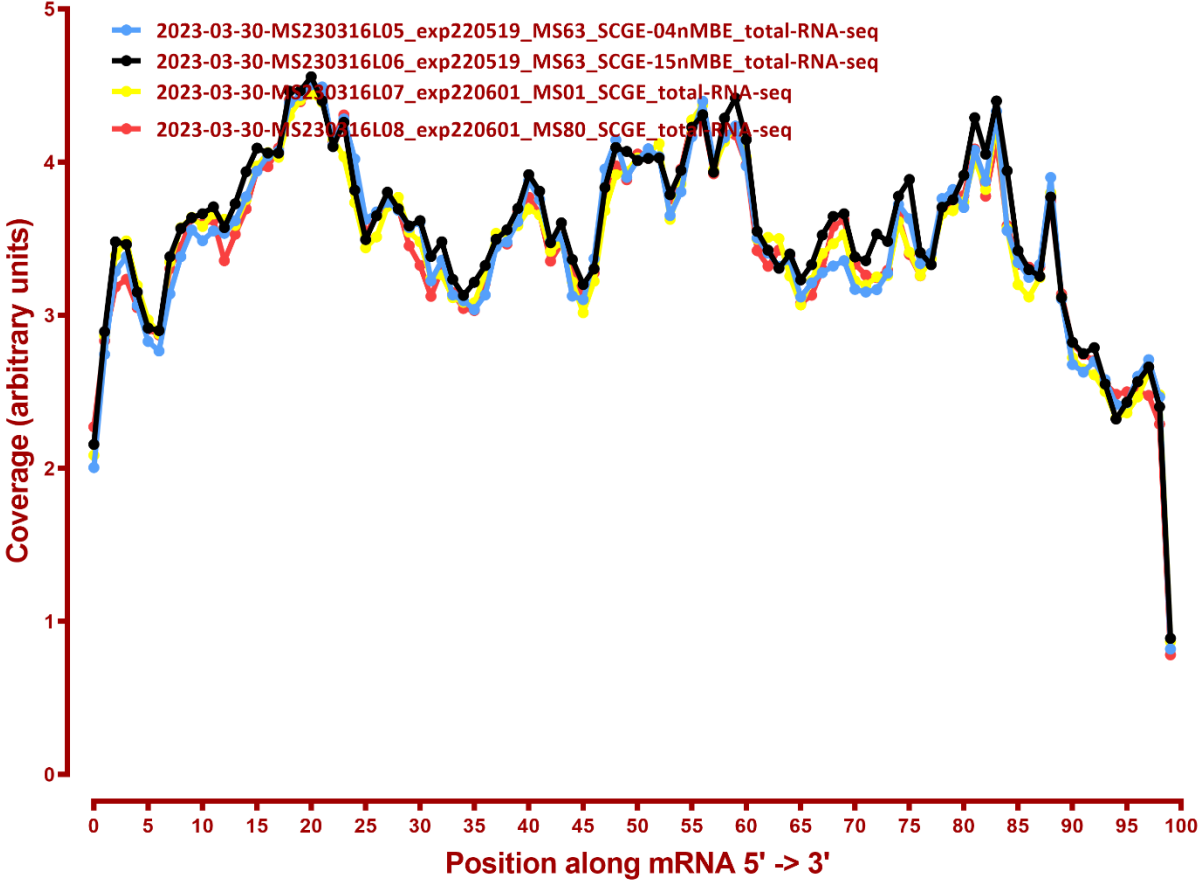
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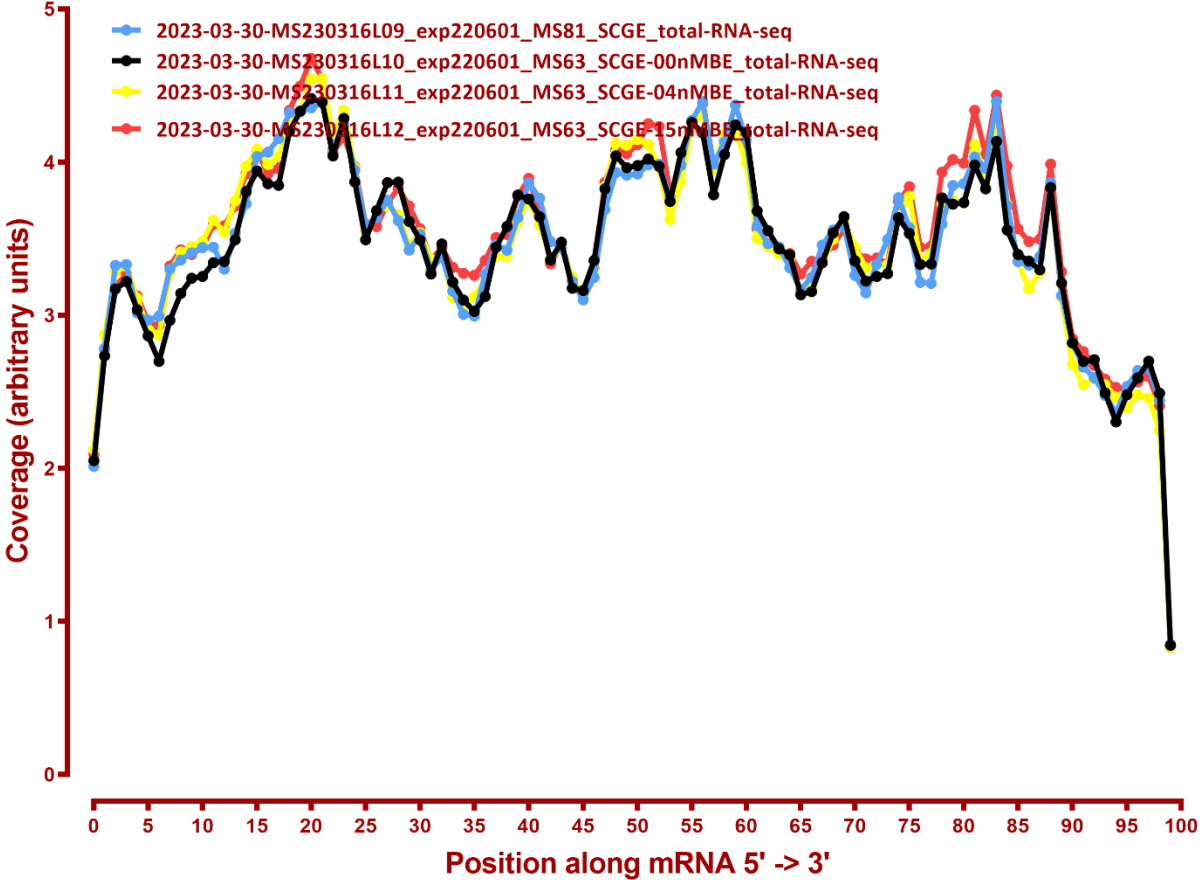
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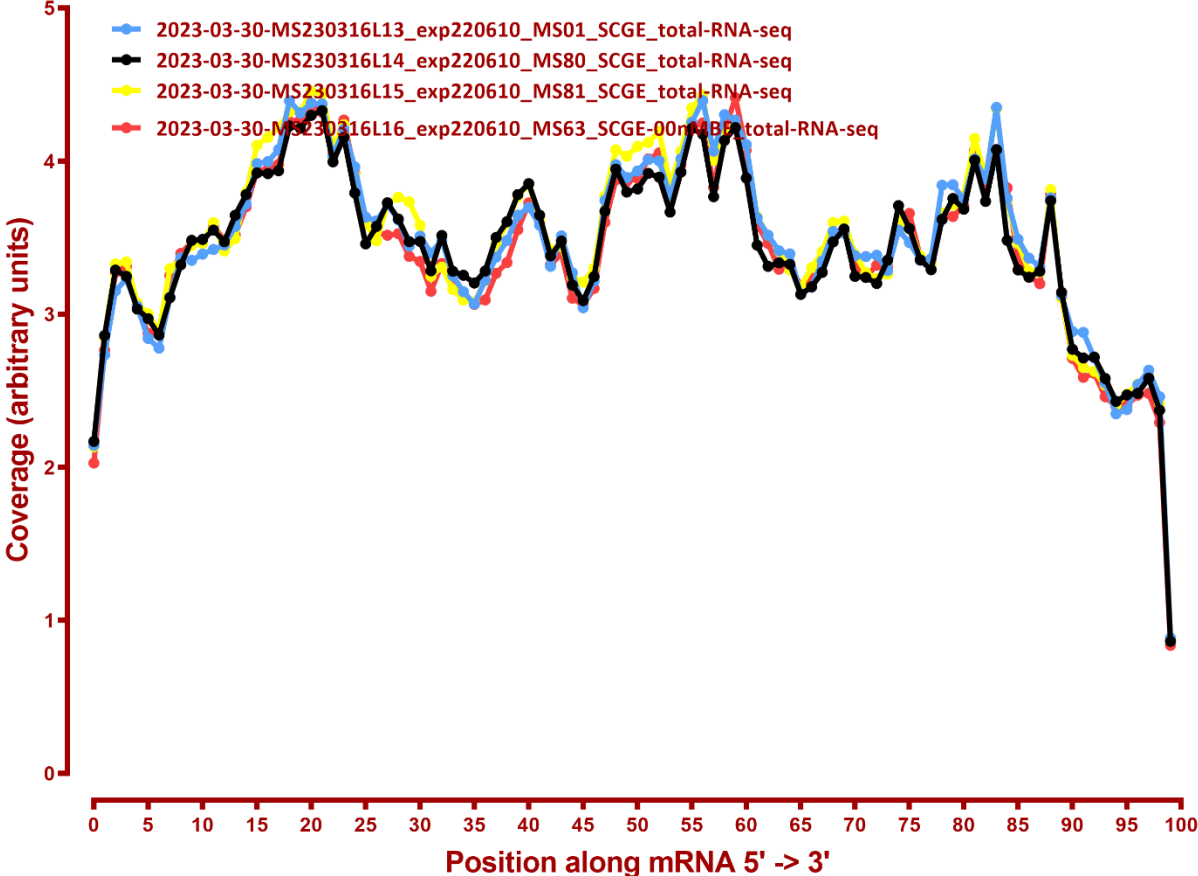
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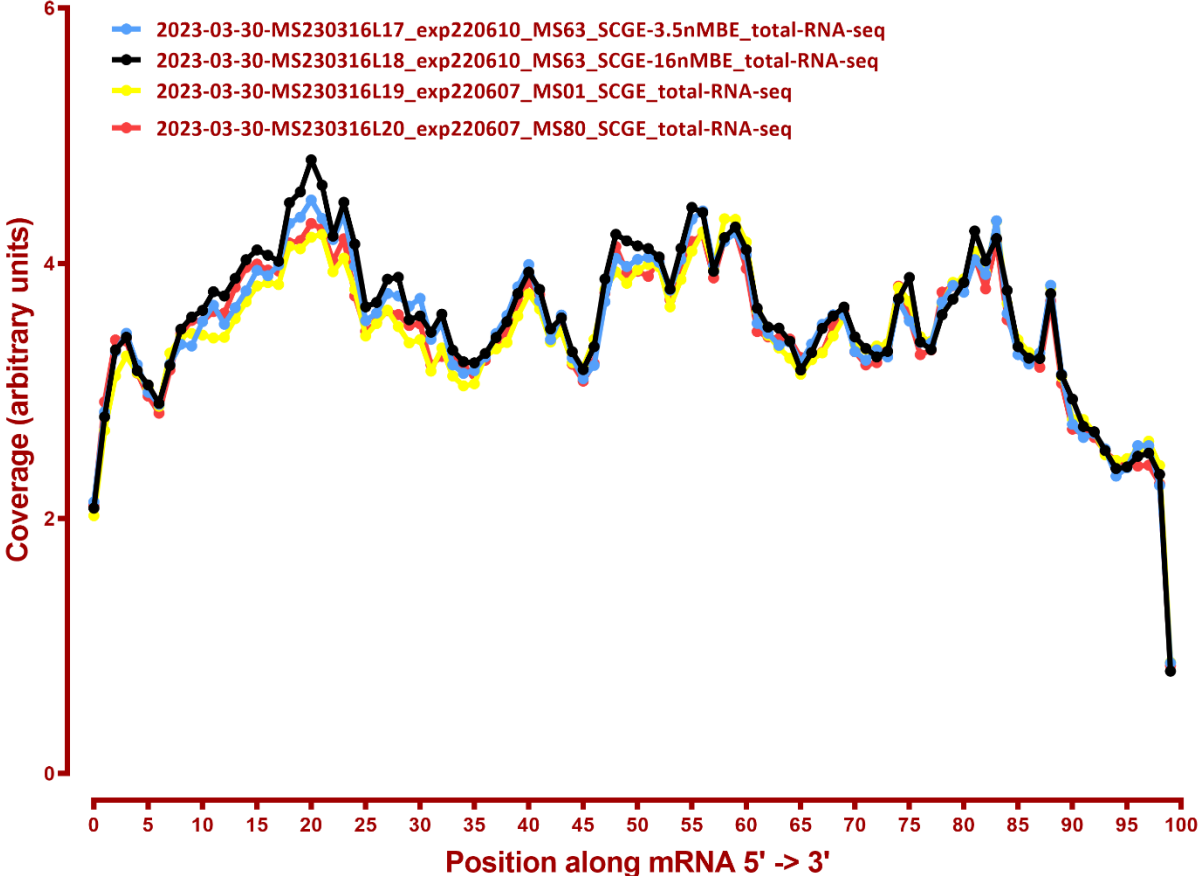
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Coverage of genes; >1000bp, sacCer3



Coverage of genes; >1000bp, sacCer3



Coverage of genes; >1000bp, sacCer3

