

MAPPING STATS

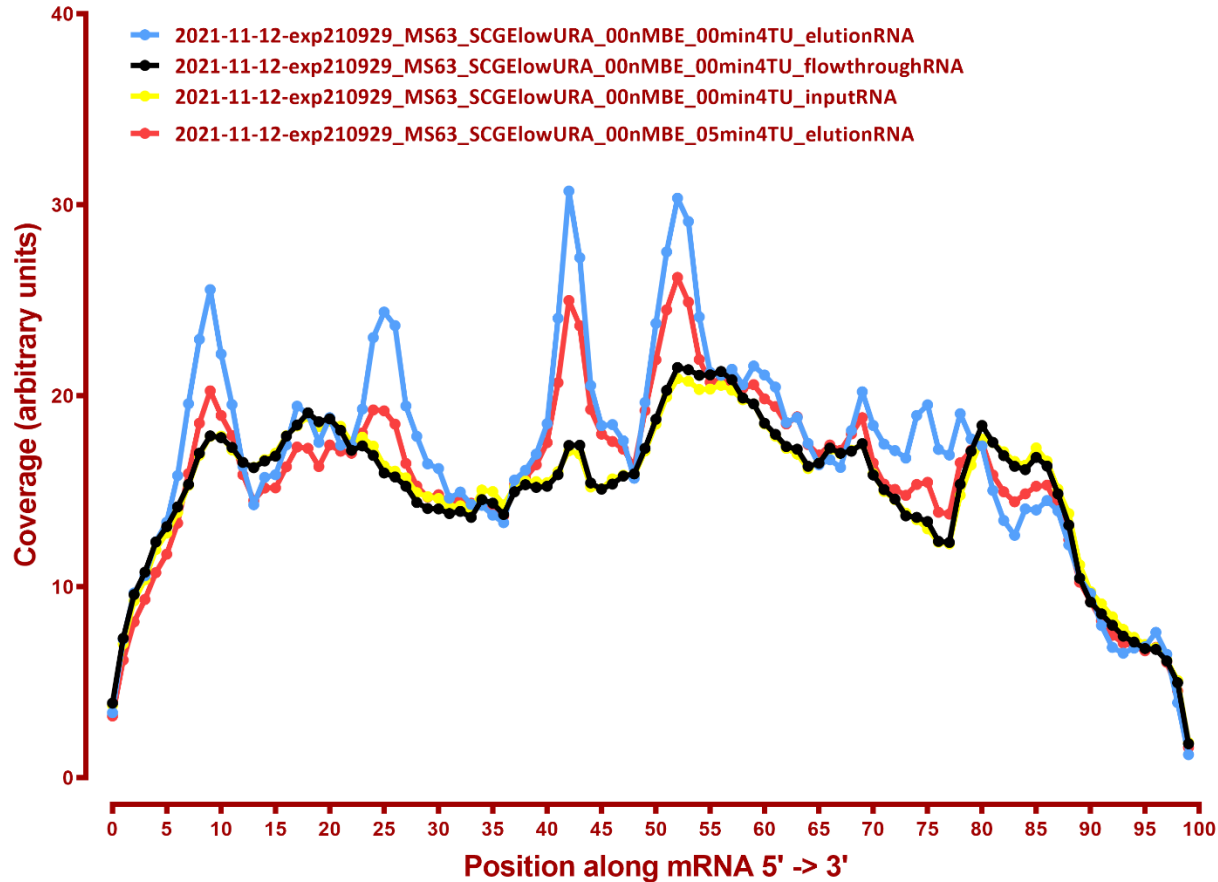
Library	Read Length	Mapping	Species	Raw fragments	Complexity	Unique	Unique Splices	Multi	Multi Splices	Fraction mapped
2021-11-12-exp210929_MS63_SCGElowURA_00nMBE_00min4TU_elutionRNA	2x150	STAR-2.5.3a	Candida	18,230,869	0.13	7,316,340	17,684	3,286,396	198,640	0.30
2021-11-12-exp210929_MS63_SCGElowURA_00nMBE_00min4TU_elutionRNA	2x150	STAR-2.5.3a	sacCer3	18,230,869	0.32	2,692,584	203,164	17,514,636	1,102,950	0.59
2021-11-12-exp210929_MS63_SCGElowURA_00nMBE_00min4TU_flowthroughRNA	2x150	STAR-2.5.3a	Candida	17,741,347	0.21	8,604,595	34,093	2,463,335	172,827	0.32
2021-11-12-exp210929_MS63_SCGElowURA_00nMBE_00min4TU_flowthroughRNA	2x150	STAR-2.5.3a	sacCer3	17,741,347	0.35	8,246,065	418,927	14,047,589	748,521	0.66
2021-11-12-exp210929_MS63_SCGElowURA_00nMBE_00min4TU_inputRNA	2x150	STAR-2.5.3a	Candida	18,304,980	0.21	9,173,096	33,446	2,566,739	176,077	0.33
2021-11-12-exp210929_MS63_SCGElowURA_00nMBE_00min4TU_inputRNA	2x150	STAR-2.5.3a	sacCer3	18,304,980	0.36	8,300,158	402,232	14,723,030	1,146,898	0.67
2021-11-12-exp210929_MS63_SCGElowURA_00nMBE_05min4TU_elutionRNA	2x150	STAR-2.5.3a	Candida	19,498,486	0.16	8,820,209	25,899	3,320,121	274,279	0.32
2021-11-12-exp210929_MS63_SCGElowURA_00nMBE_05min4TU_elutionRNA	2x150	STAR-2.5.3a	sacCer3	19,498,486	0.35	4,702,153	295,429	19,817,643	1,444,777	0.67
2021-11-12-exp210929_MS63_SCGElowURA_00nMBE_05min4TU_flowthroughRNA	2x150	STAR-2.5.3a	Candida	15,566,804	0.21	7,538,771	29,145	2,162,032	155,954	0.32
2021-11-12-exp210929_MS63_SCGElowURA_00nMBE_05min4TU_flowthroughRNA	2x150	STAR-2.5.3a	sacCer3	15,566,804	0.34	6,899,748	364,196	12,877,522	672,604	0.67
2021-11-12-exp210929_MS63_SCGElowURA_00nMBE_05min4TU_inputRNA	2x150	STAR-2.5.3a	Candida	27,883,294	0.18	13,441,478	49,796	4,004,532	320,194	0.32
2021-11-12-exp210929_MS63_SCGElowURA_00nMBE_05min4TU_inputRNA	2x150	STAR-2.5.3a	sacCer3	27,883,294	0.30	12,447,143	743,815	22,811,515	1,839,391	0.68
2021-11-12-exp210929_MS63_SCGElowURA_00nMBE_10min4TU_elutionRNA	2x150	STAR-2.5.3a	Candida	18,534,159	0.12	8,497,963	14,375	3,660,655	192,483	0.33
2021-11-12-exp210929_MS63_SCGElowURA_00nMBE_10min4TU_elutionRNA	2x150	STAR-2.5.3a	sacCer3	18,534,159	0.28	2,612,522	241,564	20,805,896	1,223,138	0.67
2021-11-12-exp210929_MS63_SCGElowURA_00nMBE_10min4TU_flowthroughRNA	2x150	STAR-2.5.3a	Candida	16,323,221	0.22	8,038,543	39,143	2,123,172	233,298	0.32
2021-11-12-exp210929_MS63_SCGElowURA_00nMBE_10min4TU_flowthroughRNA	2x150	STAR-2.5.3a	sacCer3	16,323,221	0.32	7,494,268	423,934	12,861,796	579,360	0.65
2021-11-12-exp210929_MS63_SCGElowURA_00nMBE_10min4TU_inputRNA	2x150	STAR-2.5.3a	Candida	14,900,270	0.22	7,518,164	27,066	1,965,629	145,991	0.32
2021-11-12-exp210929_MS63_SCGElowURA_00nMBE_10min4TU_inputRNA	2x150	STAR-2.5.3a	sacCer3	14,900,270	0.32	6,672,686	371,726	12,169,821	550,251	0.66
2021-11-12-exp210929_MS63_SCGElowURA_05nMBE_00min4TU_elutionRNA	2x150	STAR-2.5.3a	Candida	17,951,991	0.13	7,237,477	11,469	3,553,768	282,332	0.31
2021-11-12-exp210929_MS63_SCGElowURA_05nMBE_00min4TU_elutionRNA	2x150	STAR-2.5.3a	sacCer3	17,951,991	0.33	2,721,962	206,028	21,703,861	1,401,271	0.73
2021-11-12-exp210929_MS63_SCGElowURA_05nMBE_00min4TU_flowthroughRNA	2x150	STAR-2.5.3a	Candida	13,568,581	0.23	5,597,844	21,054	1,869,809	90,453	0.28
2021-11-12-exp210929_MS63_SCGElowURA_05nMBE_00min4TU_flowthroughRNA	2x150	STAR-2.5.3a	sacCer3	13,568,581	0.35	6,590,139	331,325	11,830,119	448,013	0.71
2021-11-12-exp210929_MS63_SCGElowURA_05nMBE_00min4TU_inputRNA	2x150	STAR-2.5.3a	Candida	21,158,405	0.20	8,878,472	28,186	3,097,538	185,006	0.29
2021-11-12-exp210929_MS63_SCGElowURA_05nMBE_00min4TU_inputRNA	2x150	STAR-2.5.3a	sacCer3	21,158,405	0.33	9,326,485	503,007	19,229,775	1,259,891	0.72
2021-11-12-exp210929_MS63_SCGElowURA_05nMBE_05min4TU_elutionRNA	2x150	STAR-2.5.3a	Candida	20,970,879	0.13	7,499,524	15,628	3,973,477	253,771	0.28
2021-11-12-exp210929_MS63_SCGElowURA_05nMBE_05min4TU_elutionRNA	2x150	STAR-2.5.3a	sacCer3	20,970,879	0.32	3,340,149	291,095	25,951,326	1,772,242	0.75

2021-11-12-exp210929_MS63_SCGElowURA_05nMBE_05min4TU_flowthroughRNA	2x150	STAR-2.5.3a	Candida	20,484,758	0.21	8,135,786	35,678	2,653,060	208,292	0.27
2021-11-12-exp210929_MS63_SCGElowURA_05nMBE_05min4TU_flowthroughRNA	2x150	STAR-2.5.3a	sacCer3	20,484,758	0.32	9,509,344	557,774	17,485,492	1,436,716	0.71
2021-11-12-exp210929_MS63_SCGElowURA_05nMBE_05min4TU_inputRNA	2x150	STAR-2.5.3a	Candida	16,382,274	0.21	6,579,470	23,470	2,149,710	175,864	0.27
2021-11-12-exp210929_MS63_SCGElowURA_05nMBE_05min4TU_inputRNA	2x150	STAR-2.5.3a	sacCer3	16,382,274	0.34	7,450,561	410,193	14,722,049	926,873	0.72
2021-11-12-exp210929_MS63_SCGElowURA_05nMBE_10min4TU_elutionRNA	2x150	STAR-2.5.3a	Candida	22,861,572	0.17	8,971,531	31,515	3,756,403	349,555	0.29
2021-11-12-exp210929_MS63_SCGElowURA_05nMBE_10min4TU_elutionRNA	2x150	STAR-2.5.3a	sacCer3	22,861,572	0.34	6,438,886	400,142	23,834,224	1,673,298	0.71
2021-11-12-exp210929_MS63_SCGElowURA_05nMBE_10min4TU_flowthroughRNA	2x150	STAR-2.5.3a	Candida	17,723,236	0.22	7,460,407	28,523	2,269,970	120,180	0.28
2021-11-12-exp210929_MS63_SCGElowURA_05nMBE_10min4TU_flowthroughRNA	2x150	STAR-2.5.3a	sacCer3	17,723,236	0.32	8,246,218	455,138	15,156,320	849,244	0.70
2021-11-12-exp210929_MS63_SCGElowURA_05nMBE_10min4TU_inputRNA	2x150	STAR-2.5.3a	Candida	17,170,014	0.22	7,179,533	25,867	2,298,340	160,358	0.28
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2021-11-12-exp210929_MS63_SCGElowURA_15nMBE_00min4TU_elutionRNA	2x150	STAR-2.5.3a	Candida	19,900,761	0.15	7,320,957	18,255	3,621,350	327,468	0.28
2021-11-12-exp210929_MS63_SCGElowURA_15nMBE_00min4TU_elutionRNA	2x150	STAR-2.5.3a	sacCer3	19,900,761	0.33	3,831,323	313,141	23,588,519	2,017,309	0.75
2021-11-12-exp210929_MS63_SCGElowURA_15nMBE_00min4TU_flowthroughRNA	2x150	STAR-2.5.3a	Candida	17,172,877	0.19	6,844,351	25,665	2,458,634	171,338	0.28
2021-11-12-exp210929_MS63_SCGElowURA_15nMBE_00min4TU_flowthroughRNA	2x150	STAR-2.5.3a	sacCer3	17,172,877	0.34	6,613,285	398,265	16,366,775	1,062,755	0.71
2021-11-12-exp210929_MS63_SCGElowURA_15nMBE_00min4TU_inputRNA	2x150	STAR-2.5.3a	Candida	16,485,337	0.19	6,702,729	20,803	2,349,763	177,623	0.28
2021-11-12-exp210929_MS63_SCGElowURA_15nMBE_00min4TU_inputRNA	2x150	STAR-2.5.3a	sacCer3	16,485,337	0.33	6,239,988	411,898	16,304,858	933,572	0.72
2021-11-12-exp210929_MS63_SCGElowURA_15nMBE_05min4TU_elutionRNA	2x150	STAR-2.5.3a	Candida	25,310,582	0.12	8,933,529	20,447	4,386,895	195,319	0.27
2021-11-12-exp210929_MS63_SCGElowURA_15nMBE_05min4TU_elutionRNA	2x150	STAR-2.5.3a	sacCer3	25,310,582	0.33	3,667,484	291,686	30,500,819	2,332,295	0.73
2021-11-12-exp210929_MS63_SCGElowURA_15nMBE_05min4TU_flowthroughRNA	2x150	STAR-2.5.3a	Candida	15,144,889	0.23	5,812,908	23,006	1,898,975	91,061	0.26
2021-11-12-exp210929_MS63_SCGElowURA_15nMBE_05min4TU_flowthroughRNA	2x150	STAR-2.5.3a	sacCer3	15,144,889	0.35	6,995,364	387,690	13,045,017	817,871	0.70
2021-11-12-exp210929_MS63_SCGElowURA_15nMBE_05min4TU_inputRNA	2x150	STAR-2.5.3a	Candida	15,977,191	0.23	6,199,947	22,819	2,153,386	121,620	0.27
2021-11-12-exp210929_MS63_SCGElowURA_15nMBE_05min4TU_inputRNA	2x150	STAR-2.5.3a	sacCer3	15,977,191	0.35	7,202,397	382,361	14,474,535	799,233	0.72
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2021-11-12-exp210929_MS63_SCGElowURA_15nMBE_10min4TU_elutionRNA	2x150	STAR-2.5.3a	sacCer3	20,698,473	0.29	3,335,804	270,862	24,574,093	1,430,229	0.72
2021-11-12-exp210929_MS63_SCGElowURA_15nMBE_10min4TU_flowthroughRNA	2x150	STAR-2.5.3a	Candida	12,984,988	0.24	5,063,821	19,435	1,642,426	67,180	0.26
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2021-11-12-exp210929_MS63_SCGElowURA_15nMBE_10min4TU_inputRNA	2x150	STAR-2.5.3a	Candida	14,045,251	0.23	5,524,461	20,881	1,912,767	101,083	0.27
2021-11-12-exp210929_MS63_SCGElowURA_15nMBE_10min4TU_inputRNA	2x150	STAR-2.5.3a	sacCer3	14,045,251	0.35	6,115,799	337,233	12,937,661	675,869	0.71

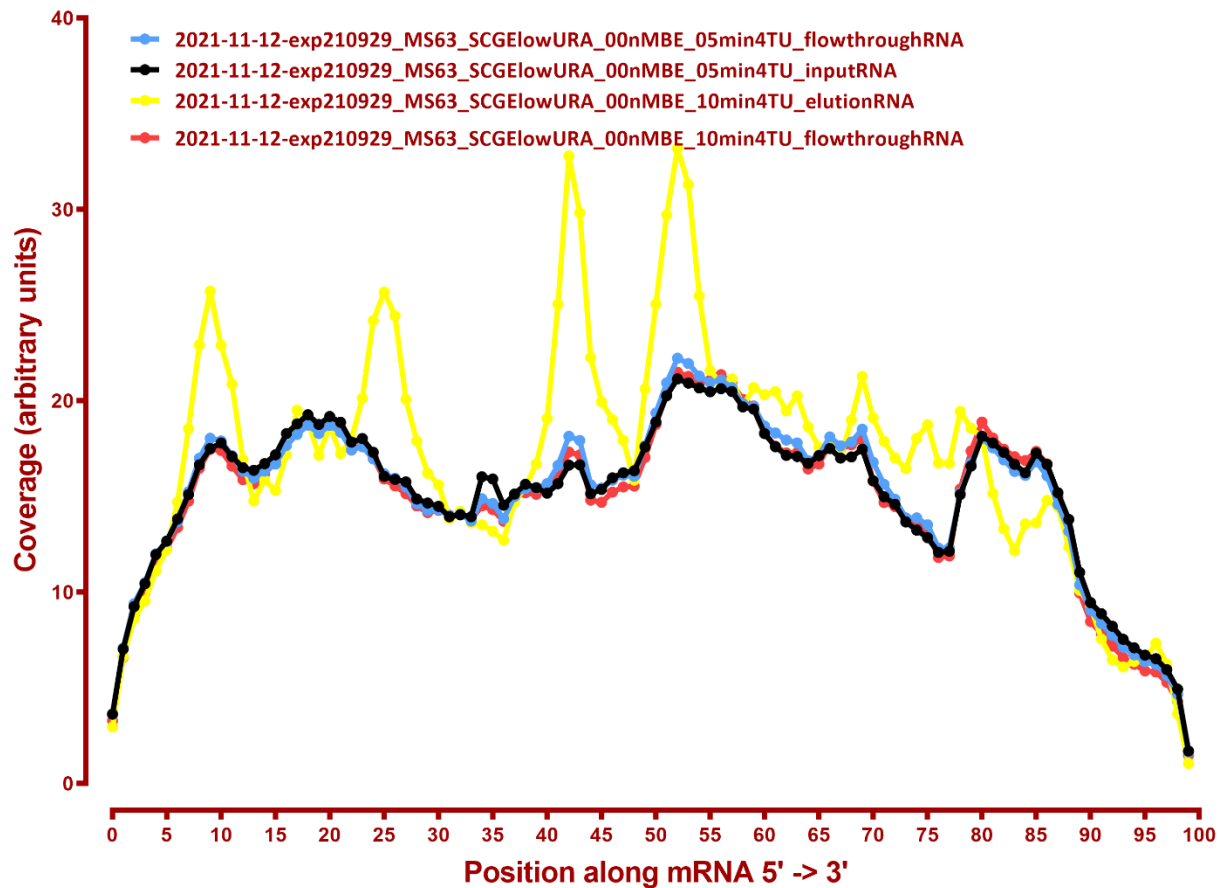
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2021-11-12-exp210929_MS63_SCGElowURA_05nMBE_00min4TU_flowthroughRNA.sacCer3	0.96	0.04	0.01
2021-11-12-exp210929_MS63_SCGElowURA_05nMBE_00min4TU_inputRNA.sacCer3	0.96	0.04	0.01
2021-11-12-exp210929_MS63_SCGElowURA_05nMBE_05min4TU_elutionRNA.sacCer3	0.97	0.03	0.00
2021-11-12-exp210929_MS63_SCGElowURA_05nMBE_05min4TU_flowthroughRNA.sacCer3	0.96	0.04	0.01
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COVERAGE ALL READS

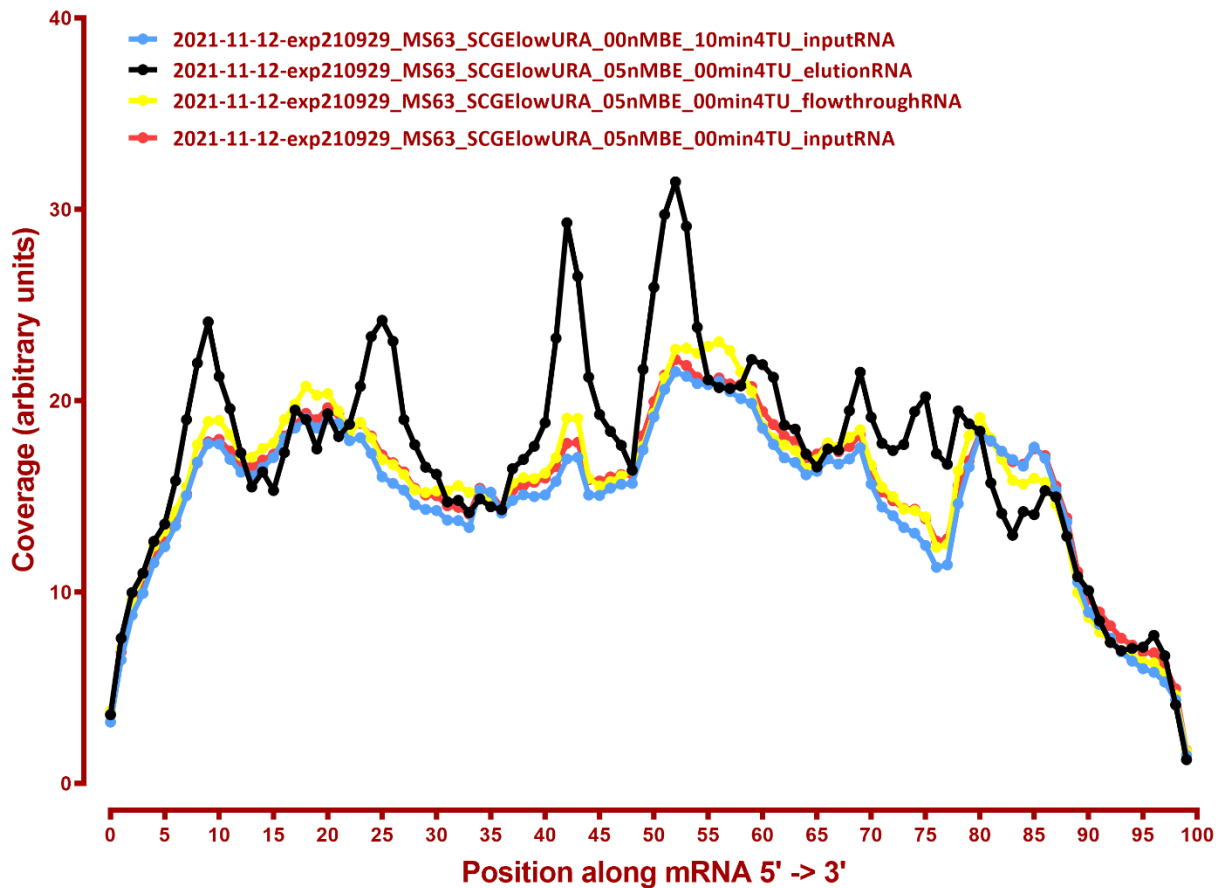
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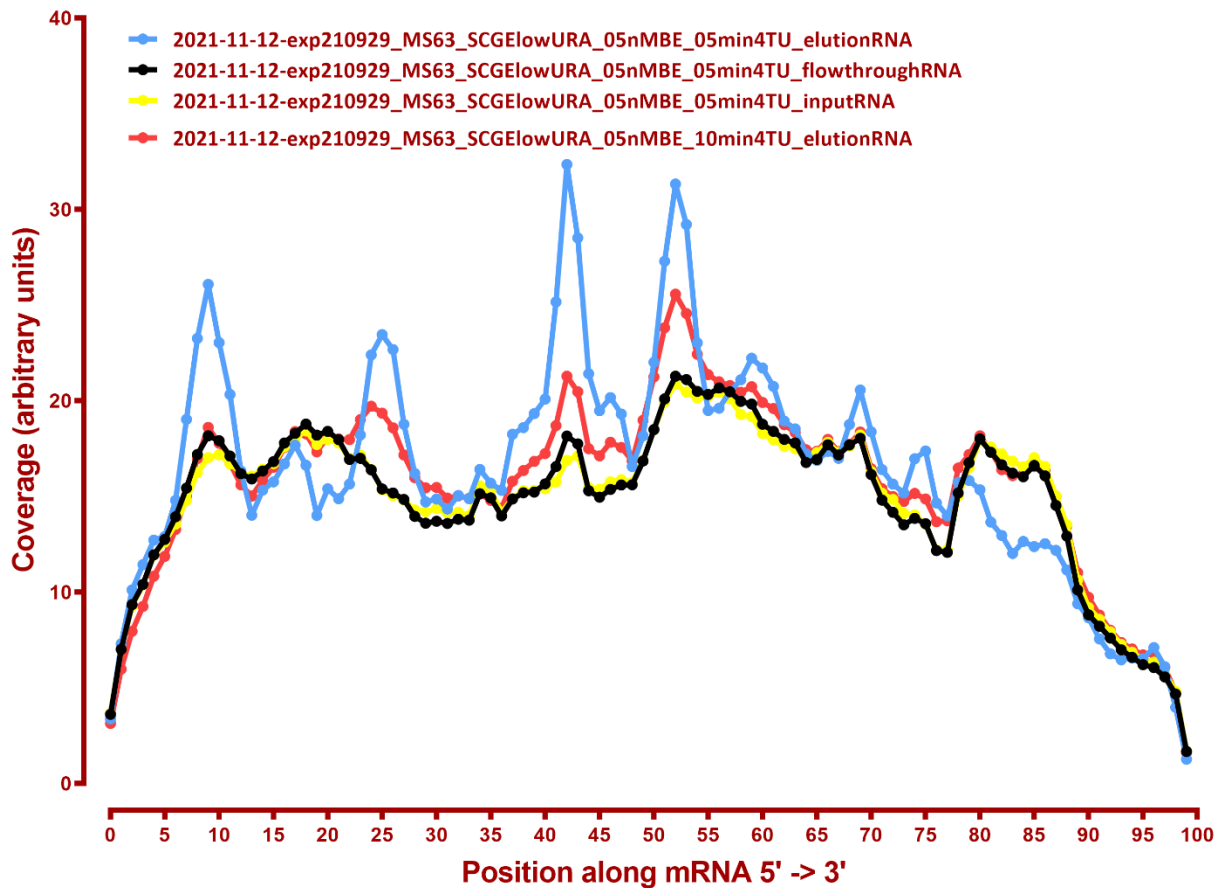
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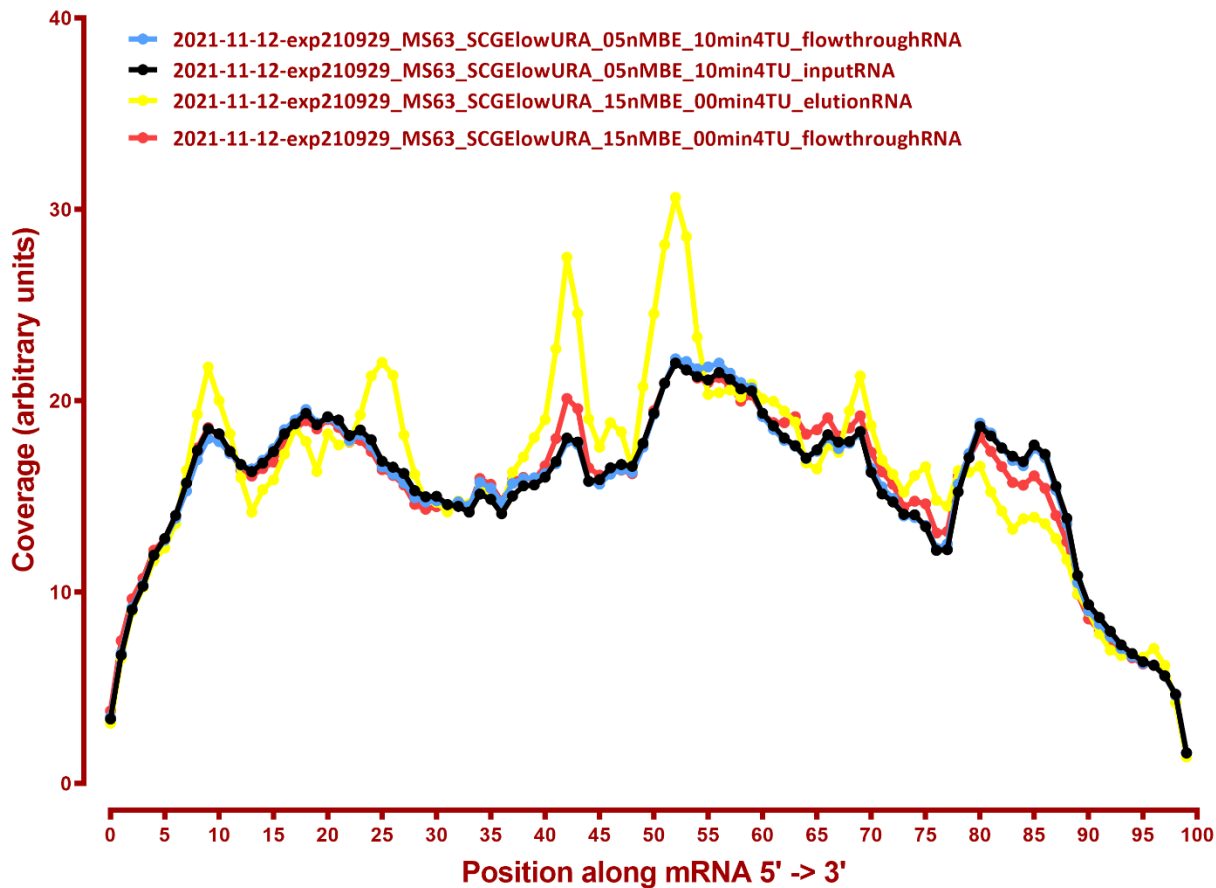
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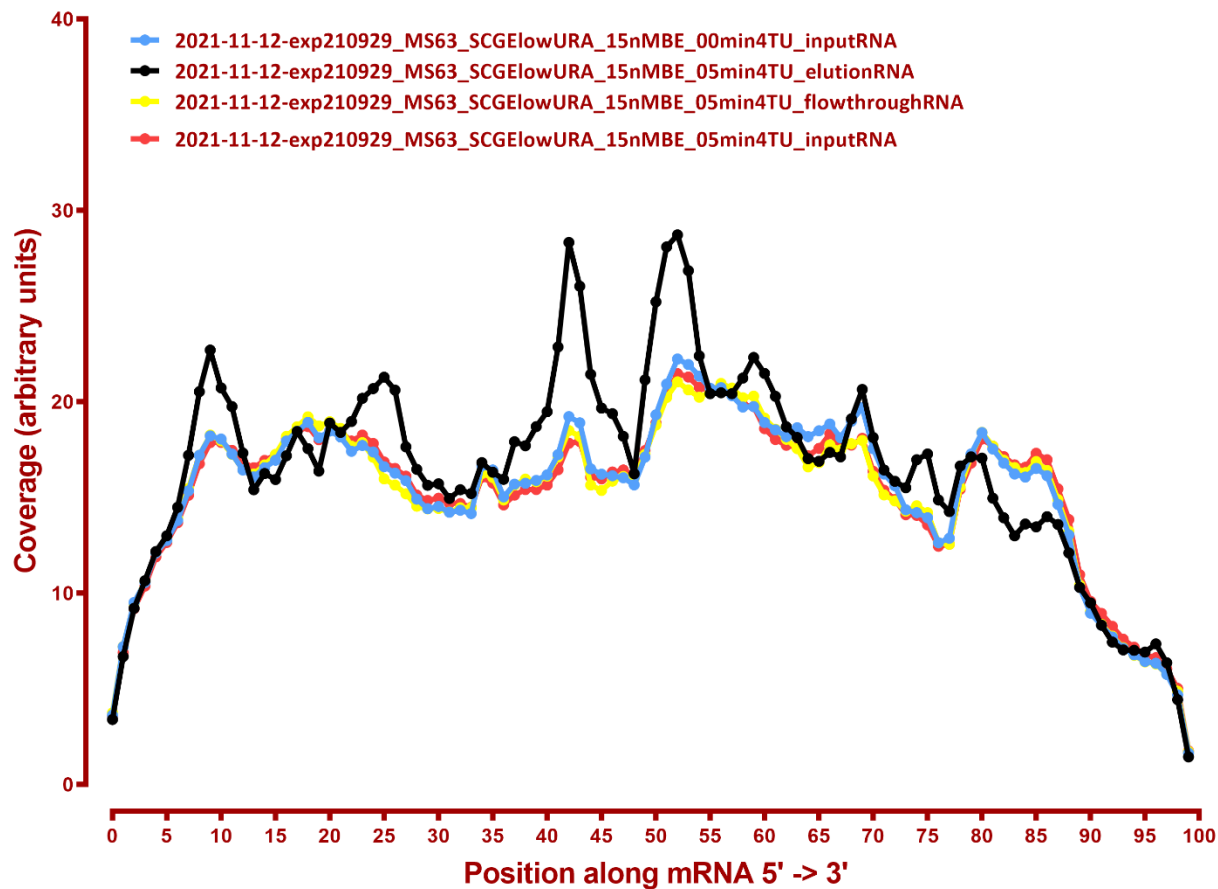
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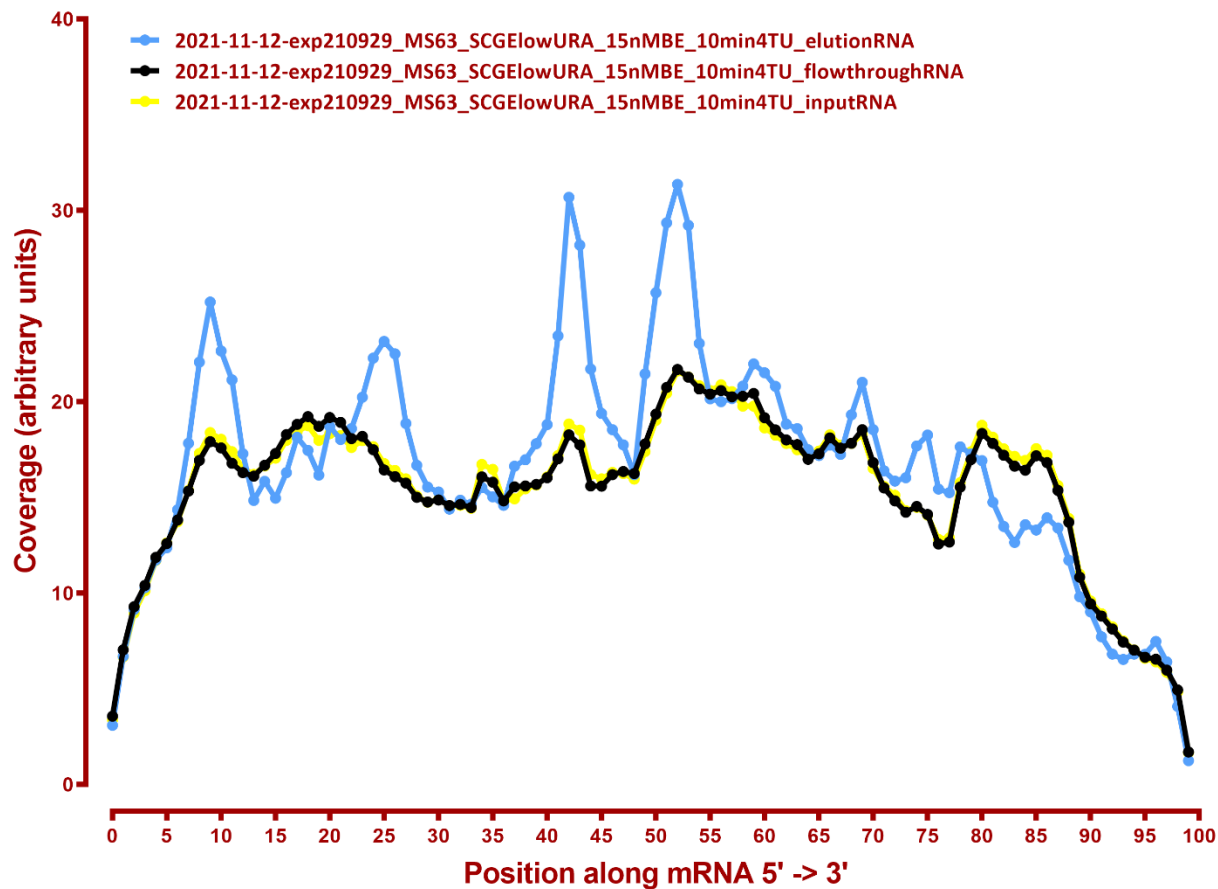
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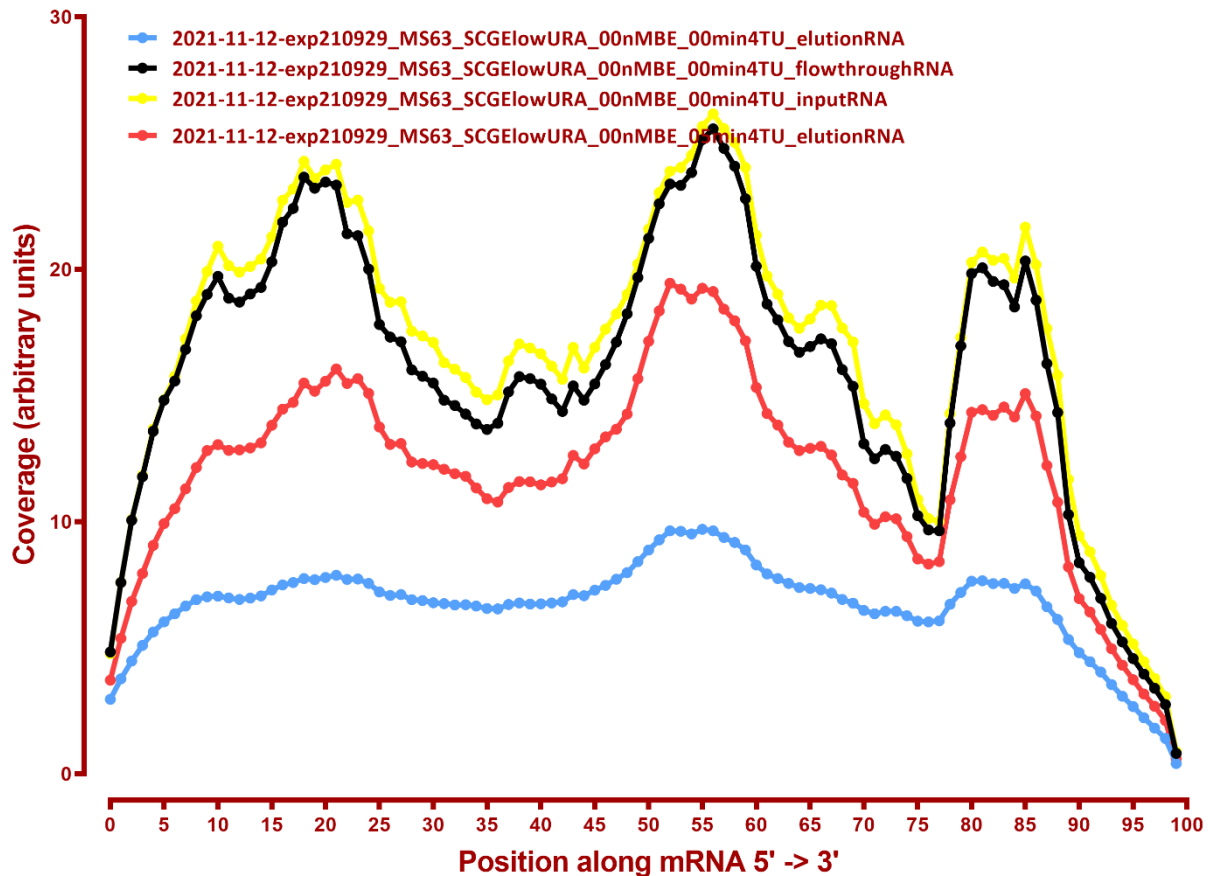


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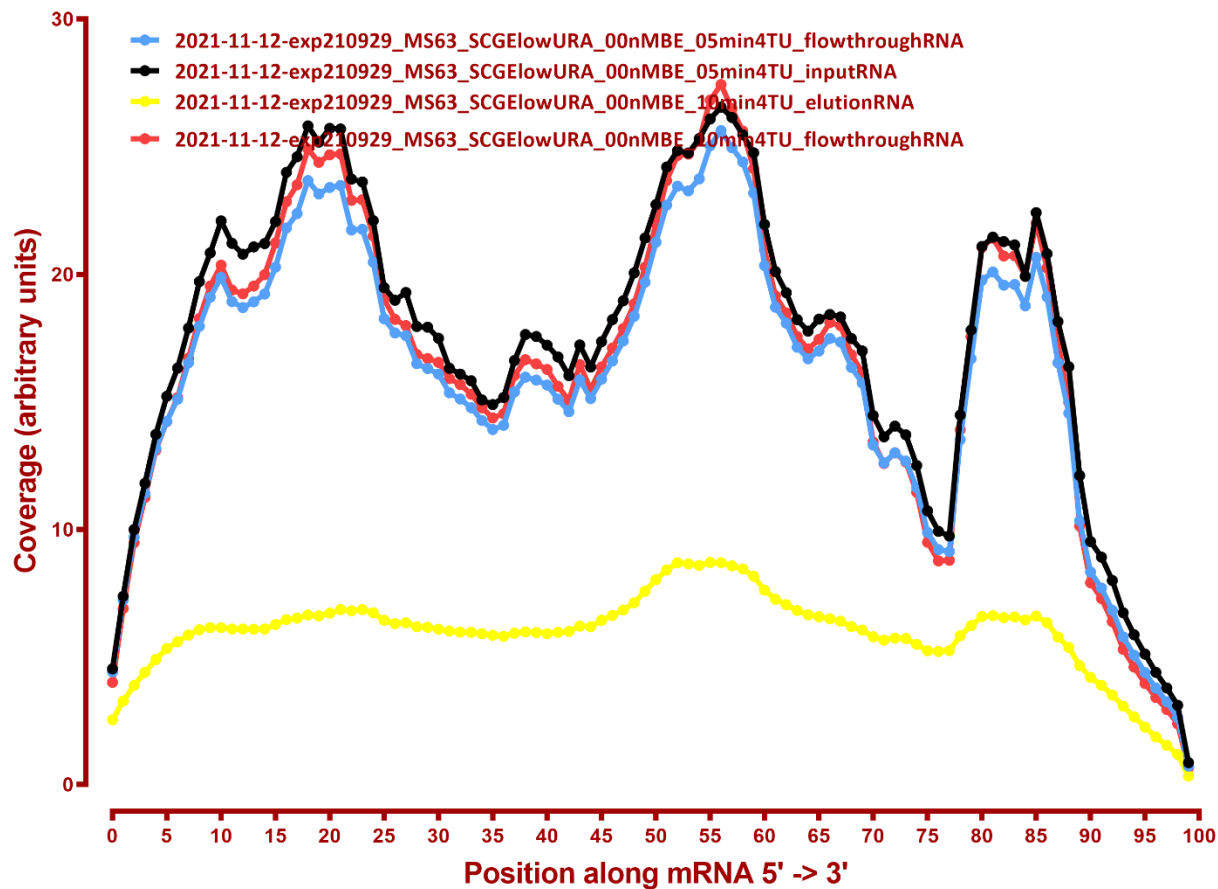


COVERAGE SENSE

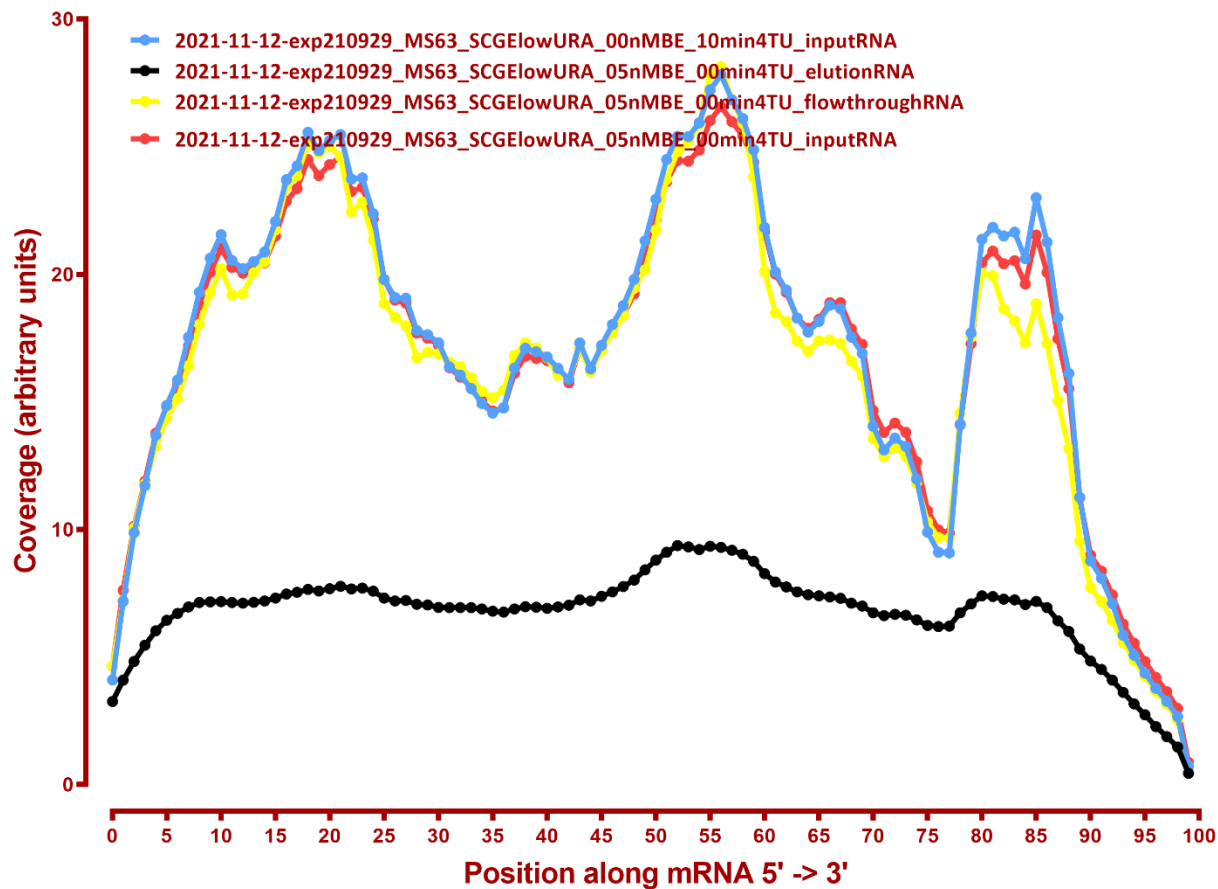
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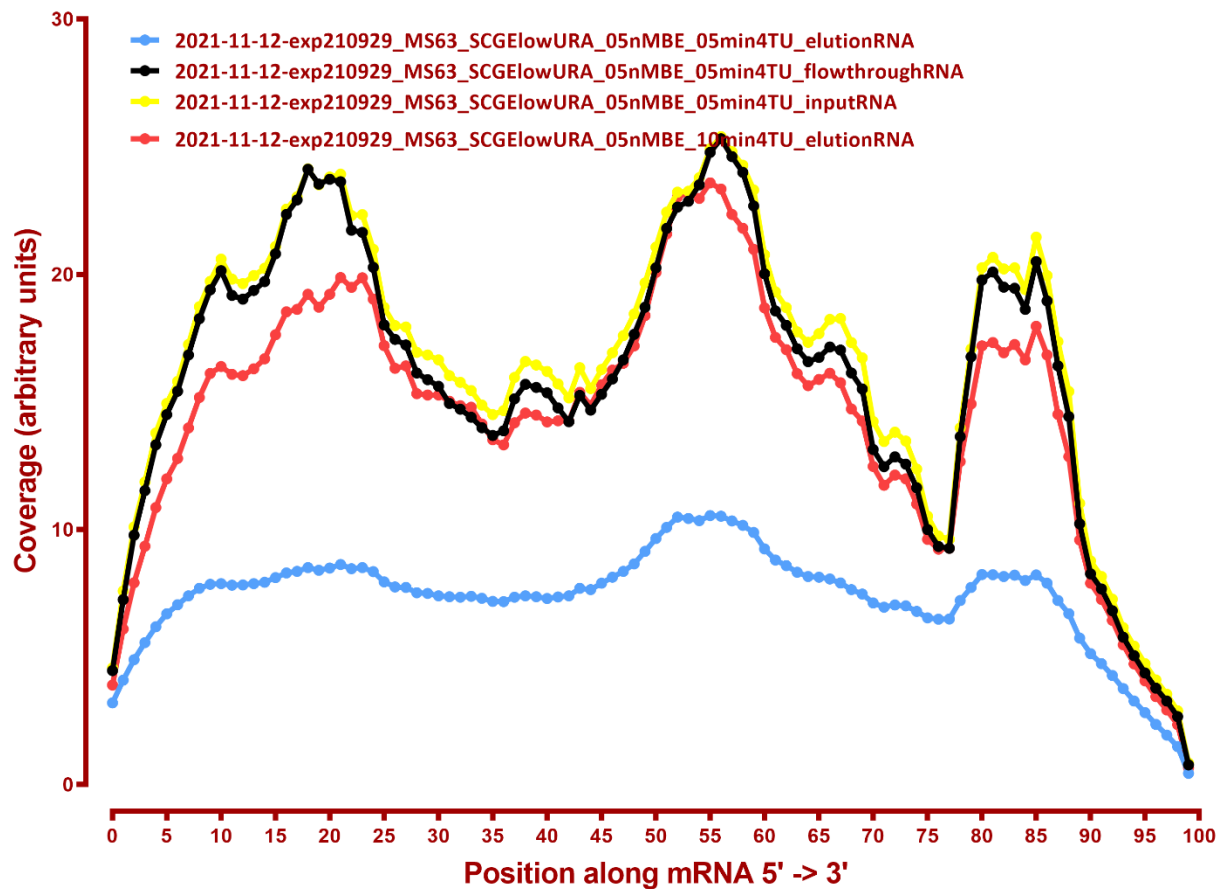
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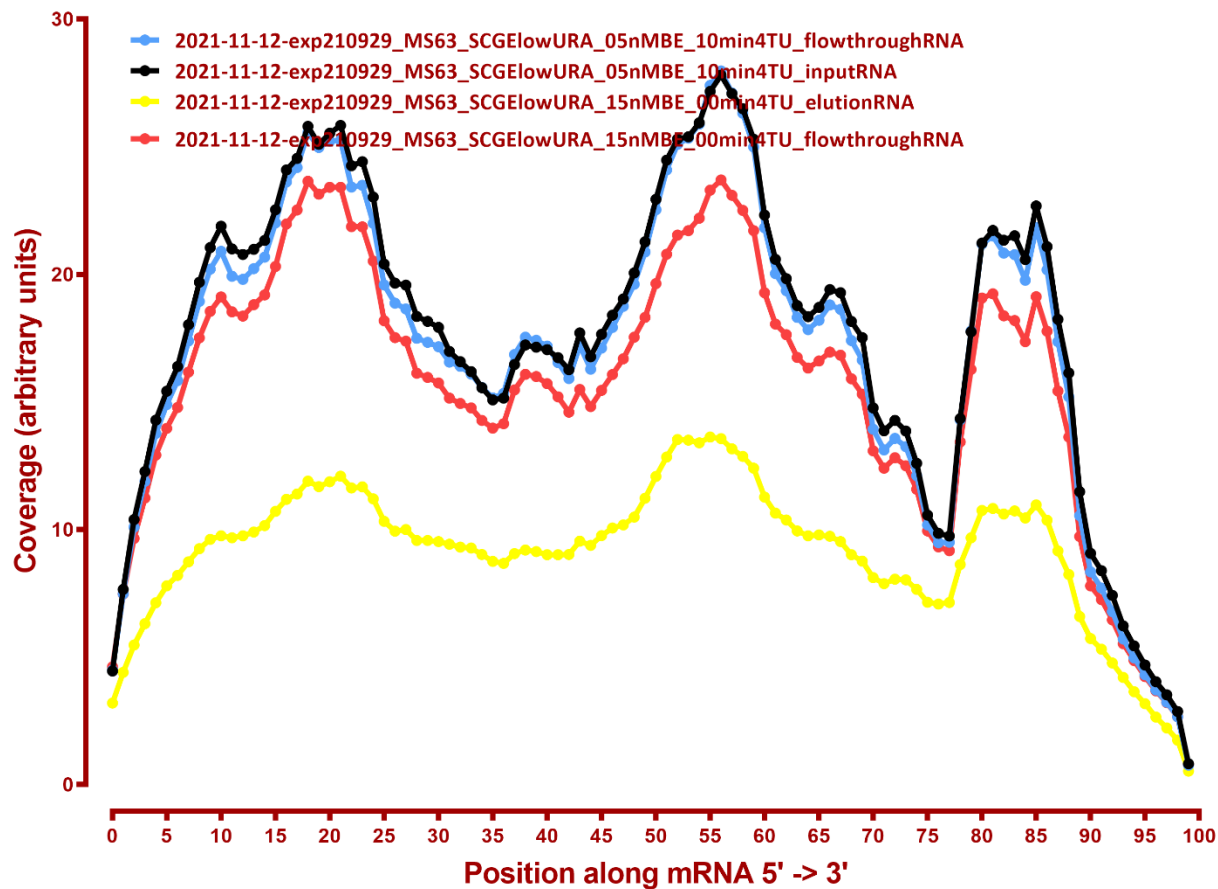
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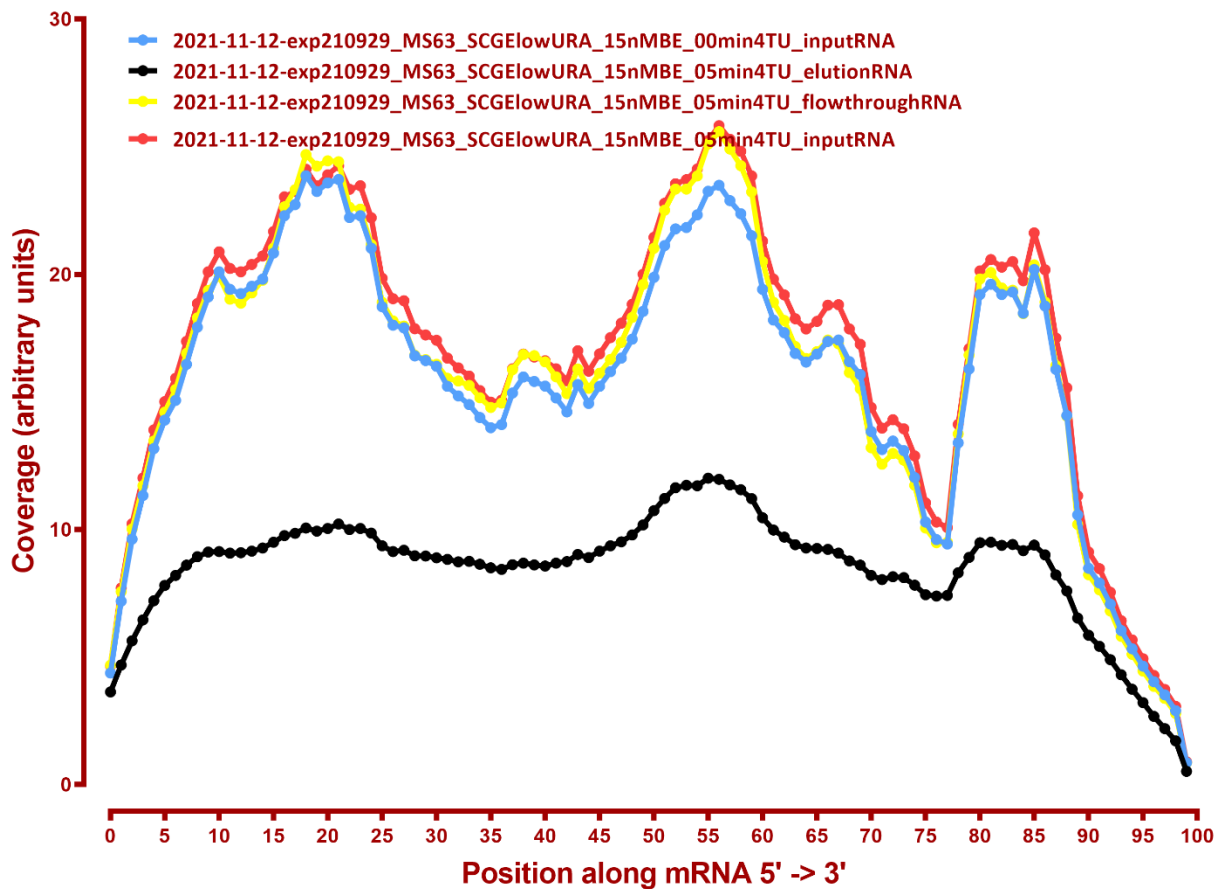
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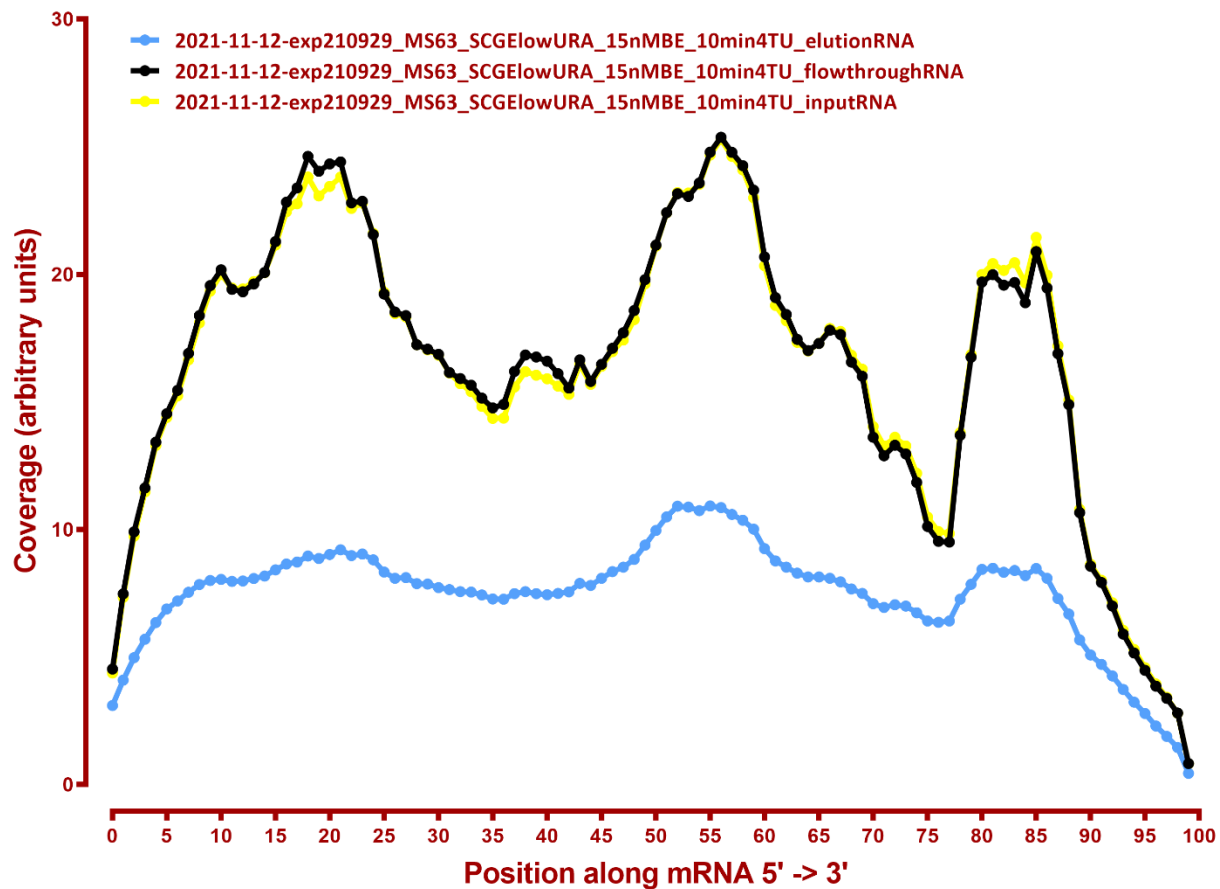
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Coverage of genes; >1000bp, sacCer3

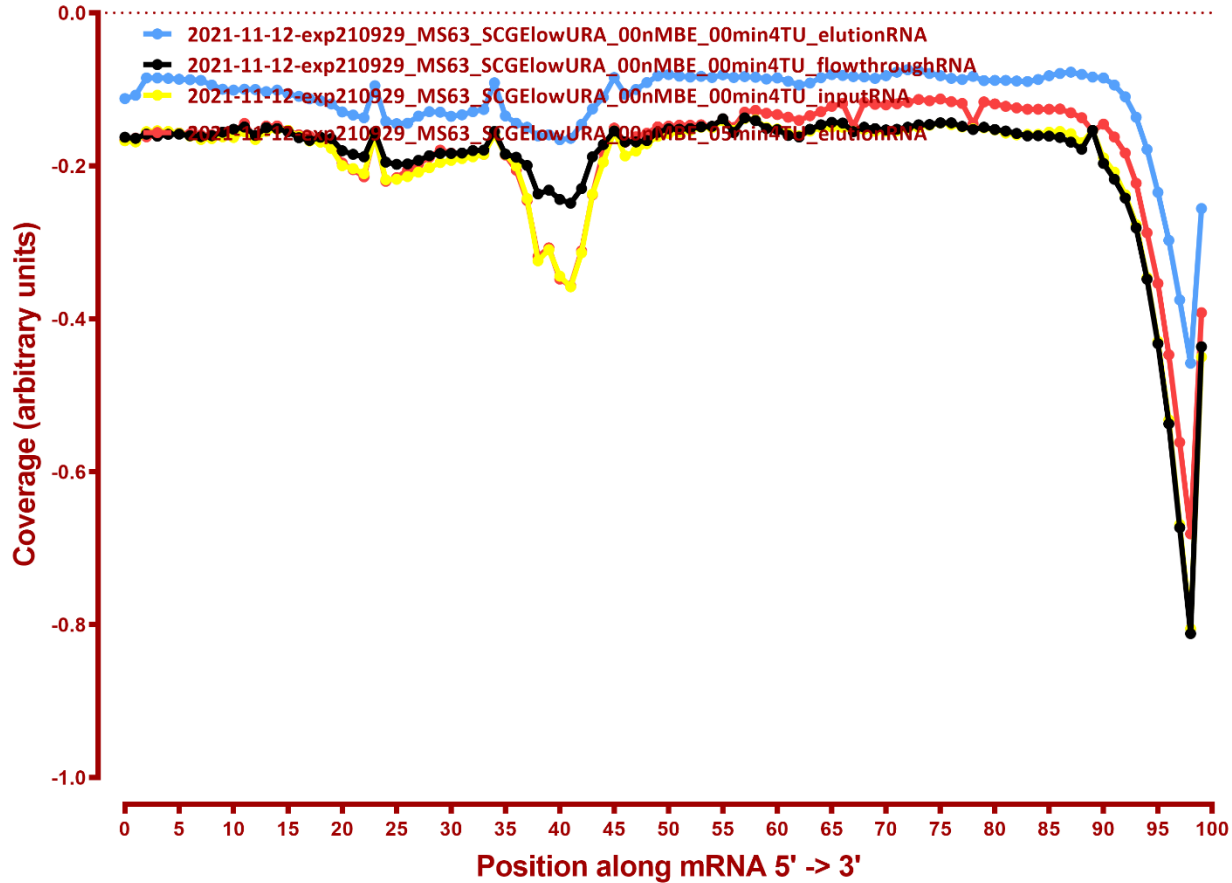


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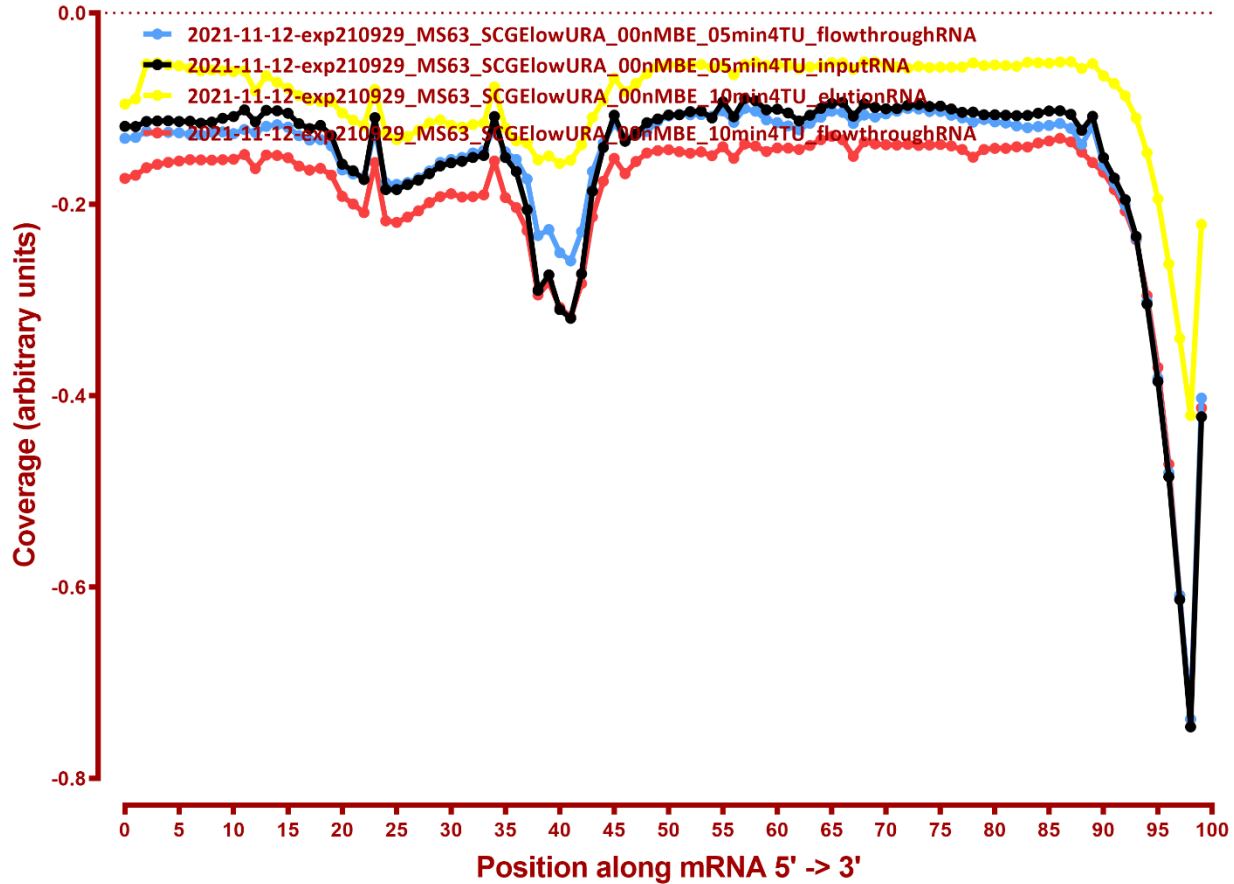


COVERAGE ANTISENSE

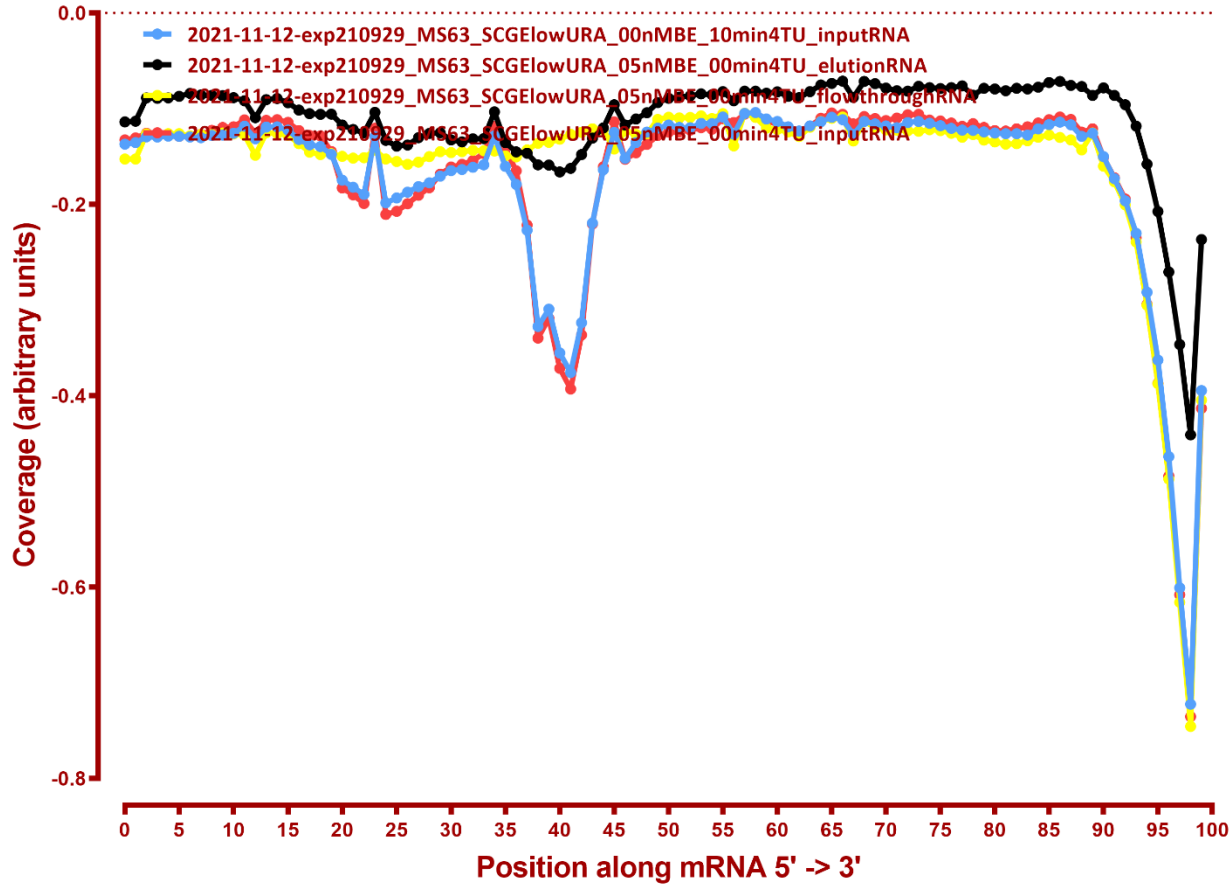
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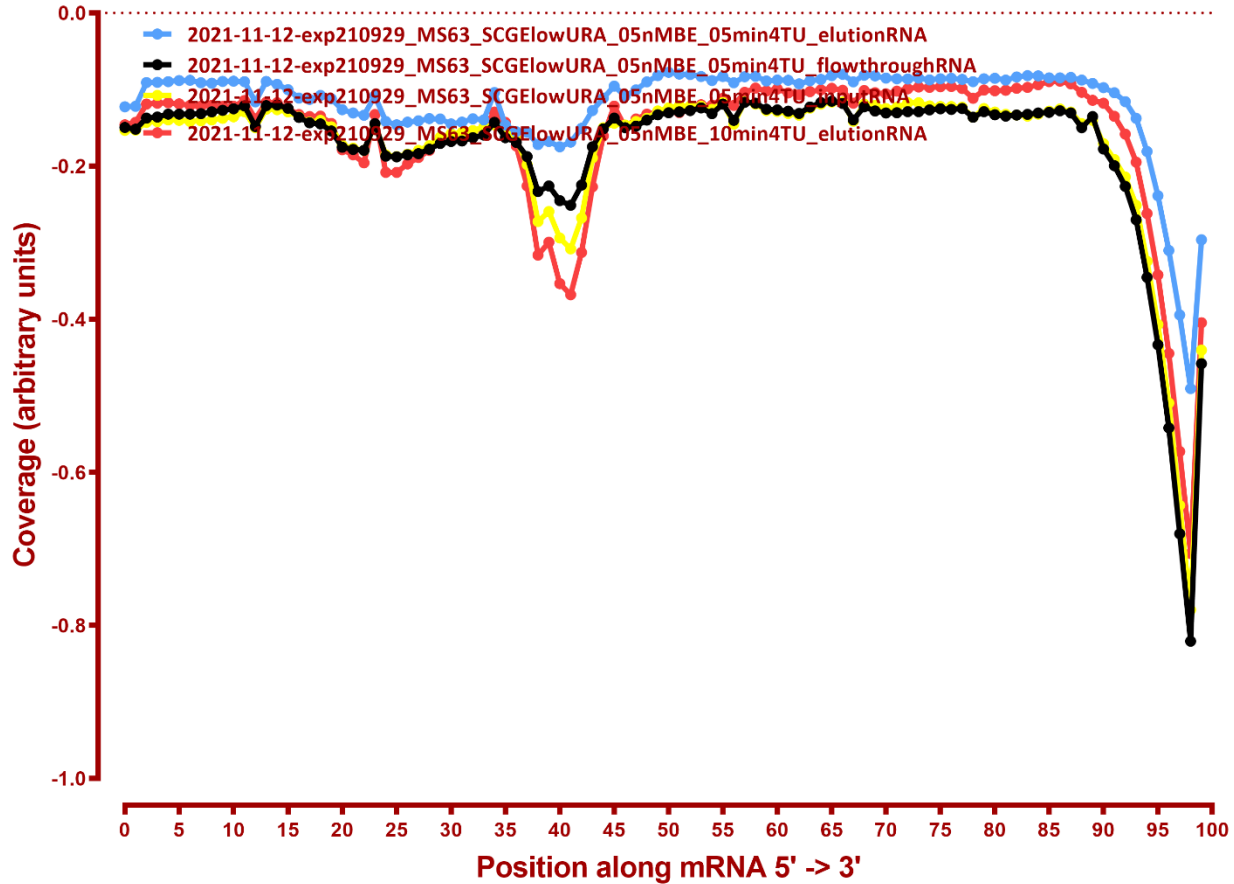
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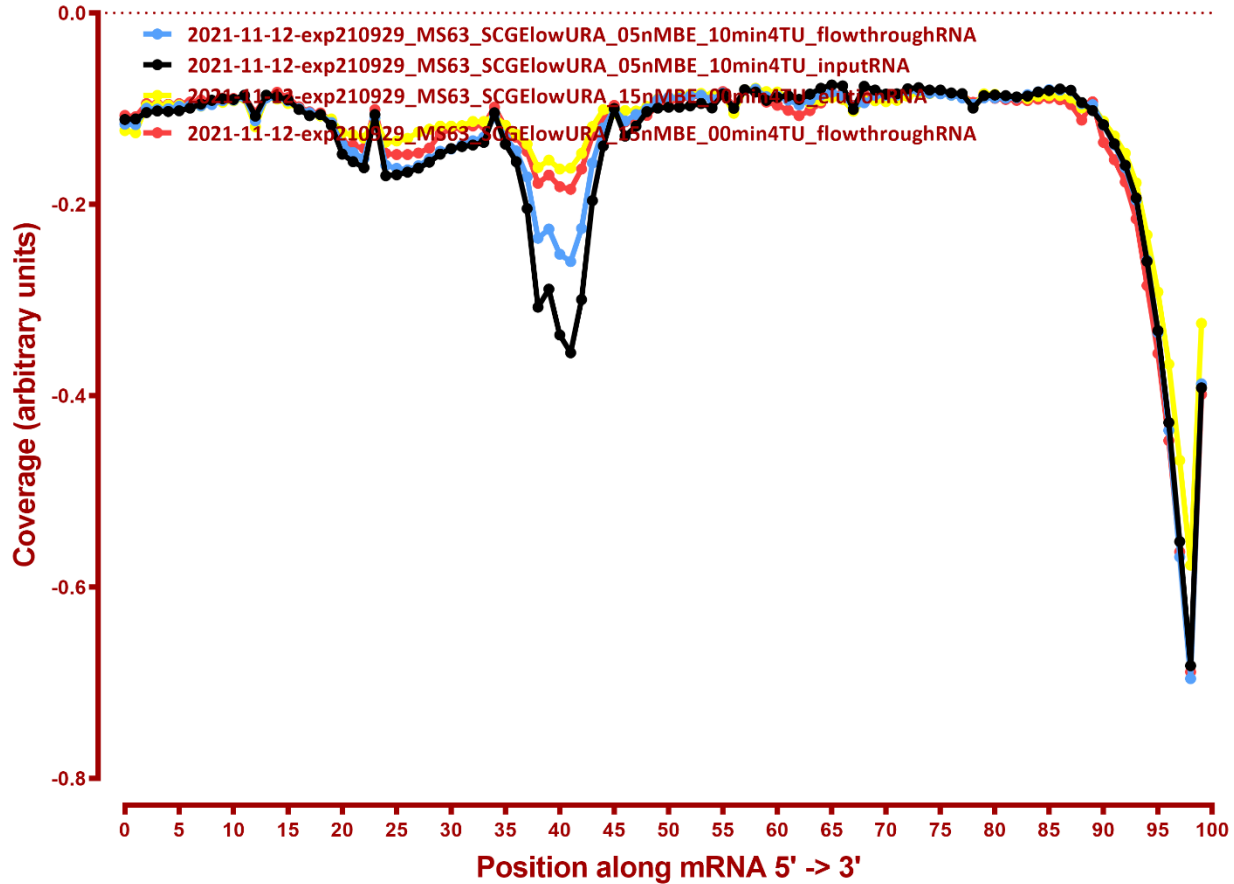
Coverage of genes; >1000bp, sacCer3



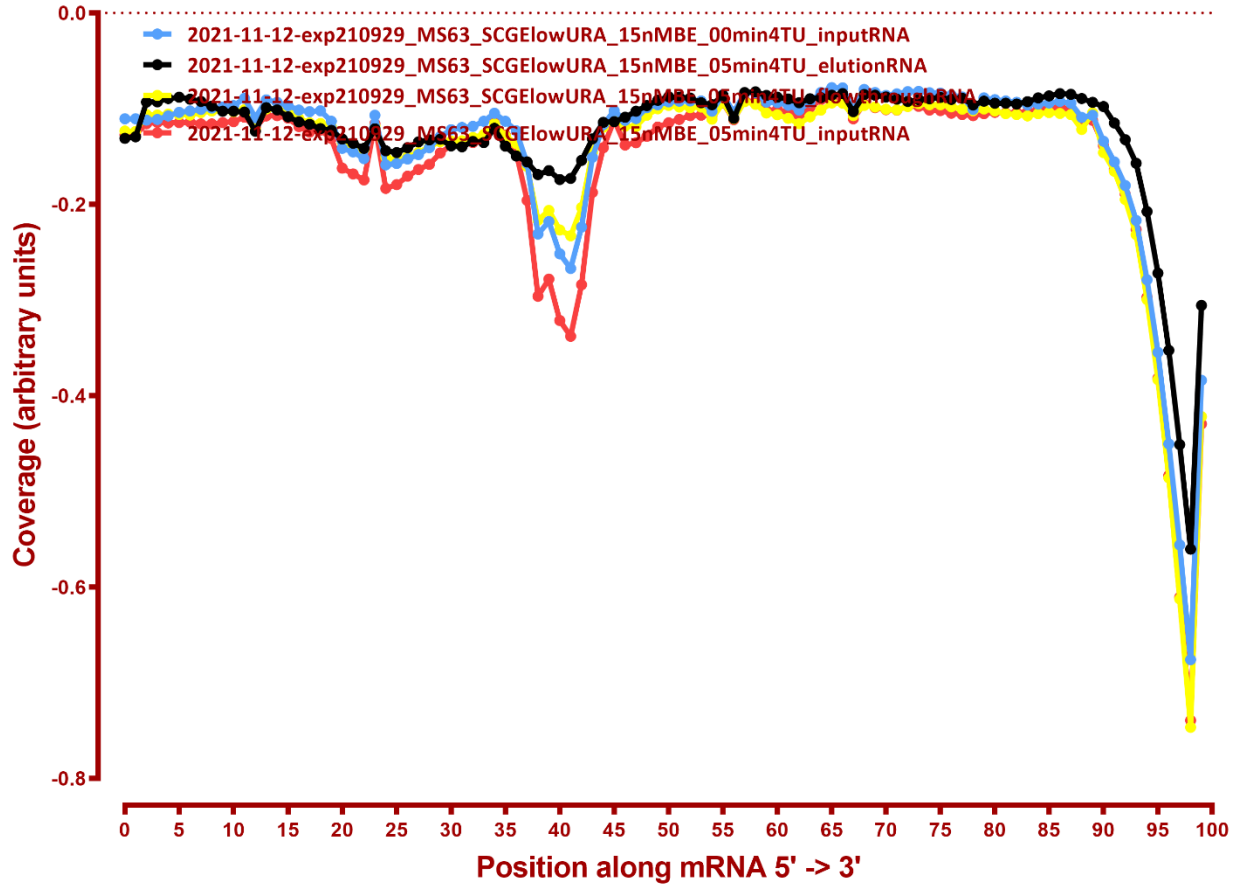
Coverage of genes; >1000bp, sacCer3



Coverage of genes; >1000bp, sacCer3



Coverage of genes; >1000bp, sacCer3



Coverage of genes; >1000bp, sacCer3

