

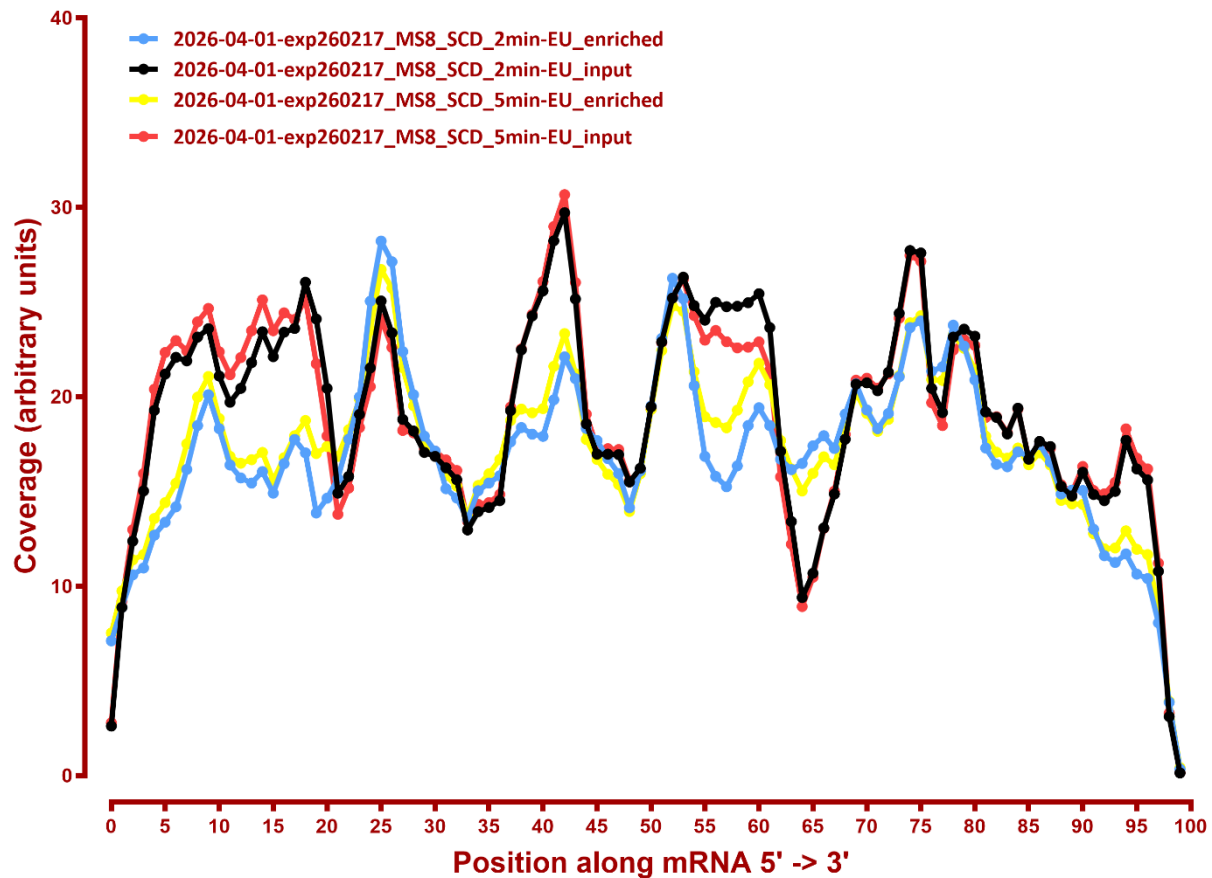
MAPPING STATS

Library	Read Length	Mapping	Species	Raw fragments	Complexity	Unique	Unique Splices	Multi	Multi Splices	Fraction mapped
2026-04-01-exp260217_MS8_SCD_2min-EU_enriched	2x150	STAR-2.5.3a	Candida	5,696,351	0.38	1,522,604	2,166	1,364,516	16,758	0.26
2026-04-01-exp260217_MS8_SCD_2min-EU_enriched	2x150	STAR-2.5.3a	sacCer3	5,696,351	0.34	2,376,679	60,057	5,474,073	190,433	0.71
2026-04-01-exp260217_MS8_SCD_2min-EU_input	2x150	STAR-2.5.3a	Candida	537,671	0.60	16,187	21	135,372	0	0.14
2026-04-01-exp260217_MS8_SCD_2min-EU_input	2x150	STAR-2.5.3a	sacCer3	537,671	0.33	237,626	3,044	705,470	10,692	0.89
2026-04-01-exp260217_MS8_SCD_5min-EU_enriched	2x150	STAR-2.5.3a	Candida	7,476,005	0.42	619,858	828	1,680,957	6,855	0.15
2026-04-01-exp260217_MS8_SCD_5min-EU_enriched	2x150	STAR-2.5.3a	sacCer3	7,476,005	0.37	3,705,785	92,613	8,086,311	327,169	0.82
2026-04-01-exp260217_MS8_SCD_5min-EU_input	2x150	STAR-2.5.3a	Candida	558,638	0.59	14,579	27	140,916	242	0.14
2026-04-01-exp260217_MS8_SCD_5min-EU_input	2x150	STAR-2.5.3a	sacCer3	558,638	0.32	247,450	3,914	736,720	7,490	0.89
2026-04-01-exp260217_MS8_SCD_no-EU_enriched	2x150	STAR-2.5.3a	Candida	4,947,610	0.34	2,900,446	6,370	1,276,295	54,103	0.43
2026-04-01-exp260217_MS8_SCD_no-EU_enriched	2x150	STAR-2.5.3a	sacCer3	4,947,610	0.18	1,295,766	39,792	3,585,045	146,757	0.51
2026-04-01-exp260217_MS8_SCD_no-EU_input	2x150	STAR-2.5.3a	Candida	629,704	0.57	19,922	16	159,332	2	0.14
2026-04-01-exp260217_MS8_SCD_no-EU_input	2x150	STAR-2.5.3a	sacCer3	629,704	0.31	280,386	6,222	823,023	12,707	0.89
2026-04-01-exp260217_MS8_SCGE_2min-EU_enriched	2x150	STAR-2.5.3a	Candida	6,103,651	0.34	2,643,779	3,699	1,367,546	47,066	0.33
2026-04-01-exp260217_MS8_SCGE_2min-EU_enriched	2x150	STAR-2.5.3a	sacCer3	6,103,651	0.41	2,524,314	52,572	4,749,673	180,877	0.61
2026-04-01-exp260217_MS8_SCGE_2min-EU_input	2x150	STAR-2.5.3a	Candida	578,267	0.57	22,672	34	142,944	450	0.14
2026-04-01-exp260217_MS8_SCGE_2min-EU_input	2x150	STAR-2.5.3a	sacCer3	578,267	0.32	271,531	3,037	717,406	13,316	0.87
2026-04-01-exp260217_MS8_SCGE_5min-EU_enriched	2x150	STAR-2.5.3a	Candida	6,574,614	0.41	972,120	2,490	1,343,596	31,030	0.18
2026-04-01-exp260217_MS8_SCGE_5min-EU_enriched	2x150	STAR-2.5.3a	sacCer3	6,574,614	0.48	3,877,018	71,608	5,763,743	212,659	0.75
2026-04-01-exp260217_MS8_SCGE_5min-EU_input	2x150	STAR-2.5.3a	Candida	565,014	0.56	23,260	14	141,617	569	0.15
2026-04-01-exp260217_MS8_SCGE_5min-EU_input	2x150	STAR-2.5.3a	sacCer3	565,014	0.34	261,746	3,622	733,084	9,466	0.89
2026-04-01-exp260217_MS8_SCGE_no-EU_enriched	2x150	STAR-2.5.3a	Candida	6,106,953	0.31	4,161,409	8,309	1,563,419	56,551	0.47
2026-04-01-exp260217_MS8_SCGE_no-EU_enriched	2x150	STAR-2.5.3a	sacCer3	6,106,953	0.17	1,501,810	52,374	4,121,229	144,839	0.48
2026-04-01-exp260217_MS8_SCGE_no-EU_input	2x150	STAR-2.5.3a	Candida	493,798	0.56	21,378	38	120,653	5	0.14
2026-04-01-exp260217_MS8_SCGE_no-EU_input	2x150	STAR-2.5.3a	sacCer3	493,798	0.36	217,434	3,020	621,279	12,479	0.86

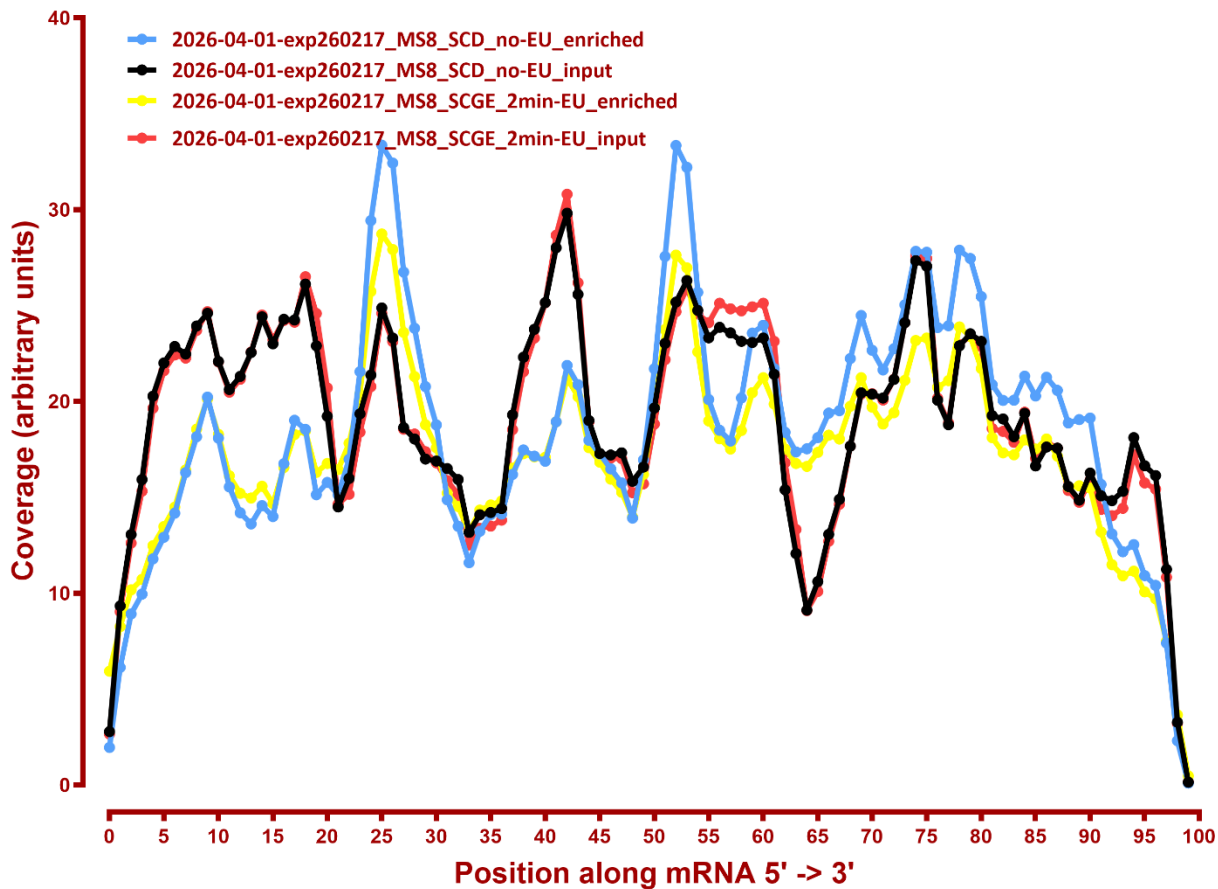
#	Exonic:	Intergenic:	Intronic:
2026-04-01-exp260217_MS8_SCD_2min-EU_enriched	0.92	0.05	0.03
2026-04-01-exp260217_MS8_SCD_2min-EU_input	0.99	0.01	0.00
2026-04-01-exp260217_MS8_SCD_5min-EU_enriched	0.93	0.04	0.02
2026-04-01-exp260217_MS8_SCD_5min-EU_input	0.99	0.01	0.00
2026-04-01-exp260217_MS8_SCD_no-EU_enriched	0.99	0.01	0.00
2026-04-01-exp260217_MS8_SCD_no-EU_input	0.99	0.01	0.00
2026-04-01-exp260217_MS8_SCGE_2min-EU_enriched	0.93	0.05	0.02
2026-04-01-exp260217_MS8_SCGE_2min-EU_input	0.99	0.01	0.00
2026-04-01-exp260217_MS8_SCGE_5min-EU_enriched	0.93	0.05	0.02
2026-04-01-exp260217_MS8_SCGE_5min-EU_input	0.99	0.01	0.00
2026-04-01-exp260217_MS8_SCGE_no-EU_enriched	0.99	0.01	0.00
2026-04-01-exp260217_MS8_SCGE_no-EU_input	0.98	0.01	0.00

COVERAGE

Coverage of genes; >1000bp, sacCer3



Coverage of genes; >1000bp, sacCer3



Coverage of genes; >1000bp, sacCer3

