

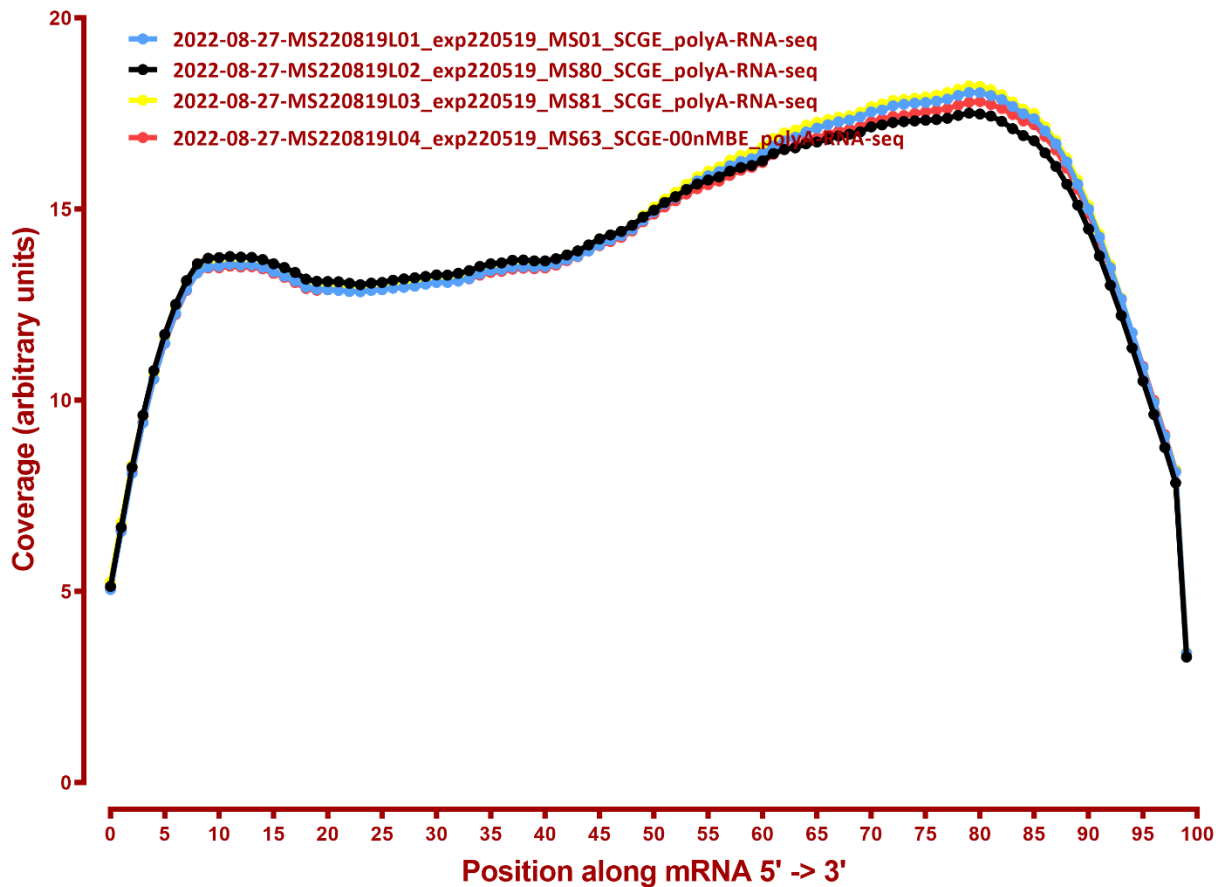
MAPPING STATS

Library	Read Length	Mapping	Species	Raw fragments	Complexity	Unique	Unique Splices	Multi	Multi Splices	Fraction mapped
2022-08-27-MS220819L01_exp220519_MS01_SCGE_polyA-RNA-seq	2x150	STAR-2.5.3a	Candida	6,464,073	0.67	2,680,927	59,343	36,800	2,482	0.22
2022-08-27-MS220819L01_exp220519_MS01_SCGE_polyA-RNA-seq	2x150	STAR-2.5.3a	sacCer3	6,464,073	0.60	8,143,354	204,260	524,978	13,482	0.69
2022-08-27-MS220819L02_exp220519_MS80_SCGE_polyA-RNA-seq	2x150	STAR-2.5.3a	Candida	8,531,681	0.64	3,673,373	80,701	42,591	3,987	0.22
2022-08-27-MS220819L02_exp220519_MS80_SCGE_polyA-RNA-seq	2x150	STAR-2.5.3a	sacCer3	8,531,681	0.57	10,422,922	254,240	651,408	16,480	0.66
2022-08-27-MS220819L03_exp220519_MS81_SCGE_polyA-RNA-seq	2x150	STAR-2.5.3a	Candida	9,109,190	0.65	3,507,164	80,884	40,291	3,463	0.20
2022-08-27-MS220819L03_exp220519_MS81_SCGE_polyA-RNA-seq	2x150	STAR-2.5.3a	sacCer3	9,109,190	0.57	11,605,712	297,524	710,727	23,709	0.69
2022-08-27-MS220819L04_exp220519_MS63_SCGE-00nMBE_polyA-RNA-seq	2x150	STAR-2.5.3a	Candida	9,301,058	0.64	3,951,007	86,539	48,676	4,898	0.22
2022-08-27-MS220819L04_exp220519_MS63_SCGE-00nMBE_polyA-RNA-seq	2x150	STAR-2.5.3a	sacCer3	9,301,058	0.57	11,552,185	277,037	747,348	18,412	0.68
2022-08-27-MS220819L05_exp220519_MS63_SCGE-04nMBE_polyA-RNA-seq	2x150	STAR-2.5.3a	Candida	8,013,346	0.65	3,209,288	72,074	40,347	3,541	0.21
2022-08-27-MS220819L05_exp220519_MS63_SCGE-04nMBE_polyA-RNA-seq	2x150	STAR-2.5.3a	sacCer3	8,013,346	0.58	10,160,396	252,850	632,343	18,129	0.69
2022-08-27-MS220819L06_exp220519_MS63_SCGE-15nMBE_polyA-RNA-seq	2x150	STAR-2.5.3a	Candida	8,862,648	0.65	3,587,102	77,542	44,441	4,575	0.21
2022-08-27-MS220819L06_exp220519_MS63_SCGE-15nMBE_polyA-RNA-seq	2x150	STAR-2.5.3a	sacCer3	8,862,648	0.58	11,349,932	265,156	666,392	17,372	0.69
2022-08-27-MS220819L07_exp220601_MS01_SCGE_polyA-RNA-seq	2x150	STAR-2.5.3a	Candida	9,119,457	0.61	5,247,385	112,763	75,771	6,123	0.30
2022-08-27-MS220819L07_exp220601_MS01_SCGE_polyA-RNA-seq	2x150	STAR-2.5.3a	sacCer3	9,119,457	0.58	10,024,865	242,545	716,762	18,822	0.60
2022-08-27-MS220819L08_exp220601_MS80_SCGE_polyA-RNA-seq	2x150	STAR-2.5.3a	Candida	9,859,082	0.59	5,992,318	127,860	82,761	9,389	0.32
2022-08-27-MS220819L08_exp220601_MS80_SCGE_polyA-RNA-seq	2x150	STAR-2.5.3a	sacCer3	9,859,082	0.56	10,524,804	241,130	749,882	15,678	0.58
2022-08-27-MS220819L09_exp220601_MS81_SCGE_polyA-RNA-seq	2x150	STAR-2.5.3a	Candida	9,142,217	0.62	5,441,082	120,896	63,463	6,127	0.31
2022-08-27-MS220819L09_exp220601_MS81_SCGE_polyA-RNA-seq	2x150	STAR-2.5.3a	sacCer3	9,142,217	0.58	9,887,480	243,338	650,353	19,059	0.59
2022-08-27-MS220819L10_exp220601_MS63_SCGE-00nMBE_polyA-RNA-seq	2x150	STAR-2.5.3a	Candida	8,011,352	0.63	4,713,134	101,284	64,242	5,810	0.30
2022-08-27-MS220819L10_exp220601_MS63_SCGE-00nMBE_polyA-RNA-seq	2x150	STAR-2.5.3a	sacCer3	8,011,352	0.60	8,710,464	213,034	620,944	16,240	0.60
2022-08-27-MS220819L11_exp220601_MS63_SCGE-04nMBE_polyA-RNA-seq	2x150	STAR-2.5.3a	Candida	8,519,693	0.63	4,725,590	106,378	47,812	5,076	0.29
2022-08-27-MS220819L11_exp220601_MS63_SCGE-04nMBE_polyA-RNA-seq	2x150	STAR-2.5.3a	sacCer3	8,519,693	0.58	9,741,804	243,876	572,259	15,487	0.62
2022-08-27-MS220819L12_exp220601_MS63_SCGE-15nMBE_polyA-RNA-seq	2x150	STAR-2.5.3a	Candida	9,368,318	0.62	4,837,533	110,743	54,506	5,874	0.27
2022-08-27-MS220819L12_exp220601_MS63_SCGE-15nMBE_polyA-RNA-seq	2x150	STAR-2.5.3a	sacCer3	9,368,318	0.58	11,084,143	264,789	637,358	17,088	0.64
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2022-08-27-MS220819L13_exp220610_MS01_SCGE_polyA-RNA-seq	2x150	STAR-2.5.3a	sacCer3	9,624,860	0.56	11,265,835	277,753	677,373	18,139	0.64
2022-08-27-MS220819L14_exp220610_MS80_SCGE_polyA-RNA-seq	2x150	STAR-2.5.3a	Candida	8,238,036	0.64	4,162,953	90,663	44,167	4,247	0.26
2022-08-27-MS220819L14_exp220610_MS80_SCGE_polyA-RNA-seq	2x150	STAR-2.5.3a	sacCer3	8,238,036	0.59	9,766,497	235,735	593,361	13,689	0.64
2022-08-27-MS220819L15_exp220610_MS81_SCGE_polyA-RNA-seq	2x150	STAR-2.5.3a	Candida	10,566,739	0.61	4,564,731	102,623	51,713	5,443	0.22
2022-08-27-MS220819L15_exp220610_MS81_SCGE_polyA-RNA-seq	2x150	STAR-2.5.3a	sacCer3	10,566,739	0.54	13,201,408	323,540	813,717	26,453	0.68
2022-08-27-MS220819L16_exp220610_MS63_SCGE-00nMBE_polyA-RNA-seq	2x150	STAR-2.5.3a	Candida	8,172,280	0.63	4,292,052	94,912	53,959	4,457	0.27
2022-08-27-MS220819L16_exp220610_MS63_SCGE-00nMBE_polyA-RNA-seq	2x150	STAR-2.5.3a	sacCer3	8,172,280	0.59	9,486,102	237,588	663,440	16,960	0.64
2022-08-27-MS220819L17_exp220610_MS63_SCGE-3.5nMBE_polyA-RNA-seq	2x150	STAR-2.5.3a	Candida	8,285,411	0.64	3,947,030	89,724	41,291	4,891	0.25
2022-08-27-MS220819L17_exp220610_MS63_SCGE-3.5nMBE_polyA-RNA-seq	2x150	STAR-2.5.3a	sacCer3	8,285,411	0.58	10,042,953	261,469	588,801	18,781	0.66
2022-08-27-MS220819L18_exp220610_MS63_SCGE-16nMBE_polyA-RNA-seq	2x150	STAR-2.5.3a	Candida	8,162,083	0.65	3,802,042	84,982	50,324	3,690	0.24
2022-08-27-MS220819L18_exp220610_MS63_SCGE-16nMBE_polyA-RNA-seq	2x150	STAR-2.5.3a	sacCer3	8,162,083	0.59	9,933,534	233,328	597,663	14,561	0.66
2022-08-27-MS220819L19_exp220607_MS01_SCGE_polyA-RNA-seq	2x150	STAR-2.5.3a	Candida	3,771,175	0.69	1,876,781	41,941	22,637	1,571	0.26
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2022-08-27-MS220819L21_exp220607_MS81_SCGE_polyA-RNA-seq	2x150	STAR-2.5.3a	Candida	8,502,377	0.64	3,800,094	82,694	39,956	4,062	0.23
2022-08-27-MS220819L21_exp220607_MS81_SCGE_polyA-RNA-seq	2x150	STAR-2.5.3a	sacCer3	8,502,377	0.58	10,438,998	257,934	650,107	22,847	0.67
2022-08-27-MS220819L22_exp220607_MS63_SCGE-00nMBE_polyA-RNA-seq	2x150	STAR-2.5.3a	Candida	7,973,137	0.64	3,775,589	84,491	54,990	3,662	0.25
2022-08-27-MS220819L22_exp220607_MS63_SCGE-00nMBE_polyA-RNA-seq	2x150	STAR-2.5.3a	sacCer3	7,973,137	0.59	9,439,830	238,974	682,773	16,339	0.65
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2022-08-27-MS220819L23_exp220607_MS63_SCGE-3.5nMBE_polyA-RNA-seq	2x150	STAR-2.5.3a	sacCer3	8,884,356	0.58	10,766,918	270,426	746,126	20,046	0.66
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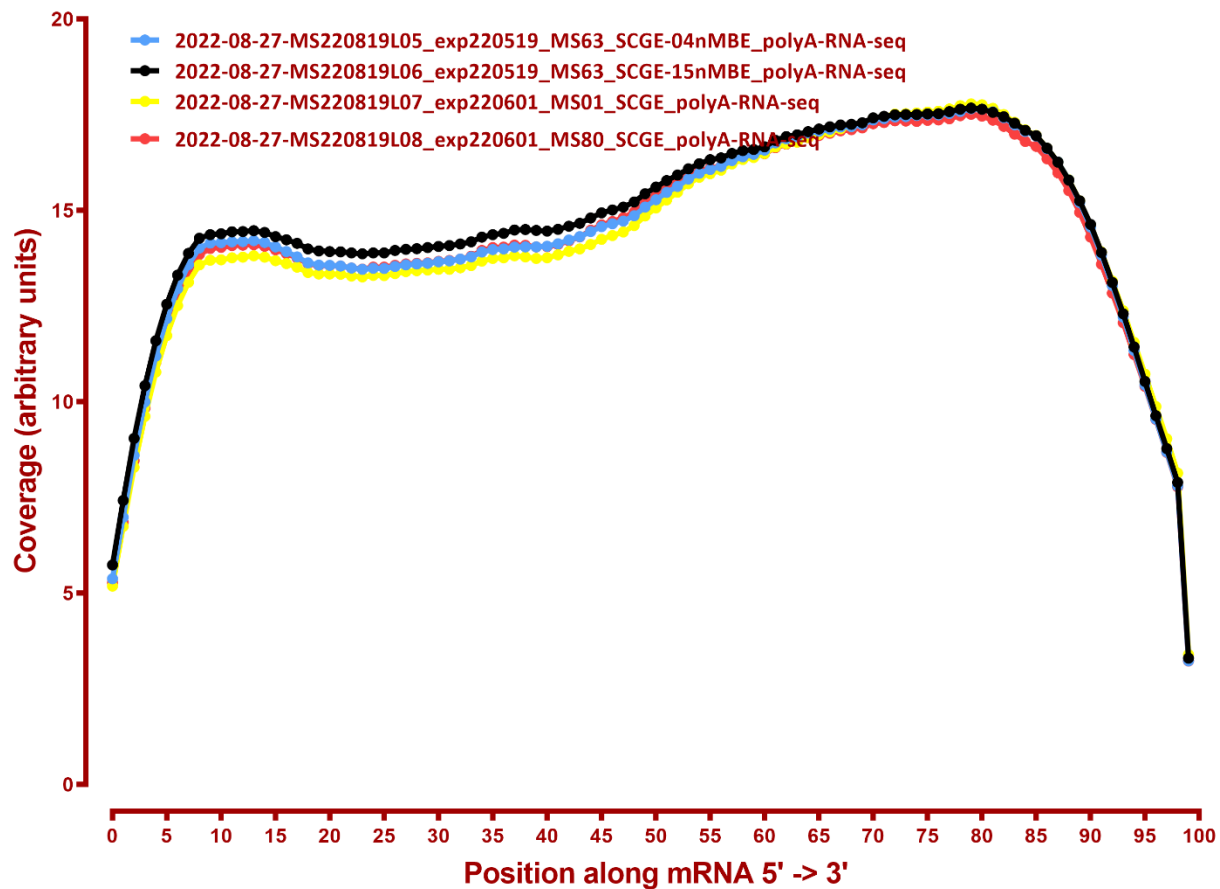
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COVERAGE ALL READS

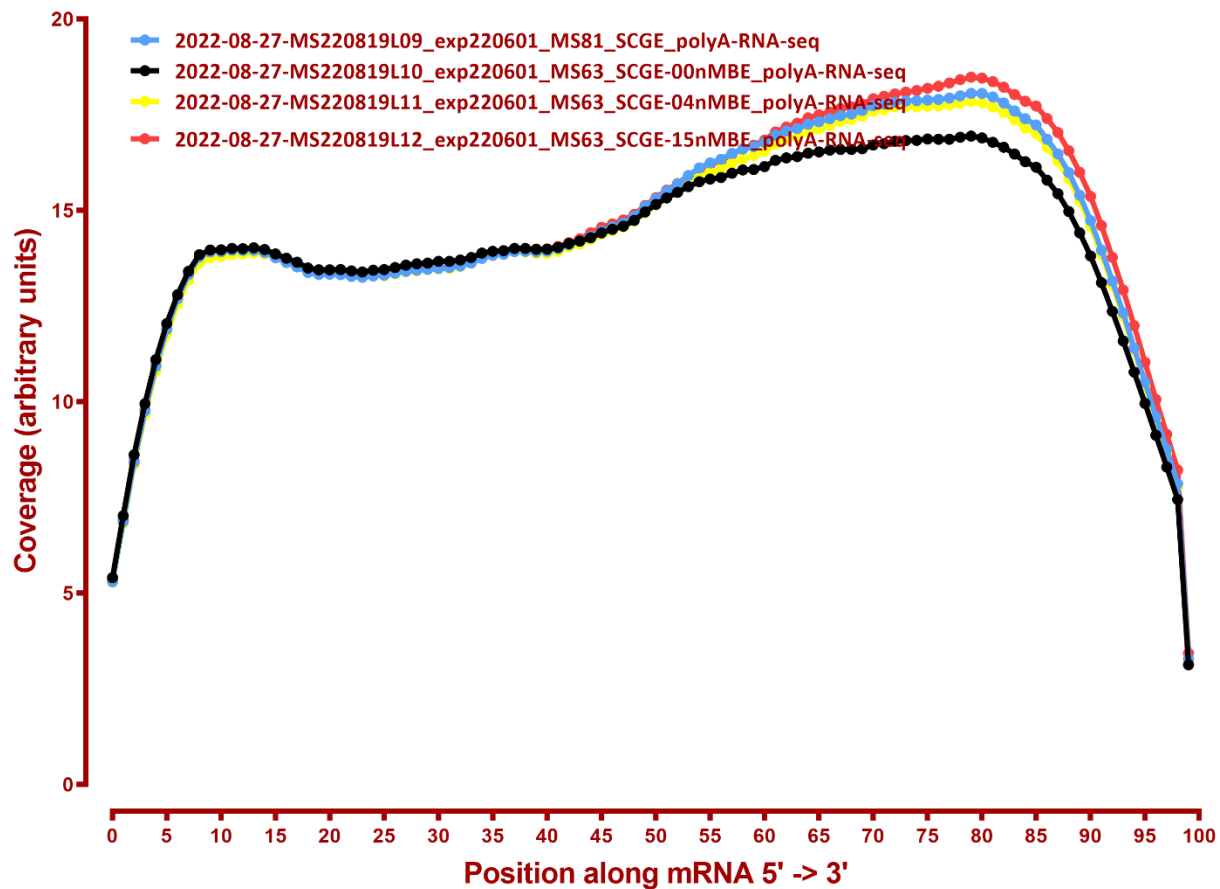
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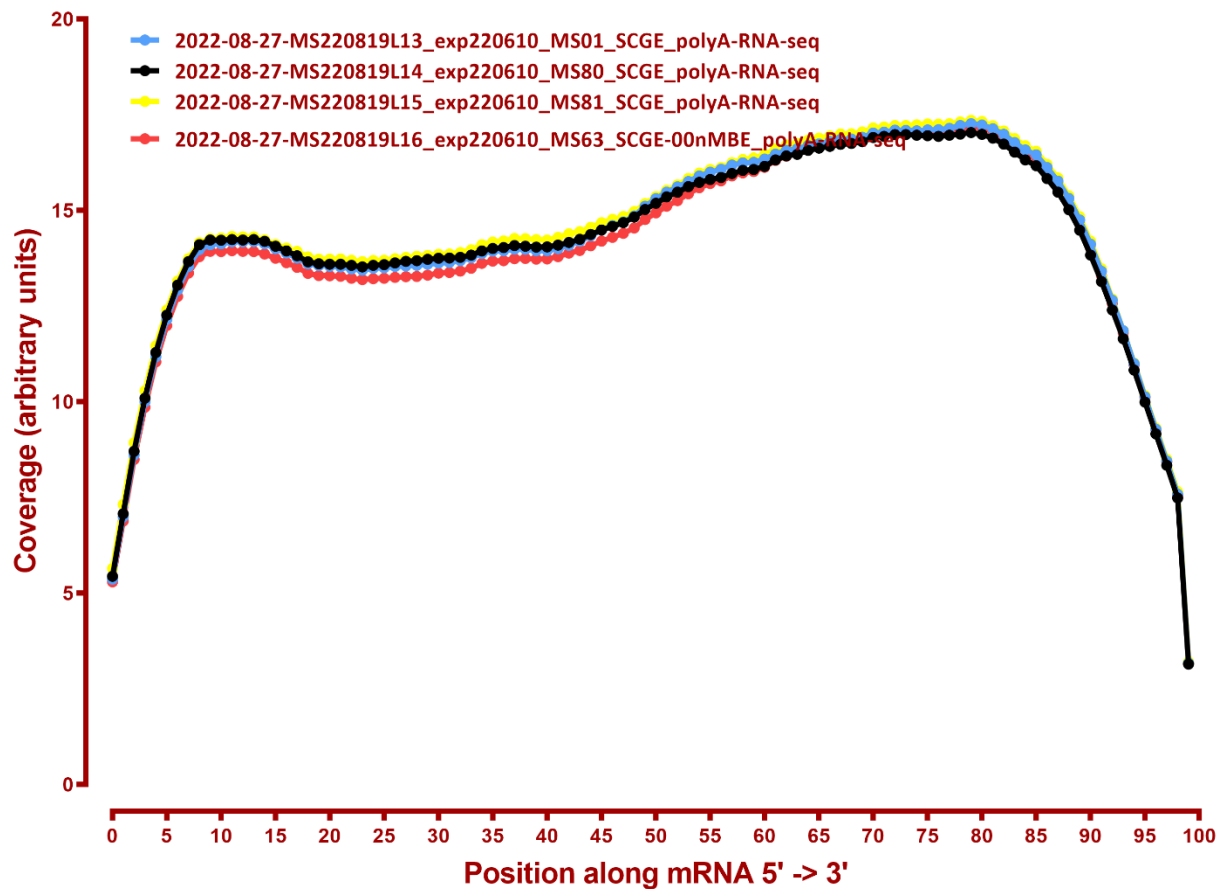
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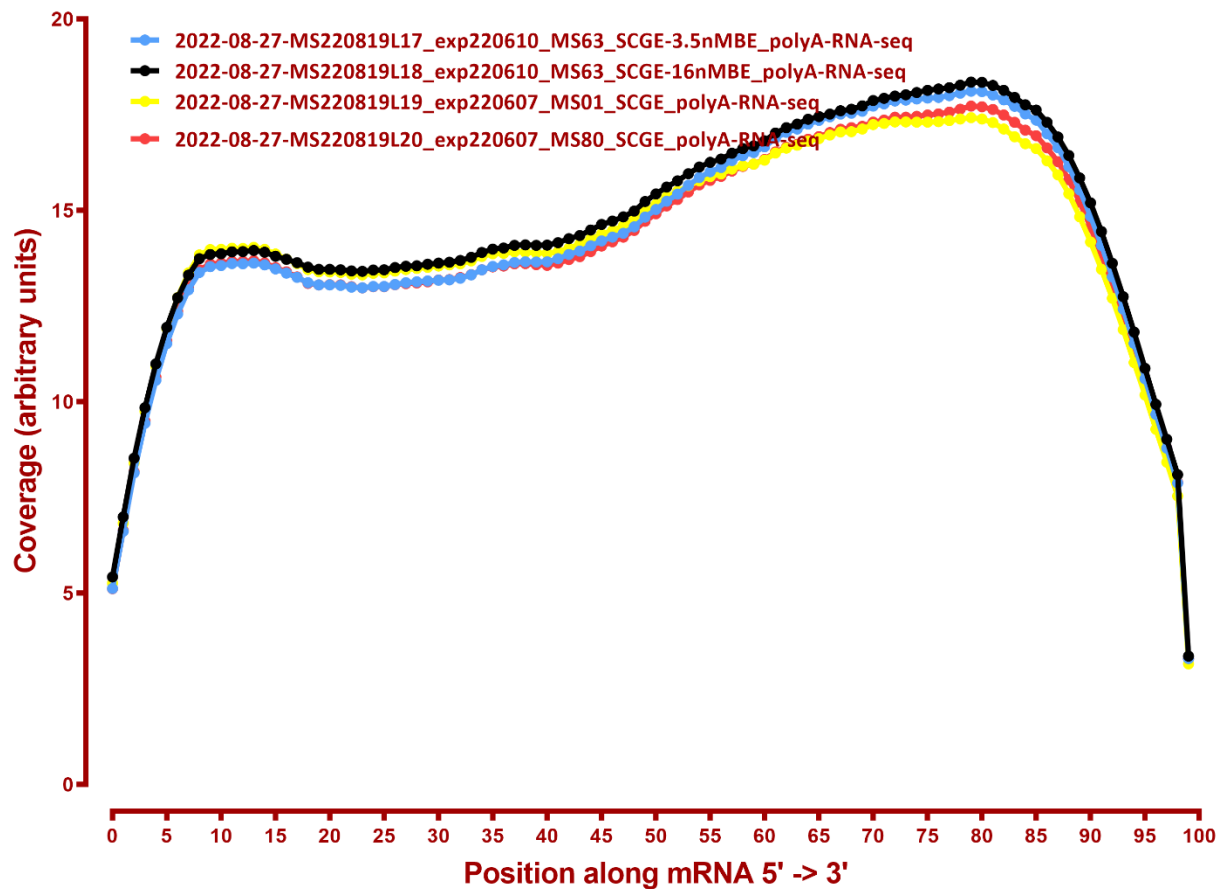
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Coverage of genes; >1000bp, sacCer3



Coverage of genes; >1000bp, sacCer3



Coverage of genes; >1000bp, sacCer3

