

MAPPING STATS

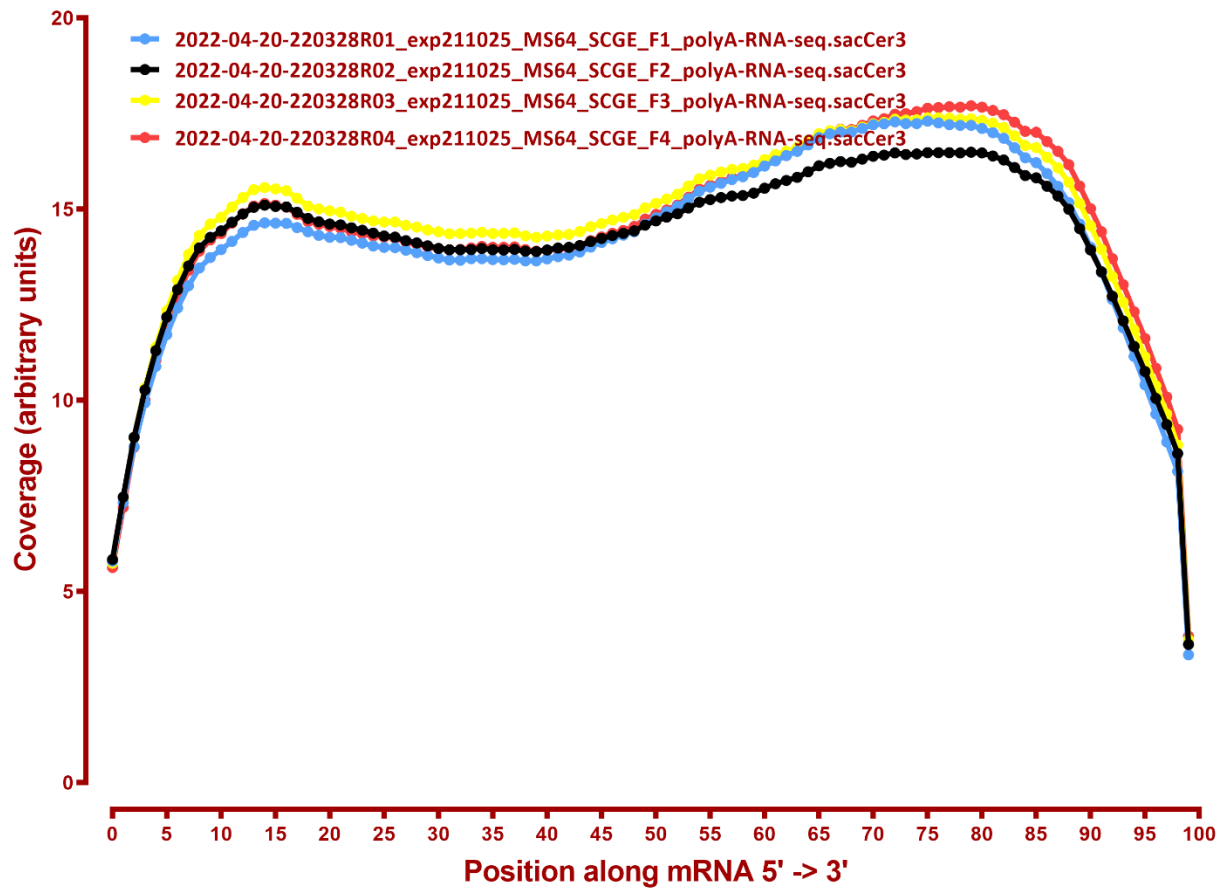
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2022-04-20-220328R01_exp211025_MS64_SCGE_F1_polyA-RNA-seq	2x150	STAR-2.5.3a	Candida	20,160,251	0.37	13,909,350	234,986	127,429	23,051	0.35
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2022-04-20-220328R03_exp211025_MS64_SCGE_F3_polyA-RNA-seq	2x150	STAR-2.5.3a	Candida	17,074,842	0.37	10,336,923	172,369	95,645	13,551	0.31
2022-04-20-220328R03_exp211025_MS64_SCGE_F3_polyA-RNA-seq	2x150	STAR-2.5.3a	sacCer3	17,074,842	0.34	21,018,692	419,164	1,105,587	42,029	0.66
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2022-04-20-220328R05_exp211025_MS64_SCGE_F5_polyA-RNA-seq	2x150	STAR-2.5.3a	Candida	17,240,352	0.38	10,584,839	169,891	93,630	14,458	0.32
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2022-04-20-220328R07_exp211025_MS64_SCGE_F7_polyA-RNA-seq	2x150	STAR-2.5.3a	Candida	18,127,013	0.36	12,560,422	198,888	116,567	17,843	0.36
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2022-04-20-220328R08_exp211021_MS67_SCGE_F1_polyA-RNA-seq	2x150	STAR-2.5.3a	Candida	17,761,738	0.29	11,429,199	154,289	95,623	13,745	0.33
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2022-04-20-220328R10_exp211021_MS67_SCGE_F3_polyA-RNA-seq	2x150	STAR-2.5.3a	Candida	18,207,050	0.37	10,379,721	170,597	87,983	11,159	0.29
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2022-04-20-220328R11_exp211021_MS67_SCGE_F4_polyA-RNA-seq	2x150	STAR-2.5.3a	Candida	17,276,252	0.37	9,281,055	150,923	81,181	12,219	0.28
2022-04-20-220328R11_exp211021_MS67_SCGE_F4_polyA-RNA-seq	2x150	STAR-2.5.3a	sacCer3	17,276,252	0.32	22,499,017	437,881	1,042,066	50,458	0.70
2022-04-20-220328R12_exp211021_MS67_SCGE_F5_polyA-RNA-seq	2x150	STAR-2.5.3a	Candida	17,480,590	0.37	9,783,980	155,428	83,270	13,828	0.29
2022-04-20-220328R12_exp211021_MS67_SCGE_F5_polyA-RNA-seq	2x150	STAR-2.5.3a	sacCer3	17,480,590	0.33	22,514,700	415,270	1,059,814	60,356	0.69
2022-04-20-220328R13_exp211021_MS67_SCGE_F6_polyA-RNA-seq	2x150	STAR-2.5.3a	Candida	16,669,629	0.38	10,025,100	158,392	87,944	12,704	0.31
2022-04-20-220328R13_exp211021_MS67_SCGE_F6_polyA-RNA-seq	2x150	STAR-2.5.3a	sacCer3	16,669,629	0.35	20,673,399	391,775	1,021,526	52,954	0.66
2022-04-20-220328R14_exp211021_MS67_SCGE_F7_polyA-RNA-seq	2x150	STAR-2.5.3a	Candida	19,224,346	0.37	11,340,599	194,895	100,759	15,749	0.30
2022-04-20-220328R14_exp211021_MS67_SCGE_F7_polyA-RNA-seq	2x150	STAR-2.5.3a	sacCer3	19,224,346	0.33	24,053,526	475,176	1,262,228	51,442	0.67

2022-04-20-220328R15_exp211028_MS64_SCGE_F1_polyA-RNA-seq	2x150	STAR-2.5.3a	Candida	19,455,026	0.34	13,855,720	219,576	119,240	17,436	0.37
2022-04-20-220328R15_exp211028_MS64_SCGE_F1_polyA-RNA-seq	2x150	STAR-2.5.3a	sacCer3	19,455,026	0.32	21,929,675	502,921	1,092,536	47,368	0.61
2022-04-20-220328R16_exp211028_MS64_SCGE_F2_polyA-RNA-seq	2x150	STAR-2.5.3a	Candida	18,173,472	0.39	11,956,371	183,033	103,489	19,787	0.34
2022-04-20-220328R16_exp211028_MS64_SCGE_F2_polyA-RNA-seq	2x150	STAR-2.5.3a	sacCer3	18,173,472	0.36	21,405,333	448,891	1,018,846	43,036	0.63
2022-04-20-220328R17_exp211028_MS64_SCGE_F3_polyA-RNA-seq	2x150	STAR-2.5.3a	Candida	16,506,488	0.41	10,247,144	151,734	85,816	13,242	0.32
2022-04-20-220328R17_exp211028_MS64_SCGE_F3_polyA-RNA-seq	2x150	STAR-2.5.3a	sacCer3	16,506,488	0.37	20,199,077	356,569	993,266	41,308	0.65
2022-04-20-220328R18_exp211028_MS64_SCGE_F4_polyA-RNA-seq	2x150	STAR-2.5.3a	Candida	18,254,554	0.39	11,688,456	170,952	103,509	13,923	0.33
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2022-04-20-220328R21_exp211028_MS64_SCGE_F7_polyA-RNA-seq	2x150	STAR-2.5.3a	Candida	19,011,639	0.39	13,098,770	199,980	130,289	18,251	0.35
2022-04-20-220328R21_exp211028_MS64_SCGE_F7_polyA-RNA-seq	2x150	STAR-2.5.3a	sacCer3	19,011,639	0.36	21,261,304	369,330	1,315,179	47,145	0.60
2022-04-20-220328R22_exp211101_MS67_SCGE_F1_polyA-RNA-seq	2x150	STAR-2.5.3a	Candida	17,741,041	0.44	10,492,167	162,763	97,499	13,799	0.30
2022-04-20-220328R22_exp211101_MS67_SCGE_F1_polyA-RNA-seq	2x150	STAR-2.5.3a	sacCer3	17,741,041	0.39	22,199,792	482,064	1,062,274	45,152	0.67
2022-04-20-220328R23_exp211101_MS67_SCGE_F2_polyA-RNA-seq	2x150	STAR-2.5.3a	Candida	18,391,156	0.37	10,301,505	162,937	97,757	12,459	0.29
2022-04-20-220328R23_exp211101_MS67_SCGE_F2_polyA-RNA-seq	2x150	STAR-2.5.3a	sacCer3	18,391,156	0.32	23,586,689	504,199	1,223,157	52,307	0.69
2022-04-20-220328R24_exp211101_MS67_SCGE_F3_polyA-RNA-seq	2x150	STAR-2.5.3a	Candida	18,222,460	0.28	10,467,645	142,175	84,571	14,253	0.29
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2022-04-20-220328R27_exp211101_MS67_SCGE_F6_polyA-RNA-seq	2x150	STAR-2.5.3a	Candida	16,998,797	0.41	9,764,070	152,290	88,056	11,410	0.29
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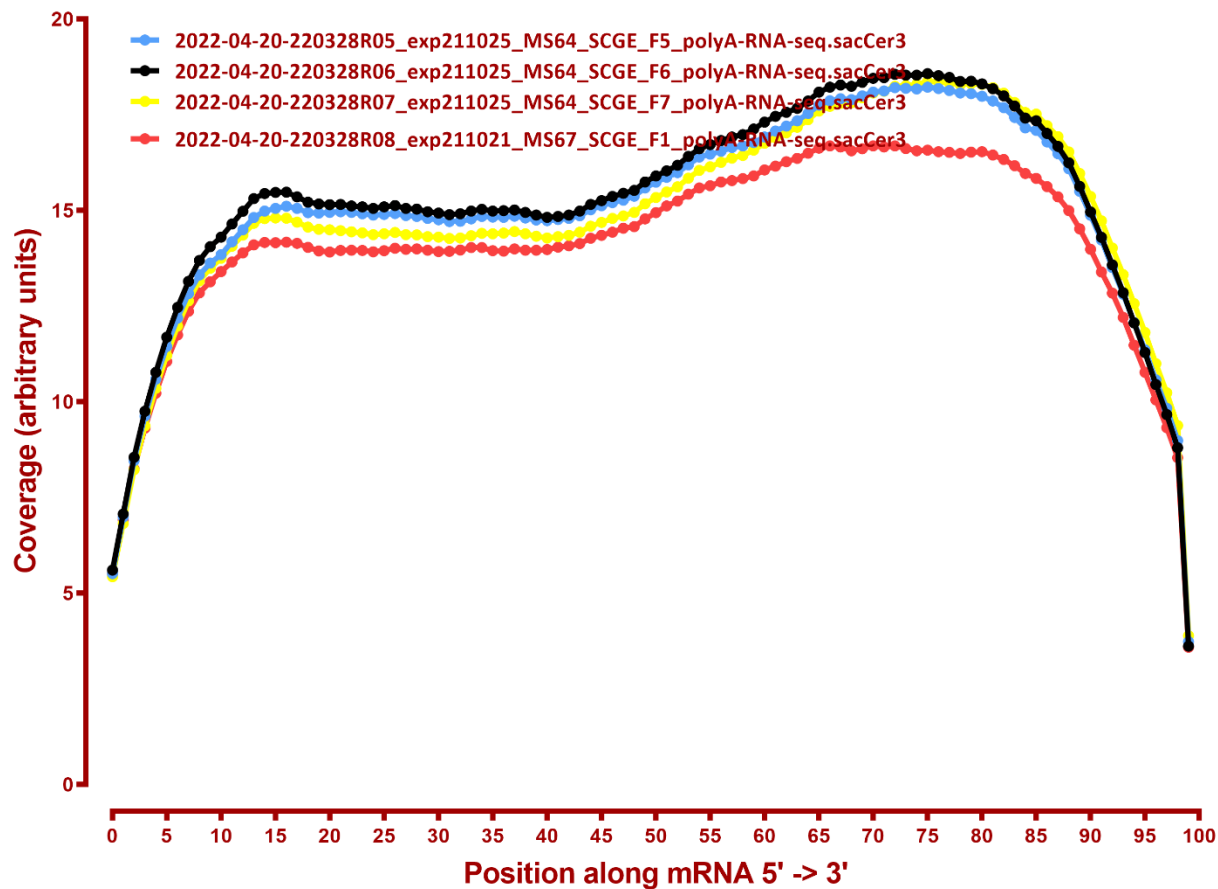
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COVERAGE ALL READS

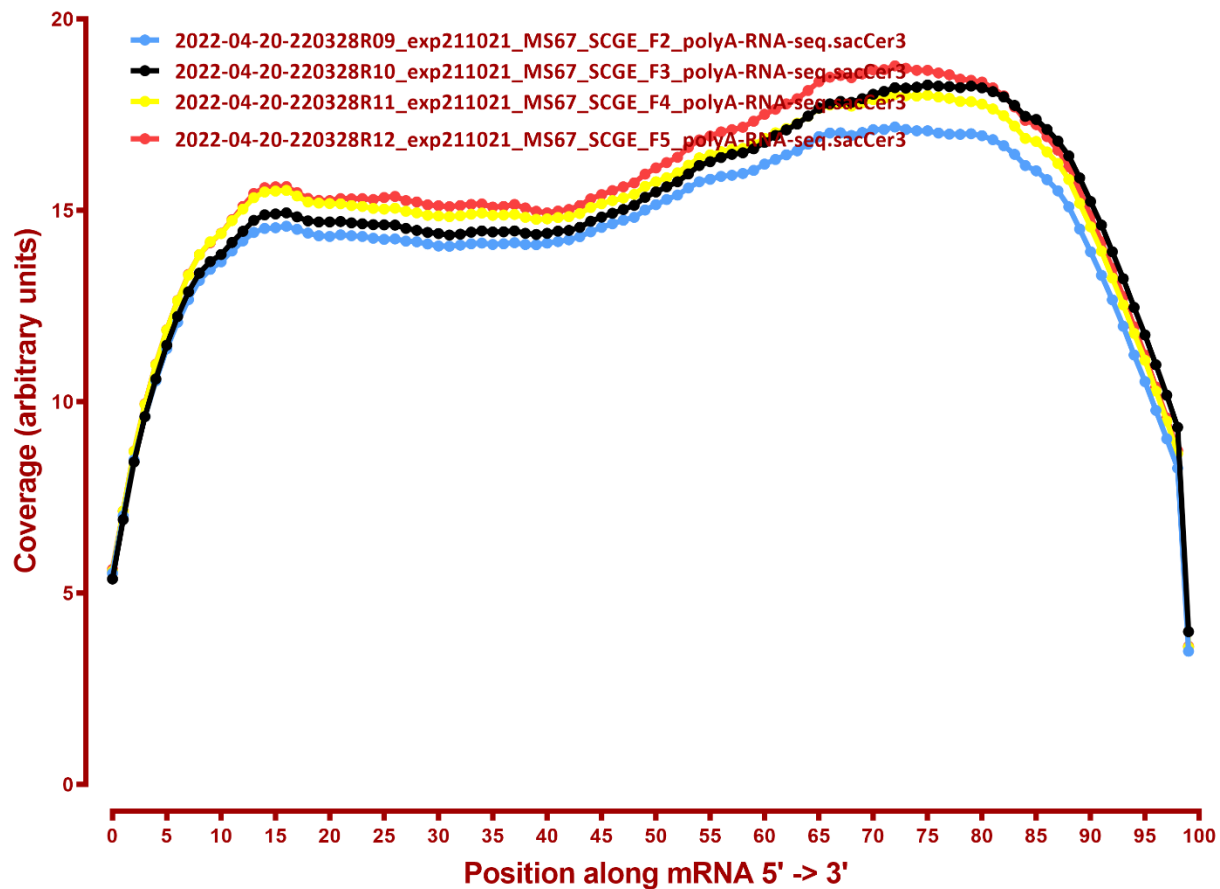
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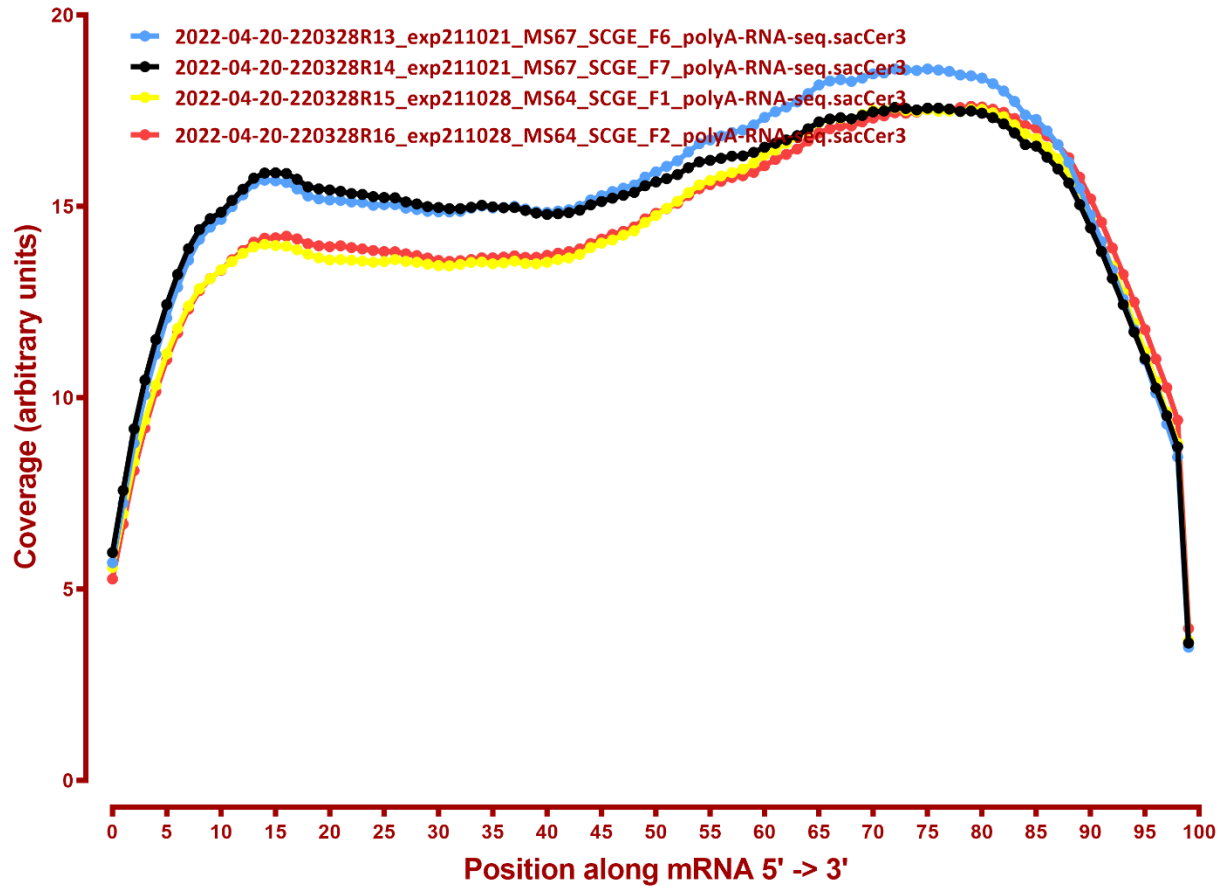
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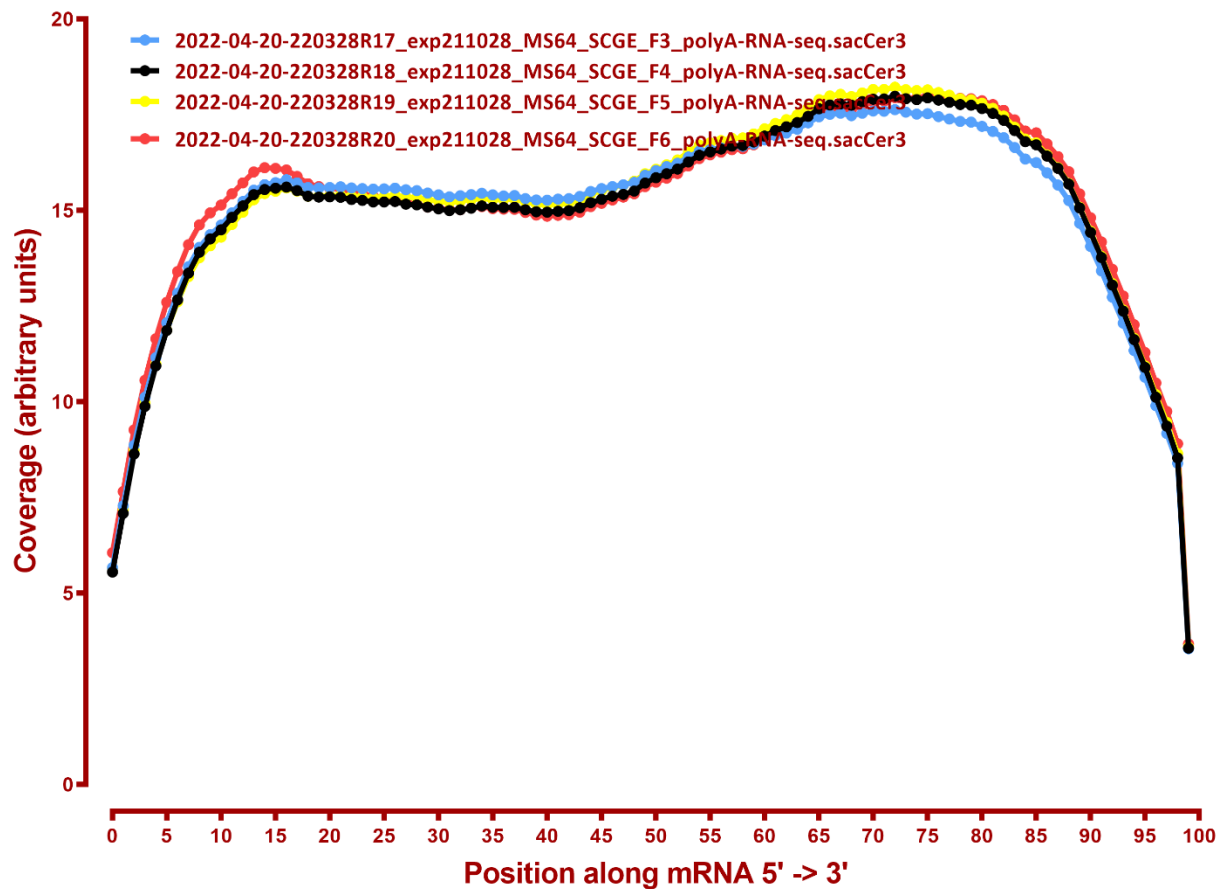
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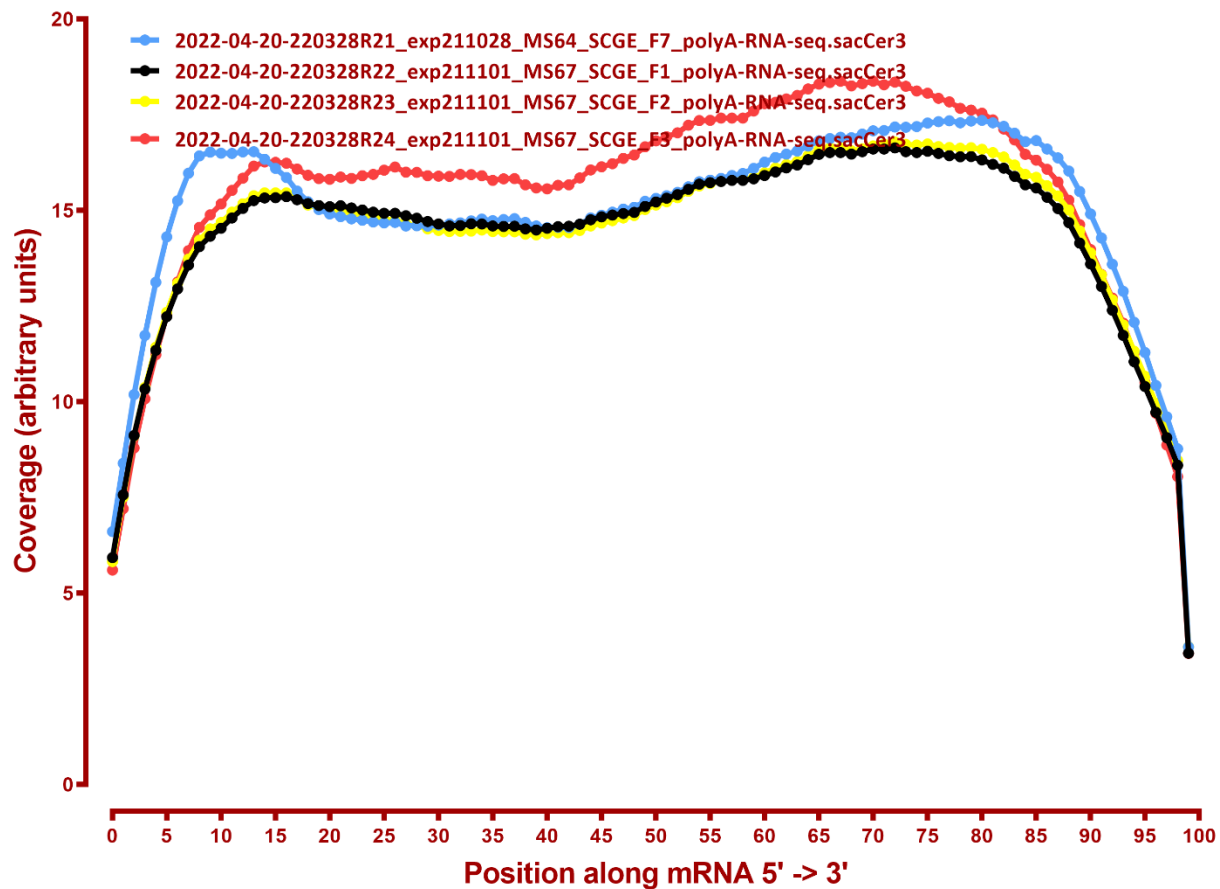
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Coverage of genes; >1000bp, sacCer3

