

MAPPING STATS

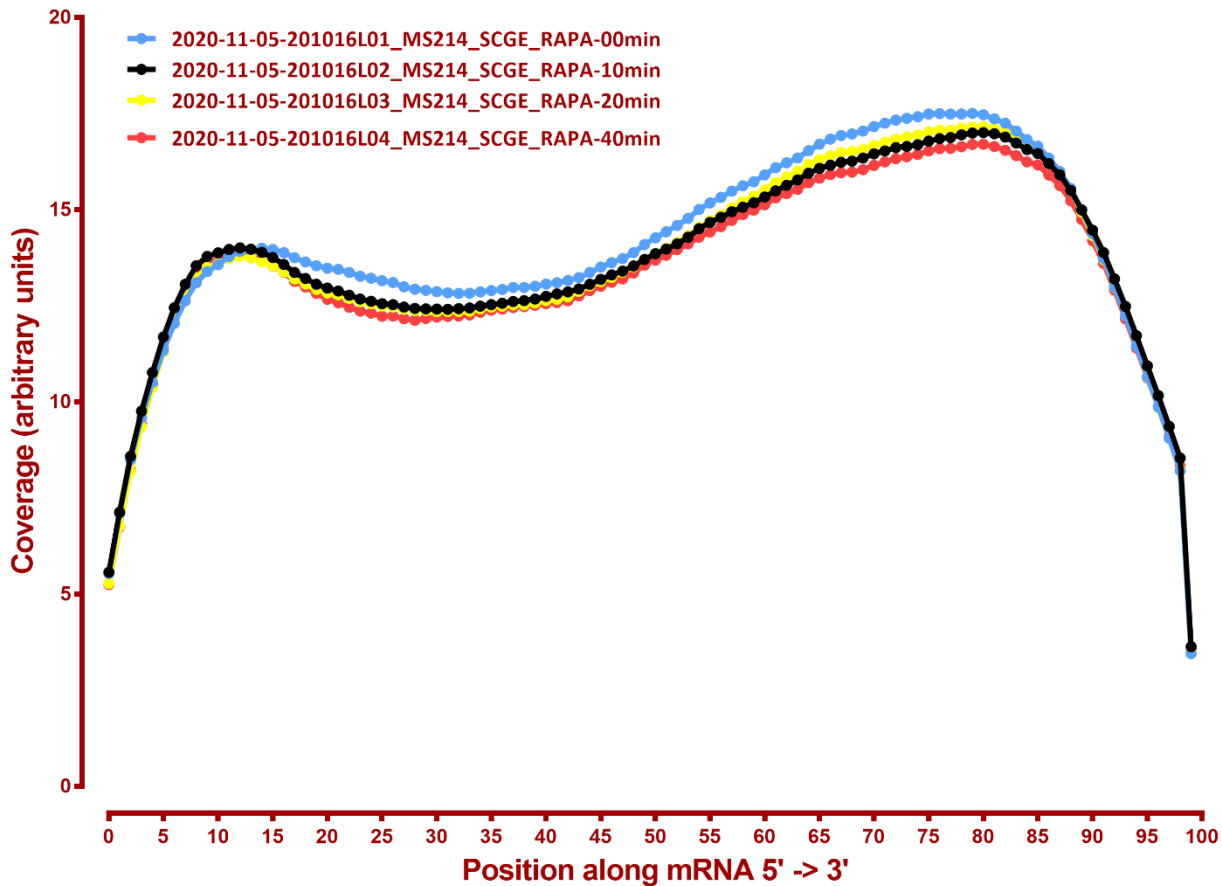
Library	Read Length	Mapping	Species	Raw fragments	Complexity	Unique	Unique Splices	Multi	Multi Splices	Fraction mapped
2019-12-03-RNAseq-exp191107-A040	2x150	STAR-2.5.3a	S_pombe	26,572,372	1.00	8	0	188	0	0.00
2019-12-03-RNAseq-exp191107-A040	2x150	STAR-2.5.3a	sacCer3	26,572,372	0.47	38,429,226	1,971,036	2,895,440	227,550	0.82
2019-12-03-RNAseq-exp191107-A050	2x150	STAR-2.5.3a	S_pombe	29,977,052	1.00	4	2	326	12	0.00
2019-12-03-RNAseq-exp191107-A050	2x150	STAR-2.5.3a	sacCer3	29,977,052	0.47	42,758,408	2,154,564	3,080,921	277,033	0.81
2019-12-03-RNAseq-exp191107-A060	2x150	STAR-2.5.3a	S_pombe	25,736,950	N/A	0	0	156	0	0.00
2019-12-03-RNAseq-exp191107-A060	2x150	STAR-2.5.3a	sacCer3	25,736,950	0.47	36,559,872	1,764,448	2,339,496	196,222	0.79
2019-12-03-RNAseq-exp191107-A070	2x150	STAR-2.5.3a	S_pombe	25,520,705	1.00	10	0	237	1	0.00
2019-12-03-RNAseq-exp191107-A070	2x150	STAR-2.5.3a	sacCer3	25,520,705	0.48	35,948,209	1,642,917	2,263,269	178,011	0.78
2020-11-05-201016L01_MS214_SCGE_RAPA-00min	2x150	STAR-2.5.3a	S_pombe	13,190,932	0.45	13,824,654	1,112,786	236,847	47,541	0.58
2020-11-05-201016L01_MS214_SCGE_RAPA-00min	2x150	STAR-2.5.3a	sacCer3	13,190,932	0.50	9,861,162	226,038	413,074	14,758	0.40
2020-11-05-201016L02_MS214_SCGE_RAPA-10min	2x150	STAR-2.5.3a	S_pombe	12,736,218	0.44	13,449,286	1,132,370	233,823	42,207	0.58
2020-11-05-201016L02_MS214_SCGE_RAPA-10min	2x150	STAR-2.5.3a	sacCer3	12,736,218	0.49	9,187,565	219,017	394,283	15,045	0.39
2020-11-05-201016L03_MS214_SCGE_RAPA-20min	2x150	STAR-2.5.3a	S_pombe	12,251,772	0.46	13,753,123	1,178,797	901,265	47,243	0.65
2020-11-05-201016L03_MS214_SCGE_RAPA-20min	2x150	STAR-2.5.3a	sacCer3	12,251,772	0.53	7,188,106	177,618	701,577	11,021	0.33
2020-11-05-201016L04_MS214_SCGE_RAPA-40min	2x150	STAR-2.5.3a	S_pombe	12,846,795	0.42	18,428,877	1,573,123	428,641	62,977	0.80
2020-11-05-201016L04_MS214_SCGE_RAPA-40min	2x150	STAR-2.5.3a	sacCer3	12,846,795	0.53	4,135,419	105,859	247,797	6,321	0.17
2020-11-05-201016L05_MS214_SCGE_RAPA-60min	2x150	STAR-2.5.3a	S_pombe	12,590,845	0.42	20,568,482	1,723,074	356,265	74,121	0.90
2020-11-05-201016L05_MS214_SCGE_RAPA-60min	2x150	STAR-2.5.3a	sacCer3	12,590,845	0.60	1,766,272	45,058	85,979	3,427	0.08
2020-11-05-201016L06_MS214_SCGE_RAPA-90min	2x150	STAR-2.5.3a	S_pombe	12,415,604	0.40	20,869,730	1,755,384	370,379	74,045	0.93
2020-11-05-201016L06_MS214_SCGE_RAPA-90min	2x150	STAR-2.5.3a	sacCer3	12,415,604	0.61	1,103,740	21,628	60,140	2,456	0.05
2020-11-05-201016L07_MS215_SCGE_RAPA-00min	2x150	STAR-2.5.3a	S_pombe	13,133,844	0.47	12,569,537	1,093,373	229,736	38,908	0.53
2020-11-05-201016L07_MS215_SCGE_RAPA-00min	2x150	STAR-2.5.3a	sacCer3	13,133,844	0.51	10,827,830	249,888	494,580	18,666	0.44
2020-11-05-201016L08_MS215_SCGE_RAPA-10min	2x150	STAR-2.5.3a	S_pombe	12,064,750	0.47	12,146,805	1,035,921	212,650	42,156	0.56
2020-11-05-201016L08_MS215_SCGE_RAPA-10min	2x150	STAR-2.5.3a	sacCer3	12,064,750	0.51	9,477,428	233,252	425,826	17,222	0.42
2020-11-05-201016L09_MS215_SCGE_RAPA-20min	2x150	STAR-2.5.3a	S_pombe	11,854,790	0.47	13,238,593	1,130,929	375,911	43,171	0.62
2020-11-05-201016L09_MS215_SCGE_RAPA-20min	2x150	STAR-2.5.3a	sacCer3	11,854,790	0.53	7,488,702	186,736	449,645	13,085	0.34
2020-11-05-201016L10_MS215_SCGE_RAPA-40min	2x150	STAR-2.5.3a	S_pombe	12,315,494	0.44	16,900,211	1,571,107	288,253	62,511	0.76
2020-11-05-201016L10_MS215_SCGE_RAPA-40min	2x150	STAR-2.5.3a	sacCer3	12,315,494	0.52	4,810,676	124,416	243,271	11,063	0.21
2020-11-05-201016L11_MS215_SCGE_RAPA-60min	2x150	STAR-2.5.3a	S_pombe	13,900,257	0.40	22,328,402	1,933,338	397,233	88,217	0.89
2020-11-05-201016L11_MS215_SCGE_RAPA-60min	2x150	STAR-2.5.3a	sacCer3	13,900,257	0.55	2,249,998	49,818	135,666	7,724	0.09
2020-11-05-201016L12_MS215_SCGE_RAPA-90min	2x150	STAR-2.5.3a	S_pombe	12,762,261	0.41	21,279,851	1,772,693	374,776	83,646	0.92
2020-11-05-201016L12_MS215_SCGE_RAPA-90min	2x150	STAR-2.5.3a	sacCer3	12,762,261	0.61	1,326,439	17,495	76,088	5,252	0.06

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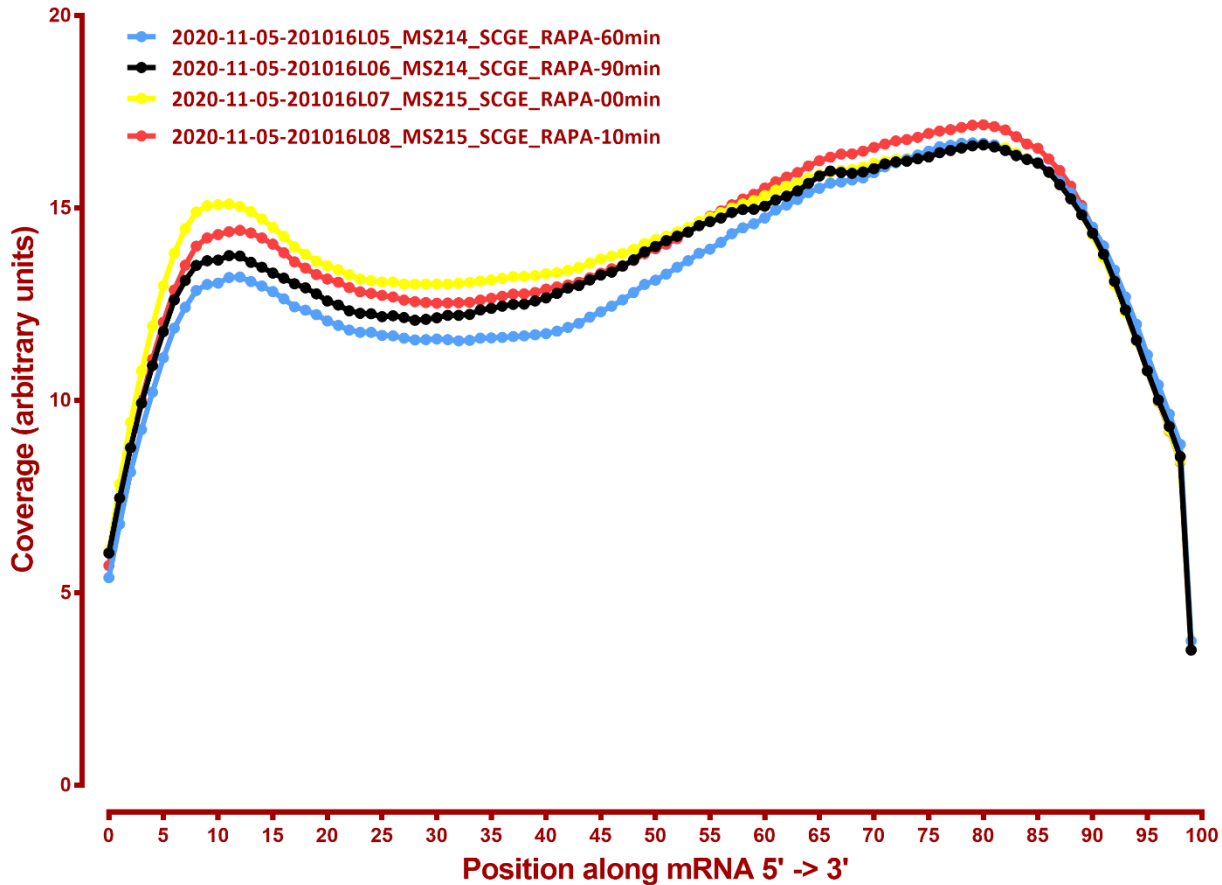
#	Exonic:	Intergenic:	Intronic:
2020-11-05-201016L01_MS214_SCGE_RAPA-00min.Schizosaccharomyces_pombe.ASM294v2.20	0.99	0.01	0.00
2020-11-05-201016L01_MS214_SCGE_RAPA-00min.sacCer3	0.91	0.09	0.00
2020-11-05-201016L02_MS214_SCGE_RAPA-10min.Schizosaccharomyces_pombe.ASM294v2.20	0.99	0.01	0.00
2020-11-05-201016L02_MS214_SCGE_RAPA-10min.sacCer3	0.91	0.09	0.00
2020-11-05-201016L03_MS214_SCGE_RAPA-20min.Schizosaccharomyces_pombe.ASM294v2.20	0.99	0.01	0.00
2020-11-05-201016L03_MS214_SCGE_RAPA-20min.sacCer3	0.91	0.09	0.00
2020-11-05-201016L04_MS214_SCGE_RAPA-40min.Schizosaccharomyces_pombe.ASM294v2.20	0.99	0.01	0.00
2020-11-05-201016L04_MS214_SCGE_RAPA-40min.sacCer3	0.91	0.09	0.00
2020-11-05-201016L05_MS214_SCGE_RAPA-60min.Schizosaccharomyces_pombe.ASM294v2.20	0.99	0.01	0.00
2020-11-05-201016L05_MS214_SCGE_RAPA-60min.sacCer3	0.90	0.10	0.00
2020-11-05-201016L06_MS214_SCGE_RAPA-90min.Schizosaccharomyces_pombe.ASM294v2.20	0.99	0.01	0.00
2020-11-05-201016L06_MS214_SCGE_RAPA-90min.sacCer3	0.90	0.10	0.00
2020-11-05-201016L07_MS215_SCGE_RAPA-00min.Schizosaccharomyces_pombe.ASM294v2.20	0.99	0.01	0.00
2020-11-05-201016L07_MS215_SCGE_RAPA-00min.sacCer3	0.91	0.09	0.00
2020-11-05-201016L08_MS215_SCGE_RAPA-10min.Schizosaccharomyces_pombe.ASM294v2.20	0.99	0.01	0.00
2020-11-05-201016L08_MS215_SCGE_RAPA-10min.sacCer3	0.91	0.09	0.00
2020-11-05-201016L09_MS215_SCGE_RAPA-20min.Schizosaccharomyces_pombe.ASM294v2.20	0.99	0.01	0.00
2020-11-05-201016L09_MS215_SCGE_RAPA-20min.sacCer3	0.91	0.09	0.00
2020-11-05-201016L10_MS215_SCGE_RAPA-40min.Schizosaccharomyces_pombe.ASM294v2.20	0.99	0.01	0.00
2020-11-05-201016L10_MS215_SCGE_RAPA-40min.sacCer3	0.90	0.09	0.00
2020-11-05-201016L11_MS215_SCGE_RAPA-60min.Schizosaccharomyces_pombe.ASM294v2.20	0.99	0.01	0.00
2020-11-05-201016L11_MS215_SCGE_RAPA-60min.sacCer3	0.90	0.10	0.00
2020-11-05-201016L12_MS215_SCGE_RAPA-90min.Schizosaccharomyces_pombe.ASM294v2.20	0.99	0.01	0.00
2020-11-05-201016L12_MS215_SCGE_RAPA-90min.sacCer3	0.90	0.10	0.00

COVERAGE QC

Coverage of genes; >1000bp, sacCer3



Coverage of genes; >1000bp, sacCer3



Coverage of genes; >1000bp, sacCer3

