

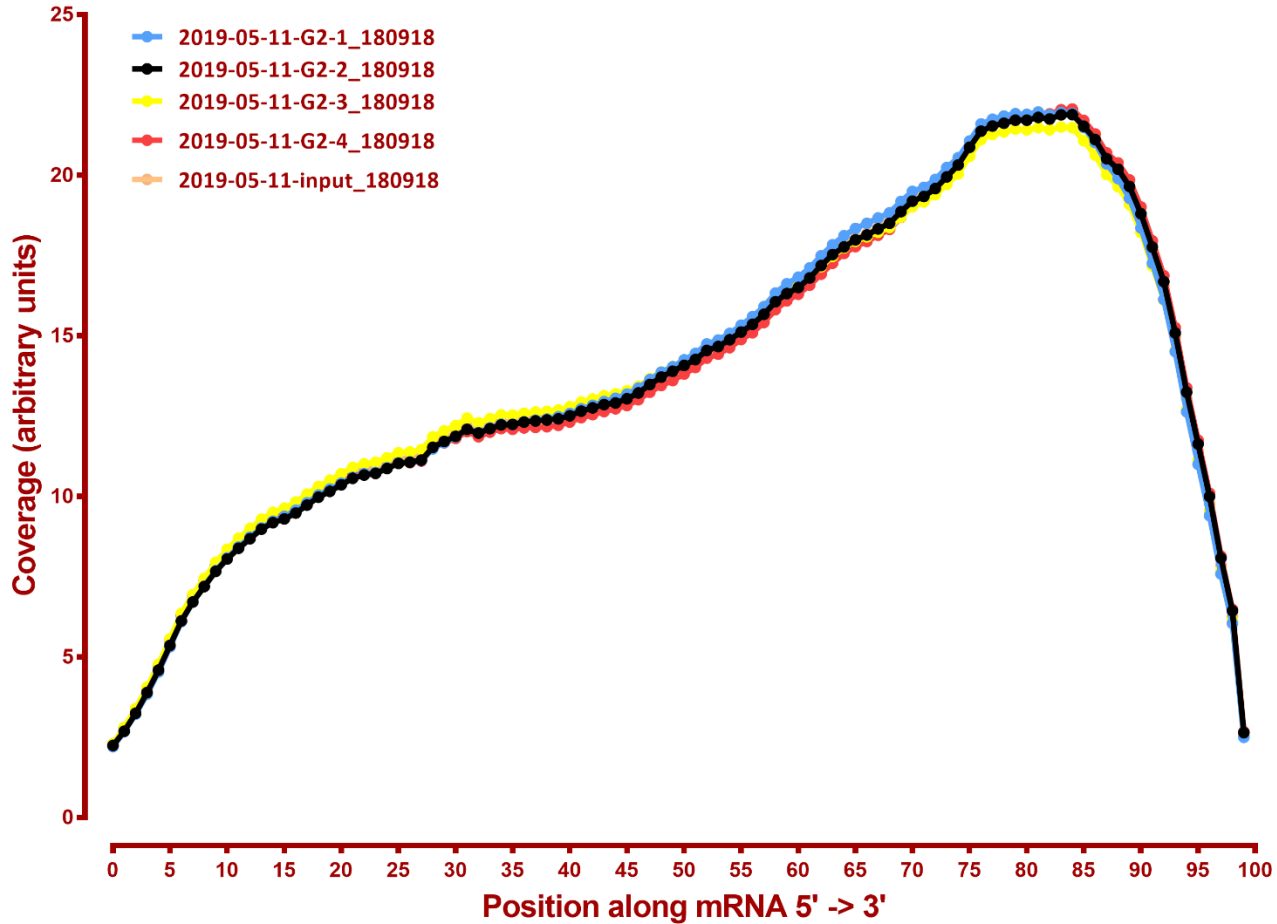
MAPPING STATS

Library	Read Length	Mapping	Species	Raw fragments	Complexity	Unique	Unique Splices	Multi	Multi Splices	Fraction mapped
2019-05-11-G2-1_180918	2x150	STAR-2.5.3a	S_pombe	39,181,969	0.19	15,142,745	1,171,441	283,926	49,030	0.21
2019-05-11-G2-1_180918	2x150	STAR-2.5.3a	sacCer3	39,181,969	0.11	54,775,326	1,478,490	2,553,231	152,519	0.75
2019-05-11-G2-2_180918	2x150	STAR-2.5.3a	S_pombe	41,029,542	0.23	14,505,185	1,106,047	318,696	54,504	0.19
2019-05-11-G2-2_180918	2x150	STAR-2.5.3a	sacCer3	41,029,542	0.13	58,401,527	1,503,025	3,149,043	171,735	0.77
2019-05-11-G2-3_180918	2x150	STAR-2.5.3a	S_pombe	40,151,563	0.24	12,724,183	970,809	268,524	43,162	0.17
2019-05-11-G2-3_180918	2x150	STAR-2.5.3a	sacCer3	40,151,563	0.13	58,694,777	1,444,187	3,380,154	175,736	0.79
2019-05-11-G2-4_180918	2x150	STAR-2.5.3a	S_pombe	43,957,042	0.23	12,213,126	954,098	253,060	40,210	0.15
2019-05-11-G2-4_180918	2x150	STAR-2.5.3a	sacCer3	43,957,042	0.12	65,937,322	1,752,956	3,779,366	222,130	0.82
2019-05-11-input_180918	2x150	STAR-2.5.3a	S_pombe	42,278,912	0.25	10,374,372	787,598	200,413	29,439	0.13
2019-05-11-input_180918	2x150	STAR-2.5.3a	sacCer3	42,278,912	0.12	64,884,206	1,760,054	3,433,700	215,180	0.83

READ DISTRIBUTION

#	Exonic:	Intergenic:	Intronic:
2019-05-11-G2-1_180918.Schizosaccharomyces_pombe.ASM294v2.20	0.98	0.02	0.00
2019-05-11-G2-1_180918.sacCer3	0.90	0.10	0.00
2019-05-11-G2-2_180918.Schizosaccharomyces_pombe.ASM294v2.20	0.98	0.02	0.00
2019-05-11-G2-2_180918.sacCer3	0.90	0.10	0.00
2019-05-11-G2-3_180918.Schizosaccharomyces_pombe.ASM294v2.20	0.98	0.02	0.00
2019-05-11-G2-3_180918.sacCer3	0.90	0.10	0.00
2019-05-11-G2-4_180918.Schizosaccharomyces_pombe.ASM294v2.20	0.98	0.02	0.00
2019-05-11-G2-4_180918.sacCer3	0.89	0.10	0.00
2019-05-11-input_180918.Schizosaccharomyces_pombe.ASM294v2.20	0.98	0.02	0.00
2019-05-11-input_180918.sacCer3	0.90	0.10	0.00

Coverage of genes; >1000bp, sacCer3



Coverage of genes; >1000bp, *S.pombe*

