

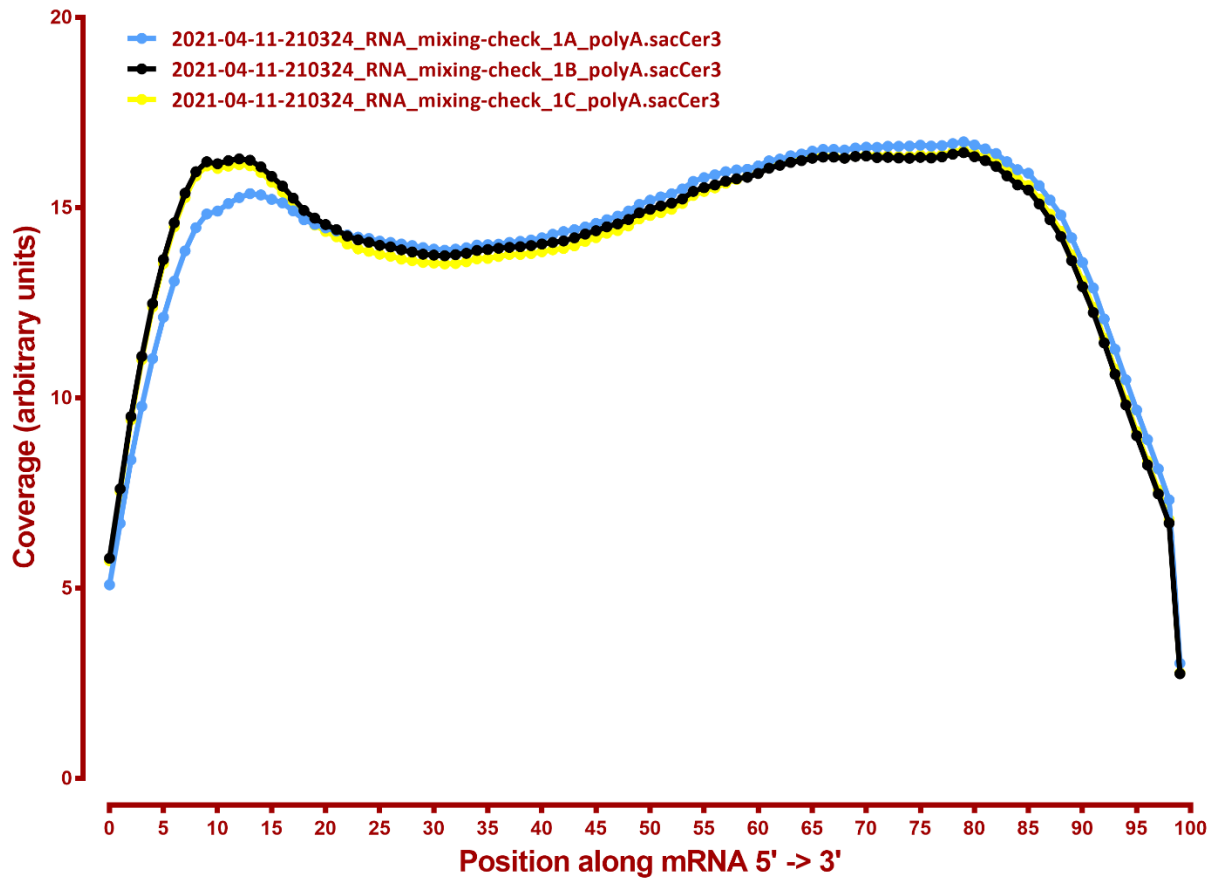
MAPPING STATS

Library	Read Length	Mapping	Species	Raw fragments	Complexity	Unique	Unique Splices	Multi	Multi Splices	Fraction mapped
2021-04-11-210324_RNA_mixing-check_1A_polyA	2x150	STAR-2.5.3a	Candida	7,574,561	0.52	5,773,295	114,871	141,942	7,616	0.40
2021-04-11-210324_RNA_mixing-check_1A_polyA	2x150	STAR-2.5.3a	sacCer3	7,574,561	0.52	7,656,923	188,673	849,942	13,842	0.57
2021-04-11-210324_RNA_mixing-check_1B_polyA	2x150	STAR-2.5.3a	Candida	6,419,037	0.58	3,107,798	69,644	78,492	3,866	0.25
2021-04-11-210324_RNA_mixing-check_1B_polyA	2x150	STAR-2.5.3a	sacCer3	6,419,037	0.53	8,295,747	223,049	725,893	13,863	0.72
2021-04-11-210324_RNA_mixing-check_1C_polyA	2x150	STAR-2.5.3a	Candida	6,627,497	0.59	1,940,639	44,839	21,919	1,885	0.15
2021-04-11-210324_RNA_mixing-check_1C_polyA	2x150	STAR-2.5.3a	sacCer3	6,627,497	0.49	10,009,914	276,286	466,731	19,311	0.81
2021-04-11-210324_RNA_mixing-check_2A_polyA	2x150	STAR-2.5.3a	Candida	6,403,480	0.55	4,775,214	109,470	71,882	7,292	0.39
2021-04-11-210324_RNA_mixing-check_2A_polyA	2x150	STAR-2.5.3a	sacCer3	6,403,480	0.54	6,813,410	187,142	506,704	11,876	0.59
2021-04-11-210324_RNA_mixing-check_2B_polyA	2x150	STAR-2.5.3a	Candida	6,787,334	0.56	3,126,533	73,847	59,092	4,008	0.24
2021-04-11-210324_RNA_mixing-check_2B_polyA	2x150	STAR-2.5.3a	sacCer3	6,787,334	0.50	9,016,855	255,579	602,881	17,065	0.73
2021-04-11-210324_RNA_mixing-check_2C_polyA	2x150	STAR-2.5.3a	Candida	5,654,272	0.61	1,578,209	36,819	66,593	1,483	0.15
2021-04-11-210324_RNA_mixing-check_2C_polyA	2x150	STAR-2.5.3a	sacCer3	5,654,272	0.51	8,261,545	239,821	768,038	16,334	0.82

#	Exonic:	Intergenic:	Intronic:
2021-04-11-210324_RNA_mixing-check_1A_polyA.sacCer3	0.93	0.07	0.00
2021-04-11-210324_RNA_mixing-check_1B_polyA.sacCer3	0.92	0.07	0.00
2021-04-11-210324_RNA_mixing-check_1C_polyA.sacCer3	0.92	0.07	0.00
2021-04-11-210324_RNA_mixing-check_2A_polyA.sacCer3	0.93	0.07	0.00
2021-04-11-210324_RNA_mixing-check_2B_polyA.sacCer3	0.92	0.07	0.00
2021-04-11-210324_RNA_mixing-check_2C_polyA.sacCer3	0.93	0.07	0.00

COVERAGE QC

Coverage of genes; >1000bp, sacCer3



Coverage of genes; >1000bp, sacCer3

