

MAPPING STATS

Library	Read Length	Mapping	Species	Raw fragments	Complexity	Unique	Unique Splices	Multi	Multi Splices	Fraction mapped
2023-05-22-exp230404_MS06_SCD-MET_20min-DMSO_polyA-RNA	2x38	STAR-2.5.3a	Candida	32,391,941	0.51	16,718,317	300,707	388,856	18,132	0.27
2023-05-22-exp230404_MS06_SCD-MET_20min-DMSO_polyA-RNA	2x38	STAR-2.5.3a	sacCer3	32,391,941	0.51	16,718,317	300,707	388,856	18,132	0.27
2023-05-22-exp230404_MS74_SCD-MET_20min-10nM-PhIAA_polyA-RNA	2x38	STAR-2.5.3a	Candida	31,456,982	0.50	17,296,552	303,170	419,812	27,008	0.29
2023-05-22-exp230404_MS74_SCD-MET_20min-10nM-PhIAA_polyA-RNA	2x38	STAR-2.5.3a	sacCer3	31,456,982	0.50	17,296,552	303,170	419,812	27,008	0.29
2023-05-22-exp230404_MS74_SCD-MET_20min-DMSO_polyA-RNA	2x38	STAR-2.5.3a	Candida	31,160,978	0.50	17,736,680	308,998	441,784	26,156	0.30
2023-05-22-exp230404_MS74_SCD-MET_20min-DMSO_polyA-RNA	2x38	STAR-2.5.3a	sacCer3	31,160,978	0.50	17,736,680	308,998	441,784	26,156	0.30
2023-05-22-exp230404_MS74_SCD-MET_40min-10nM-PhIAA_polyA-RNA	2x38	STAR-2.5.3a	Candida	29,550,414	0.50	17,938,466	305,800	446,659	19,905	0.32
2023-05-22-exp230404_MS74_SCD-MET_40min-10nM-PhIAA_polyA-RNA	2x38	STAR-2.5.3a	sacCer3	29,550,414	0.50	17,938,466	305,800	446,659	19,905	0.32
2023-05-22-exp230404_MS74_SCD-MET_60min-10nM-PhIAA_polyA-RNA	2x38	STAR-2.5.3a	Candida	30,490,856	0.50	19,707,465	320,167	507,531	21,367	0.34
2023-05-22-exp230404_MS74_SCD-MET_60min-10nM-PhIAA_polyA-RNA	2x38	STAR-2.5.3a	sacCer3	30,490,856	0.50	19,707,465	320,167	507,531	21,367	0.34

COVERAGE ALL READS

Coverage of genes; >1000bp, sacCer3

