

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	57,489,258	0.45	29,942,658	544,870	716,629	126,793	0.27
2023-05-22-exp230404_MS06_SCD-MET_20min-DMSO_polyA-RNA	2x38	STAR-2.5.3a	sacCer3	57,489,258	0.45	29,942,658	544,870	716,629	126,793	0.27
2023-05-22-exp230404_MS74_SCD-MET_20min-10nM-PhIAA_polyA-RNA	2x38	STAR-2.5.3a	Candida	55,903,611	0.44	31,013,568	550,104	769,414	59,492	0.29
2023-05-22-exp230404_MS74_SCD-MET_20min-10nM-PhIAA_polyA-RNA	2x38	STAR-2.5.3a	sacCer3	55,903,611	0.44	31,013,568	550,104	769,414	59,492	0.29
2023-05-22-exp230404_MS74_SCD-MET_20min-DMSO_polyA-RNA	2x38	STAR-2.5.3a	Candida	55,262,041	0.44	31,688,077	559,441	807,005	55,001	0.30
2023-05-22-exp230404_MS74_SCD-MET_20min-DMSO_polyA-RNA	2x38	STAR-2.5.3a	sacCer3	55,262,041	0.44	31,688,077	559,441	807,005	55,001	0.30
2023-05-22-exp230404_MS74_SCD-MET_40min-10nM-PhIAA_polyA-RNA	2x38	STAR-2.5.3a	Candida	52,488,047	0.44	32,141,665	555,709	823,291	48,793	0.32
2023-05-22-exp230404_MS74_SCD-MET_40min-10nM-PhIAA_polyA-RNA	2x38	STAR-2.5.3a	sacCer3	52,488,047	0.44	32,141,665	555,709	823,291	48,793	0.32
2023-05-22-exp230404_MS74_SCD-MET_60min-10nM-PhIAA_polyA-RNA	2x38	STAR-2.5.3a	Candida	54,206,534	0.43	35,352,601	582,057	938,648	47,902	0.34
2023-05-22-exp230404_MS74_SCD-MET_60min-10nM-PhIAA_polyA-RNA	2x38	STAR-2.5.3a	sacCer3	54,206,534	0.43	35,352,601	582,057	938,648	47,902	0.34