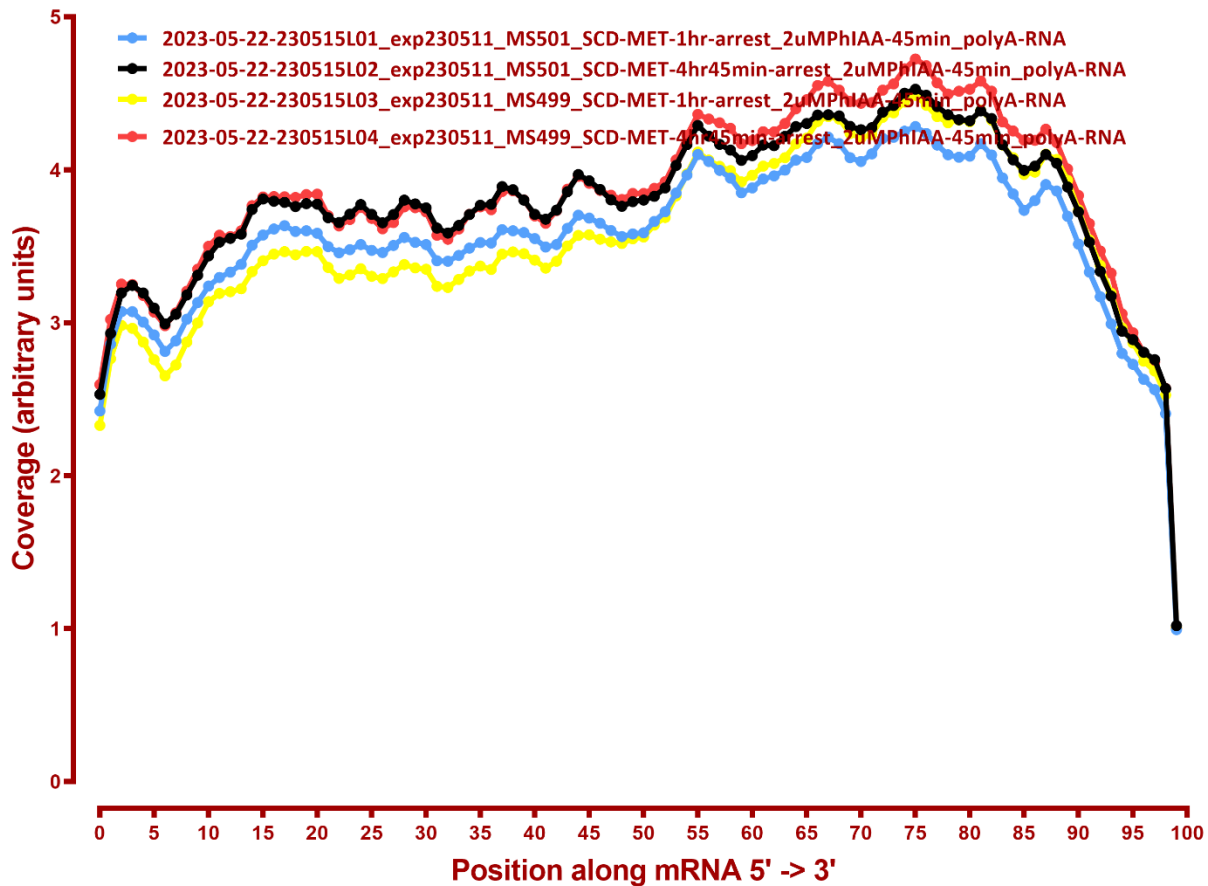


MAPPING STATS

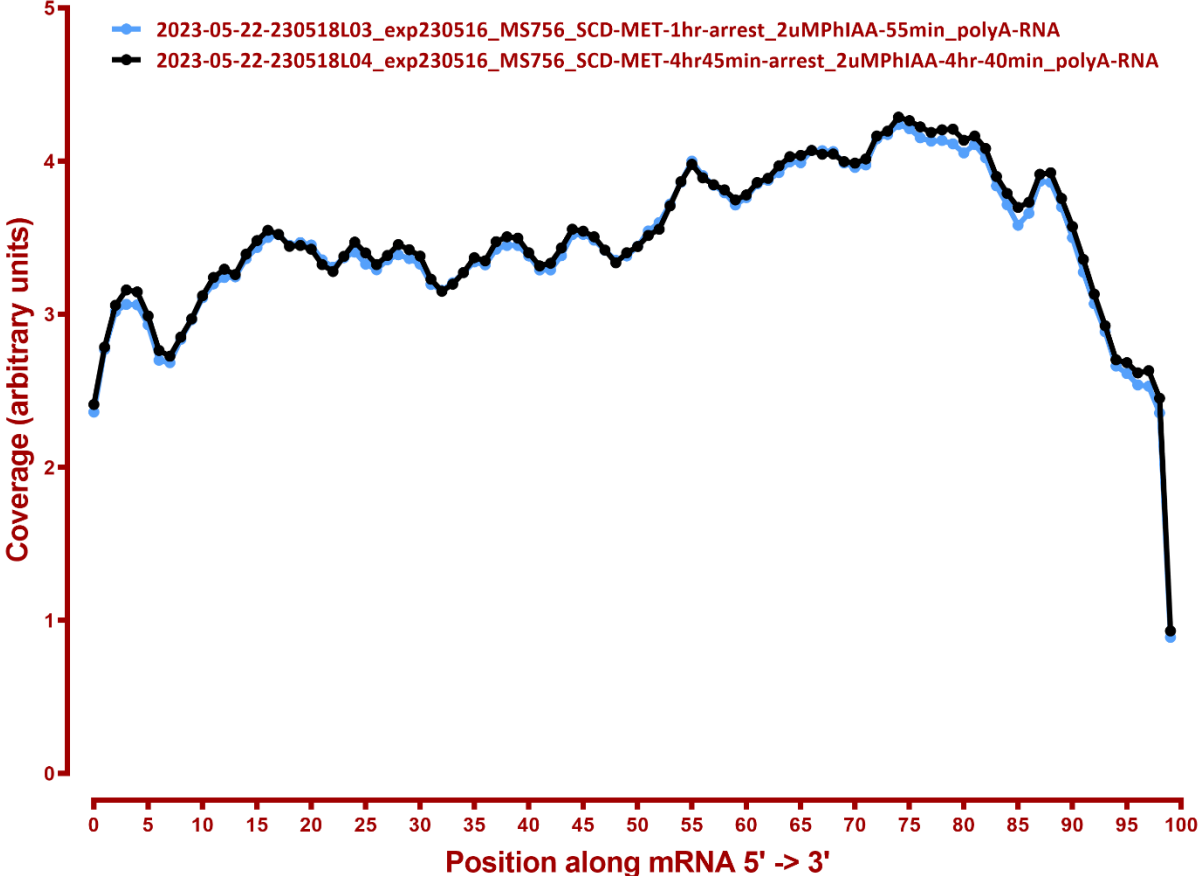
Library	Read Length	Mapping	Species	Raw fragments	Complexity	Unique	Unique Splices	Multi	Multi Splices	Fraction mapped
2023-05-22-230515L01_exp230511_MS501_SCD-MET-1hr-arrest_2uMPHIAA-45min_polyA	2x38	STAR-2.5.3a	Candida	9,591,385	0.65	140,253	3,647	5,420,163	60,229	0.29
2023-05-22-230515L01_exp230511_MS501_SCD-MET-1hr-arrest_2uMPHIAA-45min_polyA	2x38	STAR-2.5.3a	sacCer3	9,591,385	0.65	140,253	3,647	5,420,163	60,229	0.29
2023-05-22-230515L02_exp230511_MS501_SCD-MET-4hr45min-arrest_2uMPHIAA-45min	2x38	STAR-2.5.3a	Candida	9,551,610	0.64	162,449	4,159	6,221,404	68,506	0.34
2023-05-22-230515L02_exp230511_MS501_SCD-MET-4hr45min-arrest_2uMPHIAA-45min	2x38	STAR-2.5.3a	sacCer3	9,551,610	0.64	162,449	4,159	6,221,404	68,506	0.34
2023-05-22-230515L03_exp230511_MS499_SCD-MET-1hr-arrest_2uMPHIAA-45min_polyA	2x38	STAR-2.5.3a	Candida	9,600,231	0.65	133,759	3,463	5,081,202	58,586	0.27
2023-05-22-230515L03_exp230511_MS499_SCD-MET-1hr-arrest_2uMPHIAA-45min_polyA	2x38	STAR-2.5.3a	sacCer3	9,600,231	0.65	133,759	3,463	5,081,202	58,586	0.27
2023-05-22-230515L04_exp230511_MS499_SCD-MET-4hr45min-arrest_2uMPHIAA-45min	2x38	STAR-2.5.3a	Candida	9,859,680	0.64	155,947	3,339	5,418,079	61,793	0.29
2023-05-22-230515L04_exp230511_MS499_SCD-MET-4hr45min-arrest_2uMPHIAA-45min	2x38	STAR-2.5.3a	sacCer3	9,859,680	0.64	155,947	3,339	5,418,079	61,793	0.29
2023-05-22-230515L05_exp230511_MS069_SCD-MET-1hr-arrest_2uMPHIAA-45min_polyA	2x38	STAR-2.5.3a	Candida	9,923,574	0.64	148,046	3,728	5,629,252	63,806	0.29
2023-05-22-230515L05_exp230511_MS069_SCD-MET-1hr-arrest_2uMPHIAA-45min_polyA	2x38	STAR-2.5.3a	sacCer3	9,923,574	0.64	148,046	3,728	5,629,252	63,806	0.29
2023-05-22-230515L06_exp230511_MS069_SCD-MET-4hr45min-arrest_2uMPHIAA-45min	2x38	STAR-2.5.3a	Candida	9,758,801	0.64	158,008	3,388	5,952,085	67,181	0.32
2023-05-22-230515L06_exp230511_MS069_SCD-MET-4hr45min-arrest_2uMPHIAA-45min	2x38	STAR-2.5.3a	sacCer3	9,758,801	0.64	158,008	3,388	5,952,085	67,181	0.32
2023-05-22-230515L07_exp230511_MS500_SCD-MET-1hr-arrest_2uMPHIAA-45min_polyA	2x38	STAR-2.5.3a	Candida	9,532,336	0.65	145,305	3,577	5,110,016	60,064	0.28
2023-05-22-230515L07_exp230511_MS500_SCD-MET-1hr-arrest_2uMPHIAA-45min_polyA	2x38	STAR-2.5.3a	sacCer3	9,532,336	0.65	145,305	3,577	5,110,016	60,064	0.28
2023-05-22-230515L08_exp230511_MS500_SCD-MET-4hr45min-arrest_2uMPHIAA-45min	2x38	STAR-2.5.3a	Candida	9,074,058	0.66	127,856	2,728	4,316,761	49,897	0.25
2023-05-22-230515L08_exp230511_MS500_SCD-MET-4hr45min-arrest_2uMPHIAA-45min	2x38	STAR-2.5.3a	sacCer3	9,074,058	0.66	127,856	2,728	4,316,761	49,897	0.25
2023-05-22-230515L09_exp230511_MS497_SCD-MET-1hr-arrest_2uMPHIAA-45min_polyA	2x38	STAR-2.5.3a	Candida	9,153,726	0.66	115,177	2,409	4,381,137	52,389	0.25
2023-05-22-230515L09_exp230511_MS497_SCD-MET-1hr-arrest_2uMPHIAA-45min_polyA	2x38	STAR-2.5.3a	sacCer3	9,153,726	0.66	115,177	2,409	4,381,137	52,389	0.25
2023-05-22-230515L10_exp230511_MS497_SCD-MET-4hr45min-arrest_2uMPHIAA-45min	2x38	STAR-2.5.3a	Candida	9,895,688	0.64	157,854	3,680	5,598,962	69,118	0.29
2023-05-22-230515L10_exp230511_MS497_SCD-MET-4hr45min-arrest_2uMPHIAA-45min	2x38	STAR-2.5.3a	sacCer3	9,895,688	0.64	157,854	3,680	5,598,962	69,118	0.29
2023-05-22-230518L01_exp230516_MS069_SCD-MET-1hr-arrest_2uMPHIAA-55min_polyA	2x38	STAR-2.5.3a	Candida	9,578,281	0.62	172,263	4,927	7,198,260	87,252	0.39
2023-05-22-230518L01_exp230516_MS069_SCD-MET-1hr-arrest_2uMPHIAA-55min_polyA	2x38	STAR-2.5.3a	sacCer3	9,578,281	0.62	172,263	4,927	7,198,260	87,252	0.39
2023-05-22-230518L02_exp230516_MS069_SCD-MET-4hr45min-arrest_2uMPHIAA-4hr-40r	2x38	STAR-2.5.3a	Candida	11,219,243	0.60	227,220	6,188	9,084,000	111,162	0.42
2023-05-22-230518L02_exp230516_MS069_SCD-MET-4hr45min-arrest_2uMPHIAA-4hr-40r	2x38	STAR-2.5.3a	sacCer3	11,219,243	0.60	227,220	6,188	9,084,000	111,162	0.42
2023-05-22-230518L03_exp230516_MS756_SCD-MET-1hr-arrest_2uMPHIAA-55min_polyA	2x38	STAR-2.5.3a	Candida	9,355,920	0.63	184,518	3,712	6,410,243	78,555	0.36
2023-05-22-230518L03_exp230516_MS756_SCD-MET-1hr-arrest_2uMPHIAA-55min_polyA	2x38	STAR-2.5.3a	sacCer3	9,355,920	0.63	184,518	3,712	6,410,243	78,555	0.36
2023-05-22-230518L04_exp230516_MS756_SCD-MET-4hr45min-arrest_2uMPHIAA-4hr-40r	2x38	STAR-2.5.3a	Candida	9,746,642	0.62	219,800	7,180	7,760,260	93,454	0.41
2023-05-22-230518L04_exp230516_MS756_SCD-MET-4hr45min-arrest_2uMPHIAA-4hr-40r	2x38	STAR-2.5.3a	sacCer3	9,746,642	0.62	219,800	7,180	7,760,260	93,454	0.41

COVERAGE ALL READS

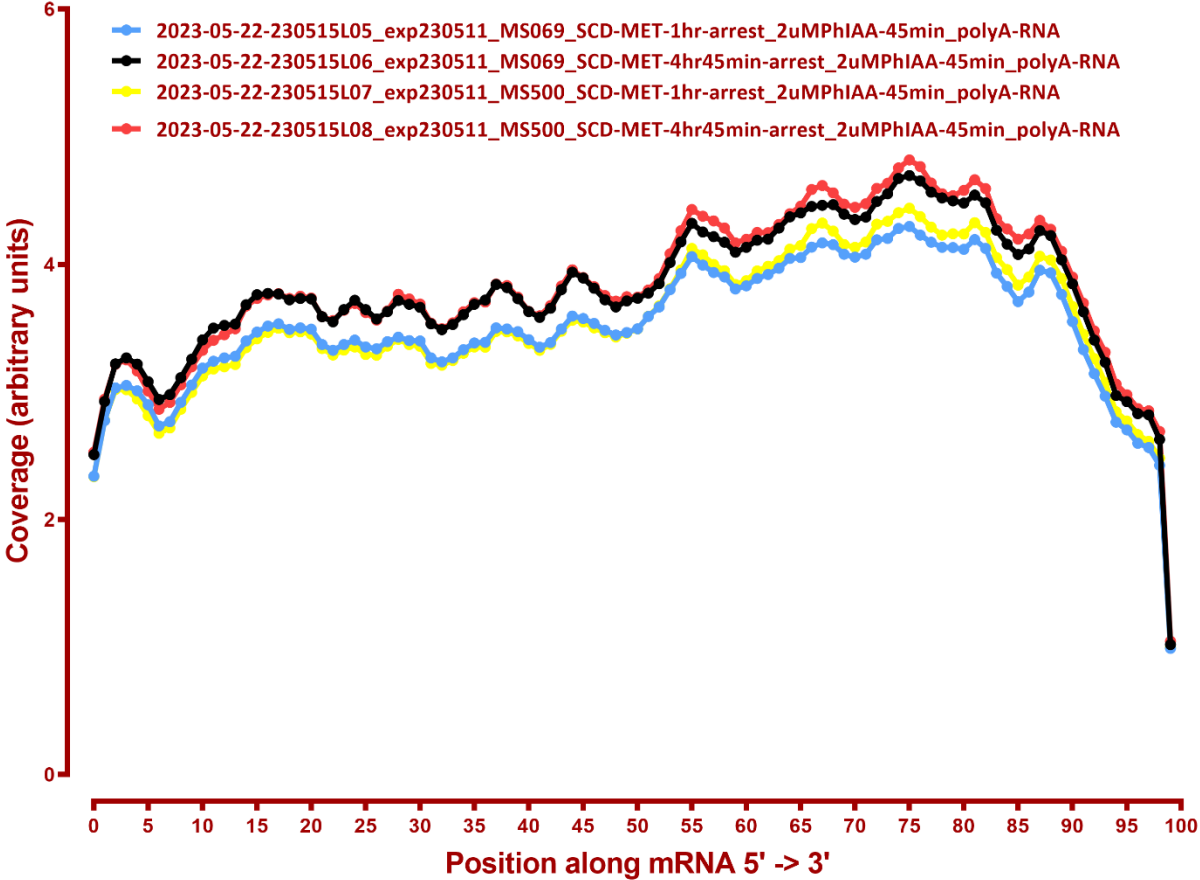
Coverage of genes; >1000bp, sacCer3



Coverage of genes; >1000bp, sacCer3



Coverage of genes; >1000bp, sacCer3



Coverage of genes; >1000bp, sacCer3

