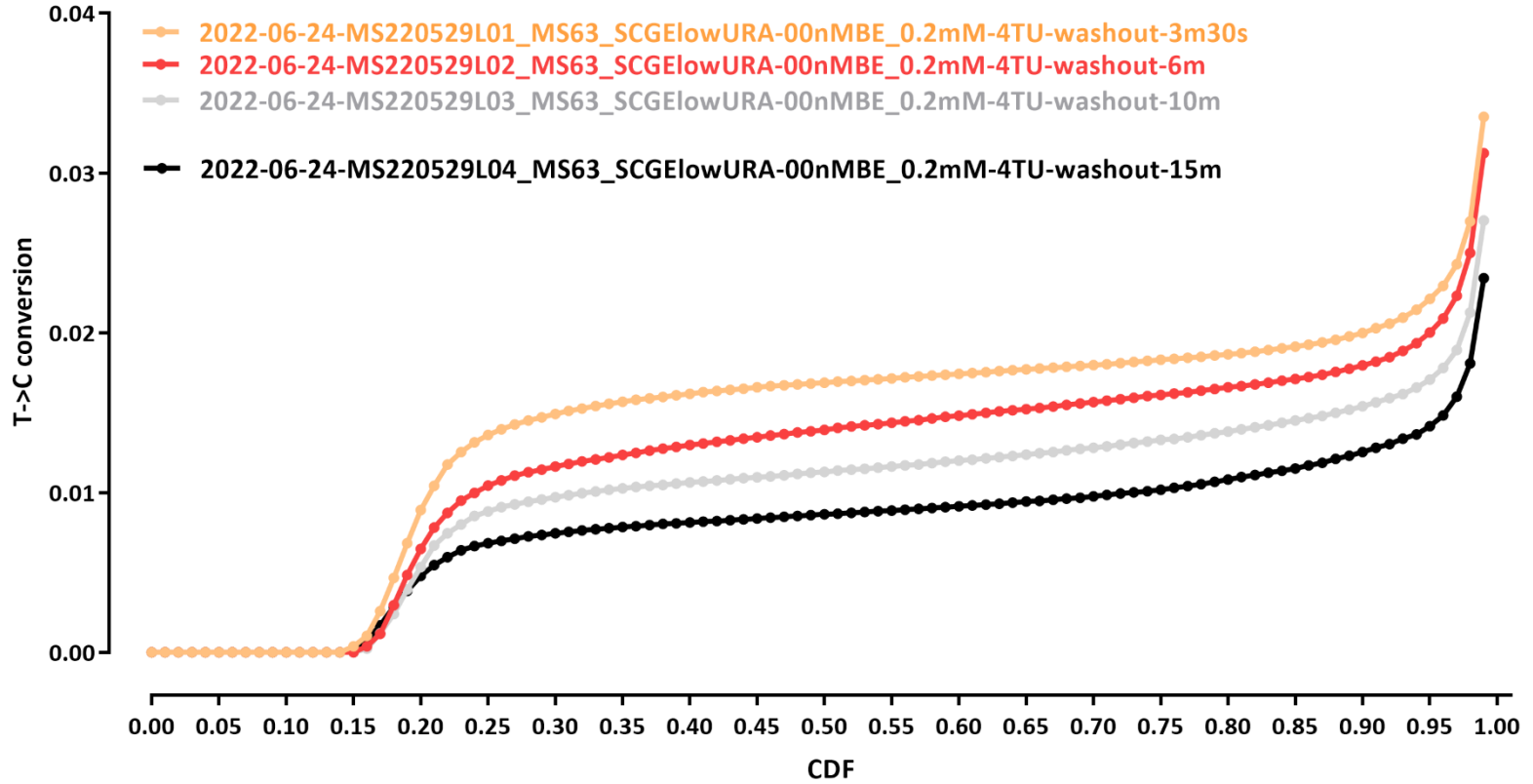


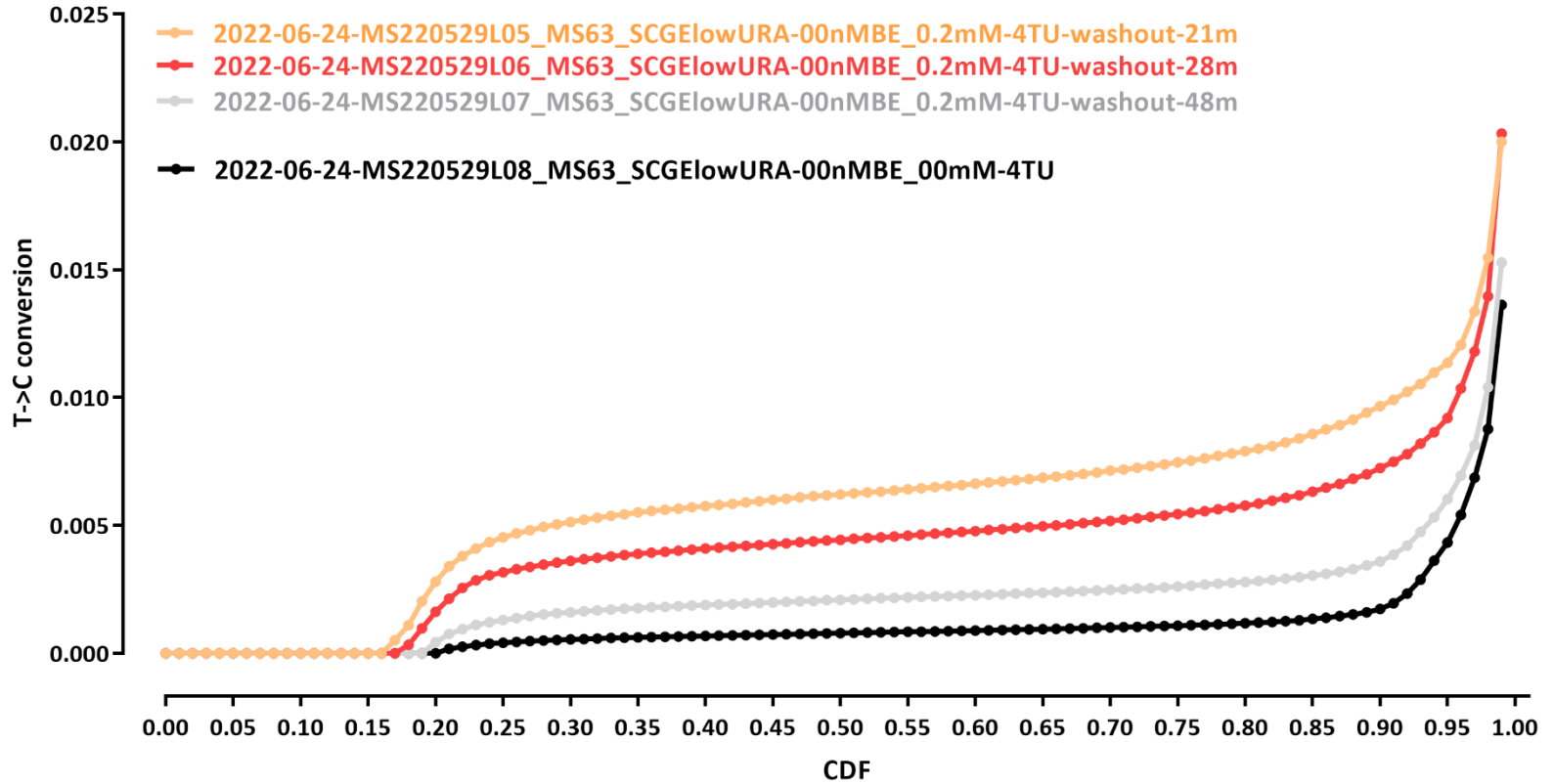
# Mapping stats

2022-06-24-SLAM-seq	Read Length	Mapping	Species	Raw fragments	Complexity	Unique	Unique Splices	Multi	Multi Splices	Fraction mapped
2022-06-24-MS220529L01_MS63_SCGElowURA-00nMBE_0.2mM-4TU-washout-3m30s	2x150	slamdunk-0.4.3, transcriptome	<b>S.cerevisiae</b>	13,419,744	0.44	22,415,175	0	23,214	0	0.84
	2x150	slamdunk-0.4.3, transcriptome								#DIV/0!
2022-06-24-MS220529L02_MS63_SCGElowURA-00nMBE_0.2mM-4TU-washout-6m	2x150	slamdunk-0.4.3, transcriptome	<b>S.cerevisiae</b>	8,796,915	0.50	14,626,595	0	15,293	0	0.83
	2x150	slamdunk-0.4.3, transcriptome								#DIV/0!
2022-06-24-MS220529L03_MS63_SCGElowURA-00nMBE_0.2mM-4TU-washout-10m	2x150	slamdunk-0.4.3, transcriptome	<b>S.cerevisiae</b>	9,727,365	0.48	16,176,555	0	15,705	0	0.83
	2x150	slamdunk-0.4.3, transcriptome								#DIV/0!
2022-06-24-MS220529L04_MS63_SCGElowURA-00nMBE_0.2mM-4TU-washout-15m	2x150	slamdunk-0.4.3, transcriptome	<b>S.cerevisiae</b>	17,003,281	0.42	28,257,507	0	27,114	0	0.83
	2x150	slamdunk-0.4.3, transcriptome								#DIV/0!
2022-06-24-MS220529L05_MS63_SCGElowURA-00nMBE_0.2mM-4TU-washout-21m	2x150	slamdunk-0.4.3, transcriptome	<b>S.cerevisiae</b>	9,485,958	0.49	15,784,667	0	14,207	0	0.83
	2x150	slamdunk-0.4.3, transcriptome								#DIV/0!
2022-06-24-MS220529L06_MS63_SCGElowURA-00nMBE_0.2mM-4TU-washout-28m	2x150	slamdunk-0.4.3, transcriptome	<b>S.cerevisiae</b>	7,516,261	0.52	12,550,706	0	11,092	0	0.84
	2x150	slamdunk-0.4.3, transcriptome								#DIV/0!
2022-06-24-MS220529L07_MS63_SCGElowURA-00nMBE_0.2mM-4TU-washout-48m	2x150	slamdunk-0.4.3, transcriptome	<b>S.cerevisiae</b>	7,667,503	0.51	12,906,648	0	10,140	0	0.84
	2x150	slamdunk-0.4.3, transcriptome								#DIV/0!
2022-06-24-MS220529L08_MS63_SCGElowURA-00nMBE_00mM-4TU	2x150	slamdunk-0.4.3, transcriptome	<b>S.cerevisiae</b>	10,767,395	0.46	18,127,839	0	14,207	0	0.84
	2x150	slamdunk-0.4.3, transcriptome								#DIV/0!
2022-06-24-MS220529L09_MS63_SCGElowURA-15nMBE_0.2mM-4TU-washout-3m30s	2x150	slamdunk-0.4.3, transcriptome	<b>S.cerevisiae</b>	6,849,347	0.52	11,544,409	0	9,908	0	0.84
	2x150	slamdunk-0.4.3, transcriptome								#DIV/0!
2022-06-24-MS220529L10_MS63_SCGElowURA-15nMBE_0.2mM-4TU-washout-6m	2x150	slamdunk-0.4.3, transcriptome	<b>S.cerevisiae</b>	6,162,655	0.53	10,412,303	0	9,010	0	0.85
	2x150	slamdunk-0.4.3, transcriptome								#DIV/0!
2022-06-24-MS220529L11_MS63_SCGElowURA-15nMBE_0.2mM-4TU-washout-10m	2x150	slamdunk-0.4.3, transcriptome	<b>S.cerevisiae</b>	8,545,483	0.49	14,398,625	0	11,658	0	0.84
	2x150	slamdunk-0.4.3, transcriptome								#DIV/0!
2022-06-24-MS220529L12_MS63_SCGElowURA-15nMBE_0.2mM-4TU-washout-15m	2x150	slamdunk-0.4.3, transcriptome	<b>S.cerevisiae</b>	6,053,761	0.53	10,176,207	0	8,650	0	0.84
	2x150	slamdunk-0.4.3, transcriptome								#DIV/0!
2022-06-24-MS220529L13_MS63_SCGElowURA-15nMBE_0.2mM-4TU-washout-21m	2x150	slamdunk-0.4.3, transcriptome	<b>S.cerevisiae</b>	6,732,247	0.52	11,307,514	0	9,417	0	0.84
	2x150	slamdunk-0.4.3, transcriptome								#DIV/0!
2022-06-24-MS220529L14_MS63_SCGElowURA-15nMBE_0.2mM-4TU-washout-28m	2x150	slamdunk-0.4.3, transcriptome	<b>S.cerevisiae</b>	5,941,387	0.52	10,020,162	0	7,685	0	0.84
	2x150	slamdunk-0.4.3, transcriptome								#DIV/0!
2022-06-24-MS220529L15_MS63_SCGElowURA-15nMBE_0.2mM-4TU-washout-48m	2x150	slamdunk-0.4.3, transcriptome	<b>S.cerevisiae</b>	5,795,427	0.52	9,819,610	0	8,162	0	0.85
	2x150	slamdunk-0.4.3, transcriptome								#DIV/0!
2022-06-24-MS220529L16_MS63_SCGElowURA-15nMBE_00mM-4TU	2x150	slamdunk-0.4.3, transcriptome	<b>S.cerevisiae</b>	5,646,662	0.53	9,570,911	0	7,946	0	0.85
	2x150	slamdunk-0.4.3, transcriptome								#DIV/0!

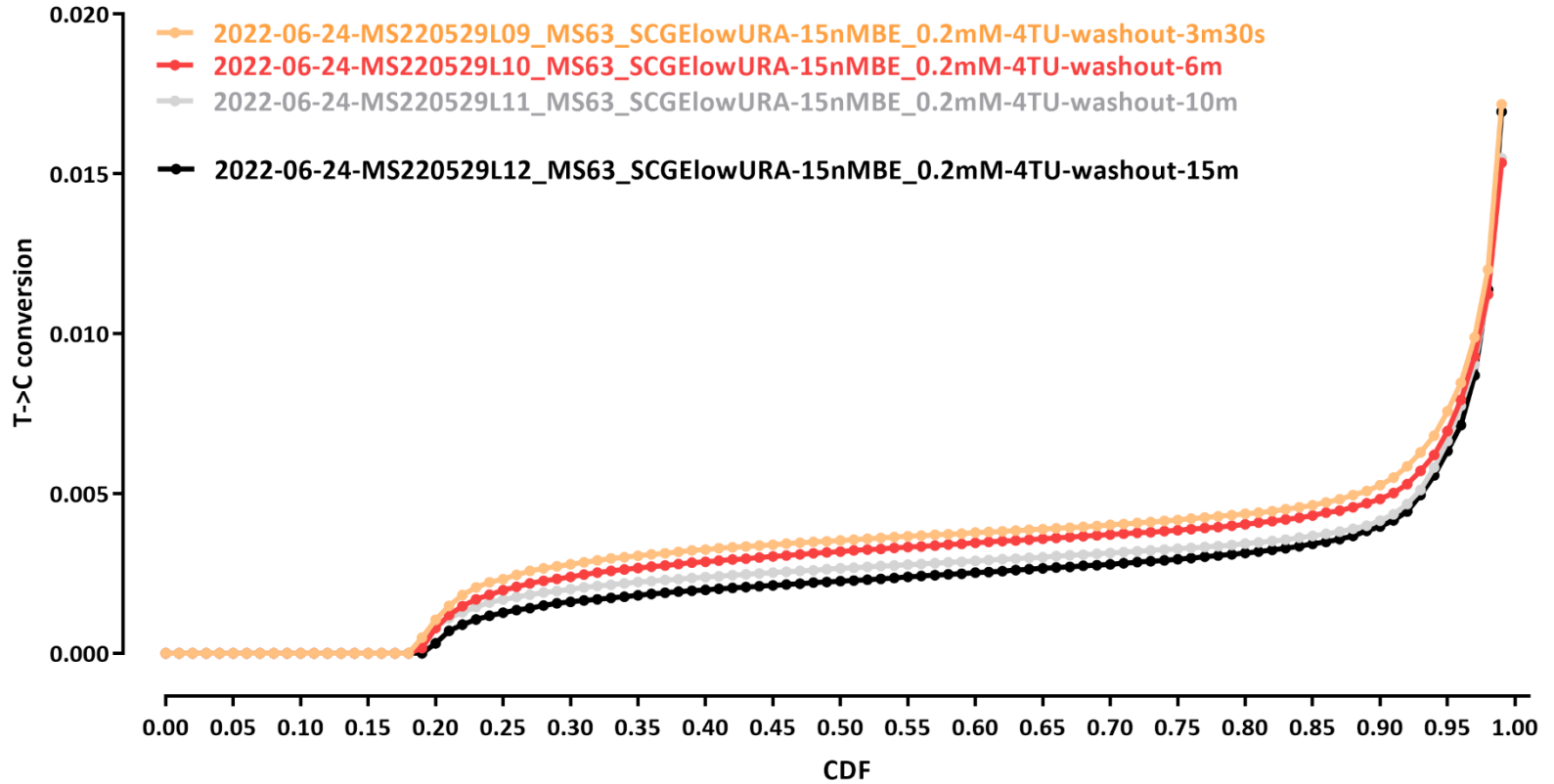
# S\_cerevisiae 1



## S\_cerevisiae 2



### S\_cerevisiae 3



### S\_cerevisiae 4

