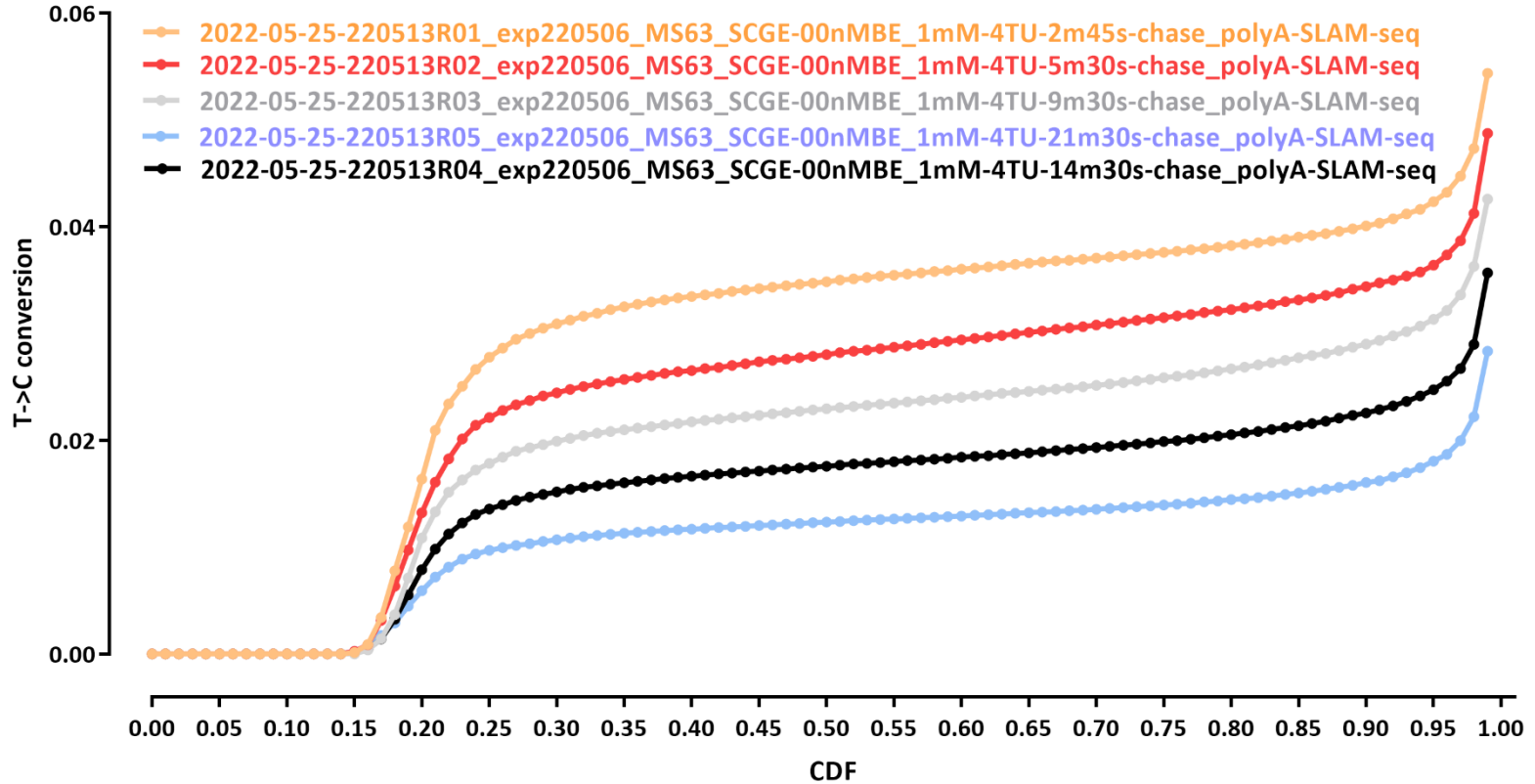


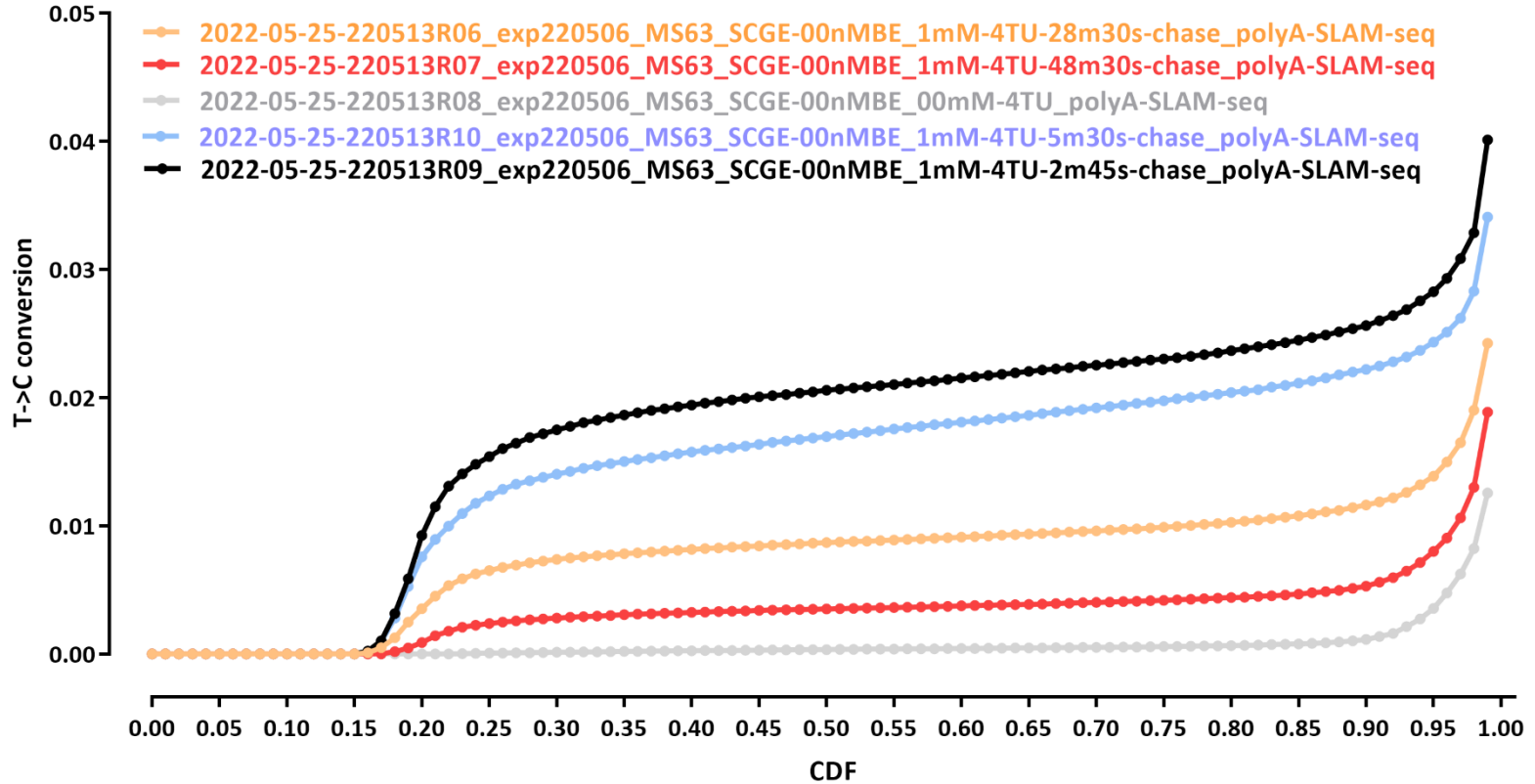
Mapping stats

Library	Read Length	Mapping	Species	Raw fragments	Complexity	Unique	Unique Splices	Multi	Multi Splices	Fraction mapped
2022-05-25-220513R01_exp220506_MS63_SCGE-00nMBE_1mM-4TU-2m45s-chase_polyA-SLAM-seq	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	27,052,848	0.10	45,541,790		65,531		0.84
	2x150	slamdunk-0.4.3, transcriptome								#DIV/0!
2022-05-25-220513R02_exp220506_MS63_SCGE-00nMBE_1mM-4TU-5m30s-chase_polyA-SLAM-seq	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	29,091,817	0.10	48,996,706		59,067		0.84
	2x150	slamdunk-0.4.3, transcriptome								#DIV/0!
2022-05-25-220513R03_exp220506_MS63_SCGE-00nMBE_1mM-4TU-9m30s-chase_polyA-SLAM-seq	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	34,491,088	0.09	44,556,204		55,652		0.65
	2x150	slamdunk-0.4.3, transcriptome								#DIV/0!
2022-05-25-220513R04_exp220506_MS63_SCGE-00nMBE_1mM-4TU-14m30s-chase_polyA-SLAM-seq	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	28,912,899	0.08	48,870,316		54,298		0.85
	2x150	slamdunk-0.4.3, transcriptome								#DIV/0!
2022-05-25-220513R05_exp220506_MS63_SCGE-00nMBE_1mM-4TU-21m30s-chase_polyA-SLAM-seq	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	31,061,202	0.10	52,427,694		51,064		0.84
	2x150	slamdunk-0.4.3, transcriptome								#DIV/0!
2022-05-25-220513R06_exp220506_MS63_SCGE-00nMBE_1mM-4TU-28m30s-chase_polyA-SLAM-seq	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	27,505,068	0.10	46,550,857		42,486		0.85
	2x150	slamdunk-0.4.3, transcriptome								#DIV/0!
2022-05-25-220513R07_exp220506_MS63_SCGE-00nMBE_1mM-4TU-48m30s-chase_polyA-SLAM-seq	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	28,992,267	0.09	49,334,562		34,239		0.85
	2x150	slamdunk-0.4.3, transcriptome								#DIV/0!
2022-05-25-220513R08_exp220506_MS63_SCGE-00nMBE_00mM-4TU_polyA-SLAM-seq	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	35,940,537	0.05	61,163,681		35,686		0.85
	2x150	slamdunk-0.4.3, transcriptome								#DIV/0!
2022-05-25-220513R09_exp220506_MS63_SCGE-00nMBE_1mM-4TU-2m45s-chase_polyA-SLAM-seq	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	32,839,435	0.06	56,111,805		68,435		0.86
	2x150	slamdunk-0.4.3, transcriptome								#DIV/0!
2022-05-25-220513R10_exp220506_MS63_SCGE-00nMBE_1mM-4TU-5m30s-chase_polyA-SLAM-seq	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	26,710,686	0.09	45,533,796		50,959		0.85
	2x150	slamdunk-0.4.3, transcriptome								#DIV/0!
2022-05-25-220513R11_exp220506_MS63_SCGE-00nMBE_1mM-4TU-9m30s-chase_polyA-SLAM-seq	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	25,708,932	0.05	43,277,209		30,795		0.84
	2x150	slamdunk-0.4.3, transcriptome								#DIV/0!
2022-05-25-220513R12_exp220506_MS63_SCGE-00nMBE_1mM-4TU-14m30s-chase_polyA-SLAM-seq	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	26,910,818	0.06	45,783,578		41,624		0.85
	2x150	slamdunk-0.4.3, transcriptome								#DIV/0!
2022-05-25-220513R13_exp220506_MS63_SCGE-00nMBE_1mM-4TU-21m30s-chase_polyA-SLAM-seq	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	27,861,269	0.05	47,381,153		36,549		0.85
	2x150	slamdunk-0.4.3, transcriptome								#DIV/0!
2022-05-25-220513R14_exp220506_MS63_SCGE-00nMBE_1mM-4TU-28m30s-chase_polyA-SLAM-seq	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	23,973,242	0.08	40,897,842		29,421		0.85
	2x150	slamdunk-0.4.3, transcriptome								#DIV/0!
2022-05-25-220513R15_exp220506_MS63_SCGE-00nMBE_1mM-4TU-48m30s-chase_polyA-SLAM-seq	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	25,342,255	0.06	43,460,025		32,607		0.86
	2x150	slamdunk-0.4.3, transcriptome								#DIV/0!
2022-05-25-220513R16_exp220506_MS63_SCGE-00nMBE_00mM-4TU_polyA-SLAM-seq	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	29,849,463	0.05	51,035,262		41,962		0.86
	2x150	slamdunk-0.4.3, transcriptome								#DIV/0!
2022-05-25-220513R17_exp220508_MS63_SCGE-00nMBE_0.2mM-4TU_polyA-SLAM-seq	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	32,117,441	0.05	54,300,526		62,443		0.85
	2x150	slamdunk-0.4.3, transcriptome								#DIV/0!

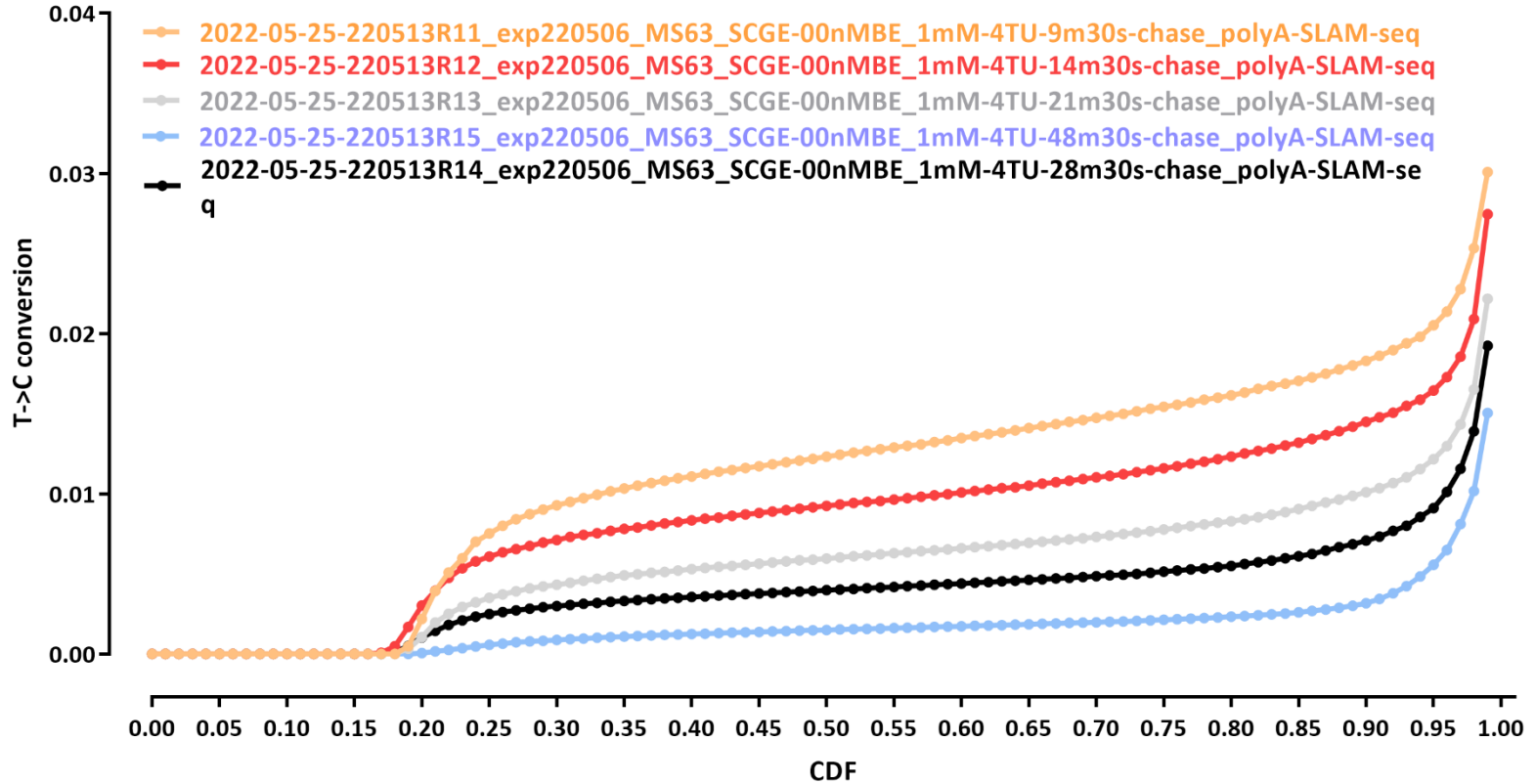
S_cerevisiae 1



S_cerevisiae 2



S_cerevisiae 3



S_cerevisiae 4

