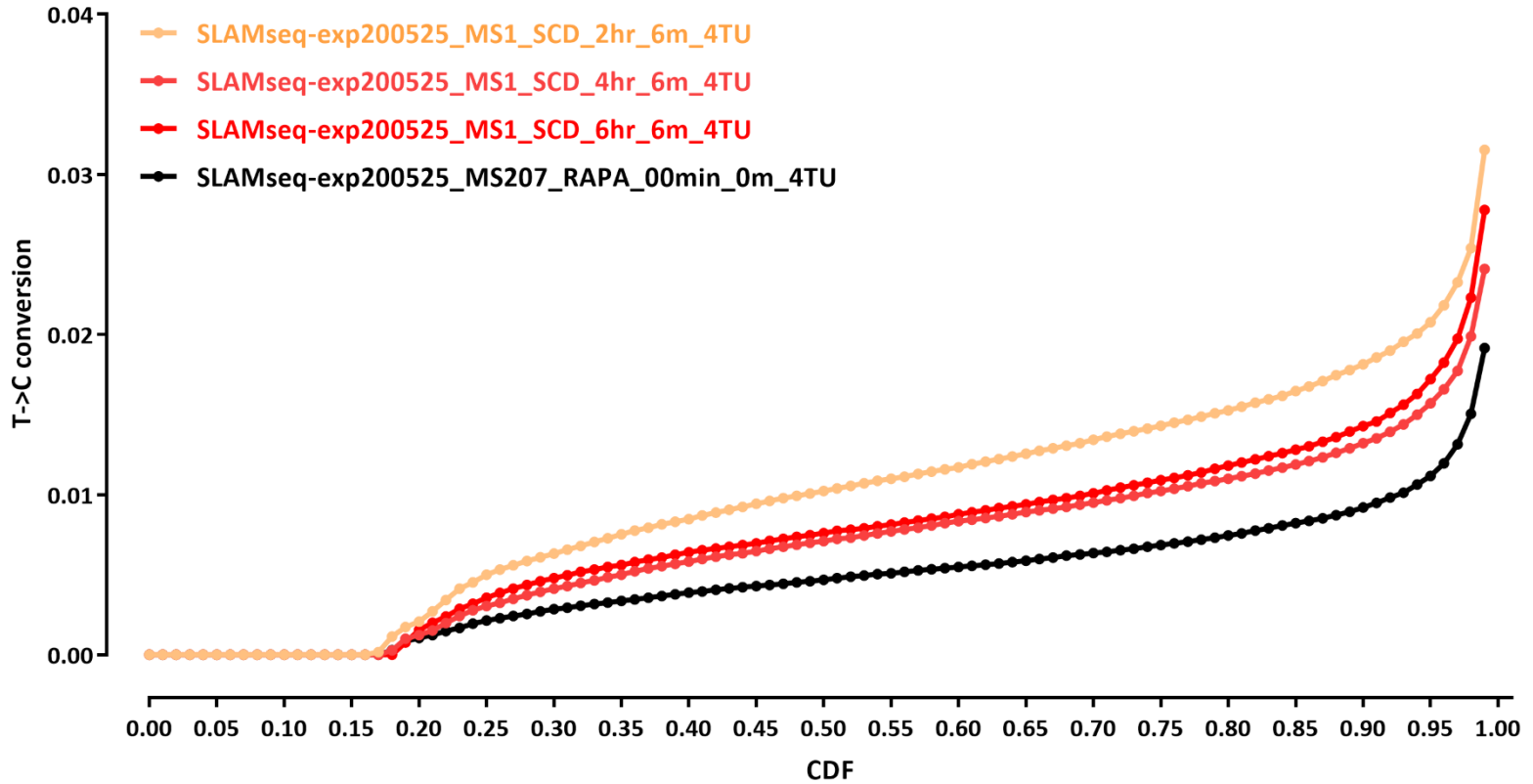


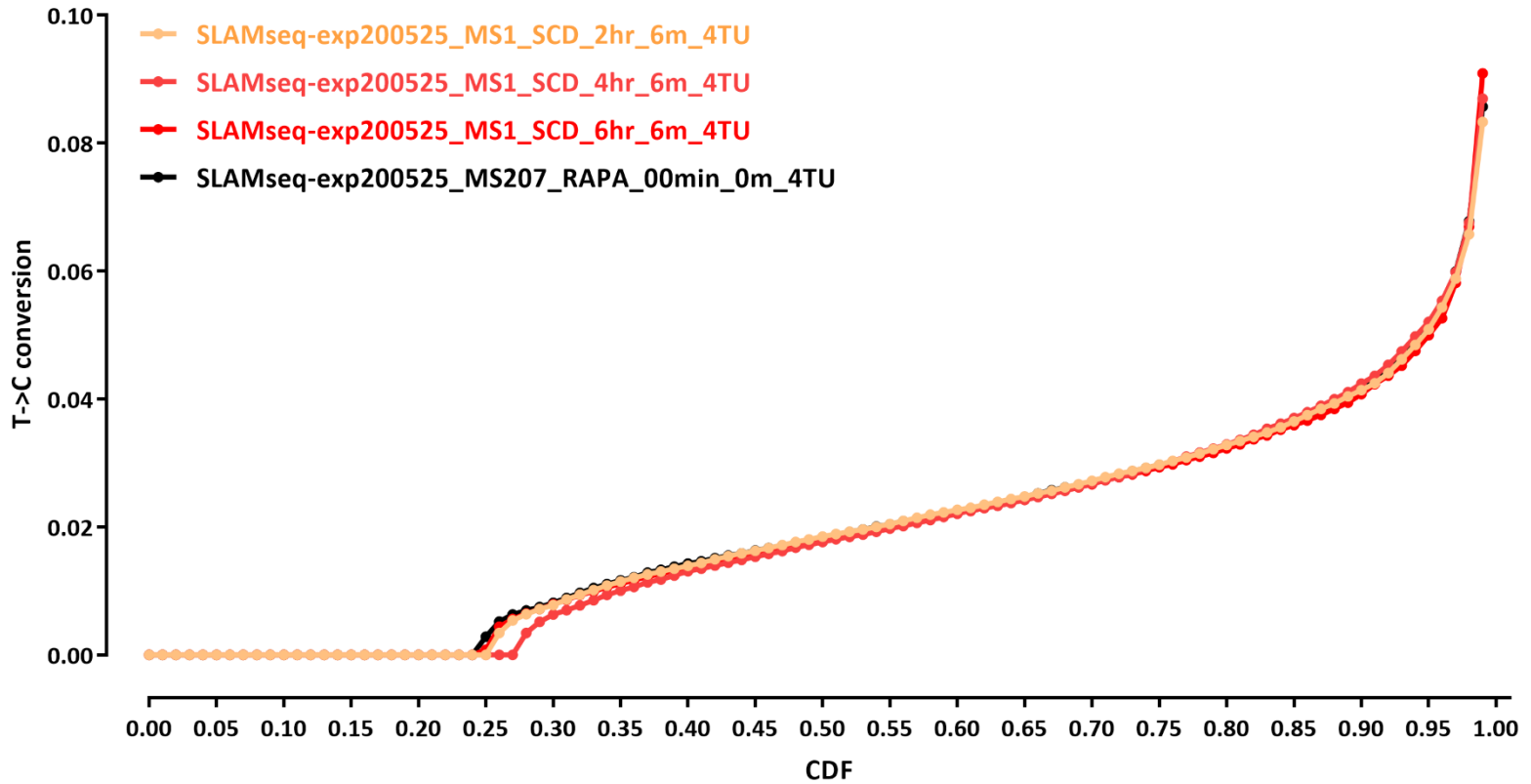
Mapping stats

Library	Read Length	Mapping	Species	Raw fragments	Complexity	Unique	Unique Splices	Multi	Multi Splices	Fraction mapped
2020-08-27-SLAMseq-exp200525_MS1_SCD_2hr_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	8,882,119	0.33	11,871,314	0	10,780	0	0.67
2020-08-27-SLAMseq-exp200525_MS1_SCD_2hr_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.pombe	8,882,119	0.45	2,962,417	0	5,380	0	0.17
2020-08-27-SLAMseq-exp200525_MS1_SCD_4hr_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	7,364,254	0.36	9,202,562	0	7,906	0	0.63
2020-08-27-SLAMseq-exp200525_MS1_SCD_4hr_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.pombe	7,364,254	0.51	1,913,456	0	3,268	0	0.13
2020-08-27-SLAMseq-exp200525_MS1_SCD_6hr_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	7,747,876	0.34	9,953,489	0	7,412	0	0.64
2020-08-27-SLAMseq-exp200525_MS1_SCD_6hr_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.pombe	7,747,876	0.46	2,868,798	0	4,590	0	0.19
2020-08-27-SLAMseq-exp200525_MS207_RAPA_00min_0m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	8,113,606	0.33	11,171,610	0	7,562	0	0.69
2020-08-27-SLAMseq-exp200525_MS207_RAPA_00min_0m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.pombe	8,113,606	0.47	2,633,956	0	4,075	0	0.16
2020-08-27-SLAMseq-exp200525_MS207_RAPA_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	8,000,441	0.33	10,942,651	0	9,644	0	0.68
2020-08-27-SLAMseq-exp200525_MS207_RAPA_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.pombe	8,000,441	0.48	2,503,426	0	4,397	0	0.16
2020-08-27-SLAMseq-exp200525_MS208_100_RAPA_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	7,206,453	0.34	9,570,767	0	7,926	0	0.66
2020-08-27-SLAMseq-exp200525_MS208_100_RAPA_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.pombe	7,206,453	0.48	2,523,327	0	4,578	0	0.18
2020-08-27-SLAMseq-exp200525_MS208_40_RAPA_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	6,827,263	0.35	8,390,156	0	4,921	0	0.61
2020-08-27-SLAMseq-exp200525_MS208_40_RAPA_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.pombe	6,827,263	0.46	3,093,771	0	4,212	0	0.23
2020-08-27-SLAMseq-exp200525_MS208_DMSO_00min_0m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	8,621,542	0.30	11,730,897	0	8,249	0	0.68
2020-08-27-SLAMseq-exp200525_MS208_DMSO_00min_0m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.pombe	8,621,542	0.44	2,922,535	0	4,494	0	0.17
2020-08-27-SLAMseq-exp200525_MS208_DMSO_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	7,737,421	0.35	10,900,004	0	10,190	0	0.71
2020-08-27-SLAMseq-exp200525_MS208_DMSO_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.pombe	7,737,421	0.51	2,083,177	0	3,899	0	0.13
2020-08-27-SLAMseq-exp200525_MS208_ON_RAPA_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	8,439,742	0.31	10,779,734	0	8,141	0	0.64
2020-08-27-SLAMseq-exp200525_MS208_ON_RAPA_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.pombe	8,439,742	0.44	3,278,712	0	5,395	0	0.19
2020-08-27-SLAMseq-exp200525_MS289_65_RAPA_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	9,476,345	0.30	2,277,187	0	1,976	0	0.12
2020-08-27-SLAMseq-exp200525_MS289_65_RAPA_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.pombe	9,476,345	0.22	13,117,629	0	22,967	0	0.69
2020-08-27-SLAMseq-exp200525_MS86_SCD_2hr_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	8,379,084	0.34	10,357,643	0	7,771	0	0.62
2020-08-27-SLAMseq-exp200525_MS86_SCD_2hr_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.pombe	8,379,084	0.44	3,567,418	0	5,495	0	0.21
2020-08-27-SLAMseq-exp200525_MS86_SCD_4hr_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	7,964,334	0.35	9,248,705	0	6,257	0	0.58
2020-08-27-SLAMseq-exp200525_MS86_SCD_4hr_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.pombe	7,964,334	0.41	3,893,501	0	6,014	0	0.24
2020-08-27-SLAMseq-exp200525_MS86_SCD_6hr_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	8,041,473	0.35	7,896,439	0	5,856	0	0.49
2020-08-27-SLAMseq-exp200525_MS86_SCD_6hr_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.pombe	8,041,473	0.37	5,367,699	0	8,247	0	0.33

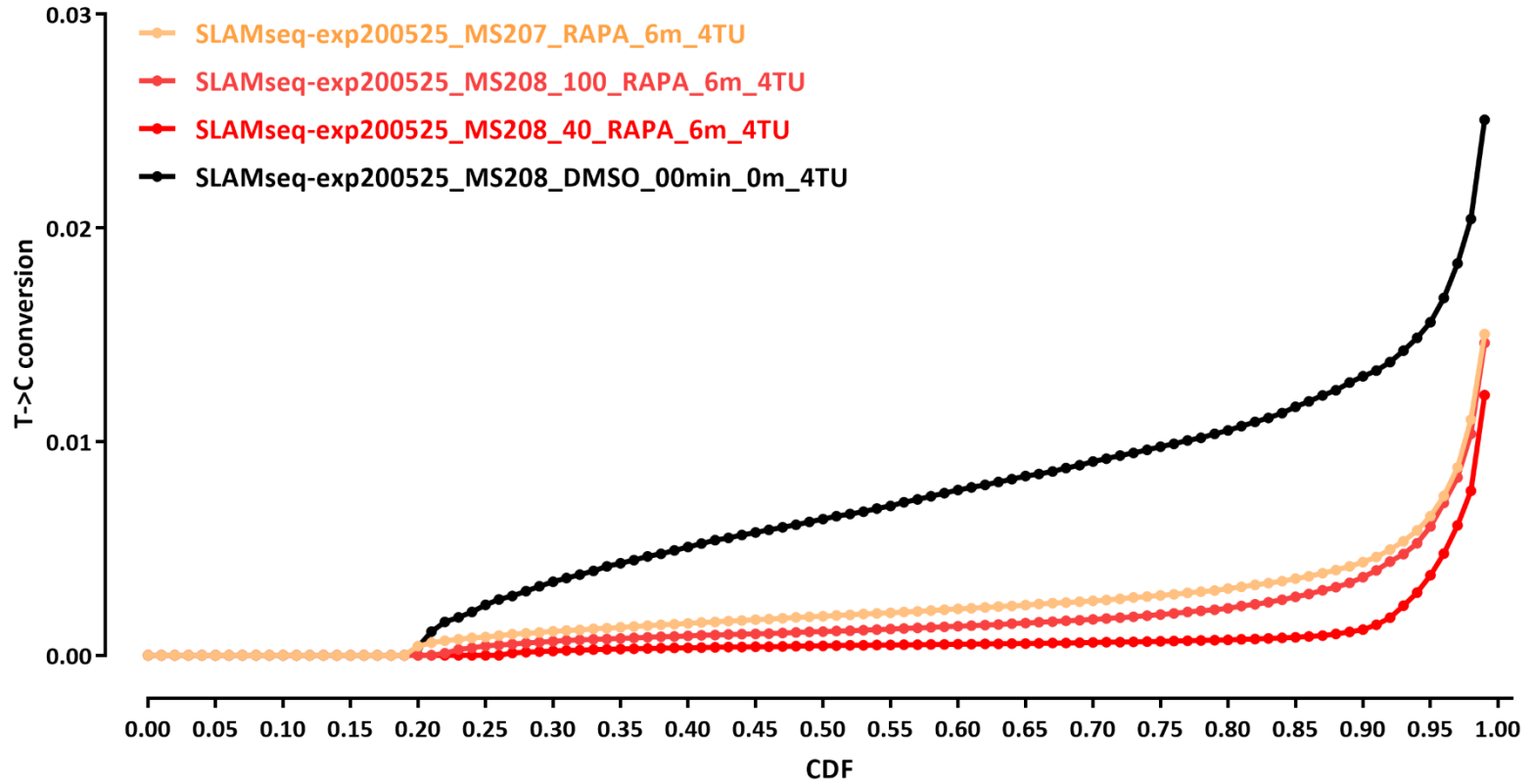
S_cerevisiae 1



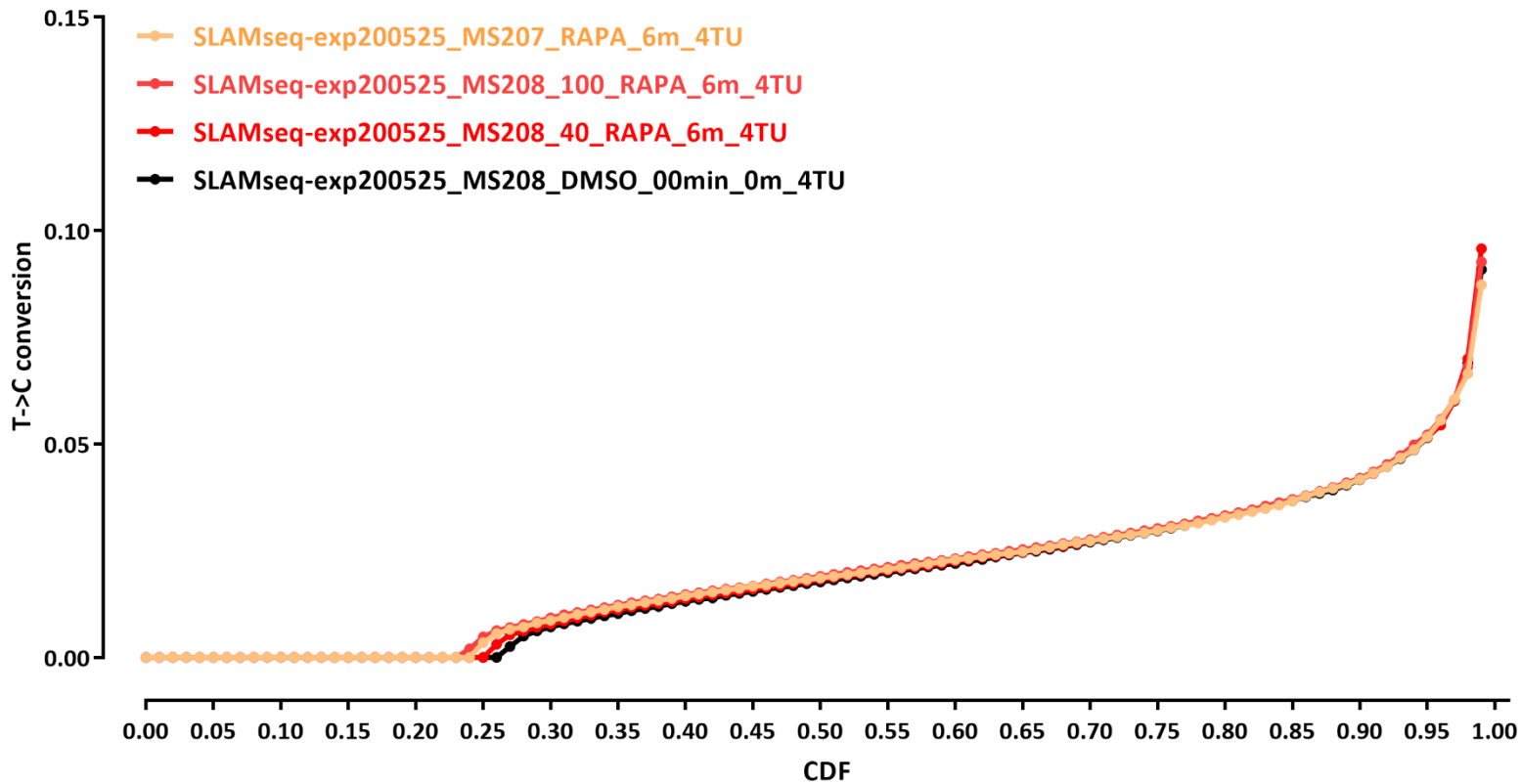
S_pombe 1



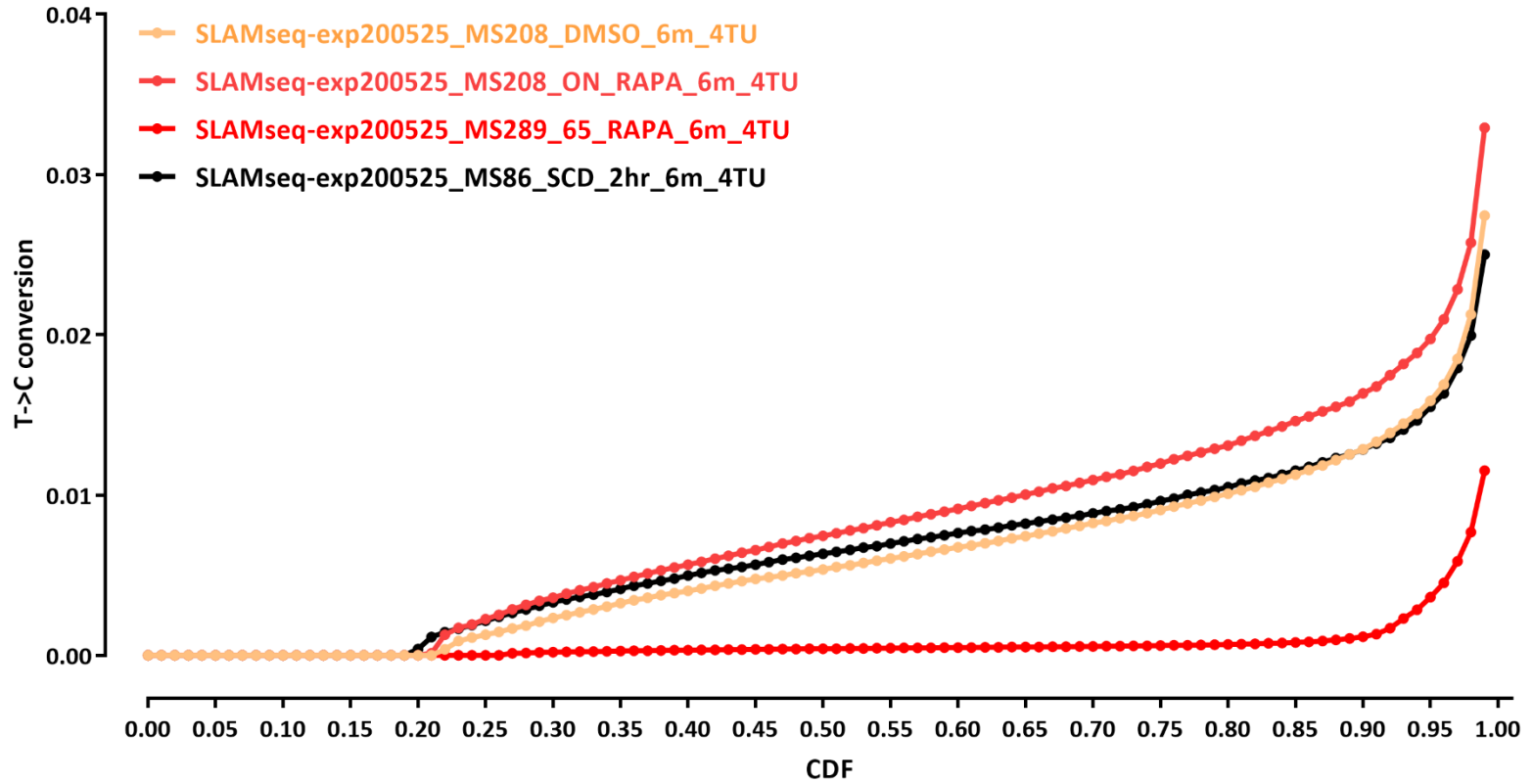
S_cerevisiae 2



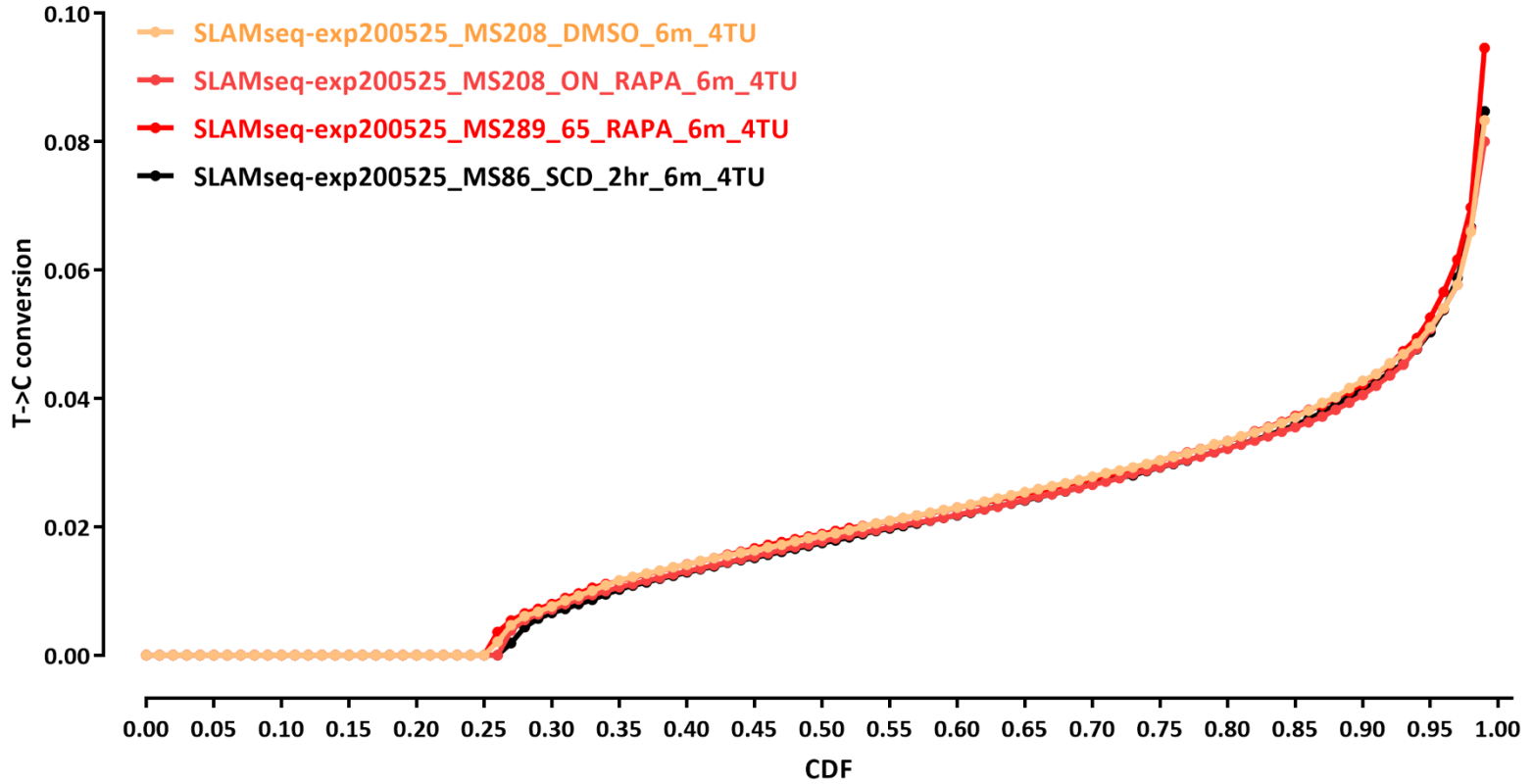
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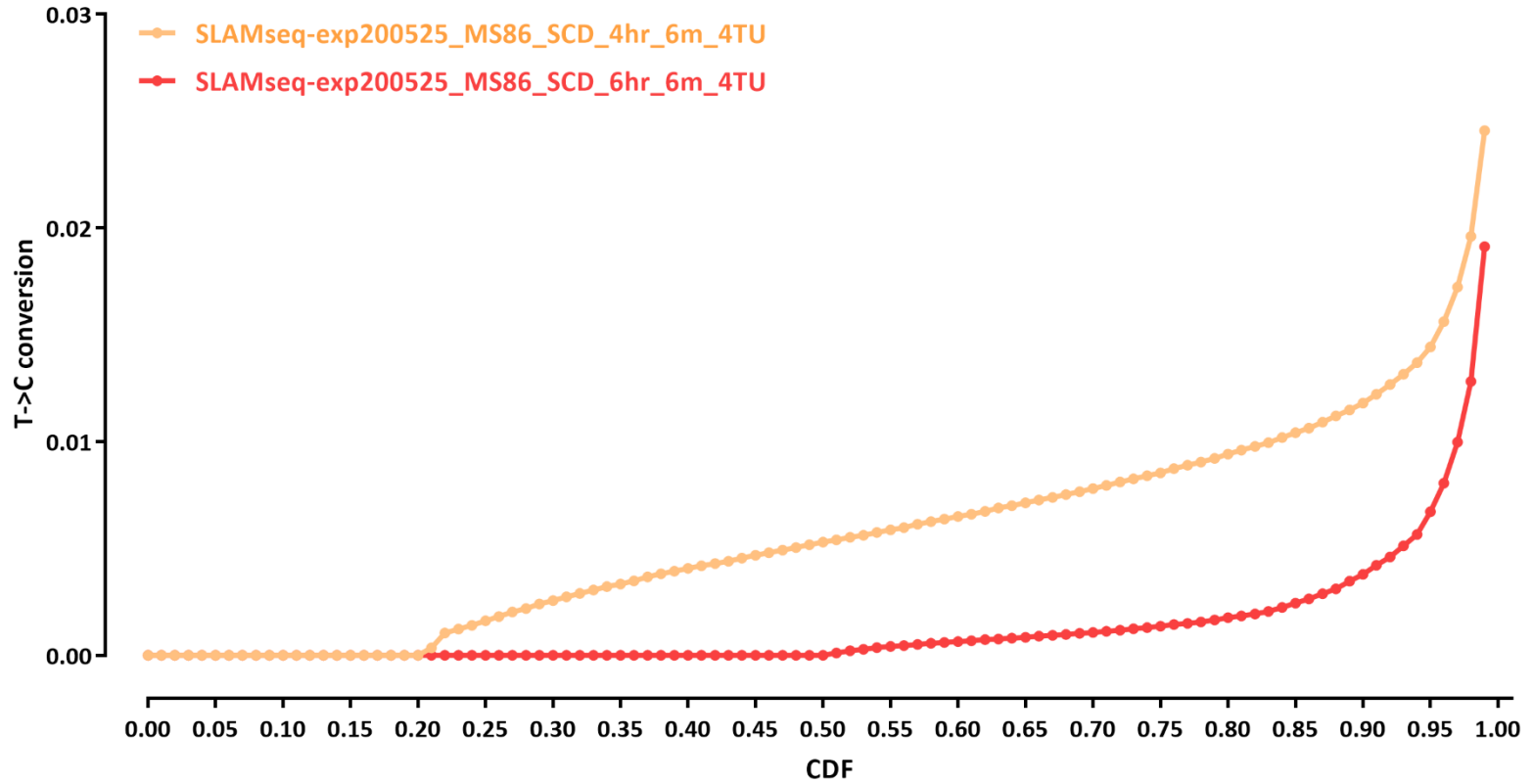
S_cerevisiae 3



S_pombe 3



S_cerevisiae 4



S_pombe 4

