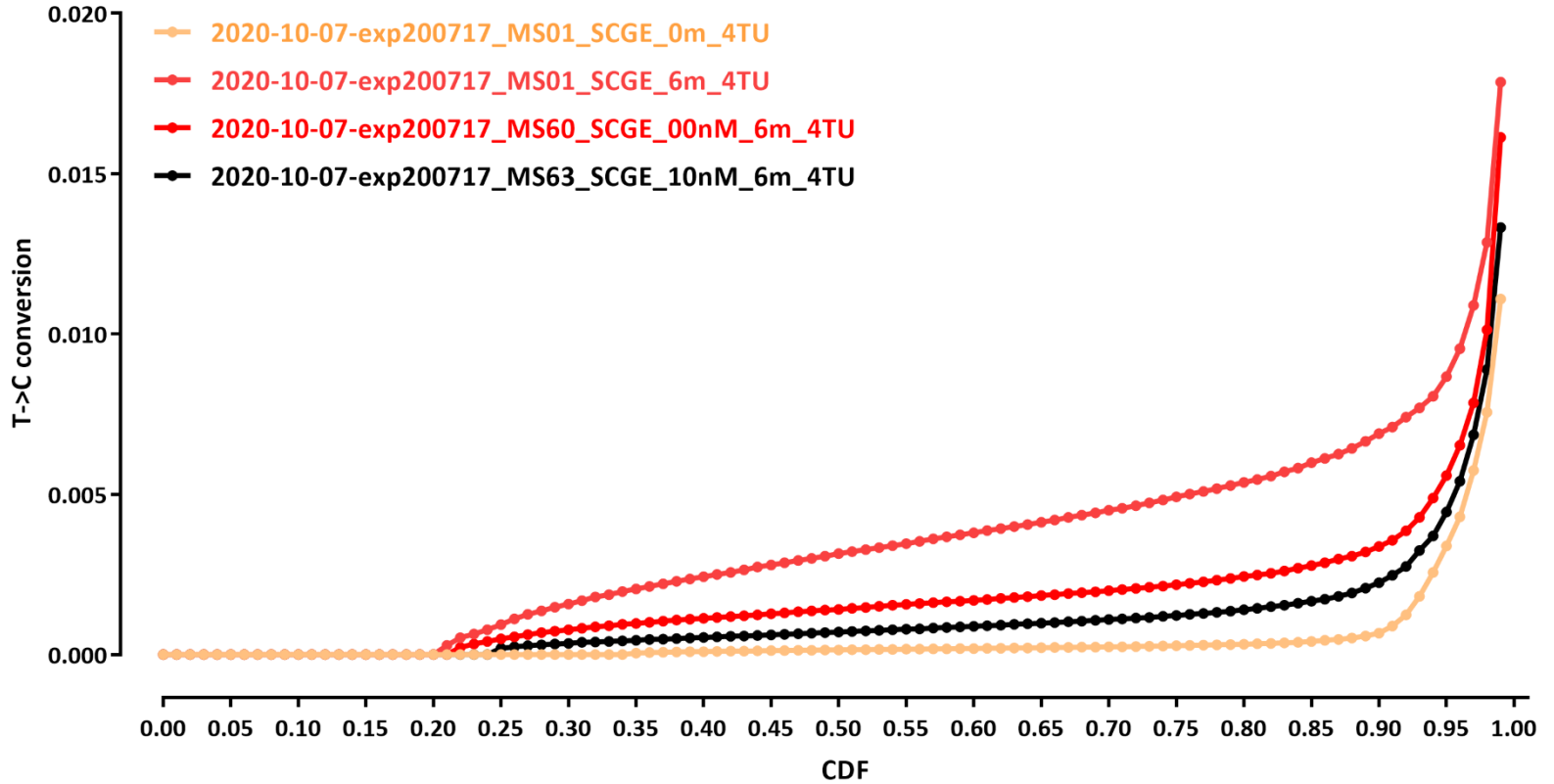


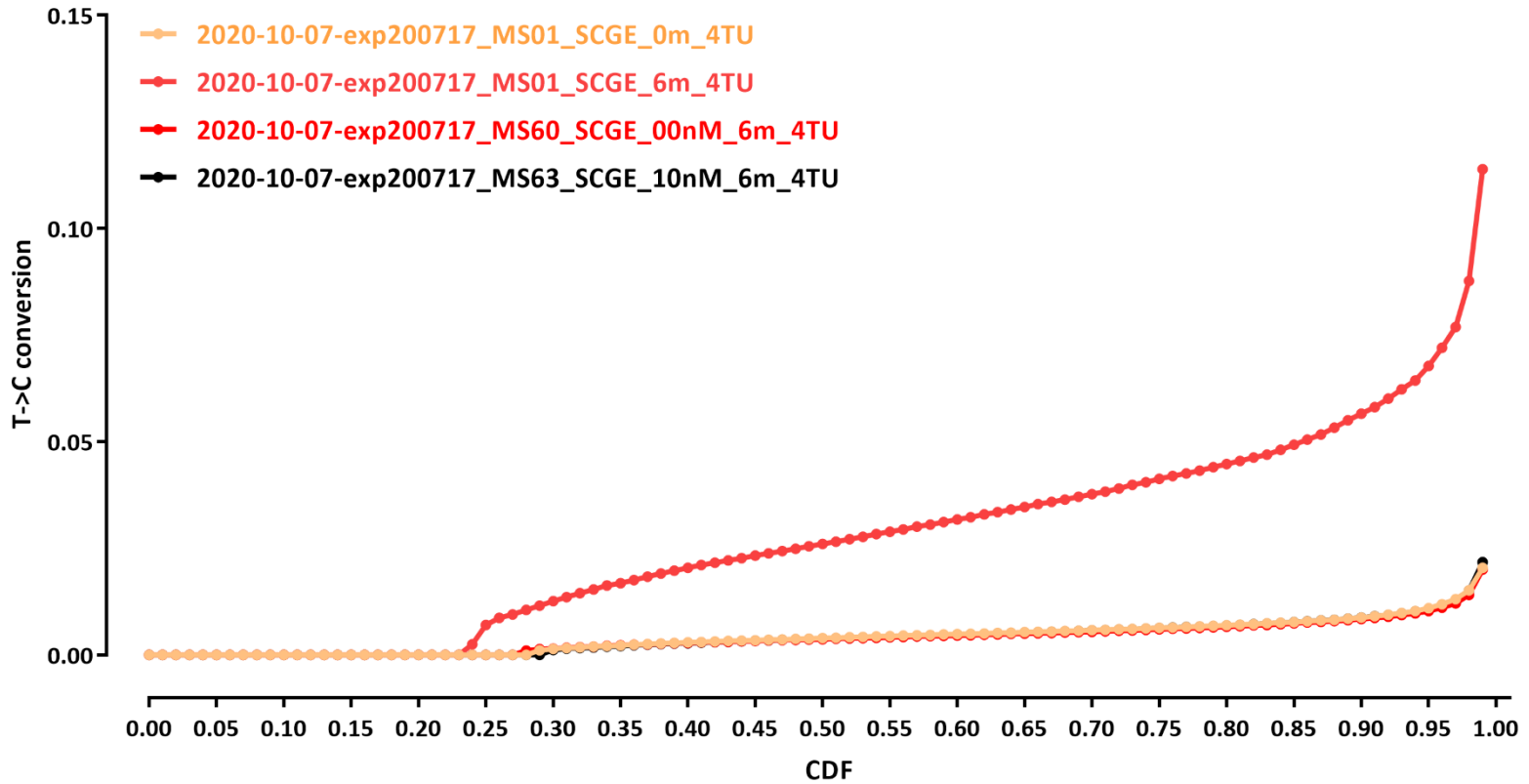
Mapping stats

Library	Read Length	Mapping	Species	Raw fragments	Complexity	Unique	Unique Splices	Multi	Multi Splices	Fraction mapped
2020-10-07-exp200717_MS01_SCGE_0m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	7,685,460	0.35	8,931,188		3,016		0.58
2020-10-07-exp200717_MS01_SCGE_0m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.pombe	7,685,460	0.42	3,782,677		4,399		0.25
2020-10-07-exp200717_MS01_SCGE_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	7,927,971	0.34	9,007,990		6,182		0.57
2020-10-07-exp200717_MS01_SCGE_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.pombe	7,927,971	0.42	4,069,786		6,668		0.26
2020-10-07-exp200717_MS60_SCGE_00nM_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	9,416,687	0.31	10,833,381		3,959		0.58
2020-10-07-exp200717_MS60_SCGE_00nM_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.pombe	9,416,687	0.40	4,671,417		5,746		0.25
2020-10-07-exp200717_MS63_SCGE_10nM_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	6,878,095	0.35	7,691,194		2,721		0.56
2020-10-07-exp200717_MS63_SCGE_10nM_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.pombe	6,878,095	0.42	3,746,446		4,328		0.27
2020-10-07-exp200717_MS63_SCGE_30nM_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	6,768,765	0.36	7,823,706		2,729		0.58
2020-10-07-exp200717_MS63_SCGE_30nM_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.pombe	6,768,765	0.43	3,512,457		4,004		0.26
2020-10-07-exp200717_MS80_SCGE_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	7,557,841	0.34	8,302,575		3,151		0.55
2020-10-07-exp200717_MS80_SCGE_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.pombe	7,557,841	0.42	3,596,585		4,307		0.24
2020-10-07-exp200717_MS81_SCGE_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	8,759,615	0.31	10,298,861		3,670		0.59
2020-10-07-exp200717_MS81_SCGE_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.pombe	8,759,615	0.40	4,146,583		5,050		0.24
2020-10-07-exp200803_MS207_SCD_40_RAPA_0m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	9,299,637	0.31	10,874,928		3,482		0.58
2020-10-07-exp200803_MS207_SCD_40_RAPA_0m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.pombe	9,299,637	0.40	4,725,119		5,501		0.25
2020-10-07-exp200803_MS207_SCD_40_RAPA_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	7,811,720	0.33	9,166,762		3,226		0.59
2020-10-07-exp200803_MS207_SCD_40_RAPA_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.pombe	7,811,720	0.43	3,770,936		4,691		0.24
2020-10-07-exp200803_MS281_SCD_40_RAPA_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	7,288,119	0.32	8,573,110		2,857		0.59
2020-10-07-exp200803_MS281_SCD_40_RAPA_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.pombe	7,288,119	0.42	3,501,312		4,101		0.24
2020-10-07-exp200803_MS282_SCD_40_RAPA_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	7,290,734	0.33	8,294,581		2,839		0.57
2020-10-07-exp200803_MS282_SCD_40_RAPA_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.pombe	7,290,734	0.43	3,405,040		3,834		0.23
2020-10-07-exp200803_MS289_SCD_40_RAPA_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	7,819,577	0.46	1,295,635		459		0.08
2020-10-07-exp200803_MS289_SCD_40_RAPA_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.pombe	7,819,577	0.29	11,341,163		13,687		0.73
2020-10-07-exp200803_MS294_SCD_40_RAPA_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	8,148,608	0.33	7,792,878		2,638		0.48
2020-10-07-exp200803_MS294_SCD_40_RAPA_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.pombe	8,148,608	0.41	4,430,873		5,038		0.27
2020-10-07-exp200803_MS299_SCD_40_RAPA_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.cerevisiae	6,630,109	0.35	7,287,722		2,615		0.55
2020-10-07-exp200803_MS299_SCD_40_RAPA_6m_4TU	2x150	slamdunk-0.4.3, transcriptome	S.pombe	6,630,109	0.43	3,610,740		4,742		0.27

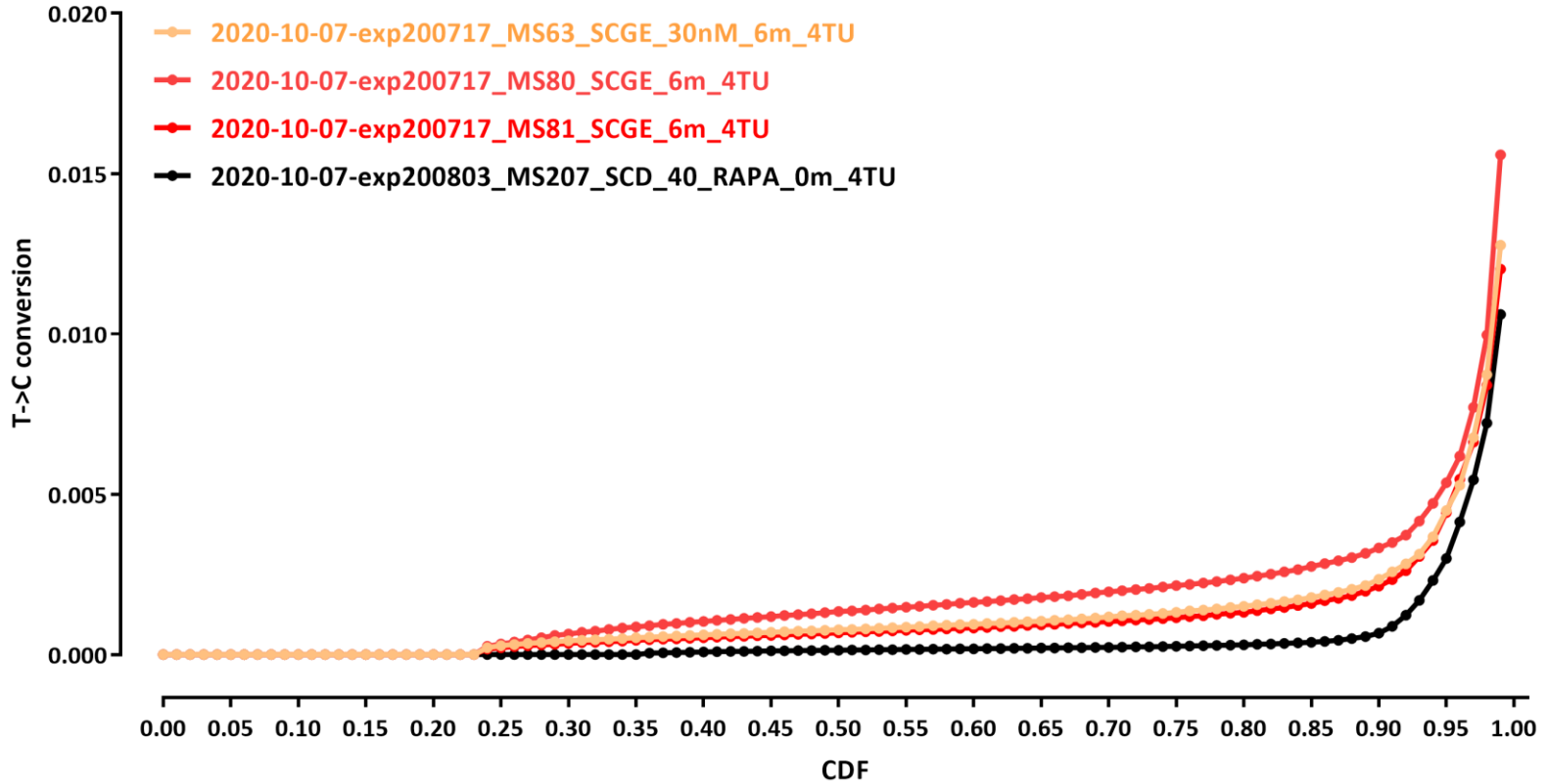
S_cerevisiae 1



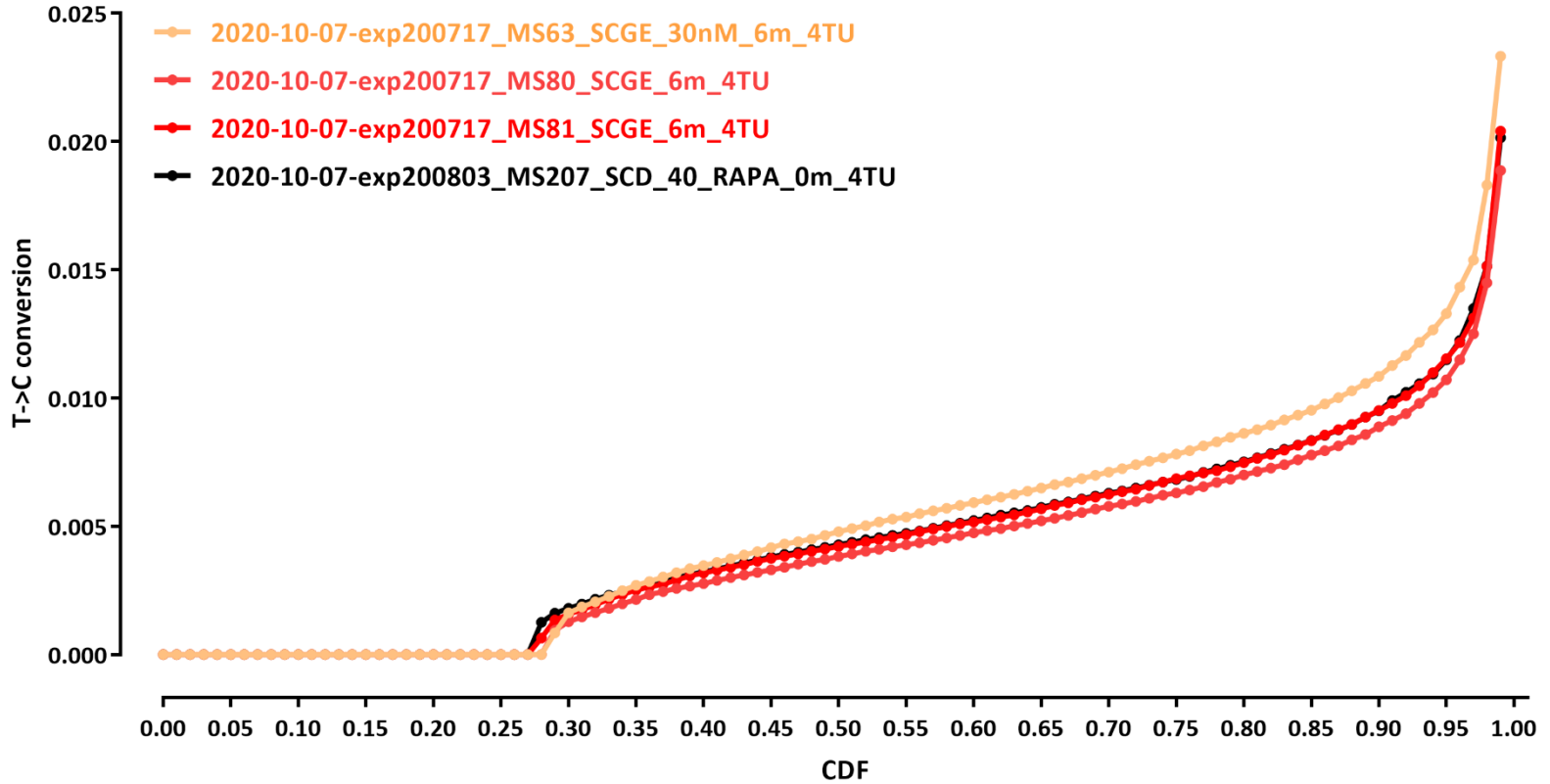
S_pombe 1



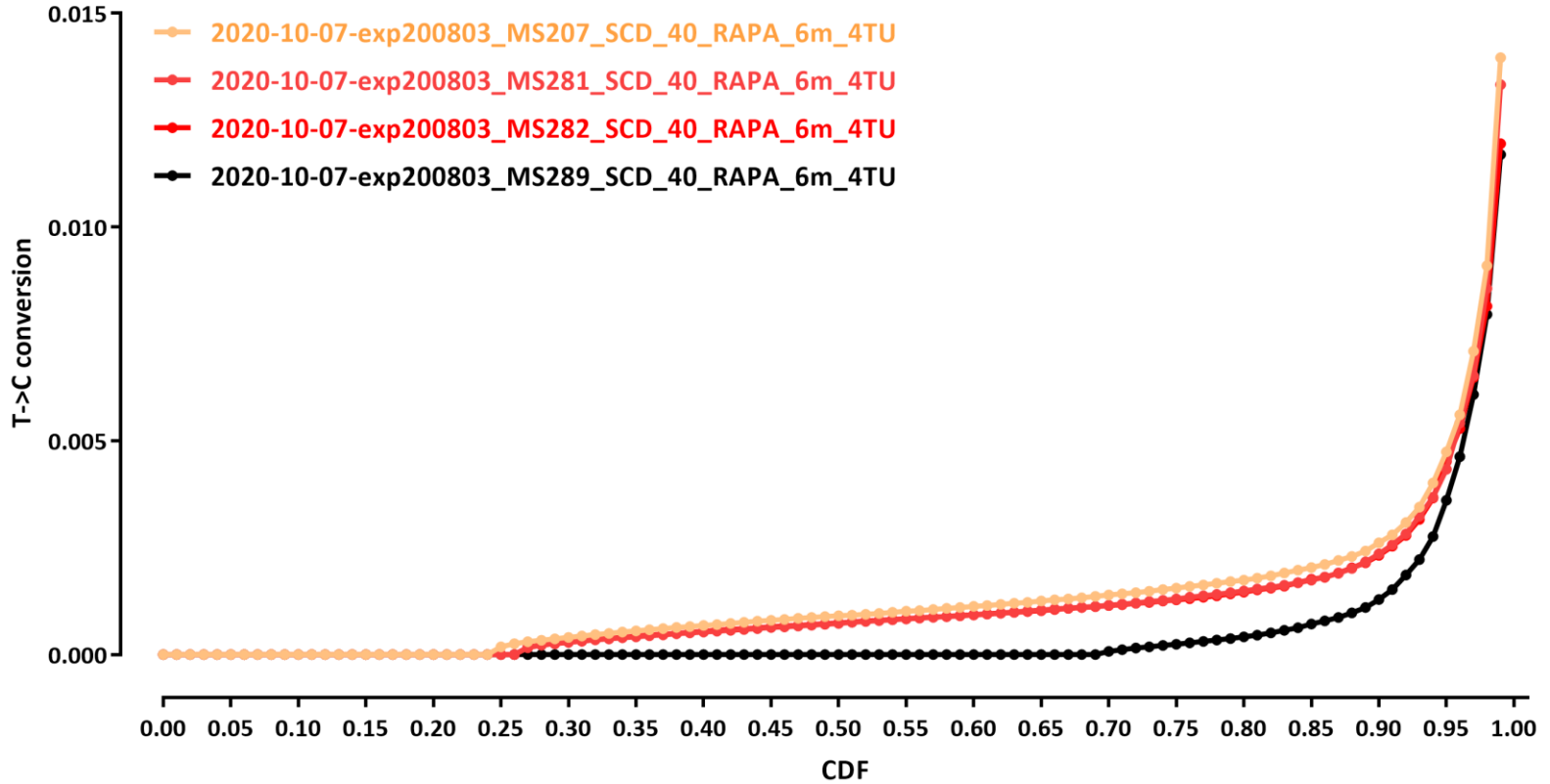
S_cerevisiae 2



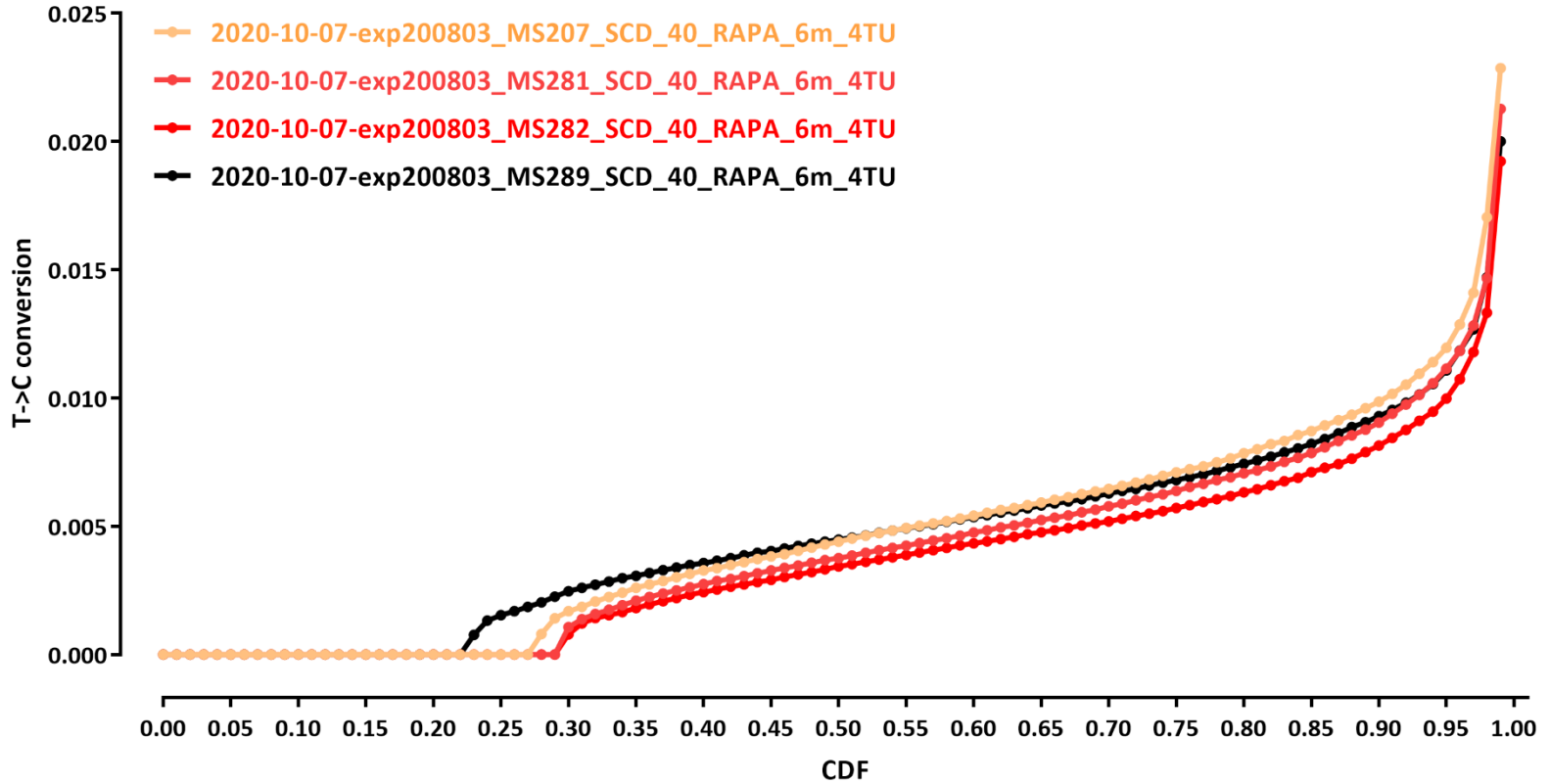
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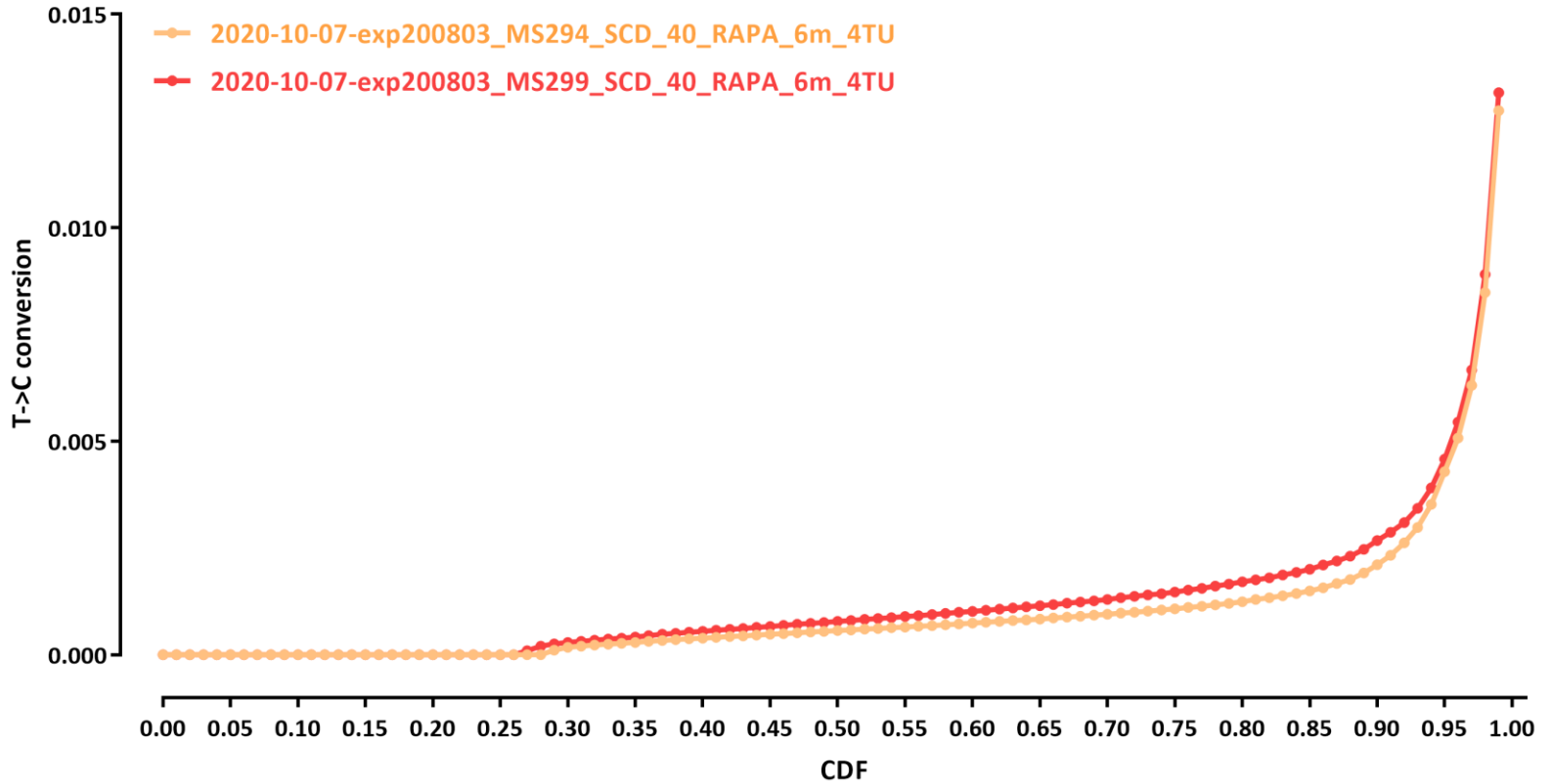
S_cerevisiae 3



S_pombe 3



S_cerevisiae 4



S_pombe 4

