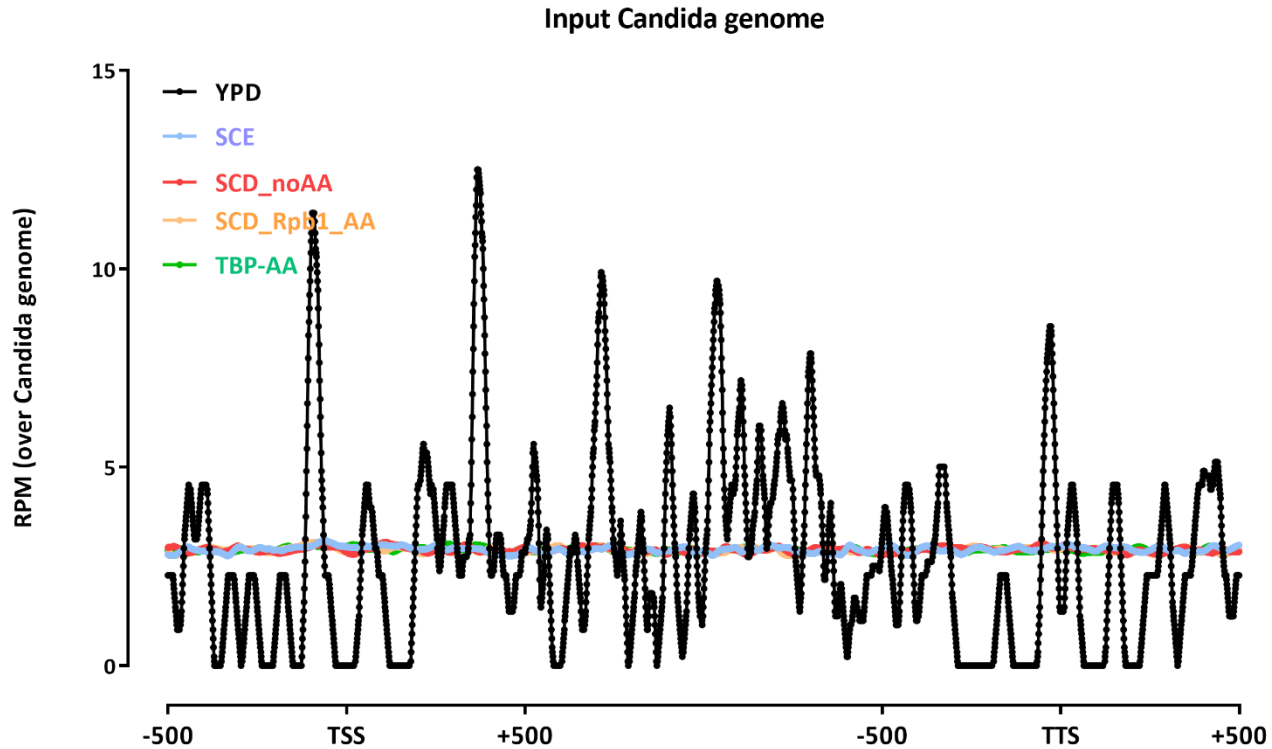


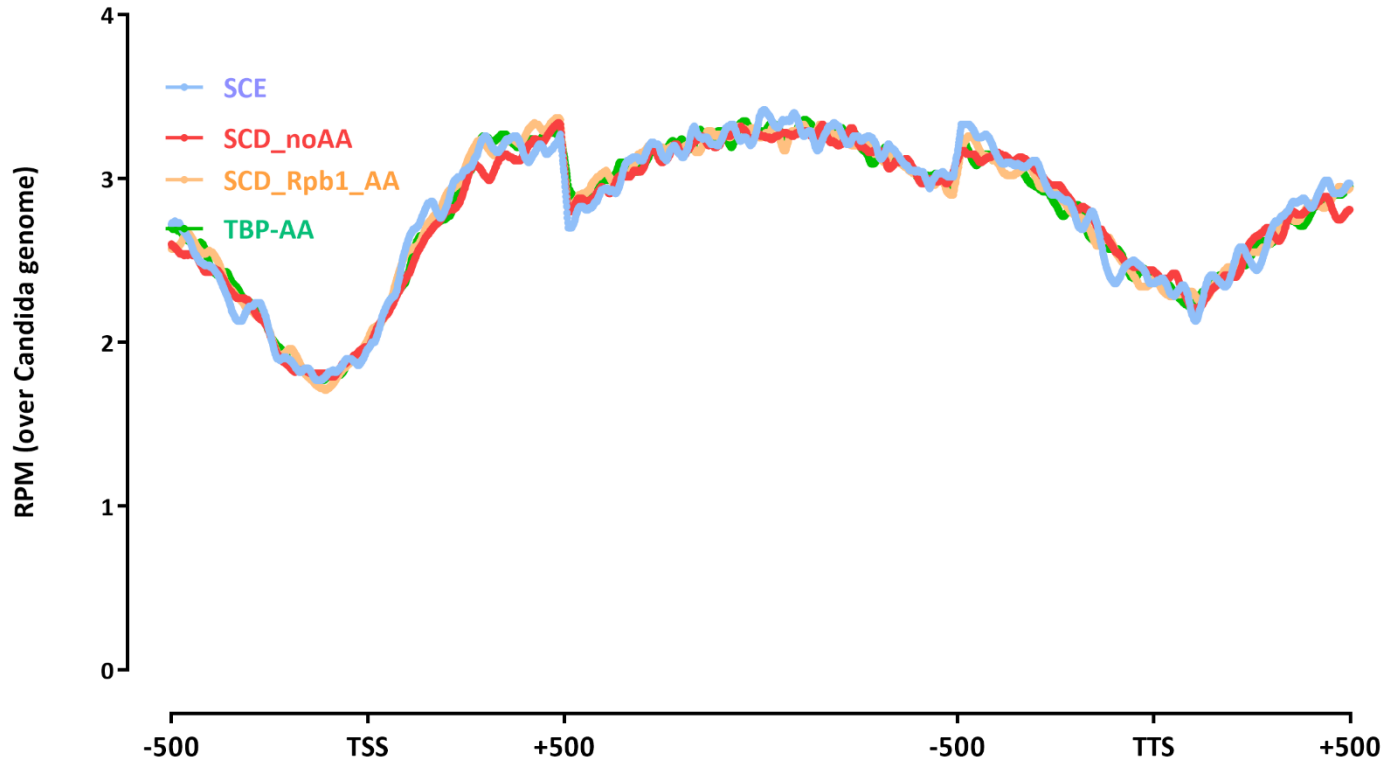
Dataset stats

Library	Raw fragments	Species	2x36mers		1x36mers		Mappability Corrected RPKM	S.cerevisiae/Candida ratio
			Unique non-chrM reads	Complexity	Unique non-chrM reads	Complexity		
2018-02-13-A-YPD-input	603,120	sacCer3	890,262	0.98	449,717	0.98	86.97	4,851.20
	603,120	Candida	162	1.00	98	1.00	0.02	
2018-02-13-B-YPD-Pol2pS2	739,280	sacCer3	1,273,346	0.95	644,537	0.94	86.97	6,251.11
	739,280	Candida	178	1.00	109	1.00	0.01	
2018-02-13-C-SCE_input	780,896	sacCer3	940,824	0.98	476,336	0.98	68.08	3.81
	780,896	Candida	260,900	0.99	132,276	0.99	17.88	
2018-02-13-D-SCE-Pol2pS2	842,765	sacCer3	1,225,412	0.96	619,794	0.96	74.11	6.08
	842,765	Candida	212,704	0.98	107,693	0.98	12.18	
2018-02-13-E-SCE-H3	778,957	sacCer3	989,158	0.98	501,680	0.98	68.38	3.88
	778,957	Candida	269,140	0.99	136,529	0.99	17.60	
2018-02-13-F-SCE-CTD	860,877	sacCer3	1,298,016	0.96	658,411	0.95	75.85	7.20
	860,877	Candida	190,596	0.99	96,702	0.99	10.54	
2018-02-13-G-YPD-CTD	566,772	sacCer3	993,618	0.95	503,125	0.95	86.96	3,092.31
	566,772	Candida	296	1.00	172	1.00	0.03	
2018-02-13-H-SCD_noAA-input	667,562	sacCer3	645,266	0.98	326,634	0.98	52.43	1.60
	667,562	Candida	424,992	0.98	215,270	0.98	32.69	
2018-02-13-I-SCD_Rpb1_AA-input	606,366	sacCer3	571,242	0.99	289,240	0.98	50.28	1.45
	606,366	Candida	416,914	0.98	211,192	0.98	34.72	
2018-02-13-J-SCD_TBP-AA-input	868,003	sacCer3	814,286	0.98	411,515	0.98	51.09	1.50
	868,003	Candida	571,646	0.98	289,140	0.98	33.96	
2018-02-13-K-SCD_noAA-H3	589,040	sacCer3	541,558	0.98	275,845	0.98	47.57	1.28
	589,040	Candida	447,944	0.99	228,581	0.98	37.29	
2018-02-13-L-SCD_Rpb1_AA-H3	716,259	sacCer3	604,192	0.98	307,202	0.98	42.11	0.99
	716,259	Candida	643,024	0.98	327,346	0.98	42.45	
2018-02-13-M-SCD_TBP-AA-H3	561,170	sacCer3	460,992	0.98	233,911	0.98	42.46	1.01
	561,170	Candida	482,094	0.99	245,237	0.98	42.11	
2018-02-13-N-YPD-Rpb3	627,400	sacCer3	1,093,878	0.95	551,792	0.94	86.93	1,611.40
	627,400	Candida	662	1.00	362	1.00	0.05	
2018-02-13-O-SCE-Rpb3	703,670	sacCer3	1,169,660	0.96	592,993	0.95	85.10	47.63
	703,670	Candida	25,890	0.99	13,162	0.99	1.79	
2018-02-13-P-SCD_noAA-CTD	1,142,307	sacCer3	1,230,878	0.95	622,354	0.95	51.42	1.53
	1,142,307	Candida	851,708	0.96	430,557	0.96	33.65	
2018-02-13-Q-SCD_Rpb1_AA-CTD	757,685	sacCer3	149,888	0.99	75,778	0.99	9.16	0.12
	757,685	Candida	1,274,360	0.95	643,900	0.94	73.62	
2018-02-13-R-SCD_TBP-AA-CTD	556,727	sacCer3	105,842	0.99	53,696	0.99	8.96	0.12
	556,727	Candida	923,868	0.95	467,501	0.95	73.81	
2018-02-13-S-SCD_noAA-Rpb3	826,610	sacCer3	1,424,848	0.94	722,292	0.93	84.21	32.14
	826,610	Candida	46,910	0.99	23,761	0.99	2.62	
2018-02-13-T-YPD-Pol2pS5_C	614,811	sacCer3	1,076,346	0.95	544,717	0.94	86.85	667.26
	614,811	Candida	1,652	0.99	863	0.99	0.13	
2018-02-13-U-YPD-Pol2pS5_D	635,448	sacCer3	1,103,120	0.95	557,787	0.94	86.75	389.73
	635,448	Candida	2,940	1.00	1,513	1.00	0.22	
2018-02-13-V-YPD-Pol2pS5_E	491,950	sacCer3	870,774	0.95	439,526	0.95	86.86	719.26
	491,950	Candida	1,254	0.99	646	1.00	0.12	
2018-02-13-W-SCD_Rpb1_AA-Rpb3	586,675	sacCer3	879,138	0.97	443,443	0.96	74.98	6.60
	586,675	Candida	140,978	0.98	71,030	0.98	11.36	
2018-02-13-X-SCD_TBP-AA-Rpb3	565,781	sacCer3	581,678	0.98	293,236	0.98	60.16	2.37
	565,781	Candida	260,278	0.97	130,775	0.96	25.38	

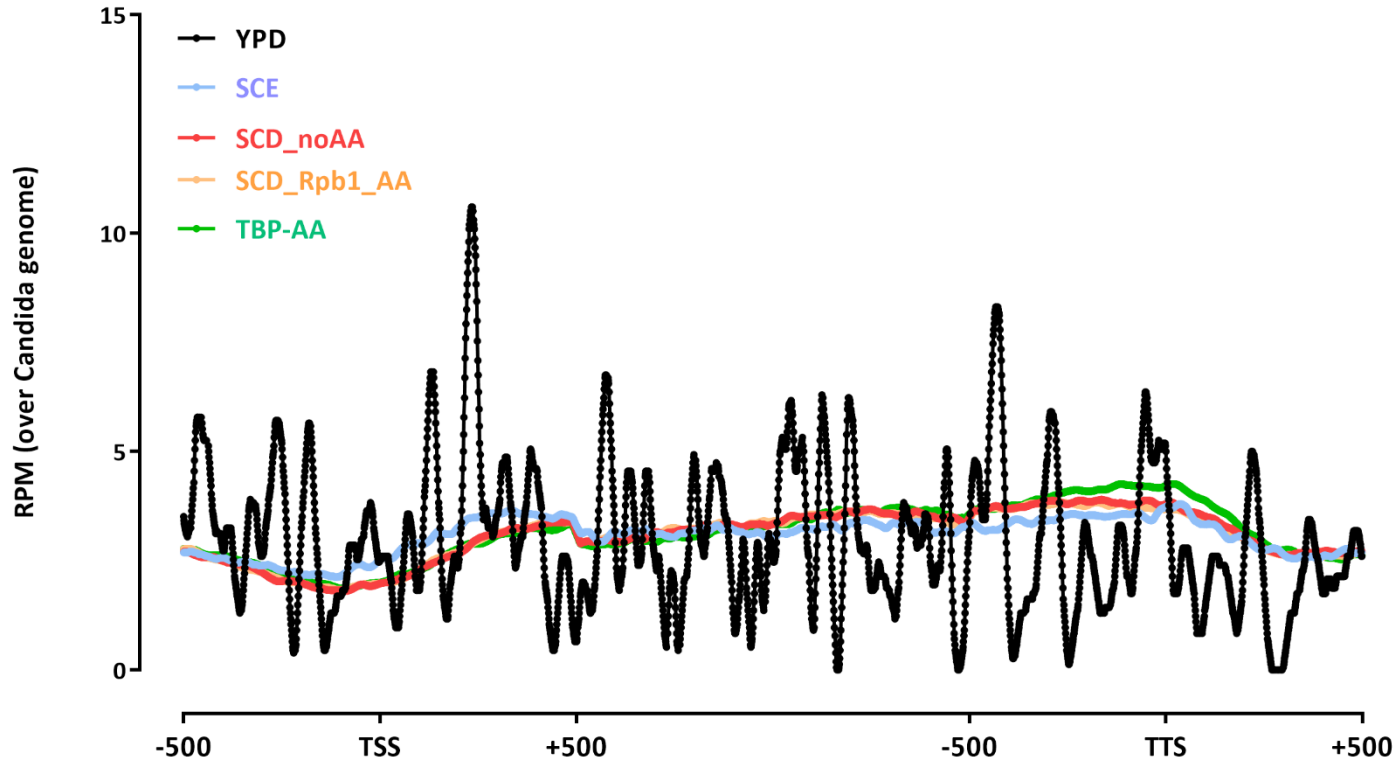
Average gene profiles without renormalization



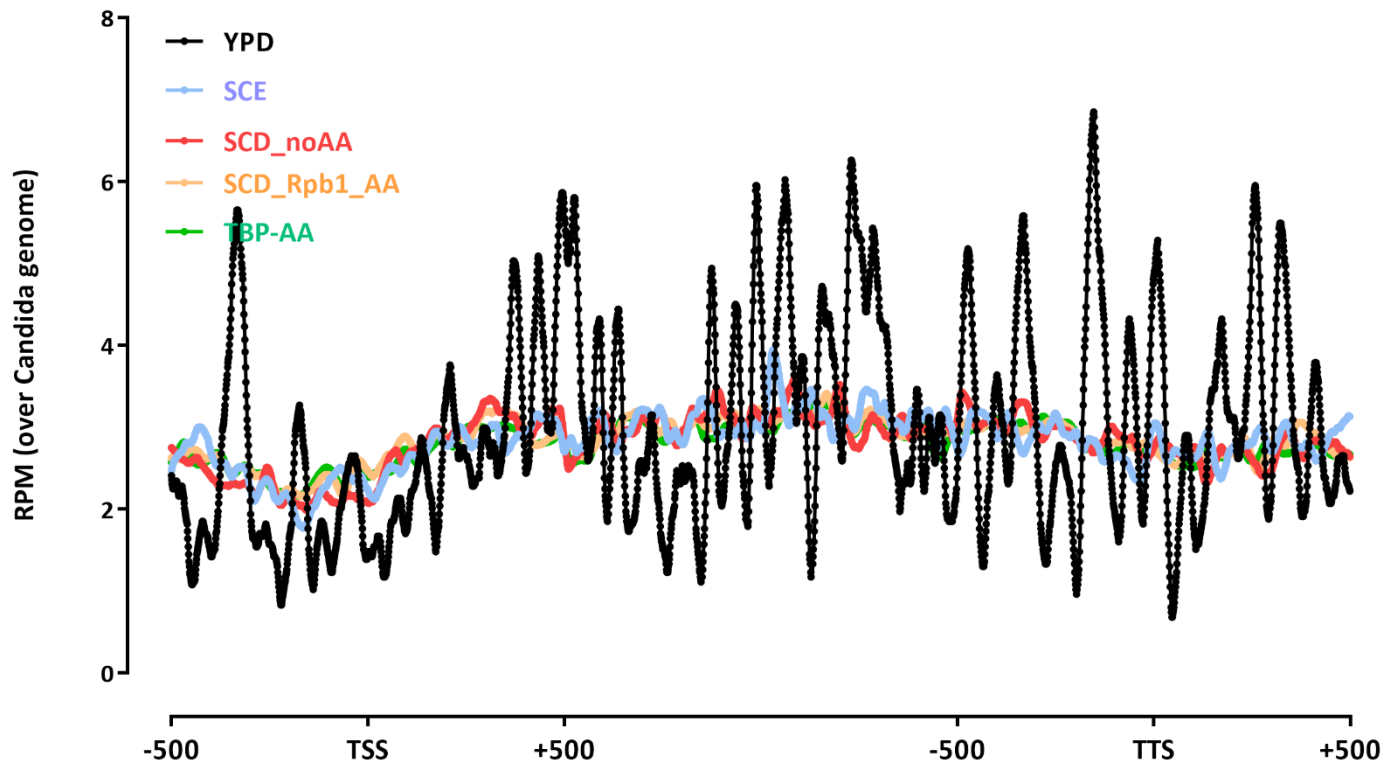
H3 Candida genome



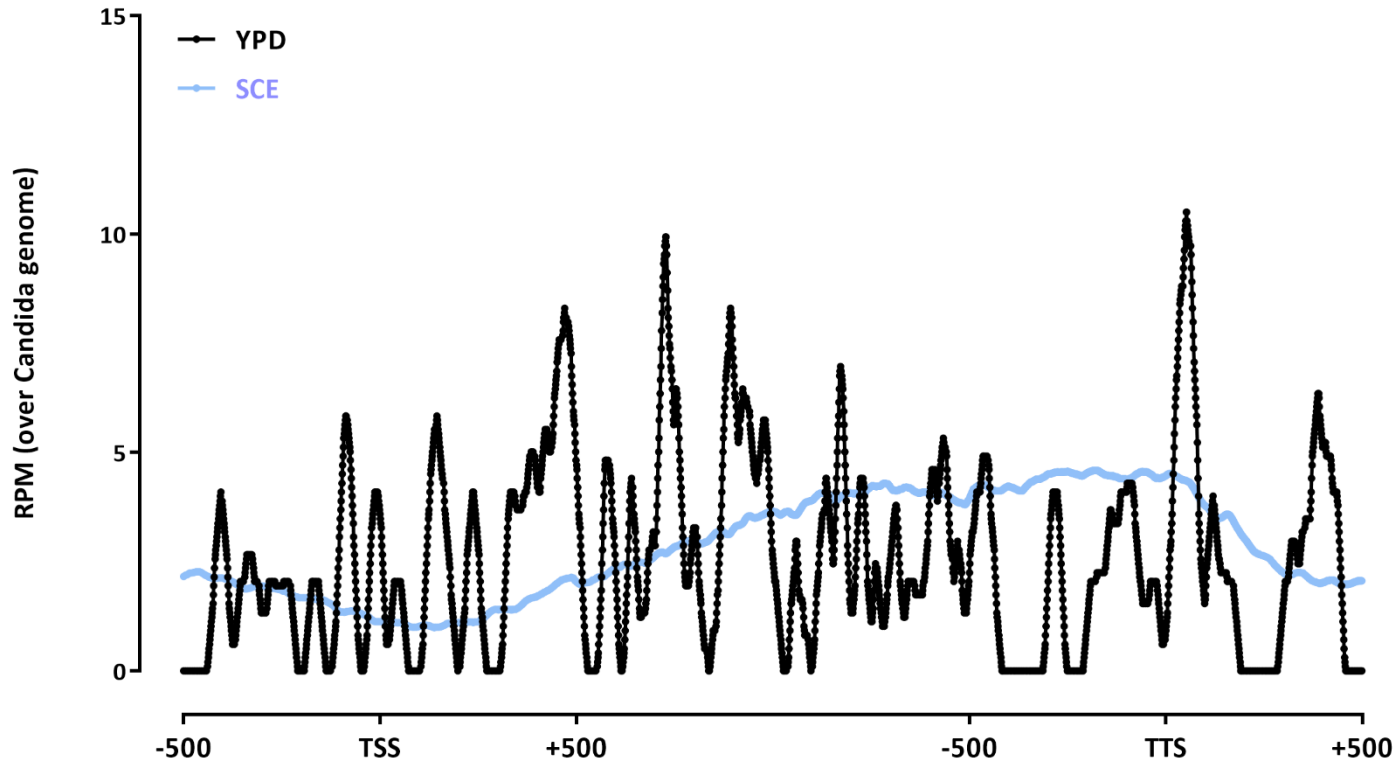
CTD Candida genome



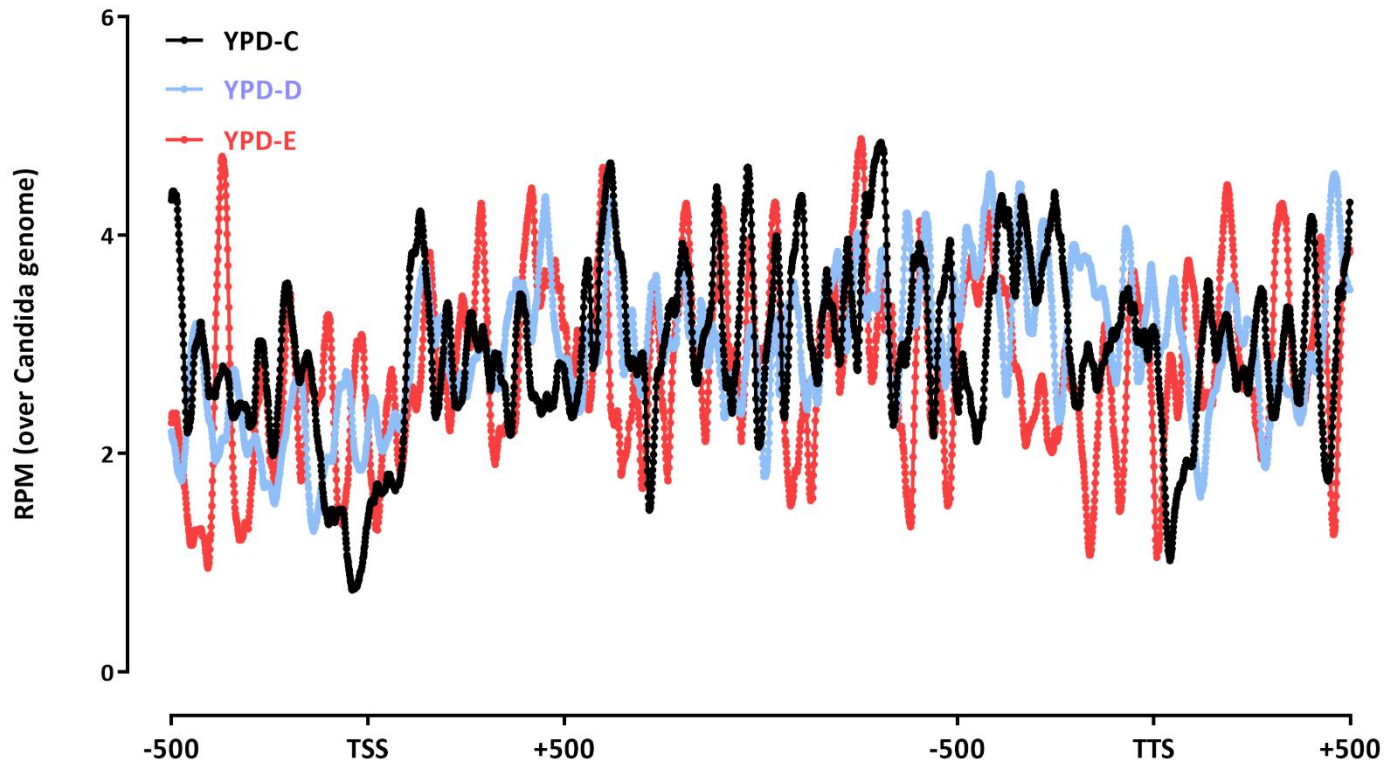
Rpb3 Candida genome



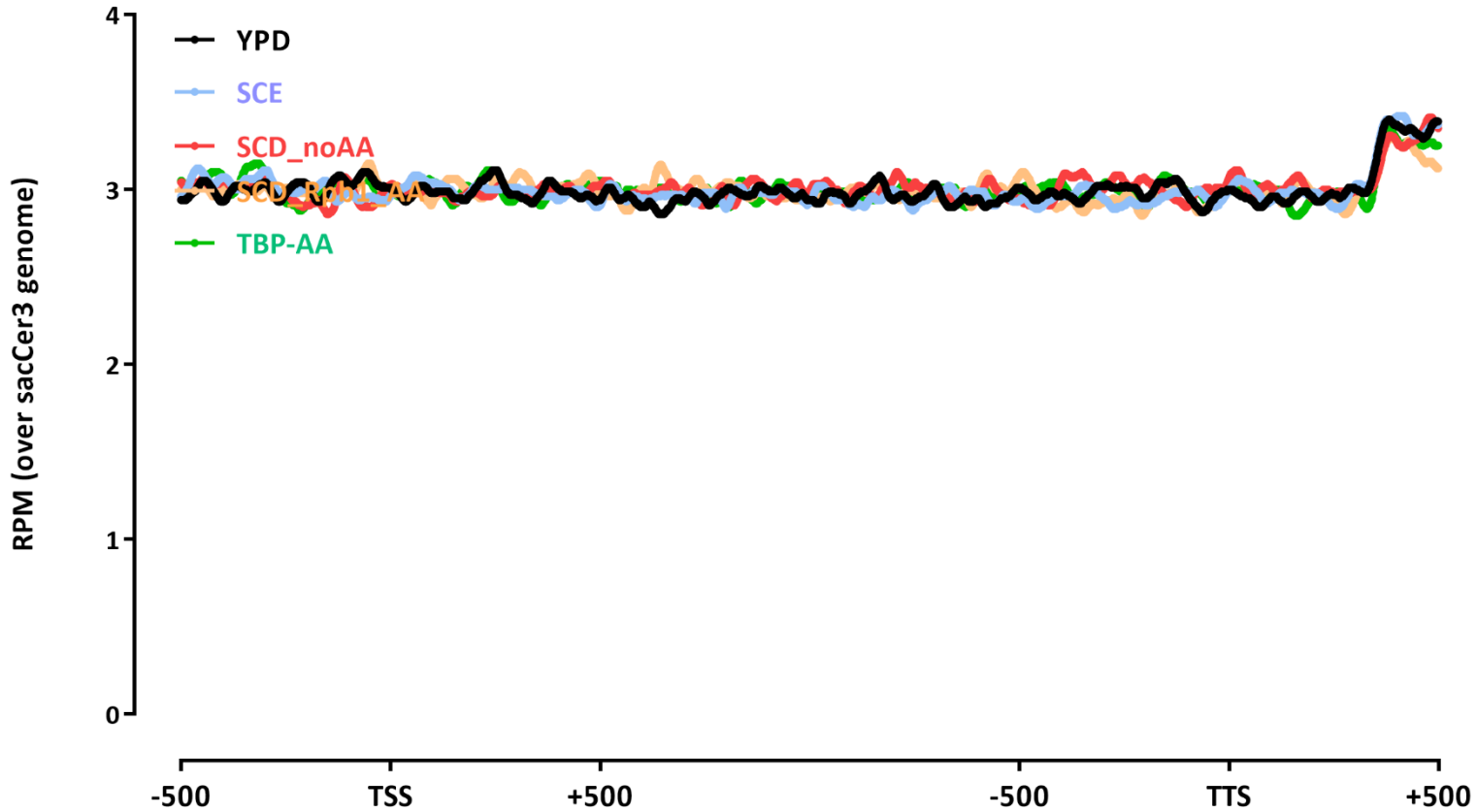
Pol2pS2 Candida genome



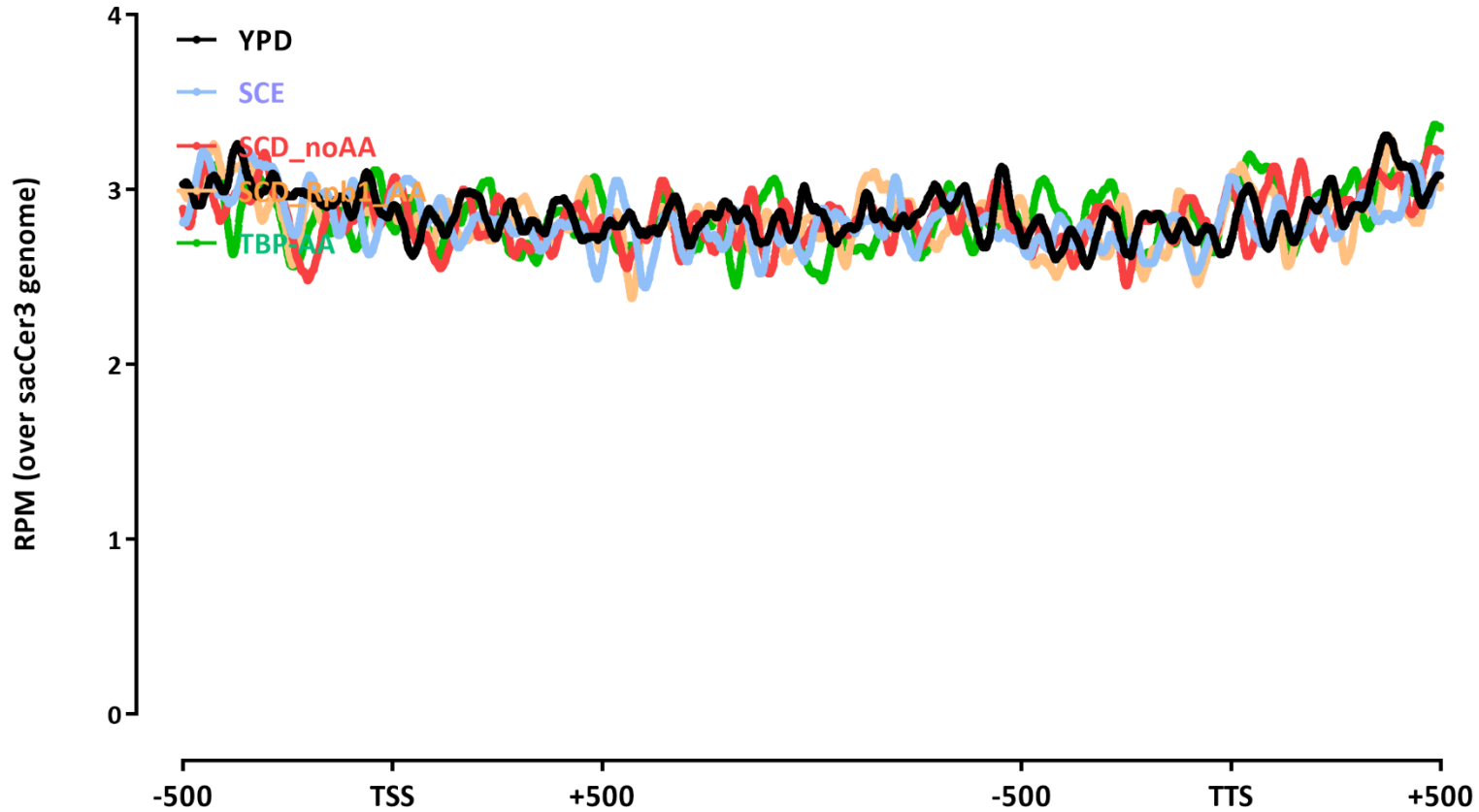
Pol2pS5 Candida genome



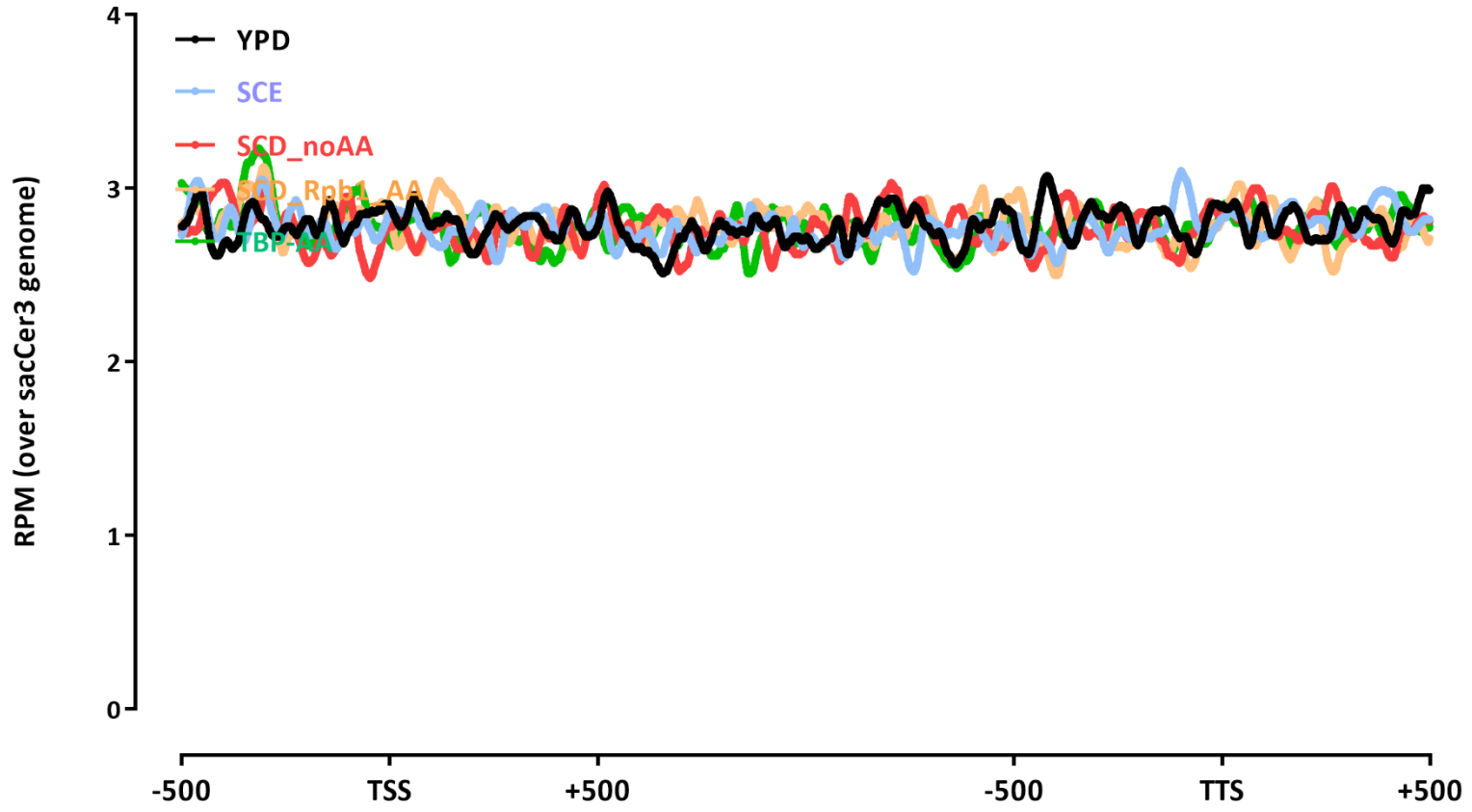
Input sacCer3 genome



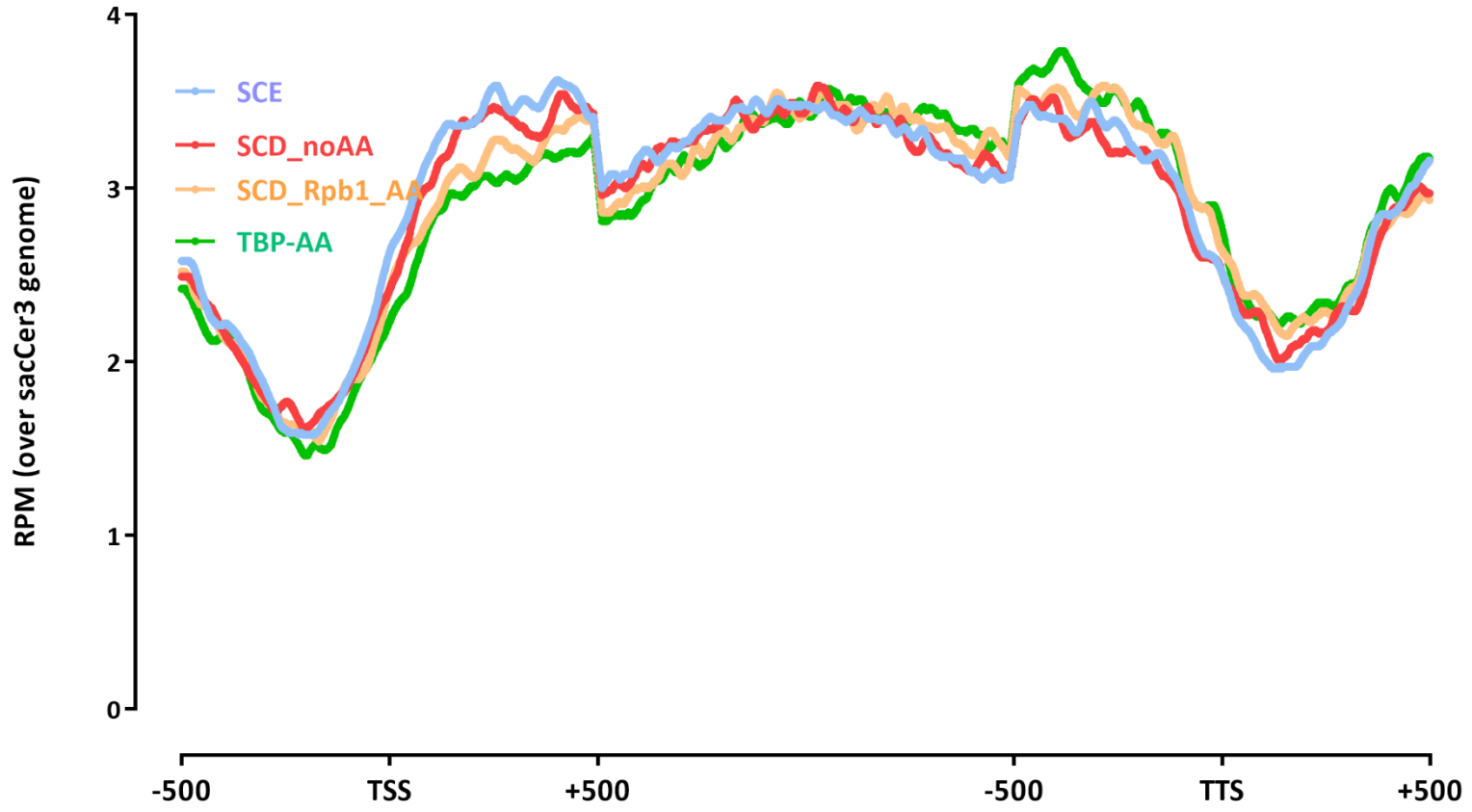
Input sacCer3 genome top expression quintile



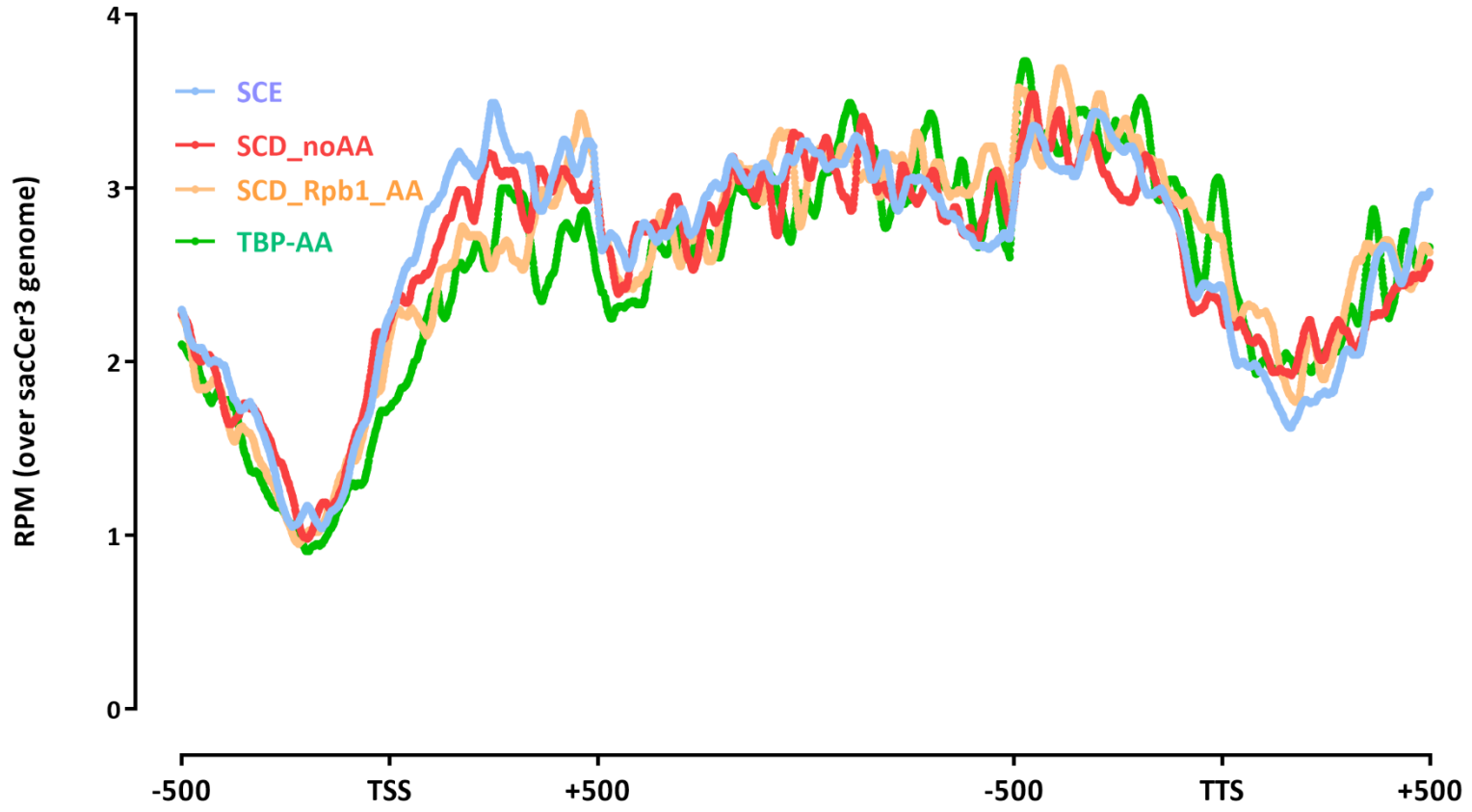
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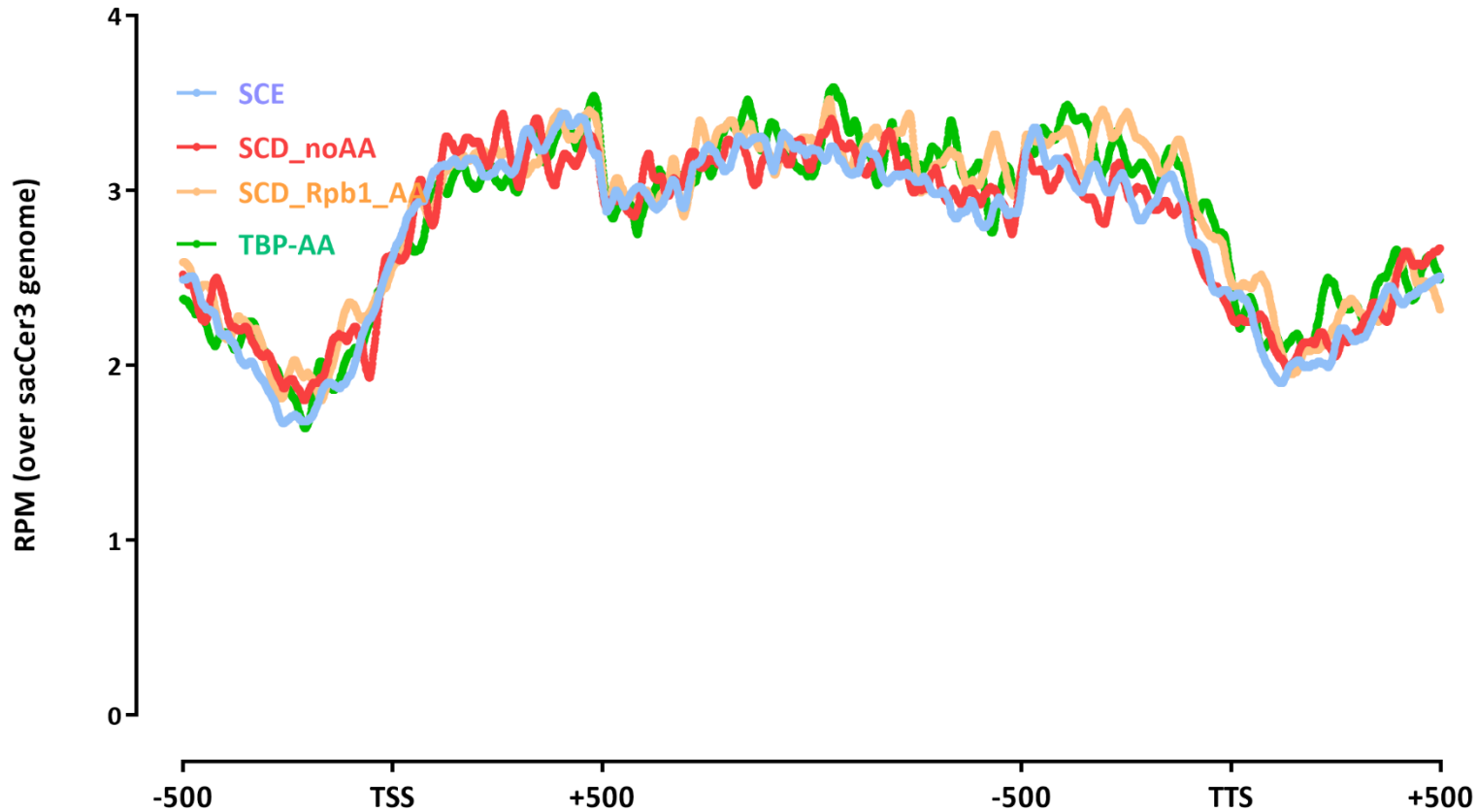
H3 sacCer3 genome



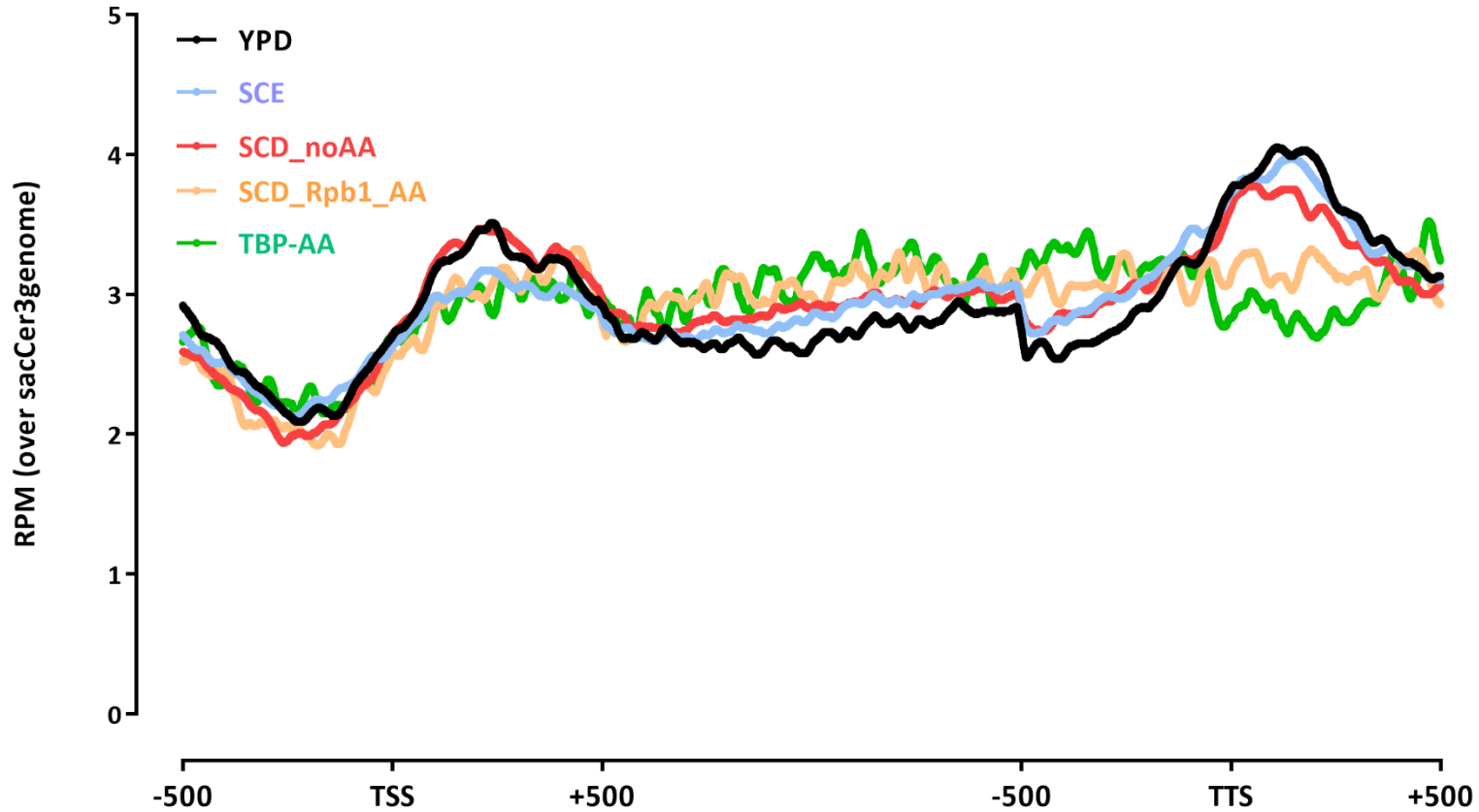
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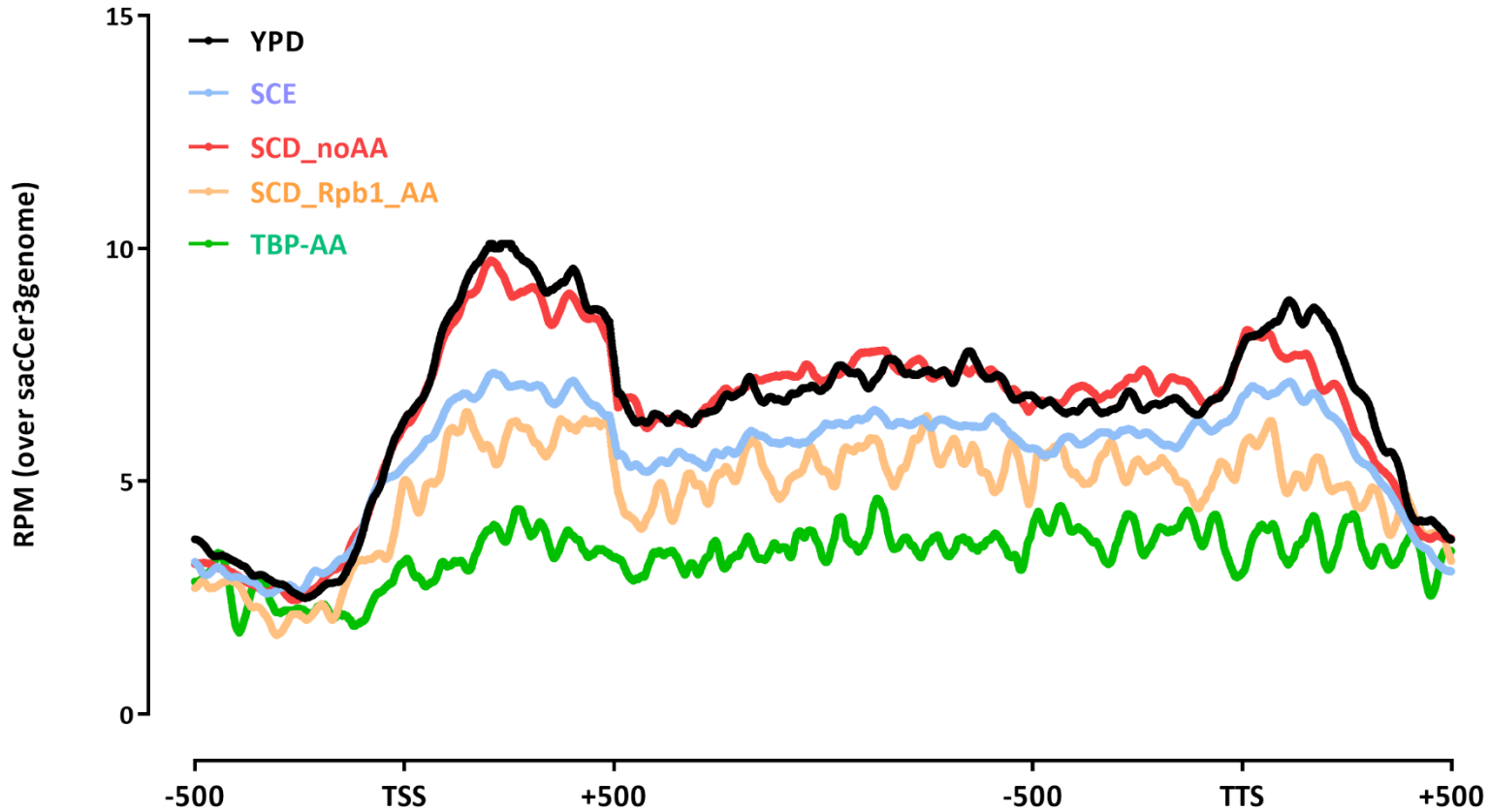
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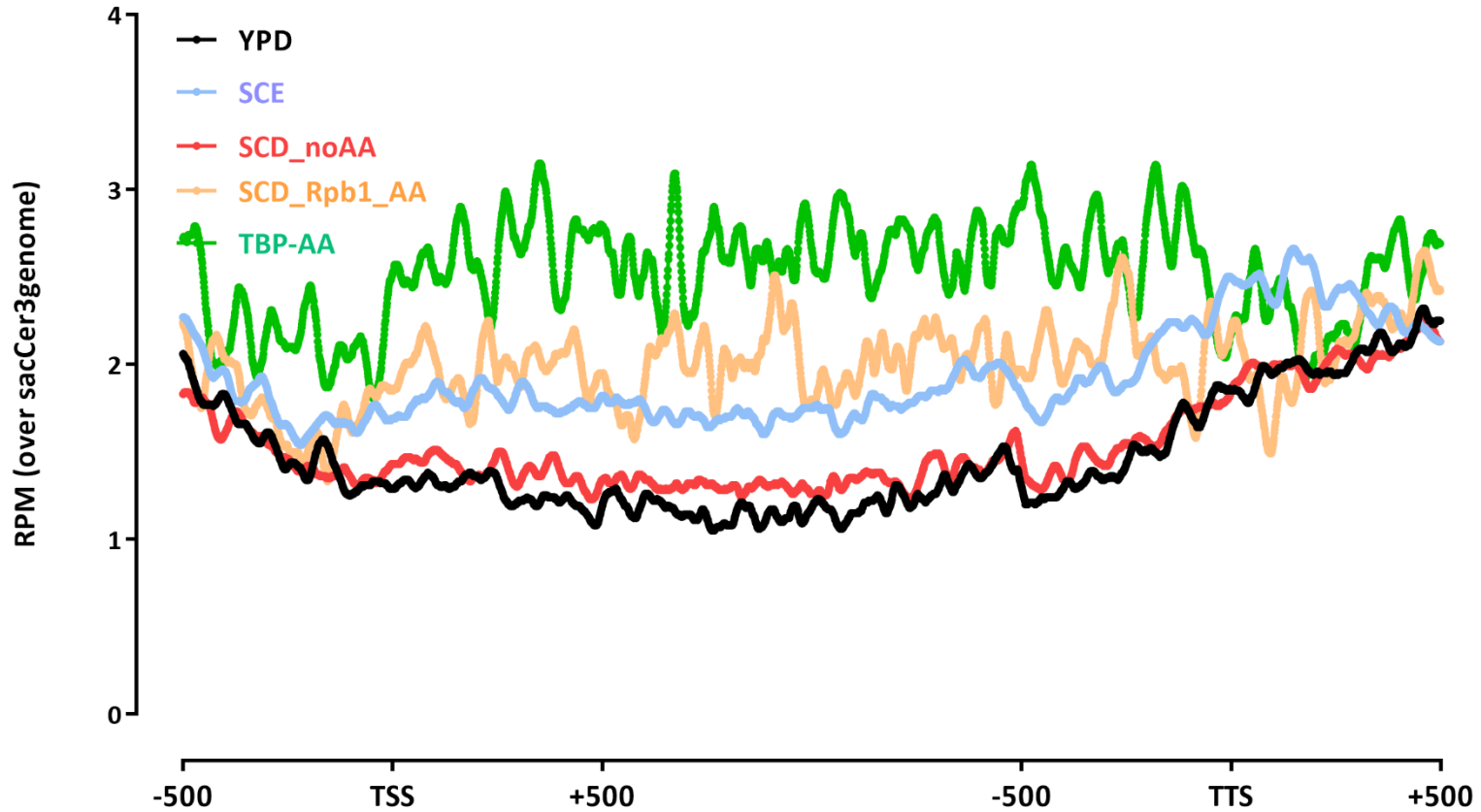
CTD sacCer3 genome



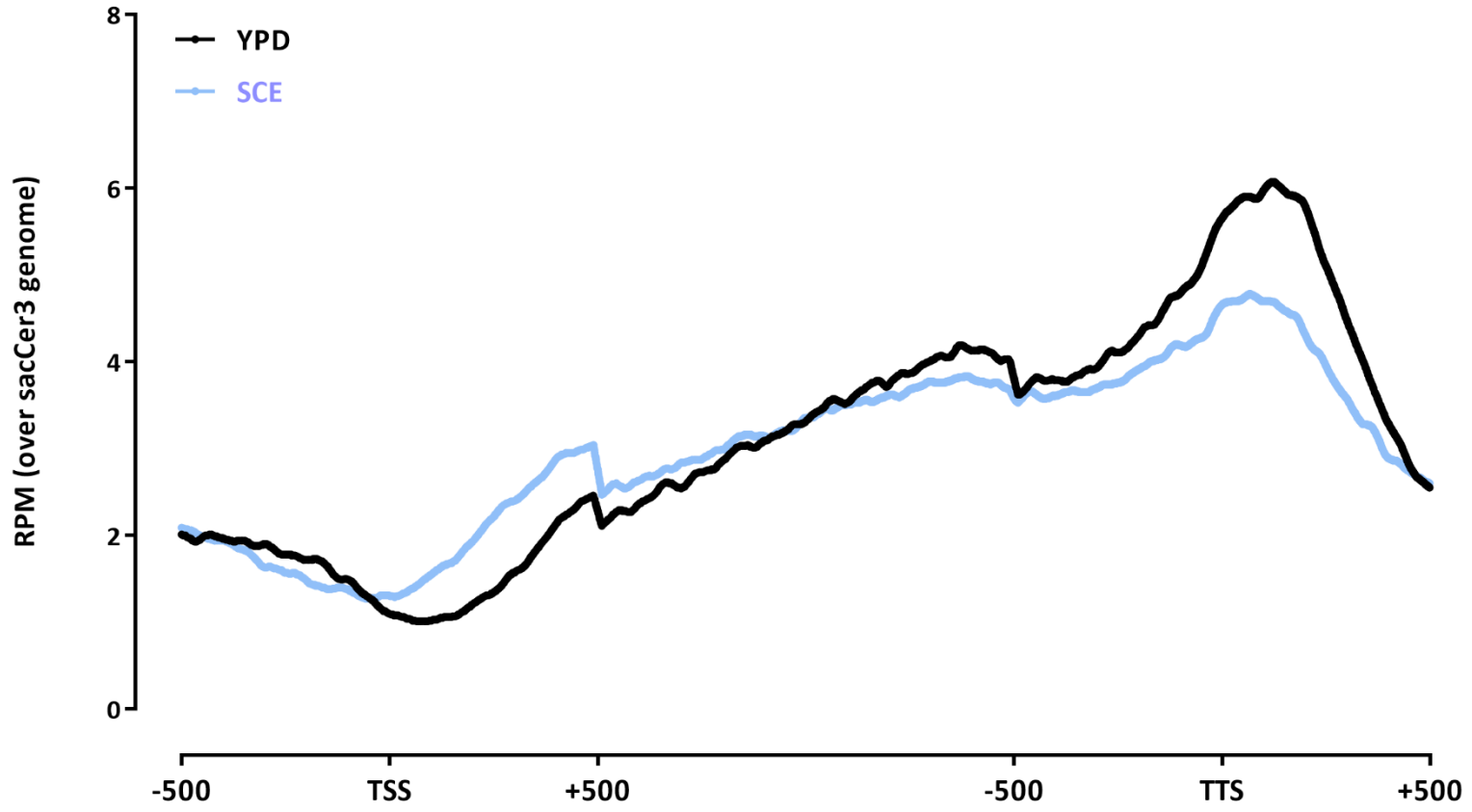
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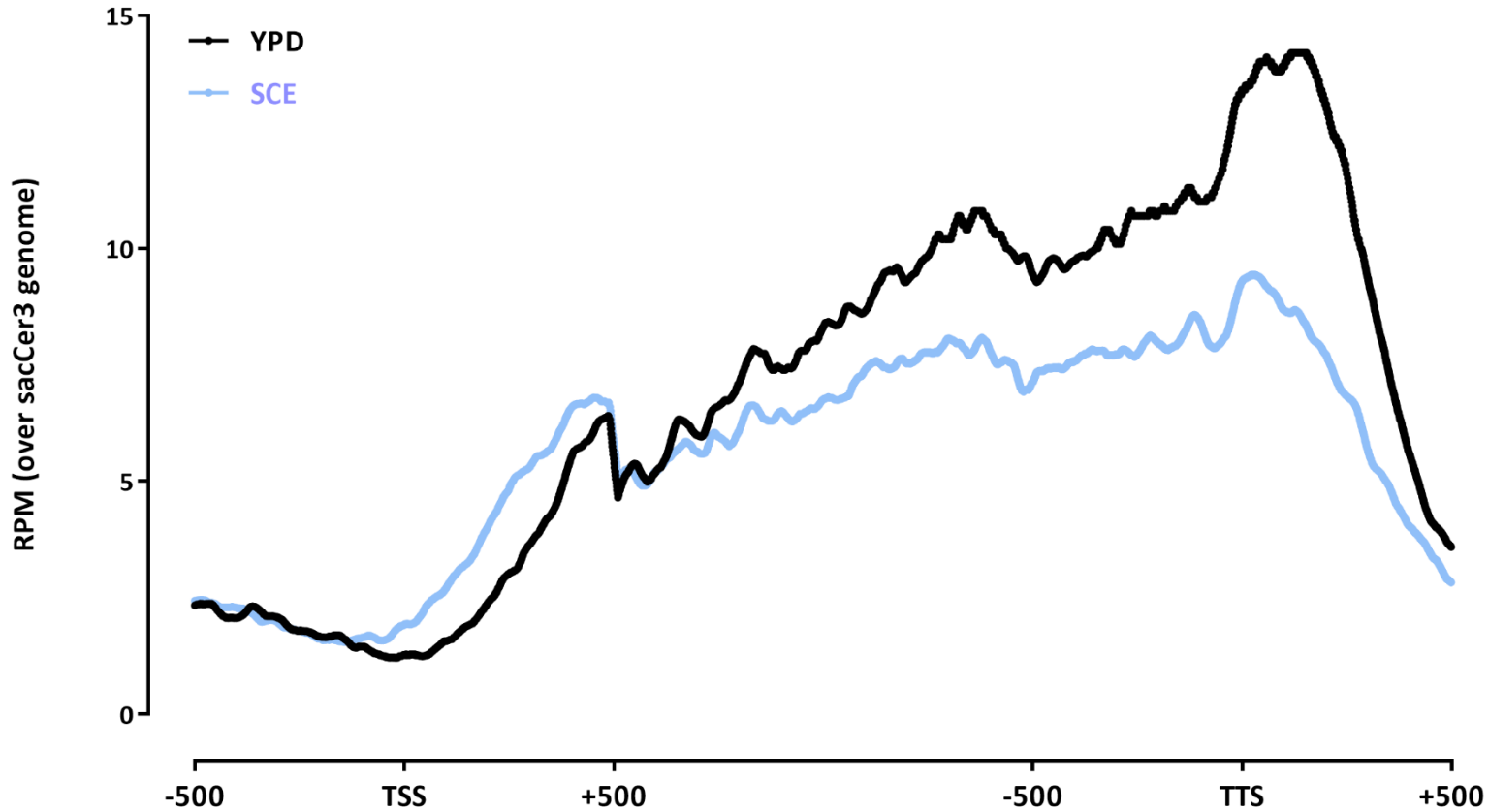
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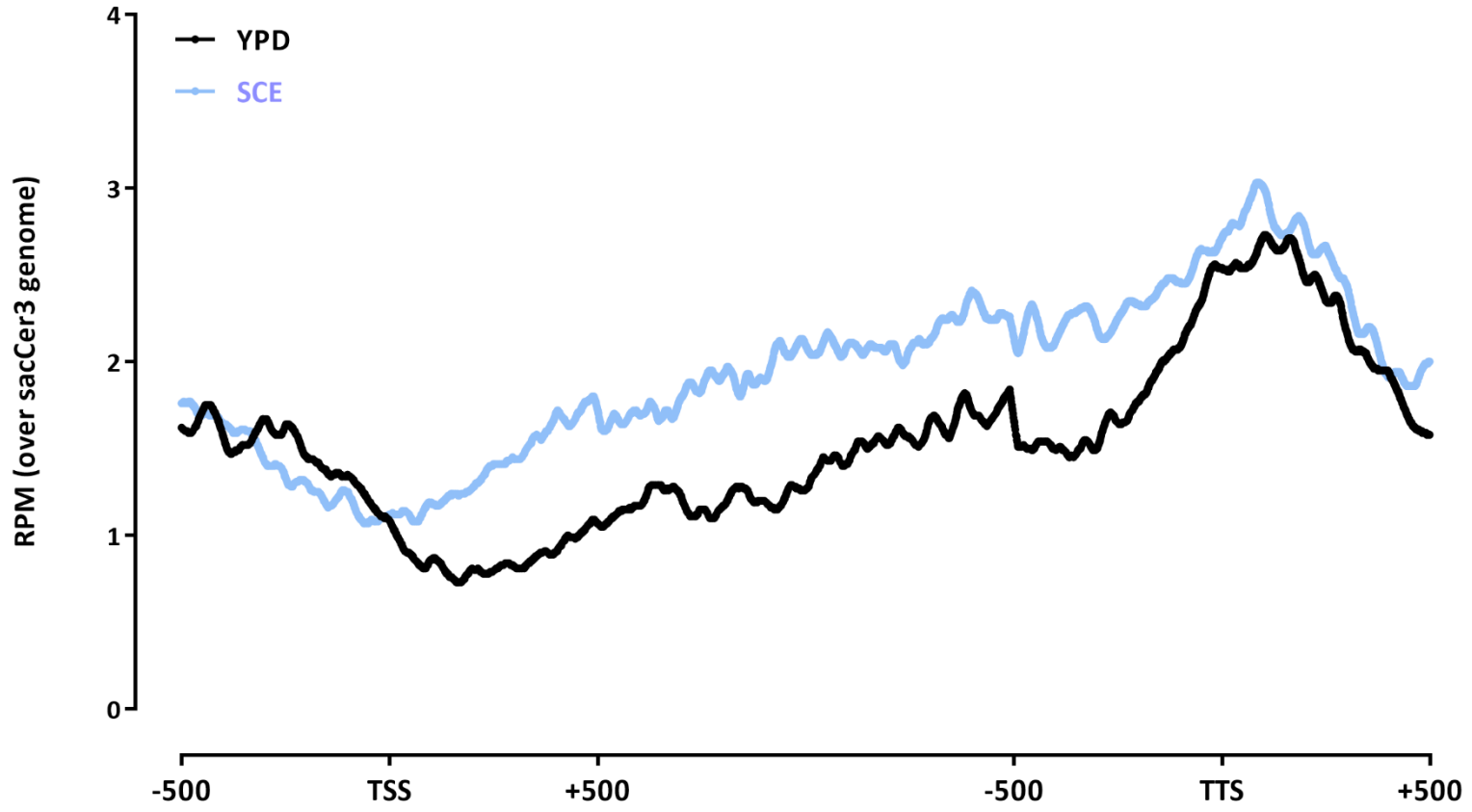
Pol2pS2 sacCer3 genome



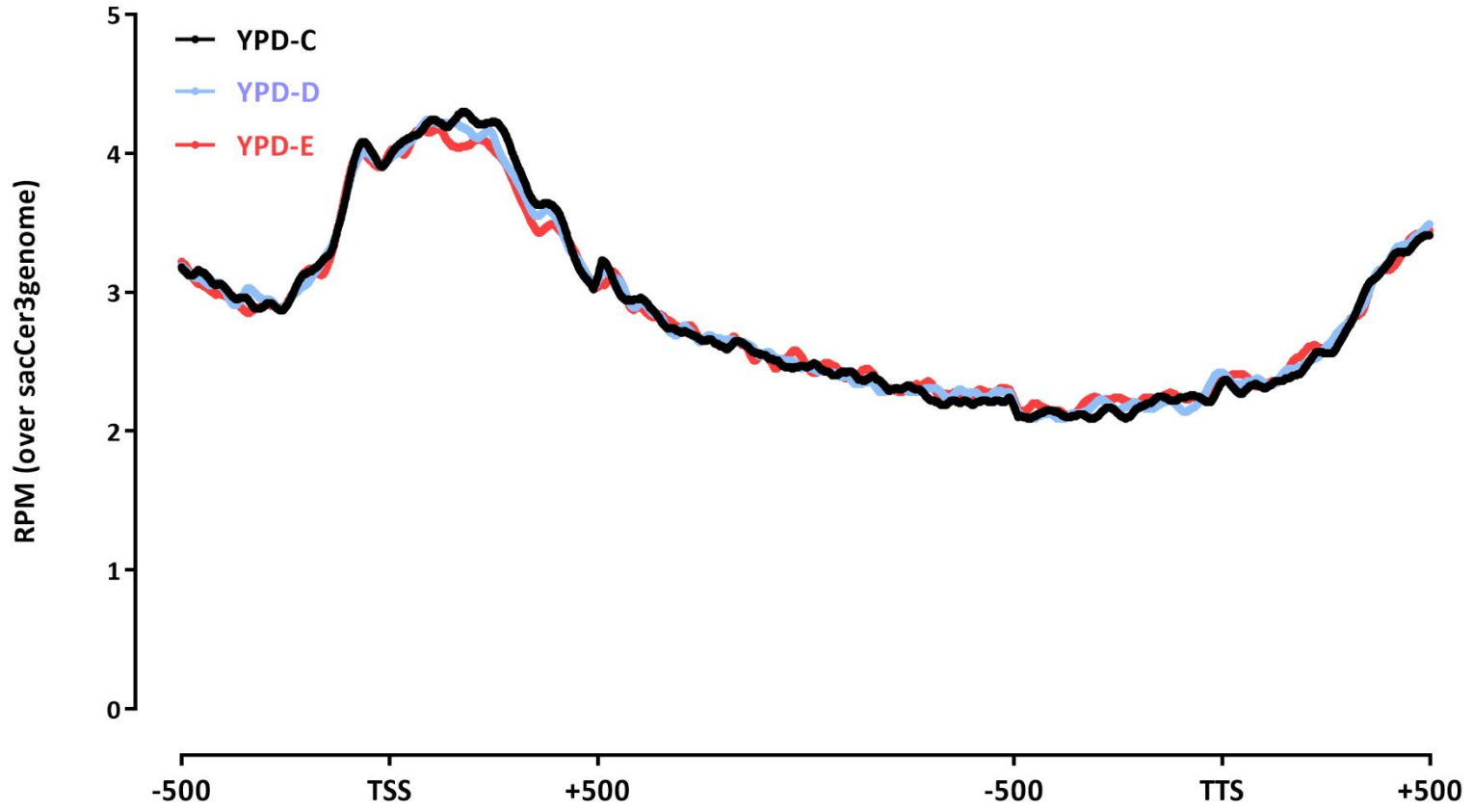
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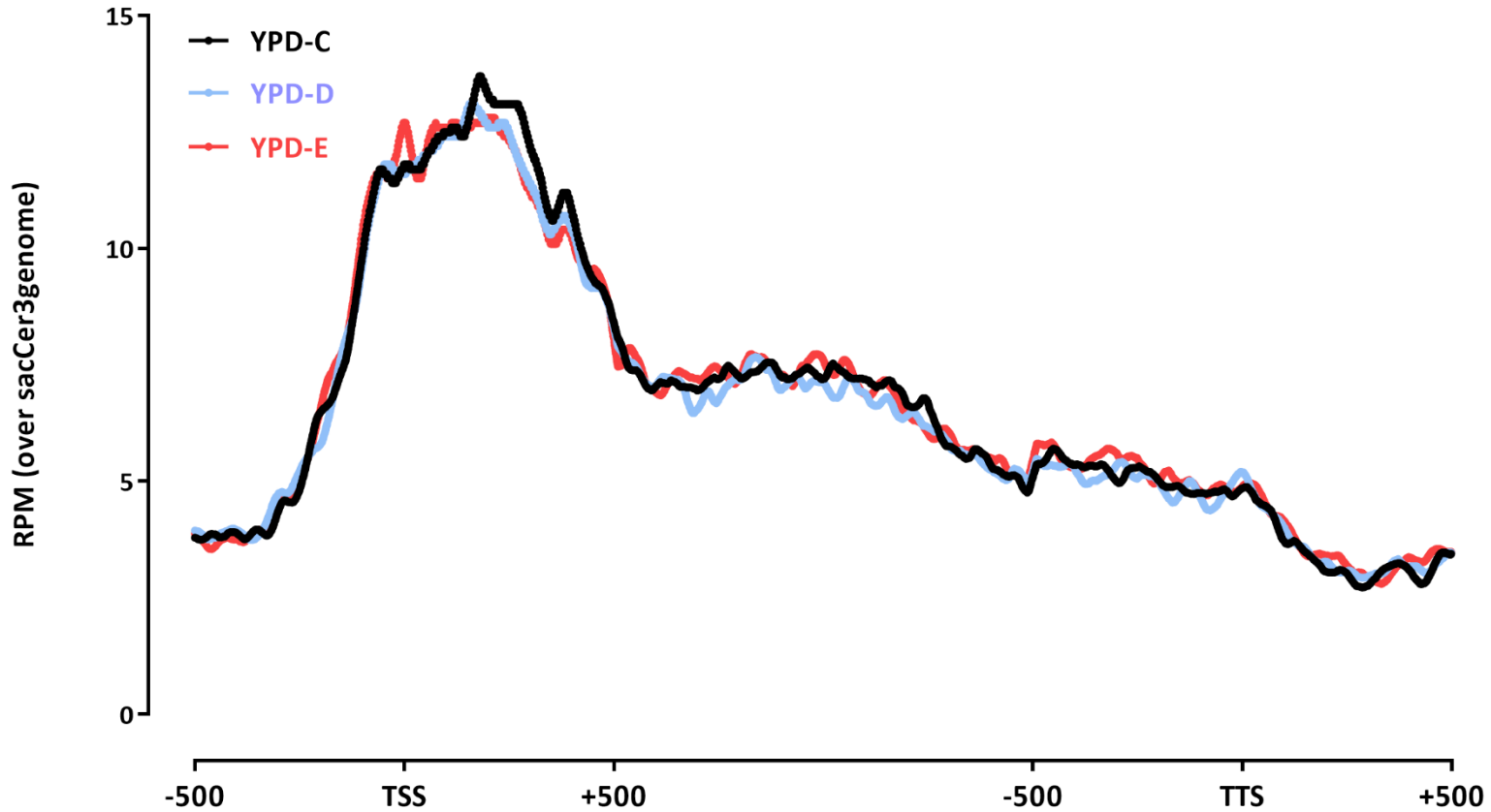
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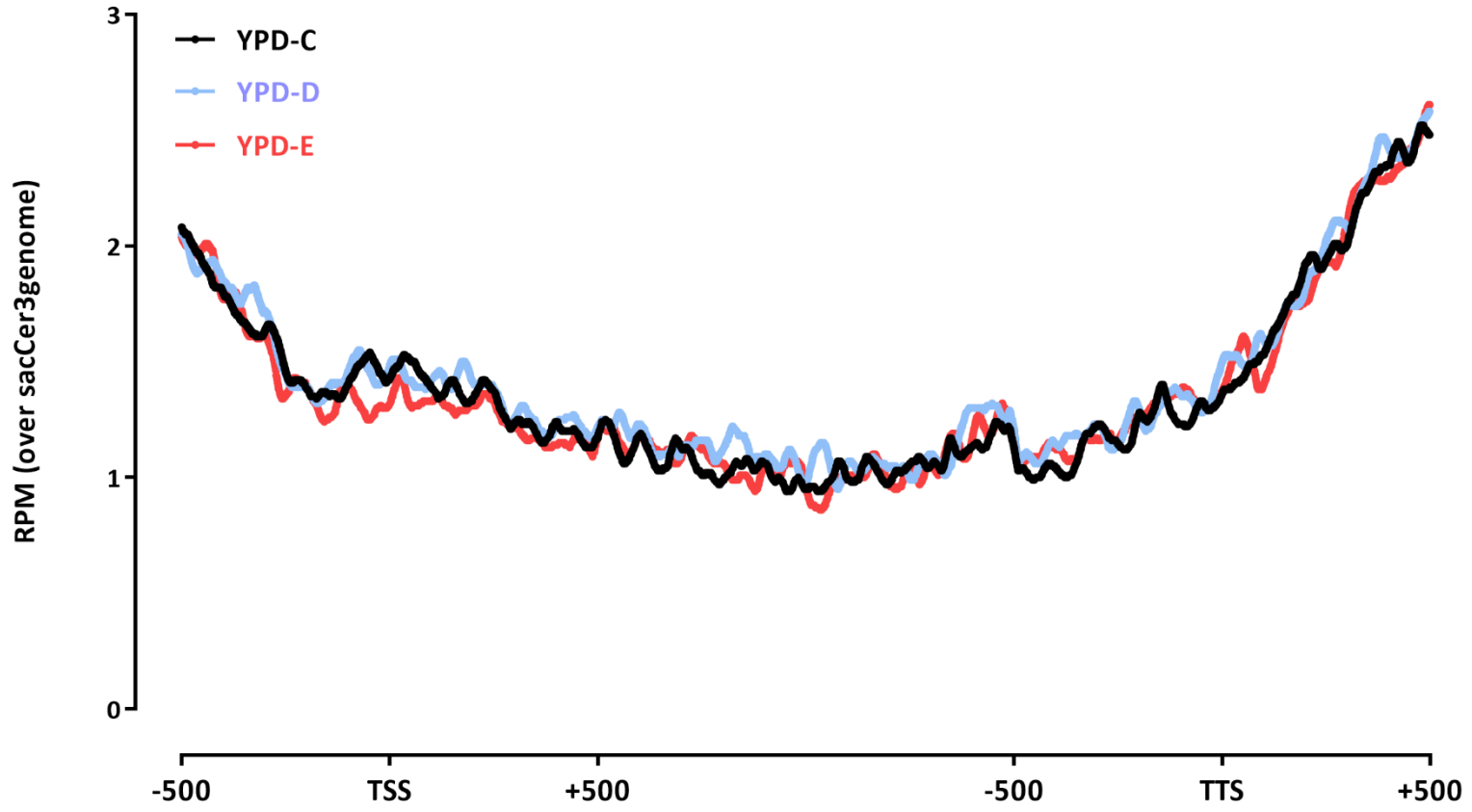
Pol2pS5 sacCer3 genome



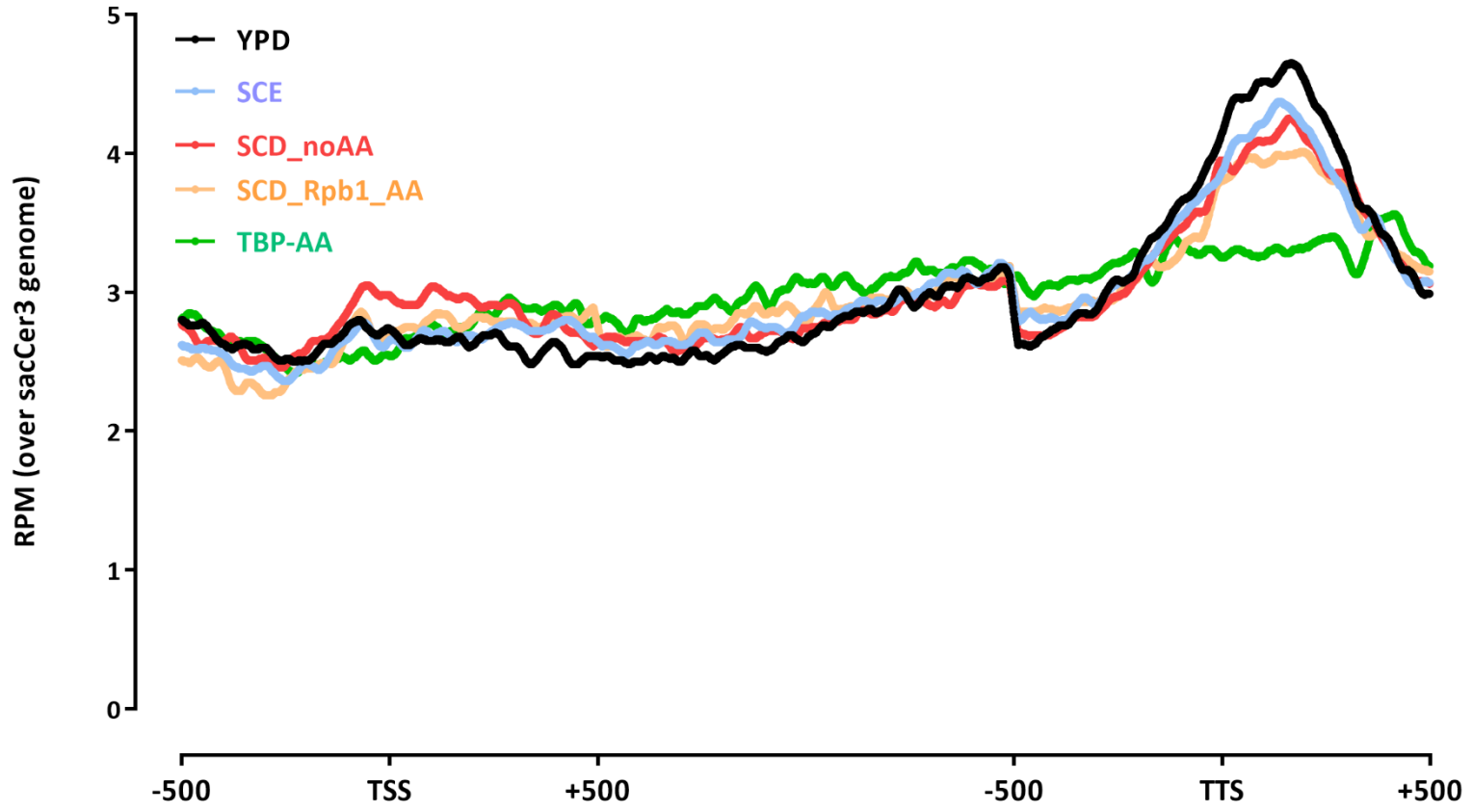
Pol2pS5 sacCer3 genome top expression quintile



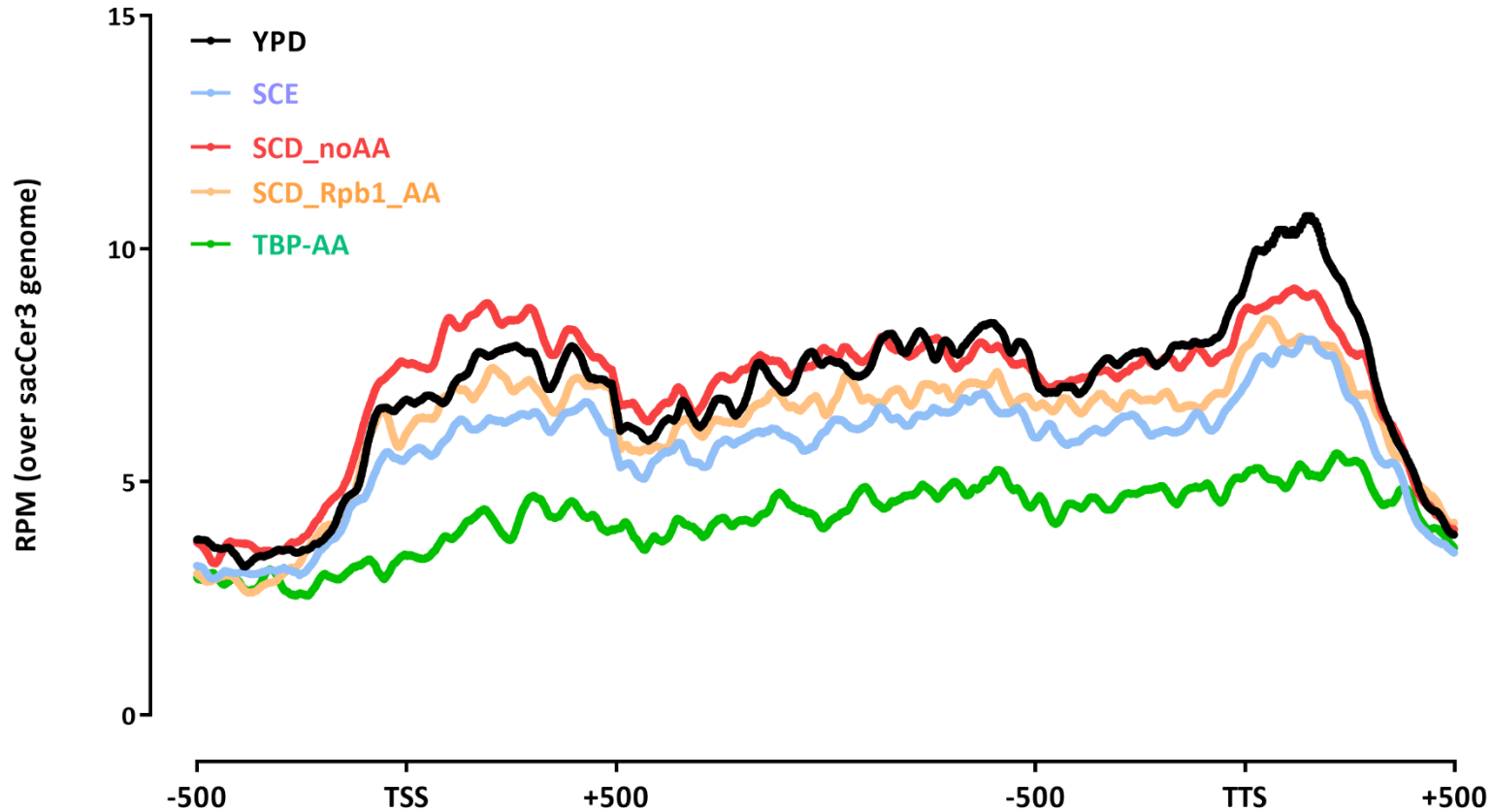
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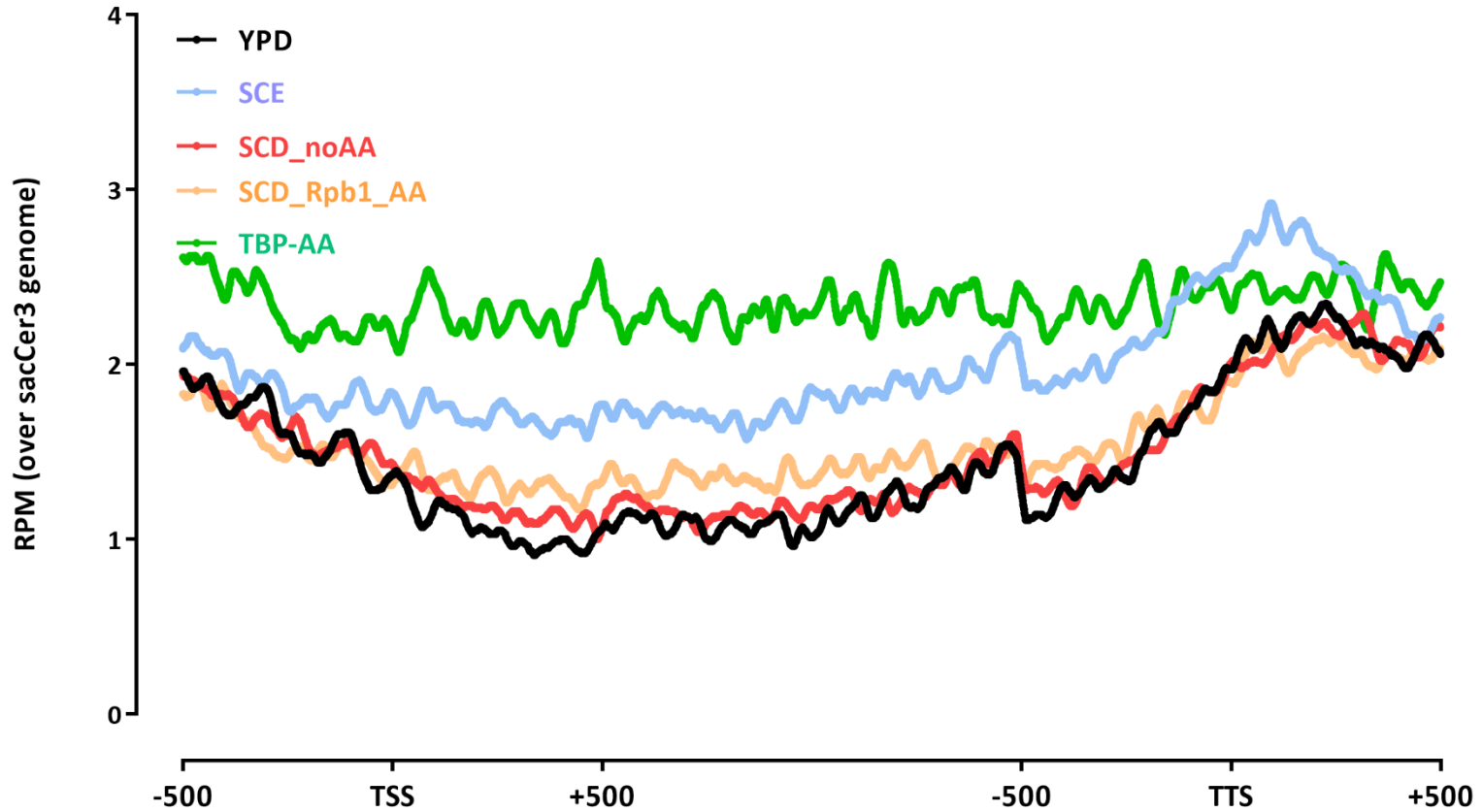
Rpb3 sacCer3 genome



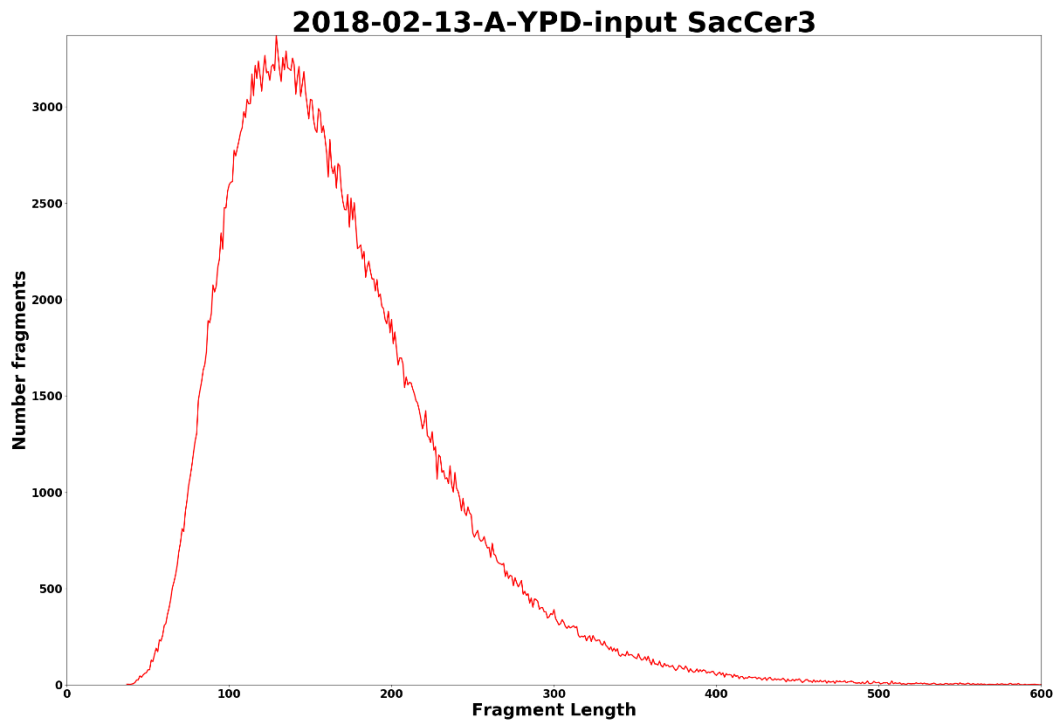
Rpb3 sacCer3 genome top expression quintile



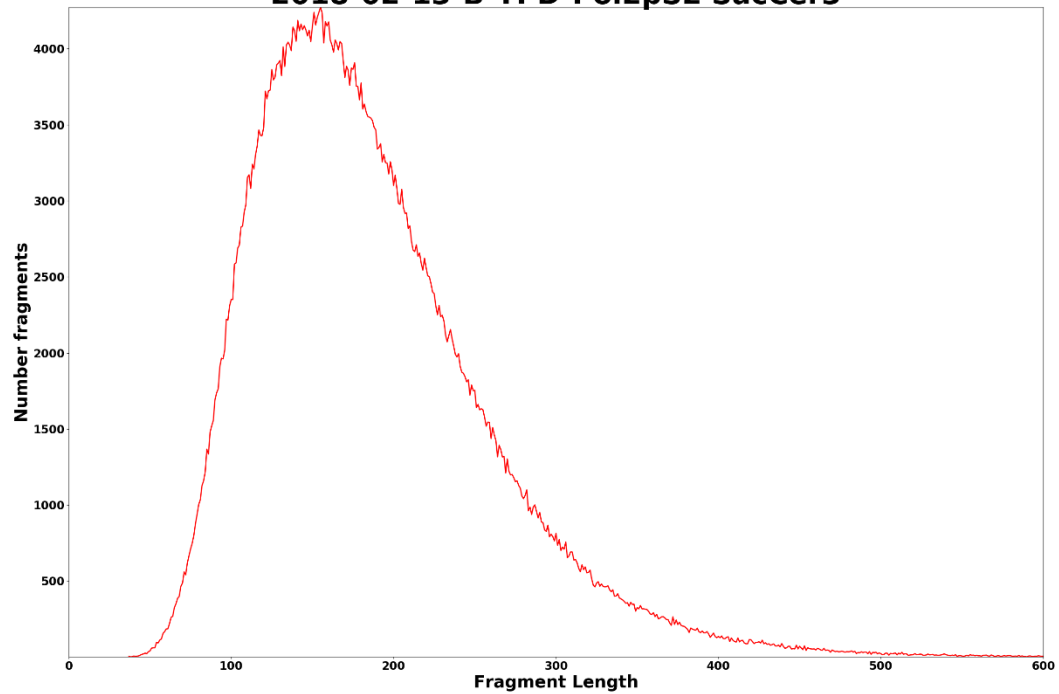
Rpb3 sacCer3 genome bottom expression quintile

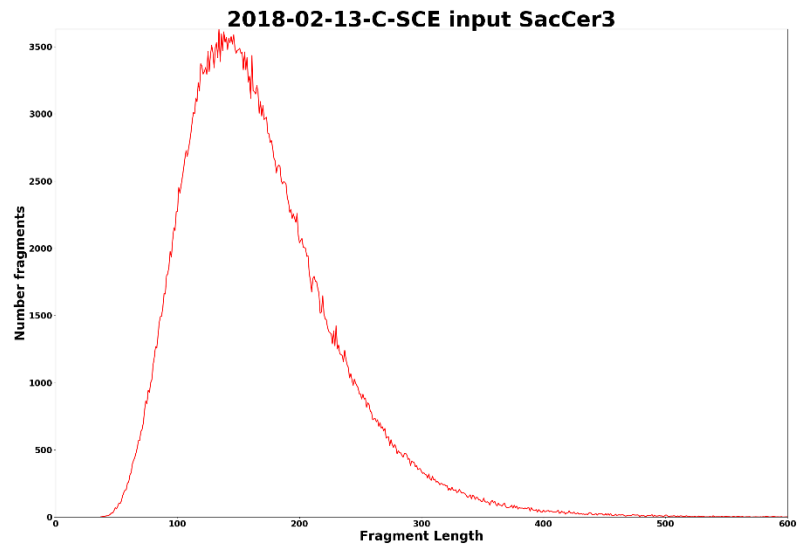
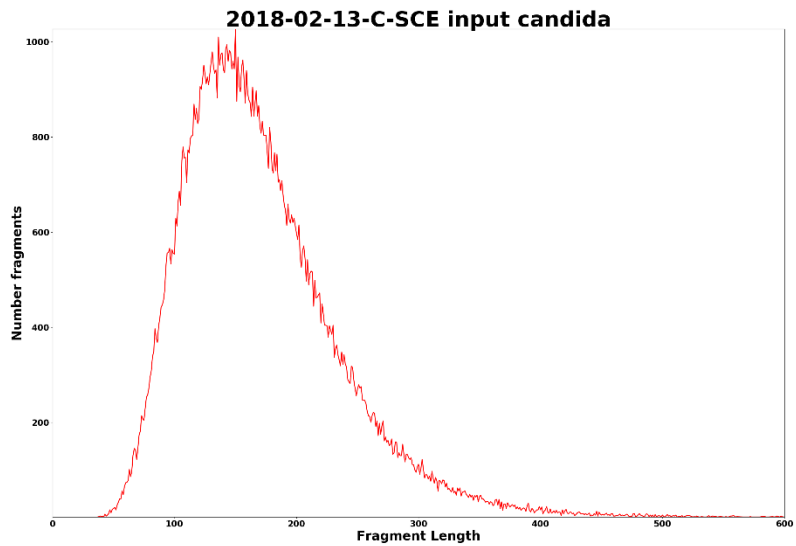


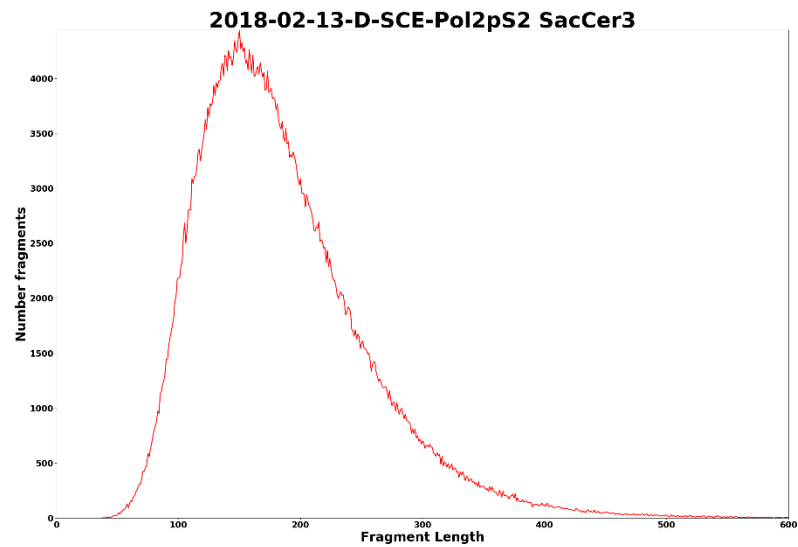
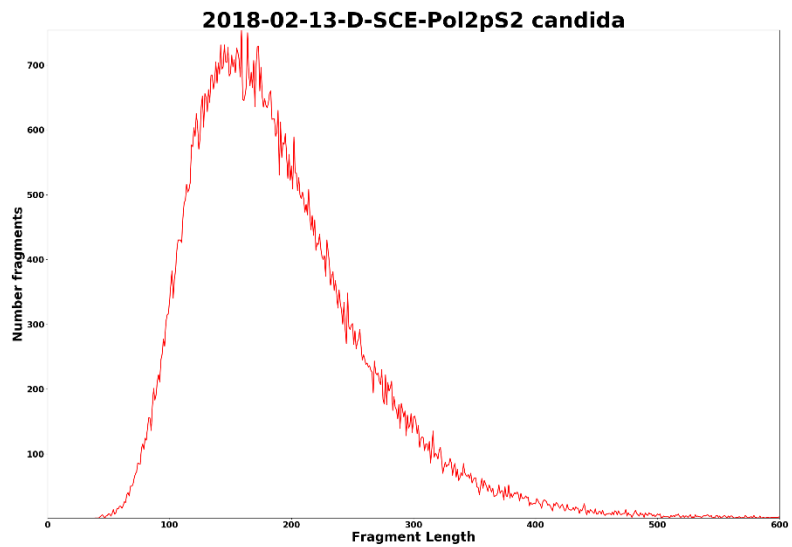
Fragment length distributions

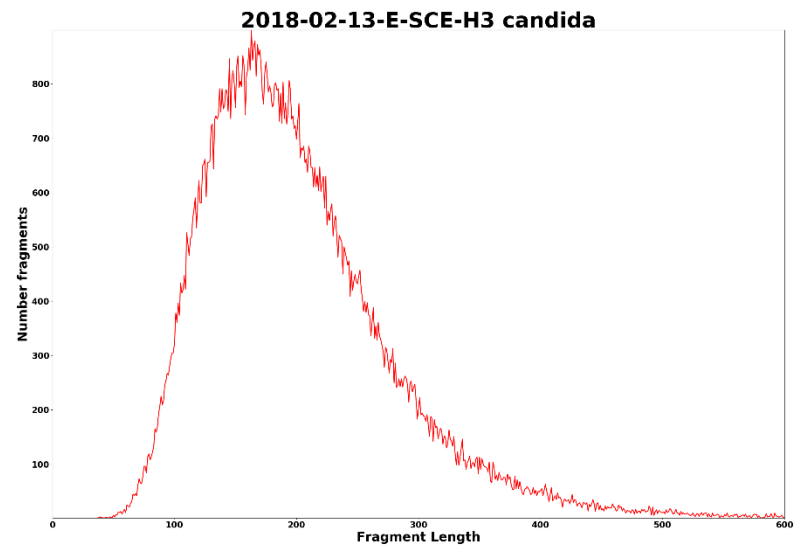
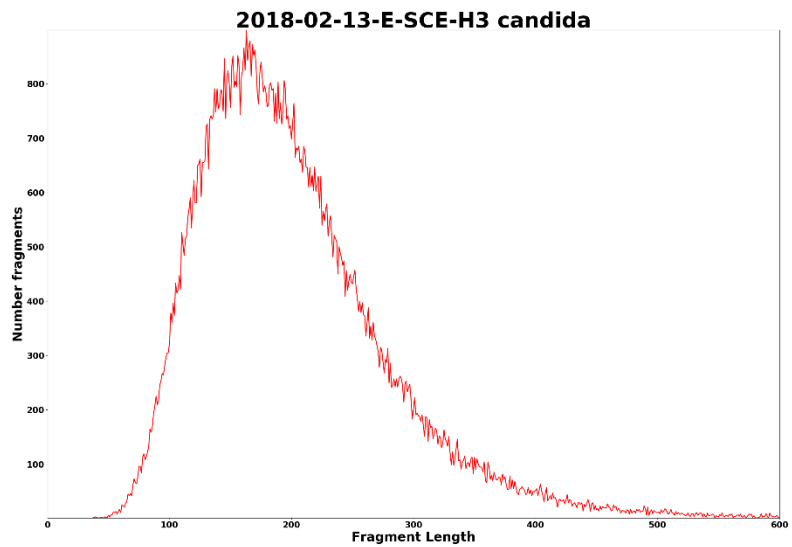


2018-02-13-B-YPD-Pol2pS2 SacCer3

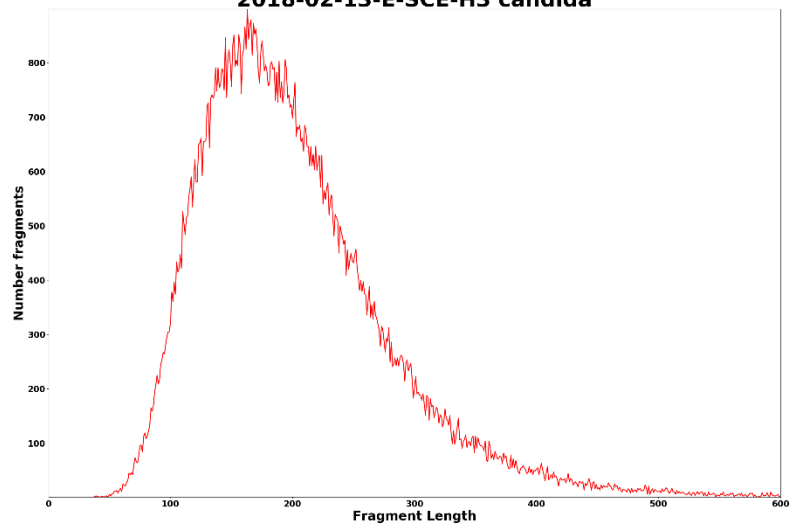




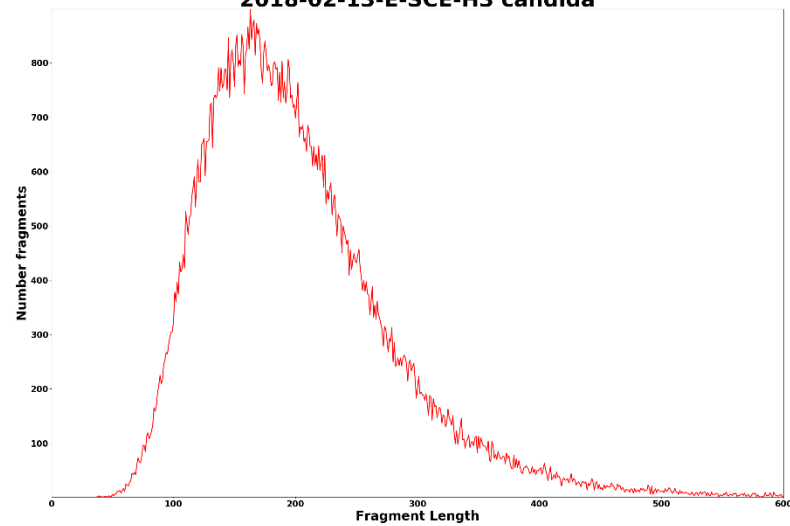


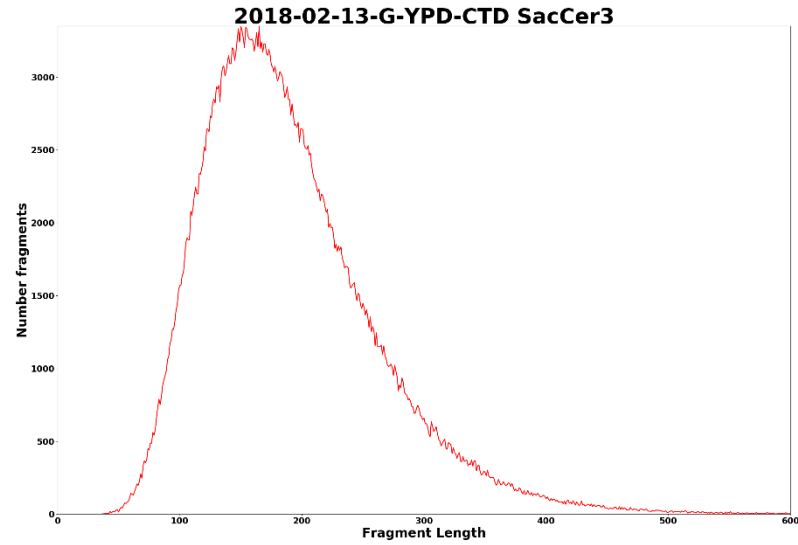


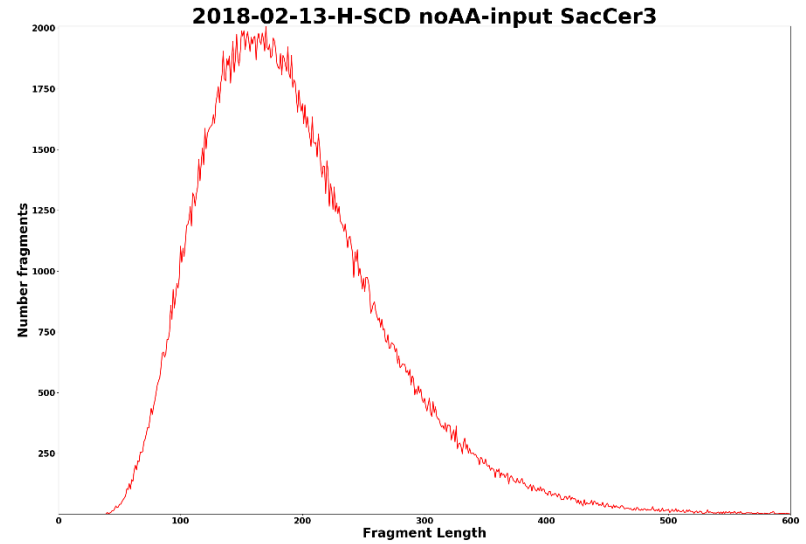
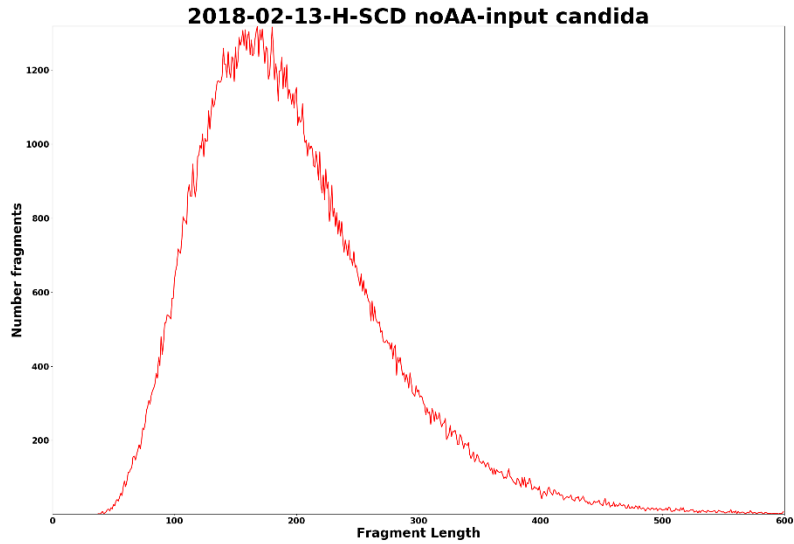
2018-02-13-E-SCE-H3 candida

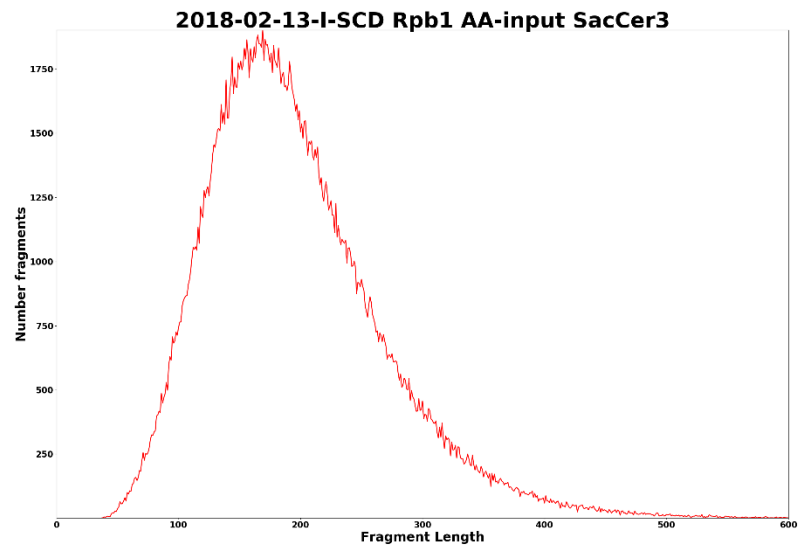
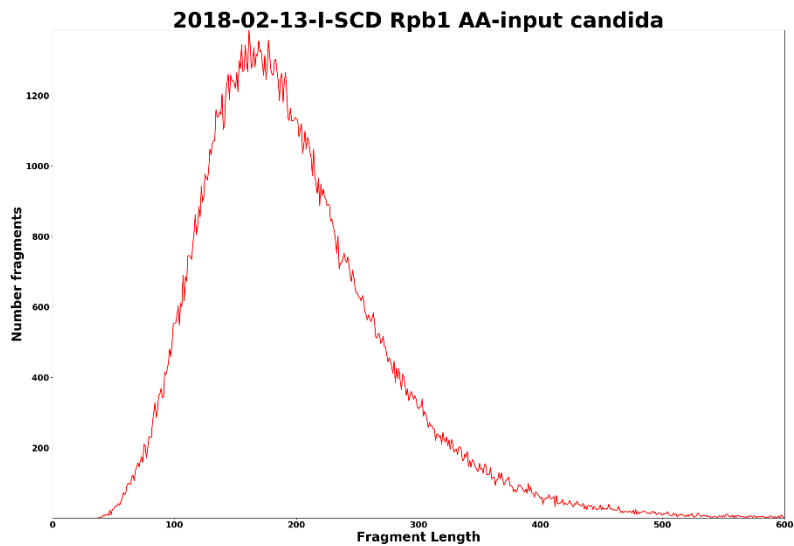


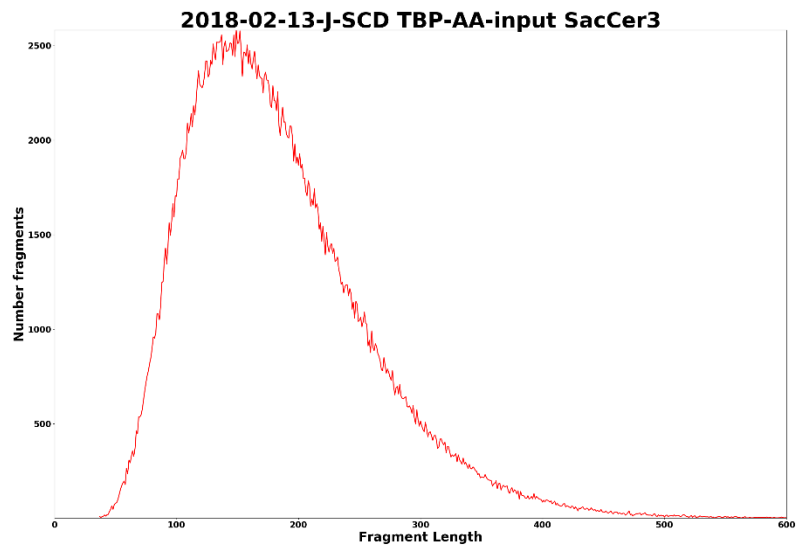
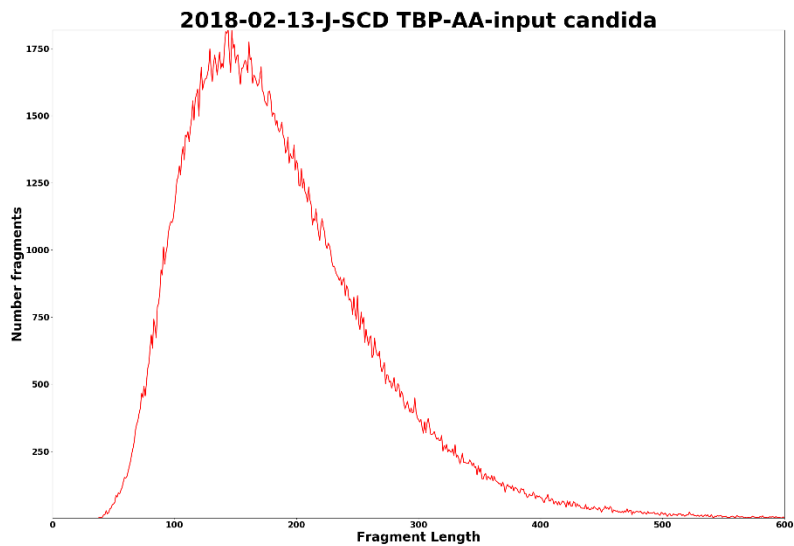
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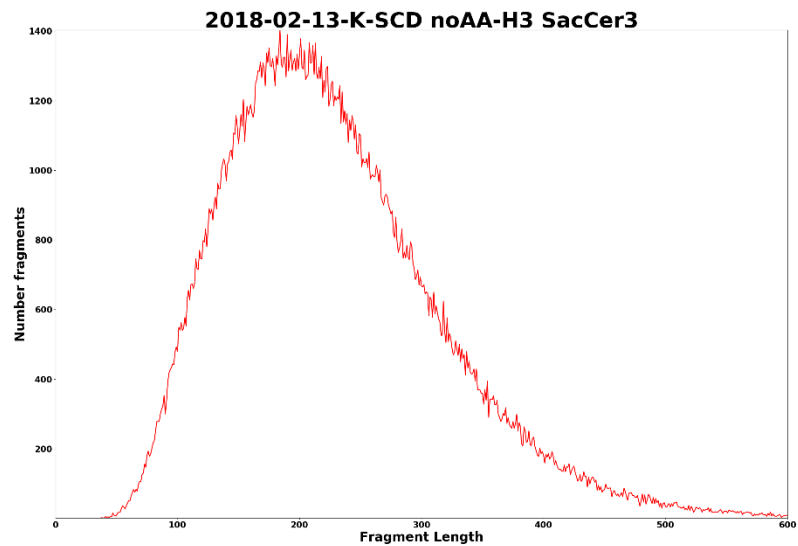
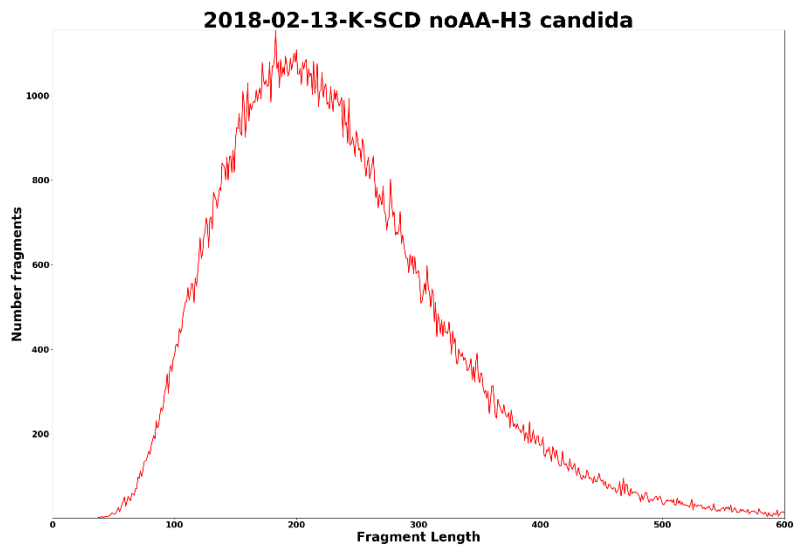


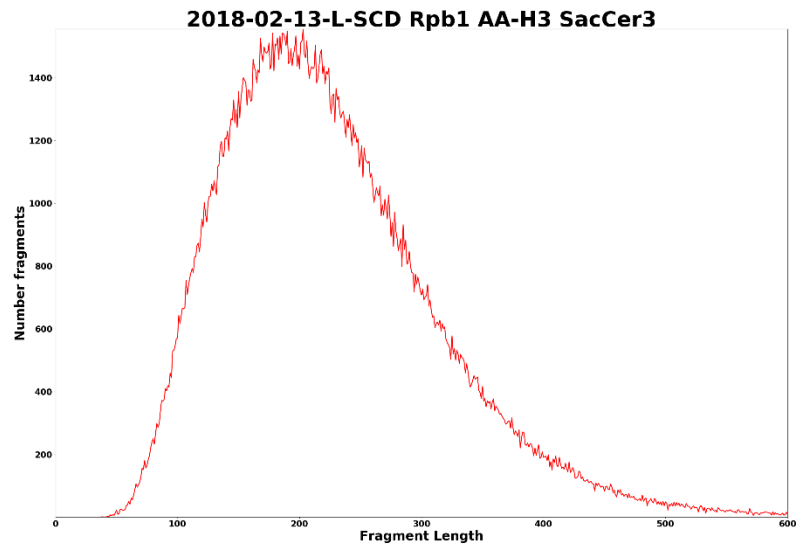
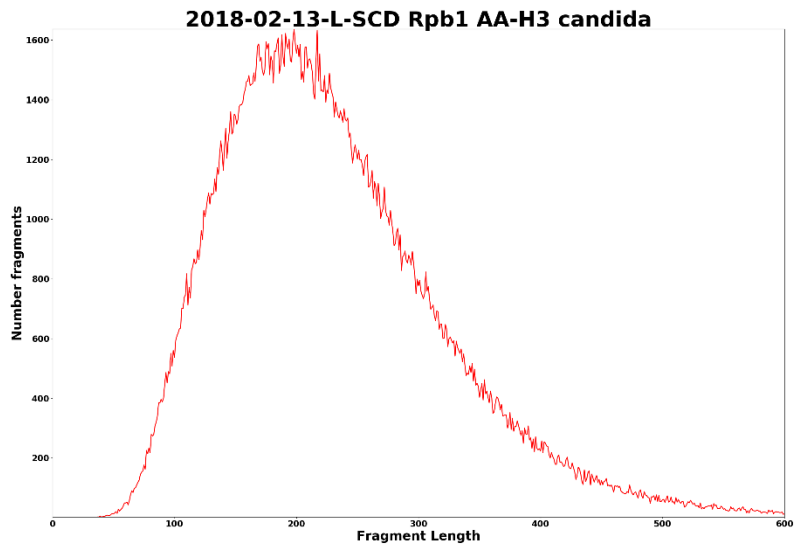




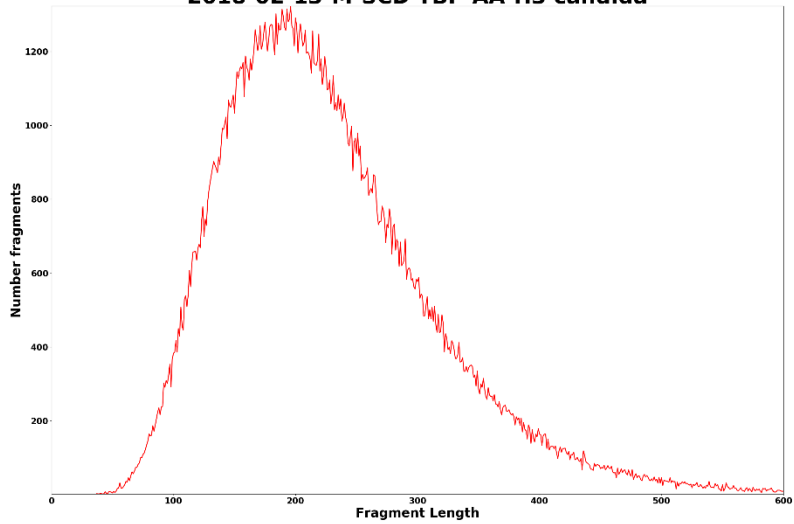




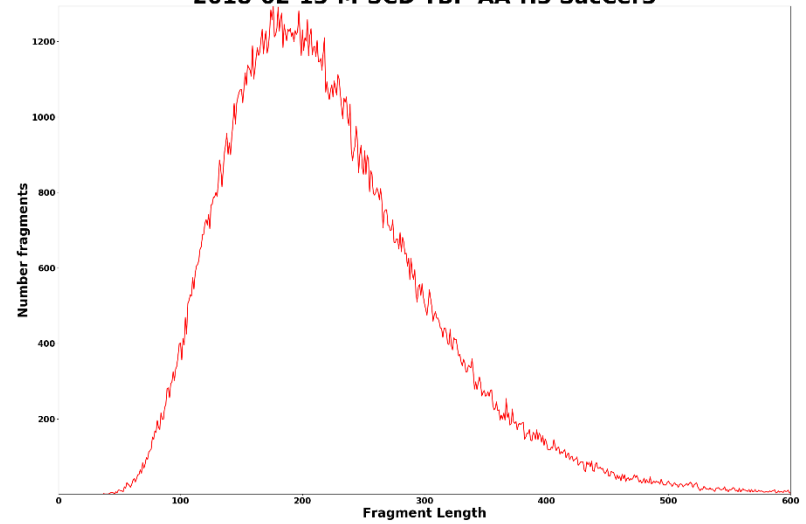




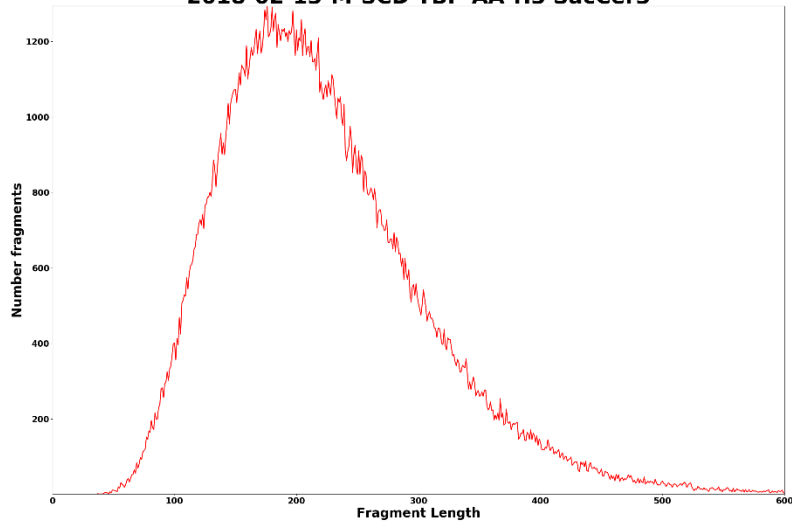
2018-02-13-M-SCD TBP-AA-H3 candida



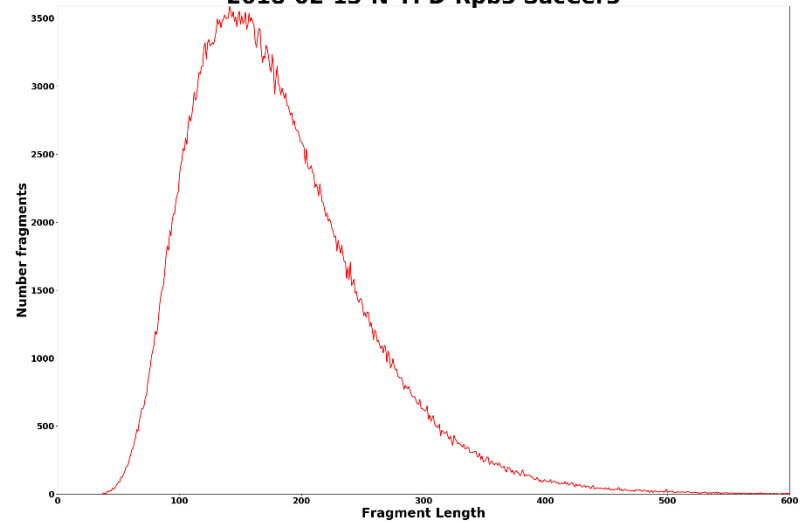
2018-02-13-M-SCD TBP-AA-H3 SacCer3

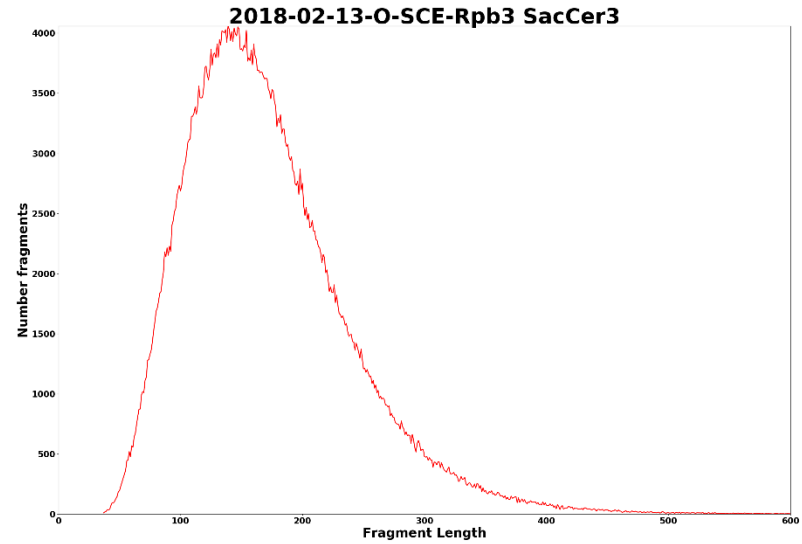
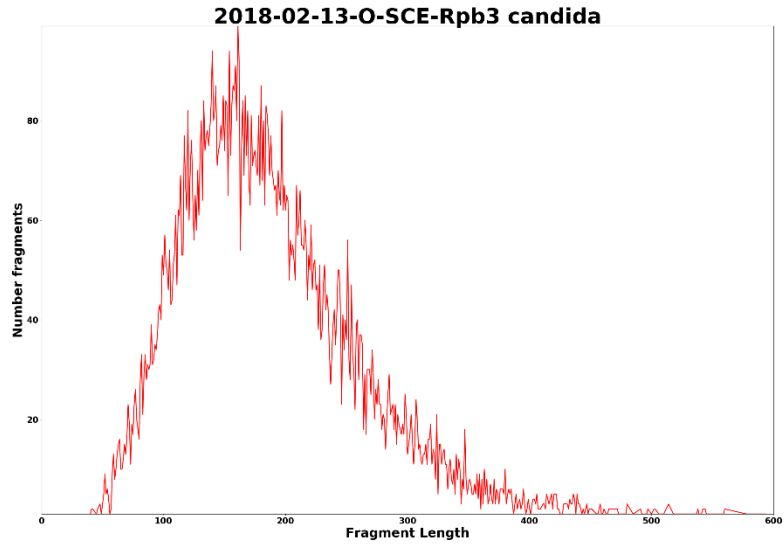


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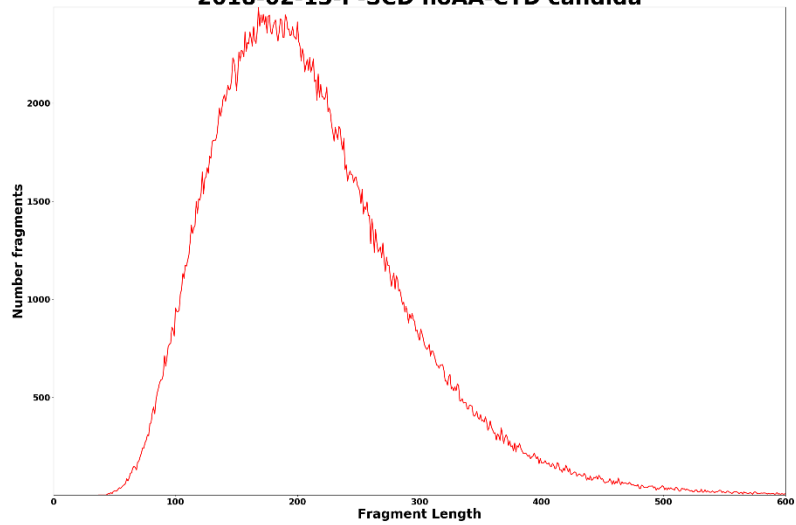


2018-02-13-N-YPD-Rpb3 SacCer3

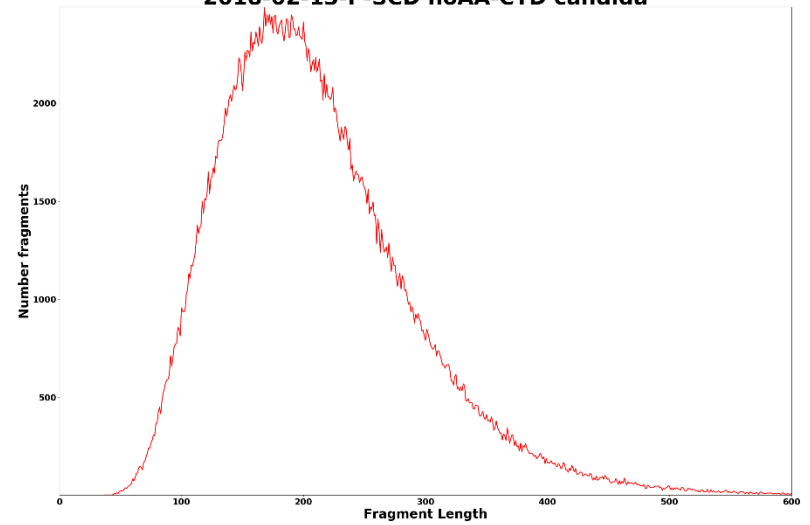




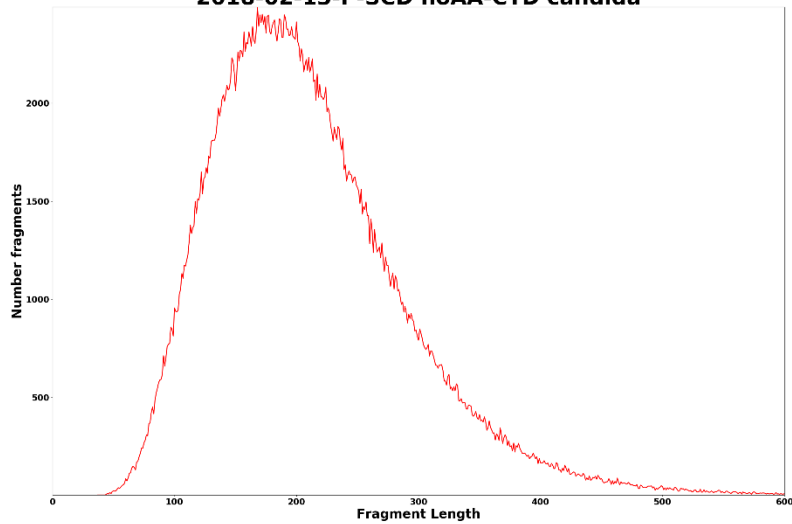
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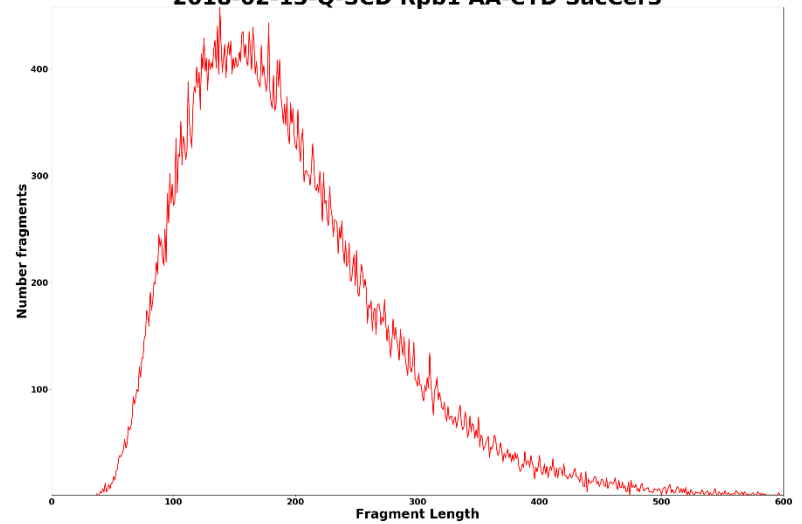
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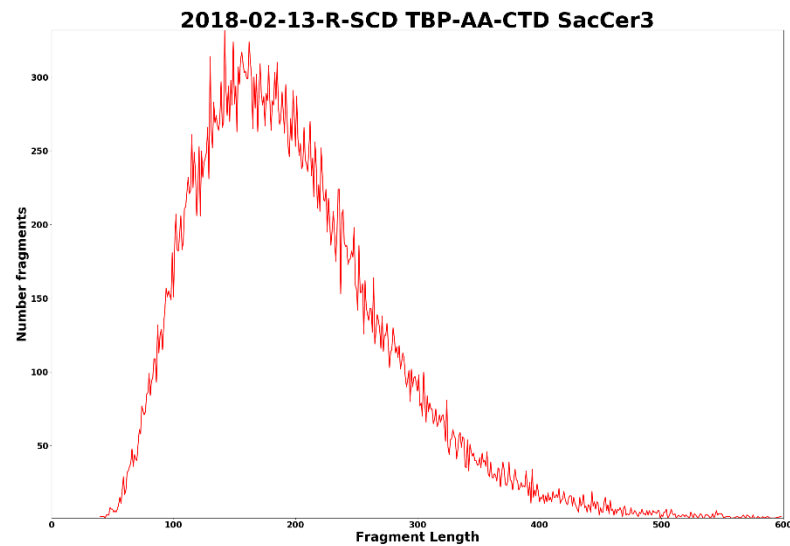
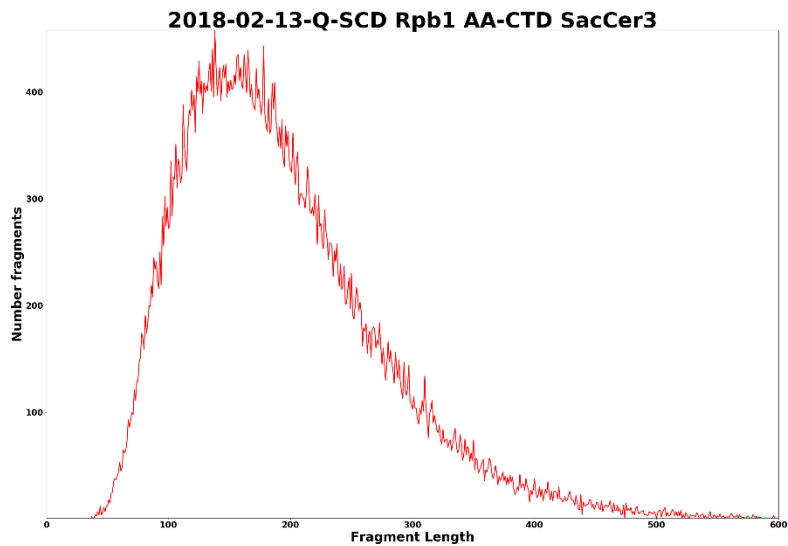


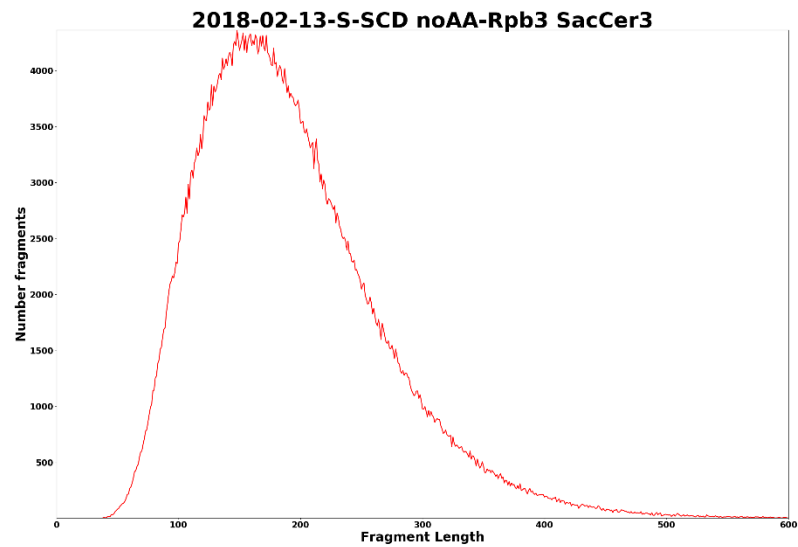
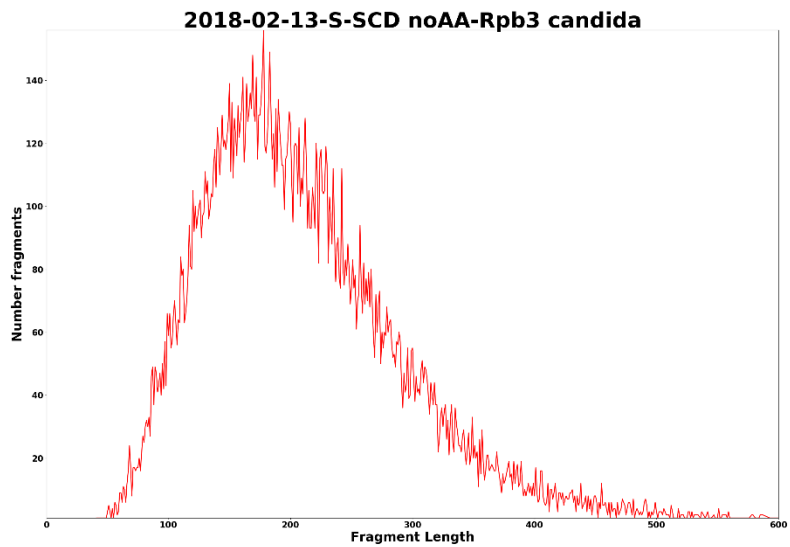
2018-02-13-P-SCD noAA-CTD candida



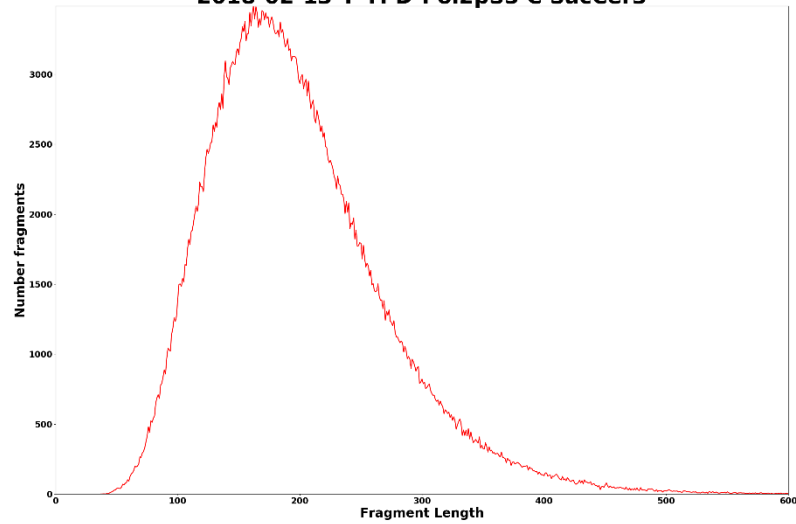
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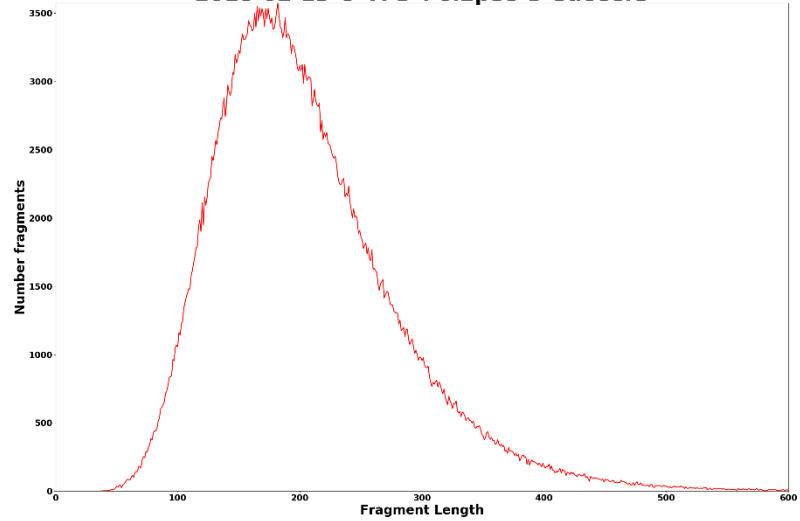




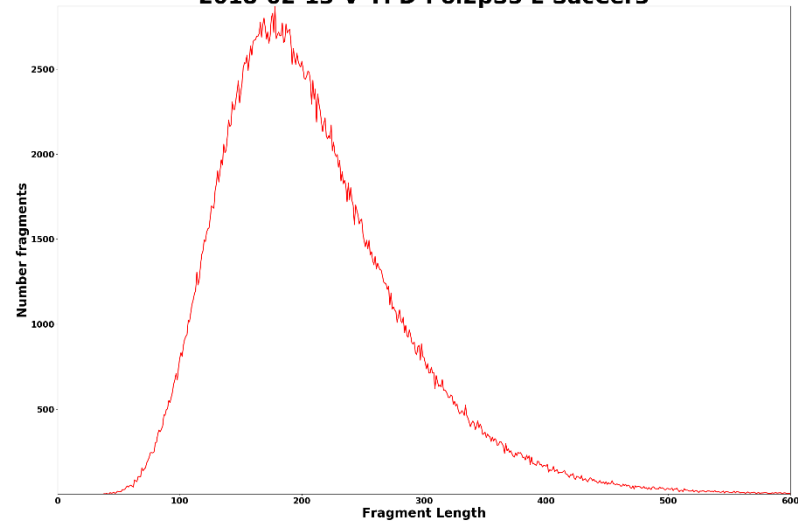
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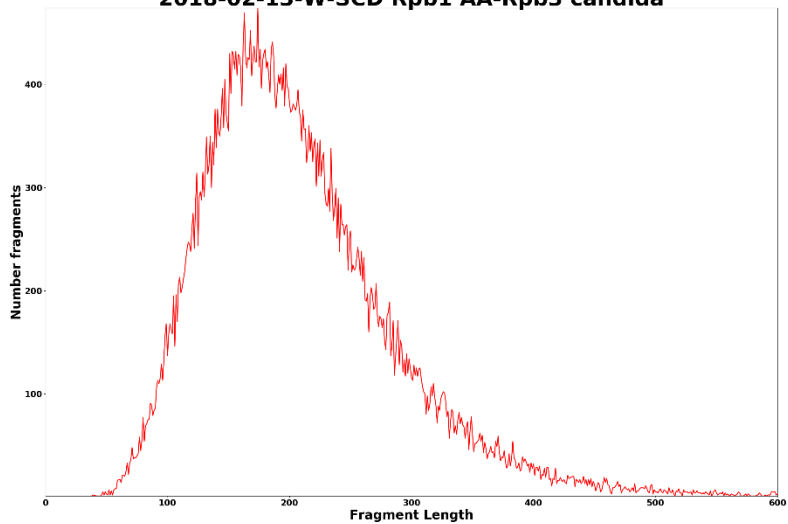
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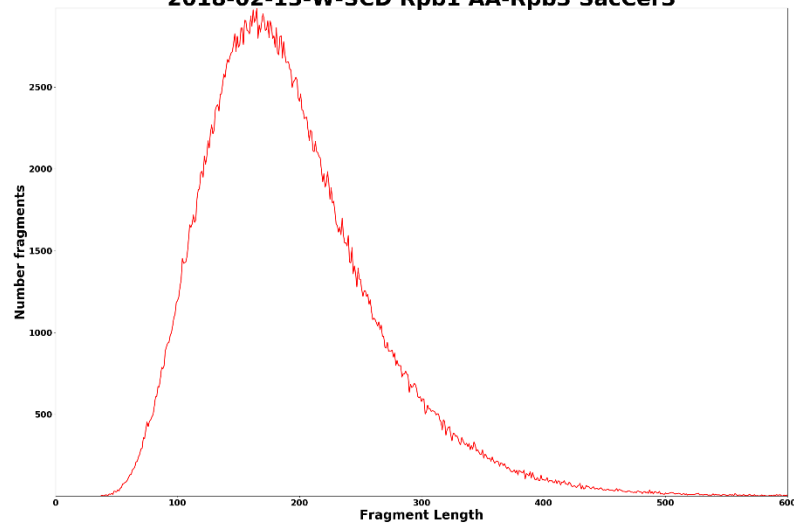
2018-02-13-V-YPD-Pol2pS5 E SacCer3



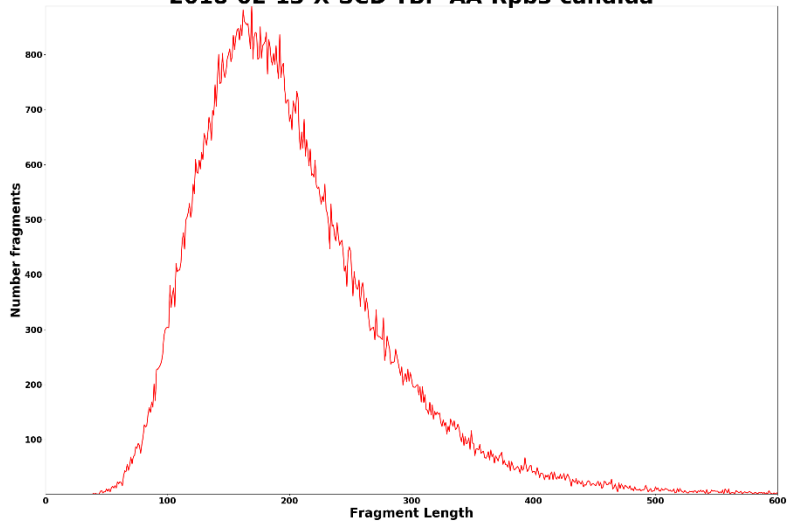
2018-02-13-W-SCD Rpb1 AA-Rpb3 candida



2018-02-13-W-SCD Rpb1 AA-Rpb3 SacCer3



2018-02-13-X-SCD TBP-AA-Rpb3 candida



2018-02-13-X-SCD TBP-AA-Rpb3 SacCer3

