

MAPPING STATS

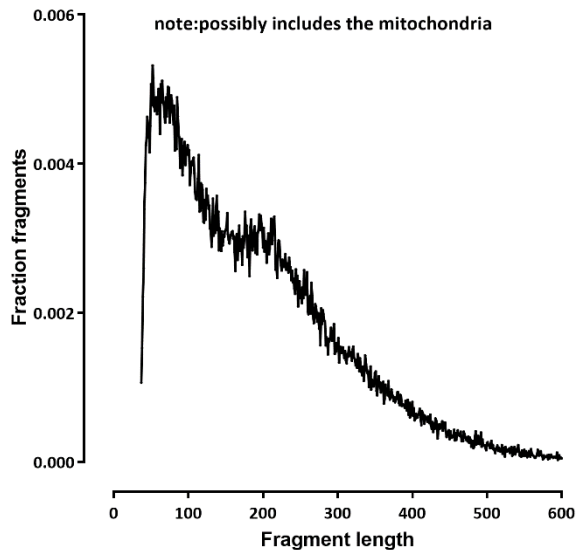
Species	Read Length	Library	Raw fragments	Unique non-chrM reads	Multi reads	Complexity	chrM reads	chrM fraction	Unique non-chrM reads after dedup	TSS ratio
<i>Saccharomyces cerevisiae</i>	2x36	L1910-GM16_Haematodinium_DVNP_rep1	309,424	222,480	276,078	0.89	26,848	0.11	214,790	1.34
<i>Saccharomyces cerevisiae</i>	2x36	L1911-GM16_Haematodinium_DVNP_rep2	330,223	181,592	225,096	0.90	36,006	0.17	175,144	1.31
<i>Saccharomyces cerevisiae</i>	2x36	L1912-GM16_Haematodinium_DVNP_ON_rep1	663,416	475,596	497,822	0.89	41,140	0.08	457,578	1.22
<i>Saccharomyces cerevisiae</i>	2x36	L1913-GM16_Haematodinium_DVNP_ON_rep2	1,451,378	1,124,098	970,800	0.86	149,174	0.12	1,076,188	1.22
<i>Saccharomyces cerevisiae</i>	2x36	L1914-GM17_vehcile_DVNP_rep1	2,558,102	1,707,020	1,719,616	0.82	77,856	0.04	1,606,590	1.24
<i>Saccharomyces cerevisiae</i>	2x36	L1915-GM17_vehcile_DVNP_rep2	1,878,565	1,314,798	1,485,492	0.83	46,964	0.03	1,237,010	1.26
<i>Saccharomyces cerevisiae</i>	2x36	L1916-GM17_vehcile_DVNP_ON_rep1	3,129,685	2,208,962	2,492,610	0.78	88,536	0.04	2,033,532	1.28
<i>Saccharomyces cerevisiae</i>	2x36	L1917-GM17_vehcile_DVNP_ON_rep2	1,047,863	801,626	886,612	0.85	48,132	0.06	760,264	1.27
<i>Saccharomyces cerevisiae</i>	2x36	L1918-GM18_rep1	110,389	59,148	93,722	0.92	8,526	0.13	57,344	1.26
<i>Saccharomyces cerevisiae</i>	2x36	L1919-GM18_rep2	2,873,420	1,966,504	2,493,850	0.78	88,596	0.04	1,811,840	1.29
<i>Saccharomyces cerevisiae</i>	2x36	L1920-GM19_rep1	5,333,495	3,649,732	4,121,402	0.73	193,994	0.05	3,314,988	1.34
<i>Saccharomyces cerevisiae</i>	2x36	L1921-GM19_rep2	2,366,205	1,743,602	1,395,362	0.83	67,710	0.04	1,647,678	1.24

MAPPING STATS, JOINT MAPPING

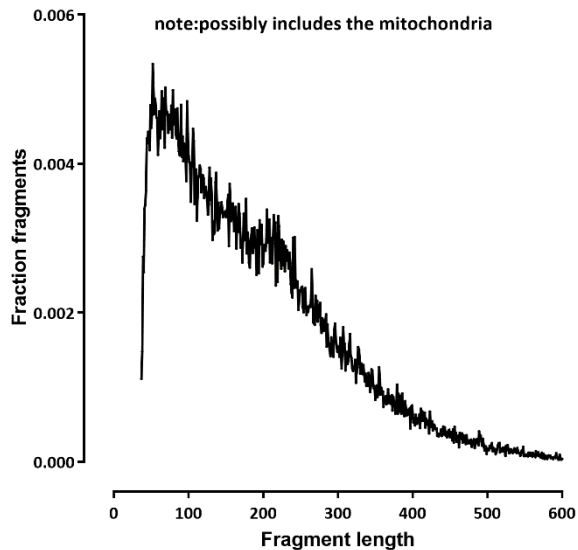
Species	Read Length	Library	Raw fragments	Unique non-chrM reads	Multi reads	Complexity	chrM reads	chrM fraction	Unique non-chrM reads after dedup	TSS ratio
sacCer3+Candida, sacCer3	2x36	L1910-GM16_Haematodinium_DVNP_rep1	309,424	222,340	160,970	0.89				1.44
sacCer3+Candida, sacCer3	2x36	L1911-GM16_Haematodinium_DVNP_rep2	330,223	181,330	129,218	0.90				1.37
sacCer3+Candida, sacCer3	2x36	L1912-GM16_Haematodinium_DVNP_ON_rep1	663,416	475,278	291,474	0.89				1.26
sacCer3+Candida, sacCer3	2x36	L1913-GM16_Haematodinium_DVNP_ON_rep2	1,451,378	1,123,364	572,376	0.86				1.26
sacCer3+Candida, sacCer3	2x36	L1914-GM17_vehcile_DVNP_rep1	2,558,102	1,705,458	969,192	0.82				1.31
sacCer3+Candida, sacCer3	2x36	L1915-GM17_vehcile_DVNP_rep2	1,878,565	1,313,802	840,602	0.83				1.33
sacCer3+Candida, sacCer3	2x36	L1916-GM17_vehcile_DVNP_ON_rep1	3,129,685	2,207,368	1,425,440	0.78				1.35
sacCer3+Candida, sacCer3	2x36	L1917-GM17_vehcile_DVNP_ON_rep2	1,047,863	801,230	512,378	0.85				1.32
sacCer3+Candida, sacCer3	2x36	L1918-GM18_rep1	110,389	59,088	53,200	0.92				1.29
sacCer3+Candida, sacCer3	2x36	L1919-GM18_rep2	2,873,420	1,965,254	1,417,210	0.78				1.36
sacCer3+Candida, sacCer3	2x36	L1920-GM19_rep1	5,333,495	3,646,892	2,353,228	0.73				1.41
sacCer3+Candida, sacCer3	2x36	L1921-GM19_rep2	2,366,205	1,741,988	814,778	0.83				1.28
sacCer3+Candida, Candida	2x36	L1910-GM16_Haematodinium_DVNP_rep1	309,424	60,374	3,002	0.94				1.74
sacCer3+Candida, Candida	2x36	L1911-GM16_Haematodinium_DVNP_rep2	330,223	171,520	9,584	0.91				1.71
sacCer3+Candida, Candida	2x36	L1912-GM16_Haematodinium_DVNP_ON_rep1	663,416	243,766	13,454	0.92				1.51
sacCer3+Candida, Candida	2x36	L1913-GM16_Haematodinium_DVNP_ON_rep2	1,451,378	511,568	22,304	0.90				1.54
sacCer3+Candida, Candida	2x36	L1914-GM17_vehcile_DVNP_rep1	2,558,102	1,240,952	47,924	0.82				1.82
sacCer3+Candida, Candida	2x36	L1915-GM17_vehcile_DVNP_rep2	1,878,565	650,948	25,790	0.85				1.93
sacCer3+Candida, Candida	2x36	L1916-GM17_vehcile_DVNP_ON_rep1	3,129,685	1,028,996	36,790	0.80				1.96
sacCer3+Candida, Candida	2x36	L1917-GM17_vehcile_DVNP_ON_rep2	1,047,863	231,076	9,950	0.90				1.95
sacCer3+Candida, Candida	2x36	L1918-GM18_rep1	110,389	44,296	2,630	0.95				1.74
sacCer3+Candida, Candida	2x36	L1919-GM18_rep2	2,873,420	859,634	40,806	0.85				1.73
sacCer3+Candida, Candida	2x36	L1920-GM19_rep1	5,333,495	2,007,476	84,132	0.77				1.84
sacCer3+Candida, Candida	2x36	L1921-GM19_rep2	2,366,205	1,207,780	50,006	0.84				1.67

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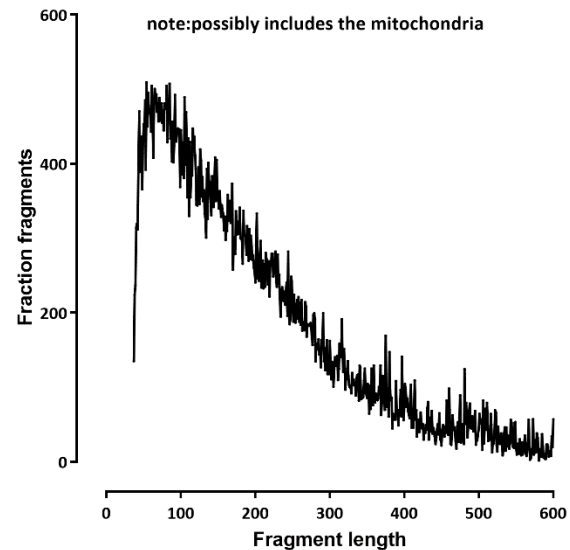
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L1910-GM16_Haematodinium_DVNP_rep1 sacCer3 all

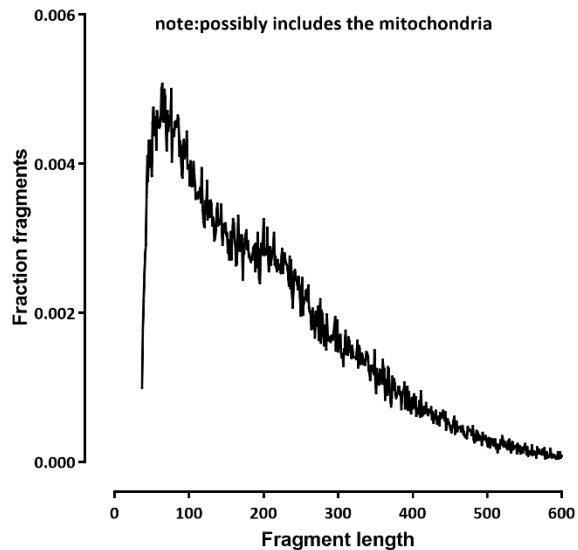


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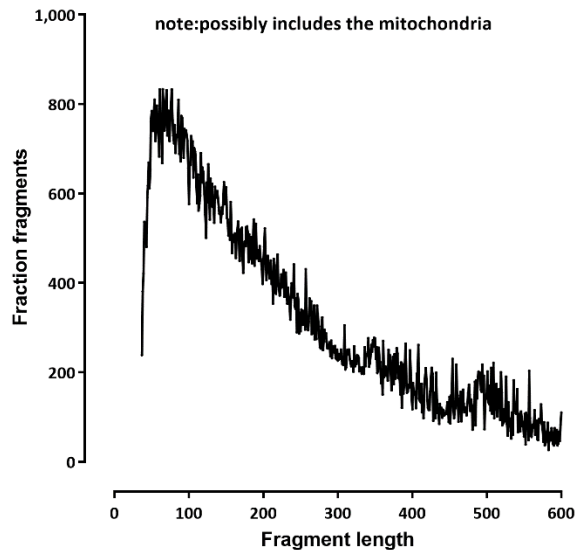


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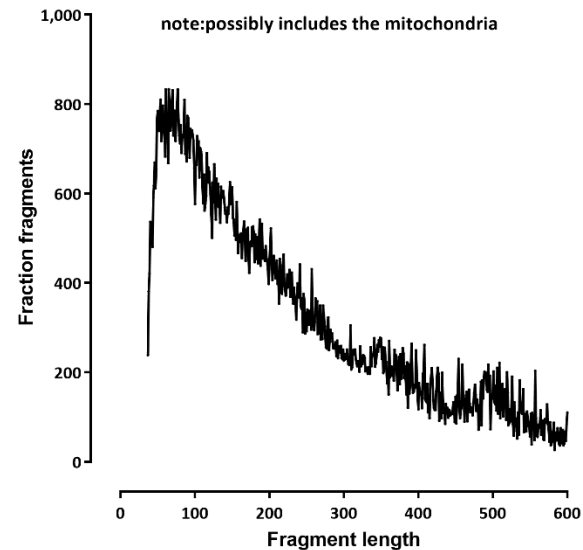
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L1911-GM16_Haematodinium_DVNP_rep2 Candida all

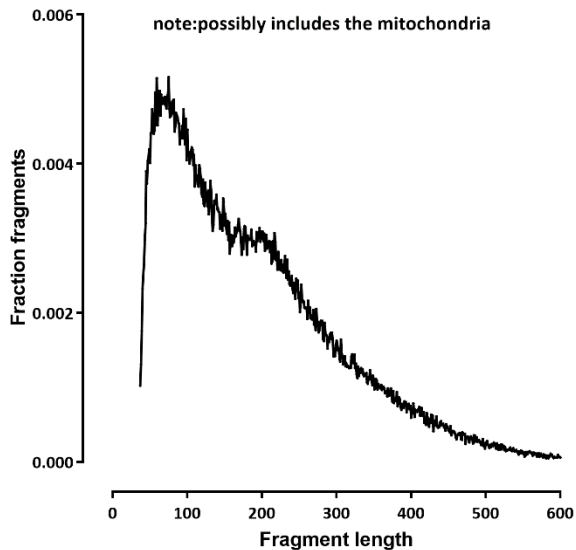


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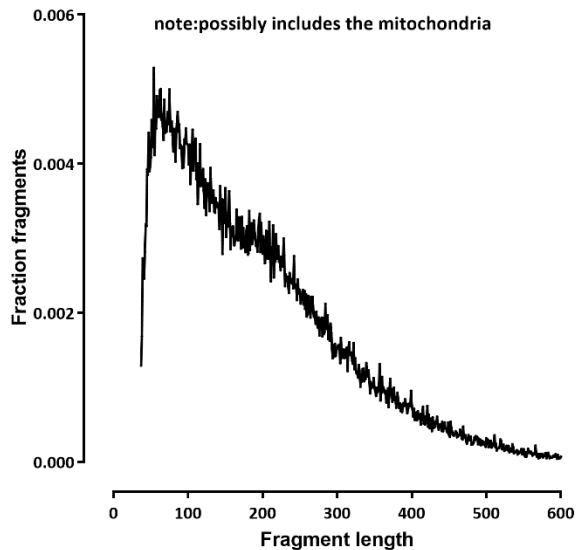


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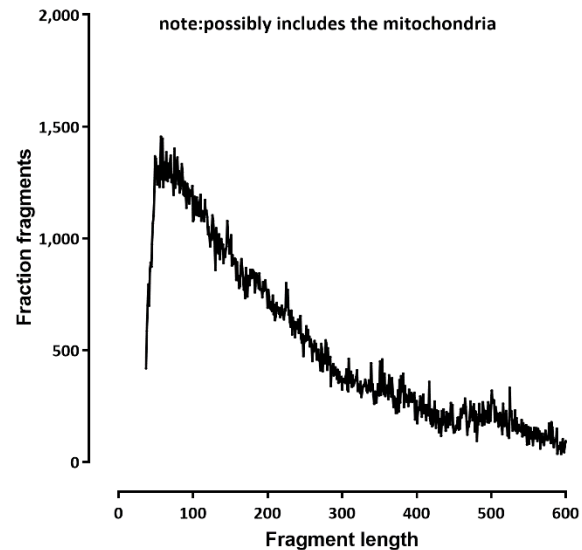
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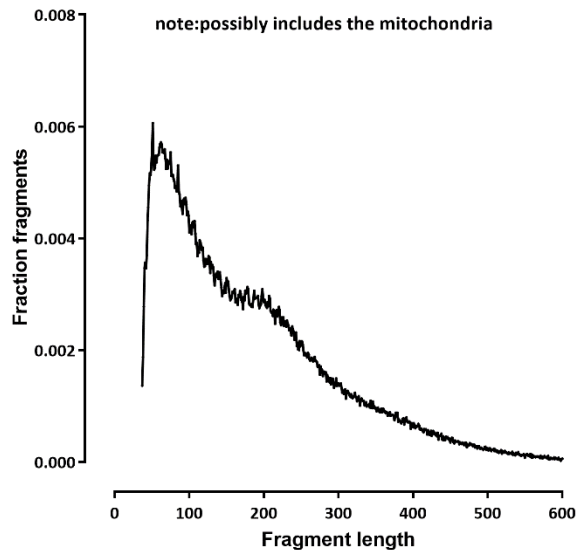


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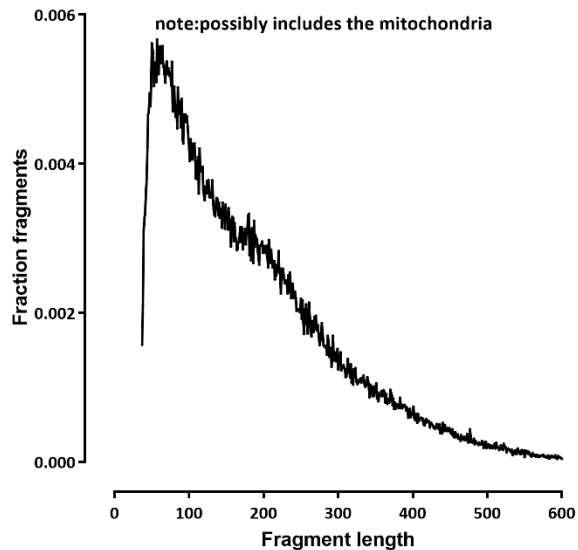


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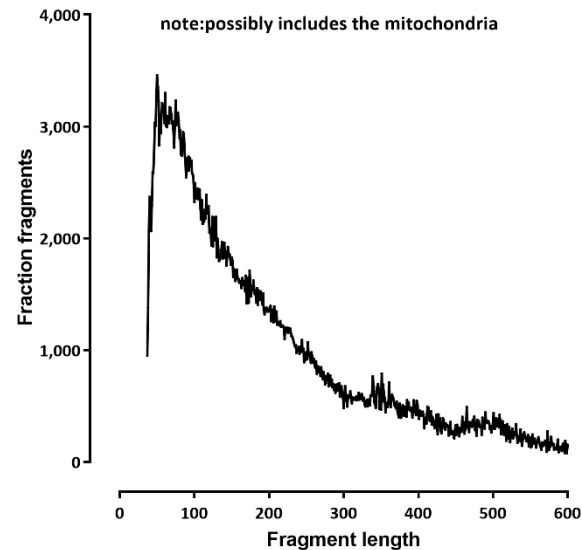
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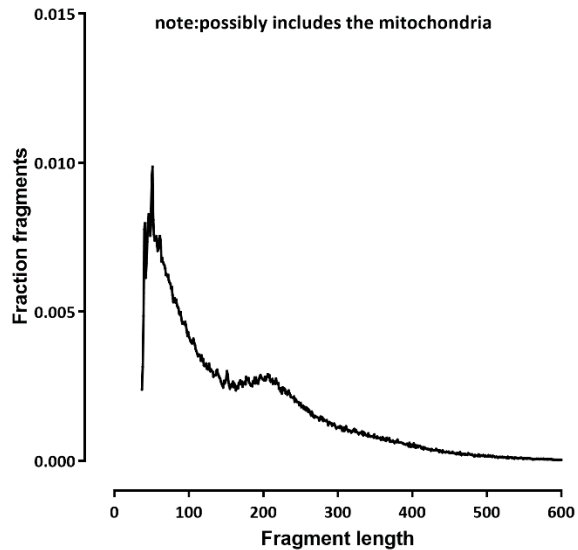


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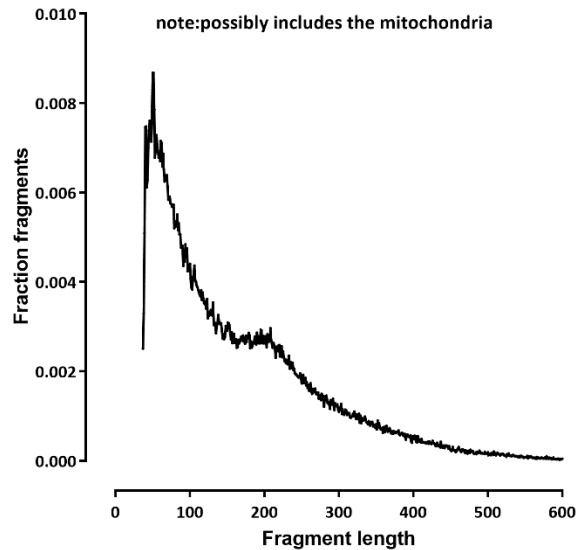


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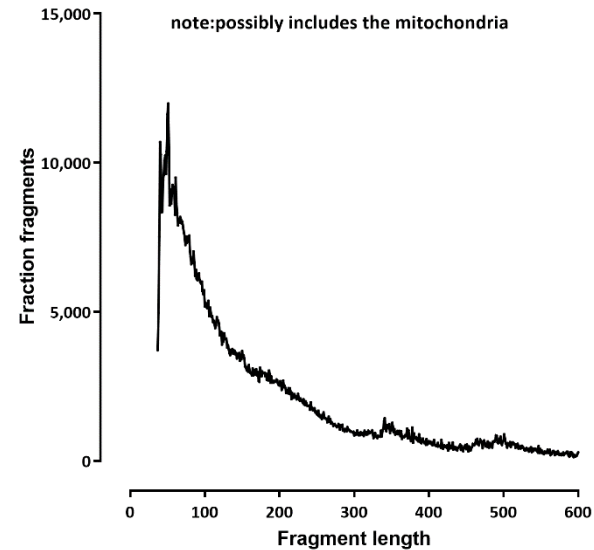
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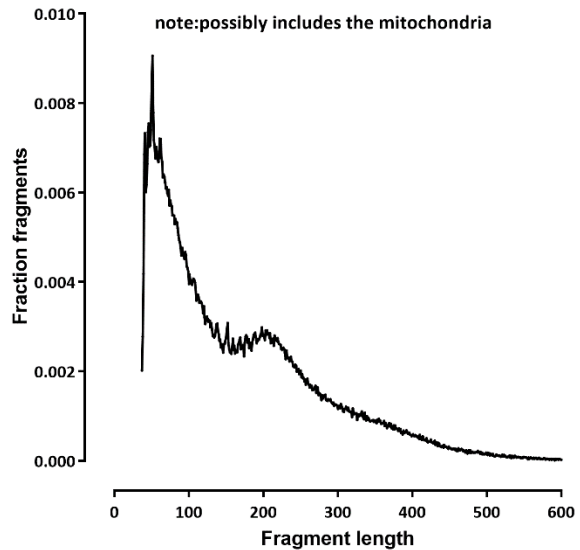


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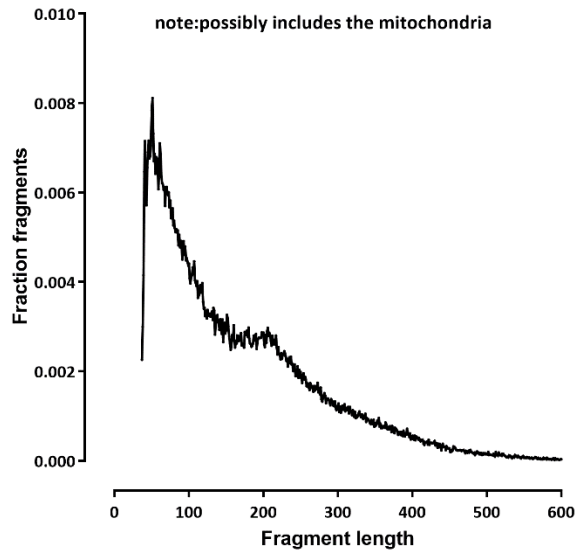


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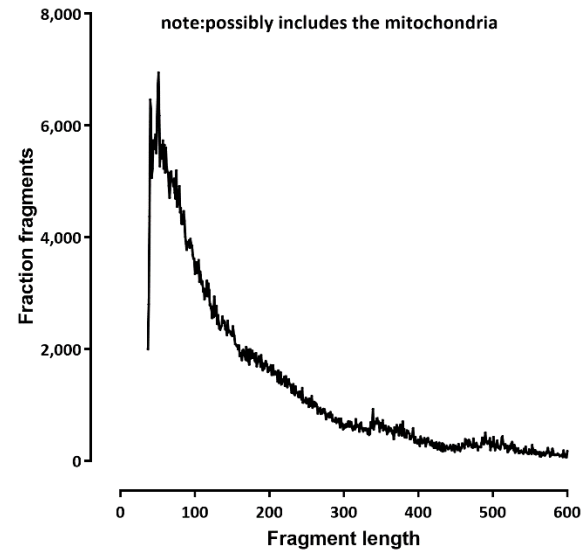
L1915-GM17_vehcile_DVNP_rep2 sacCer3 unique



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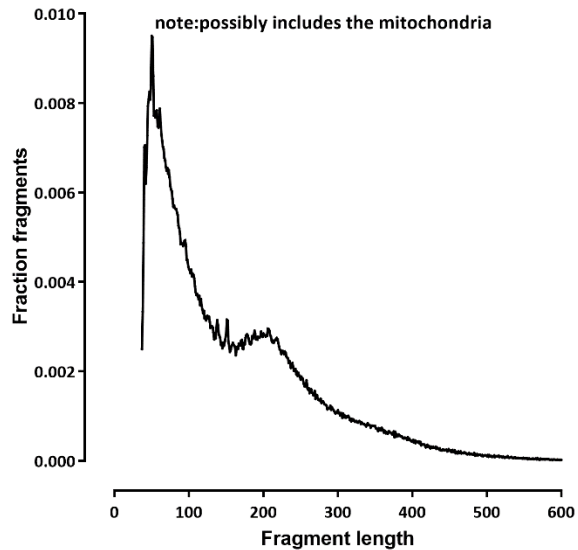


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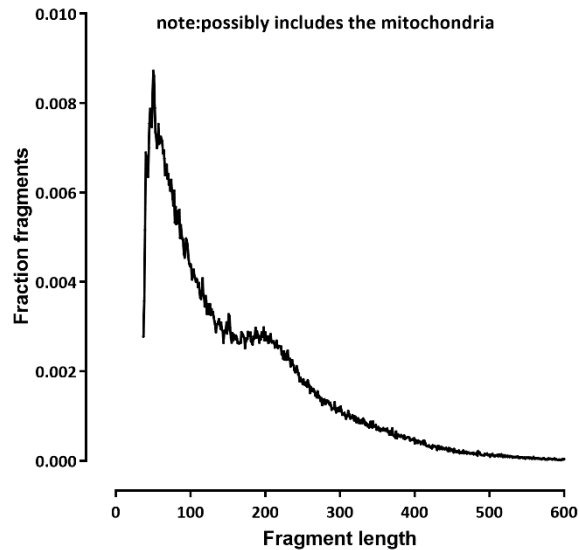


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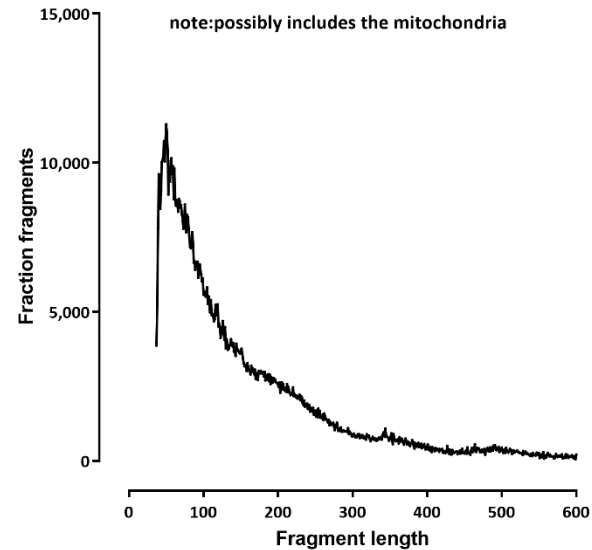
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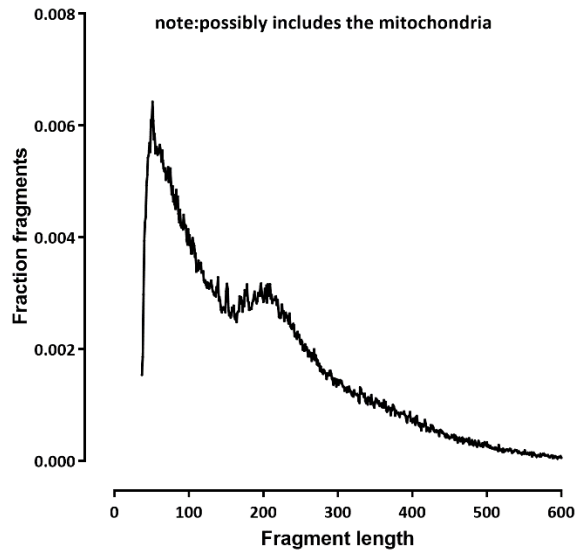


L1916-GM17_vehcile_DVNP_ON_rep1 Candida all

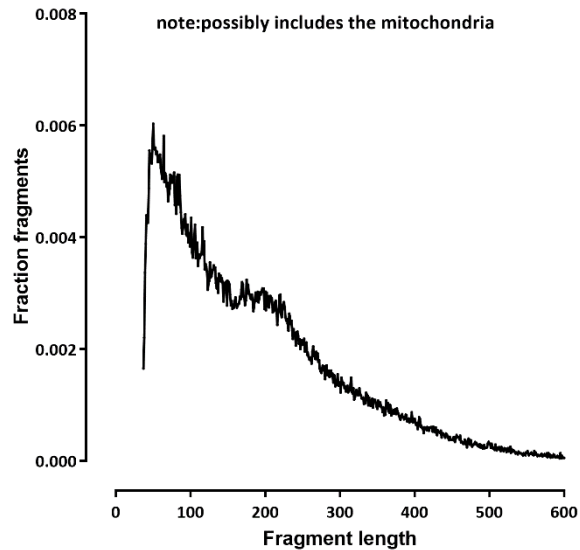


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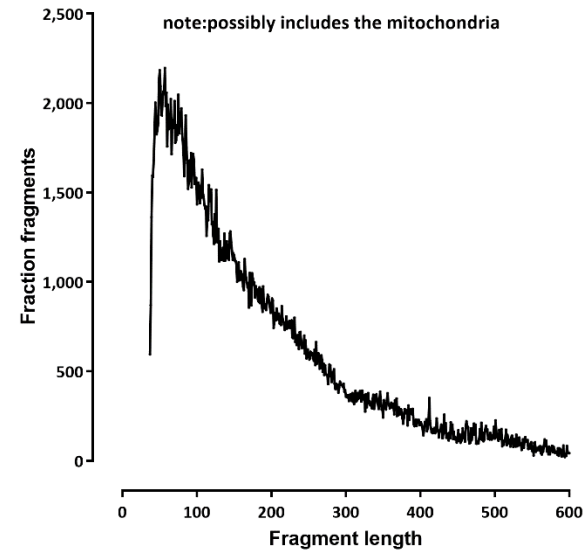
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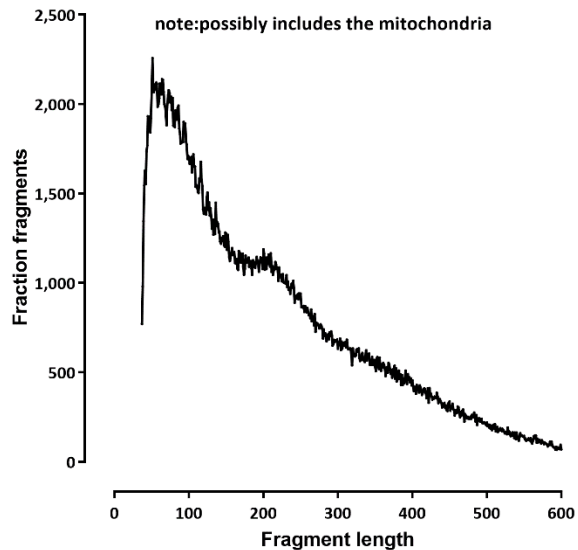


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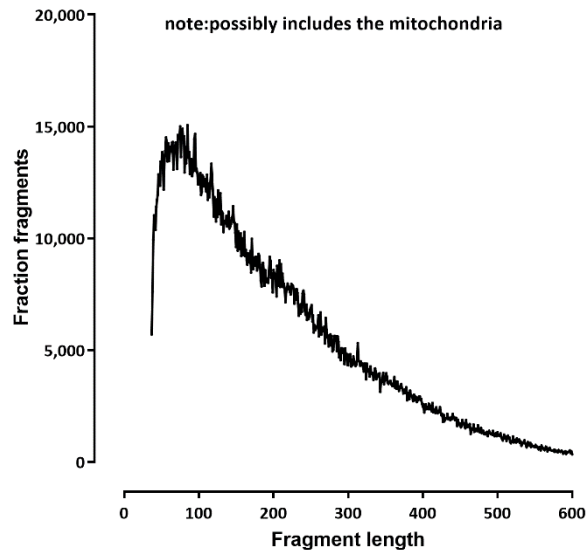


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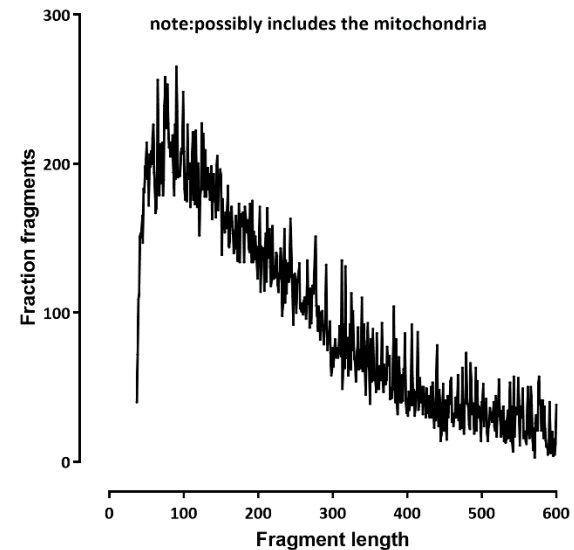
L1918-GM18_rep1 sacCer3 unique



L1918-GM18_rep1 sacCer3 all



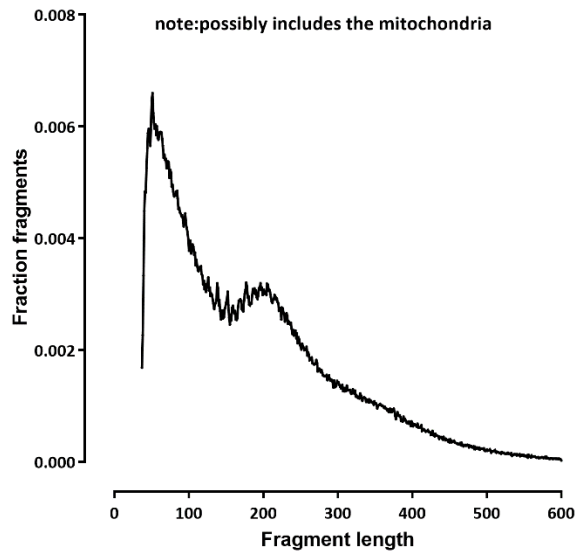
L1918-GM18_rep1 Candida all



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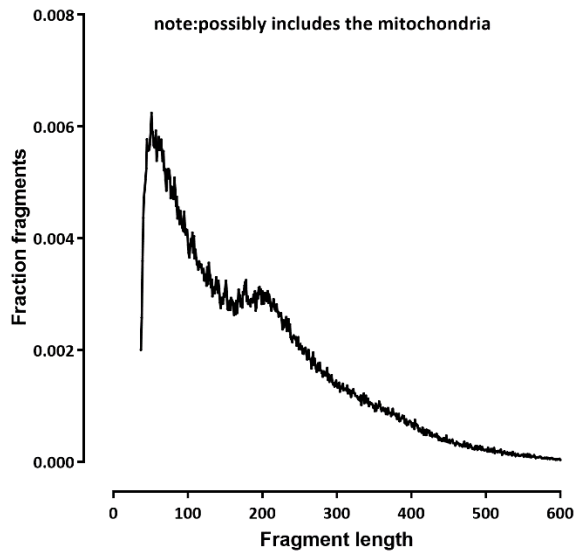
L1919-GM18_rep2 sacCer3 unique

note:possibly includes the mitochondria



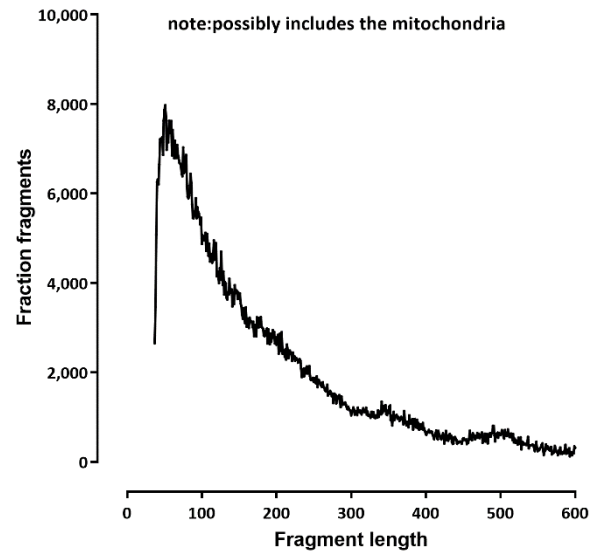
L1919-GM18_rep2 sacCer3 all

note:possibly includes the mitochondria



L1919-GM18_rep2 Candida all

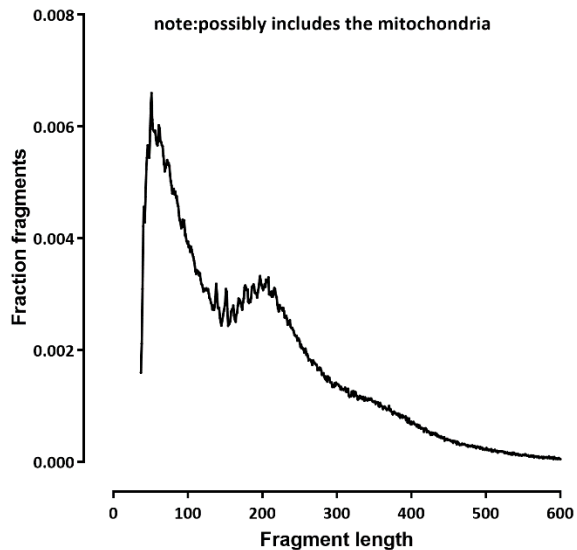
note:possibly includes the mitochondria



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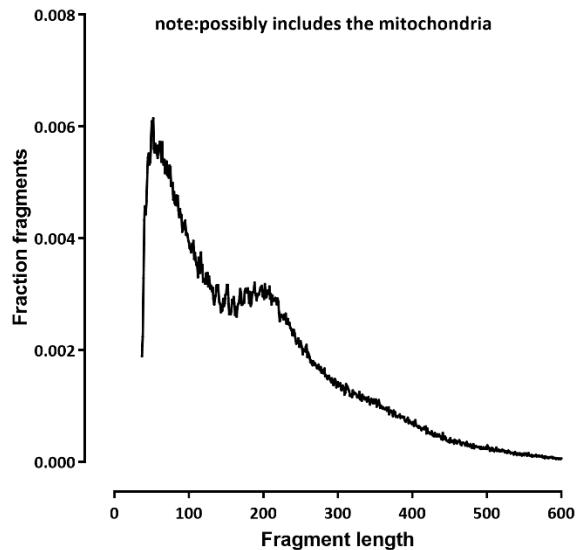
L1920-GM19_rep1 sacCer3 unique

note:possibly includes the mitochondria



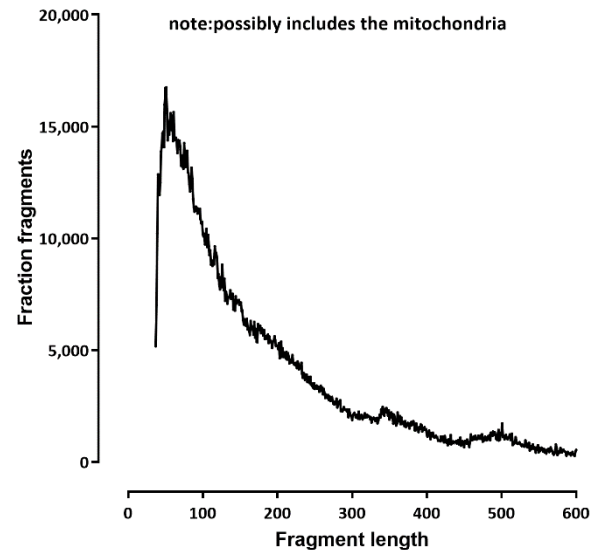
L1920-GM19_rep1 sacCer3 all

note:possibly includes the mitochondria



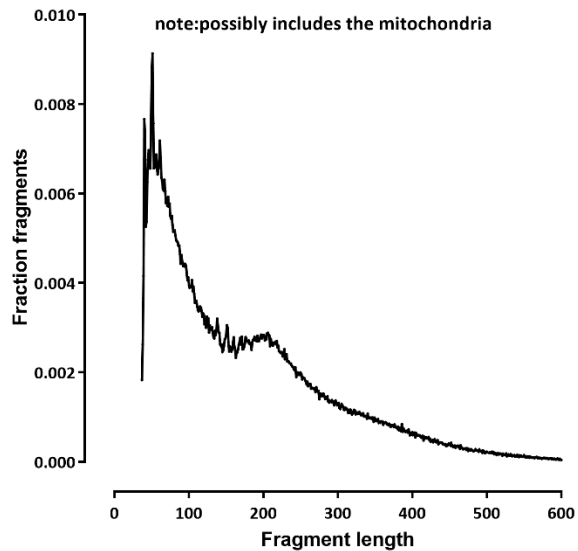
L1920-GM19_rep1 Candida all

note:possibly includes the mitochondria

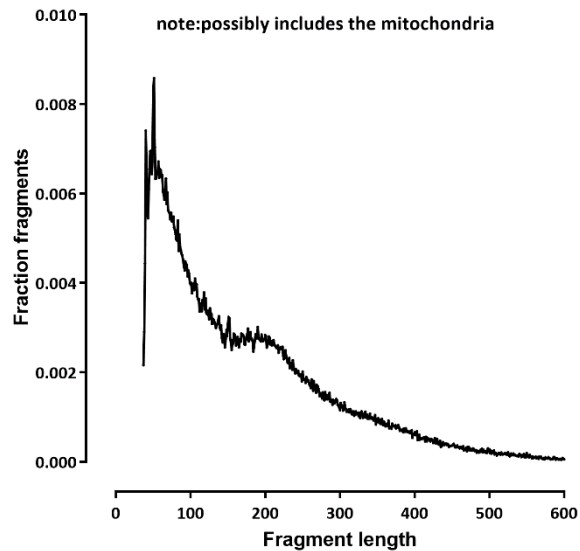


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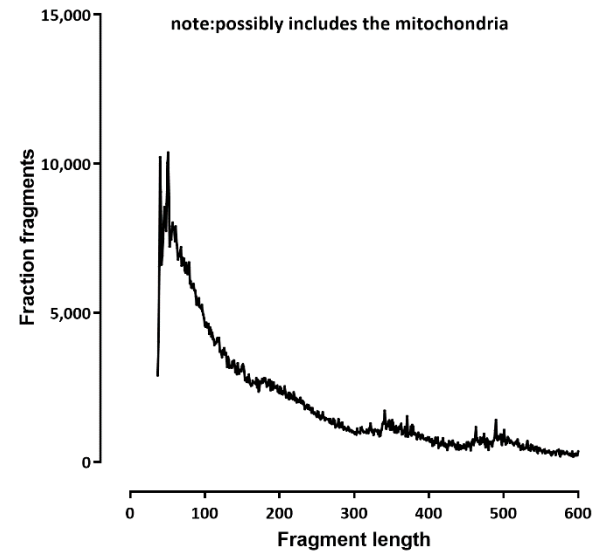
L1921-GM19_rep2 sacCer3 unique



L1921-GM19_rep2 sacCer3 all

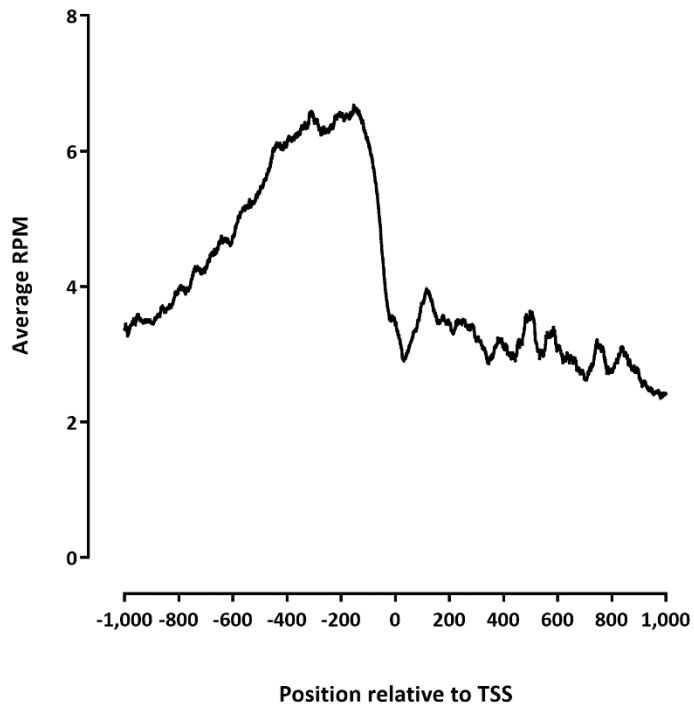


L1921-GM19_rep2 Candida all

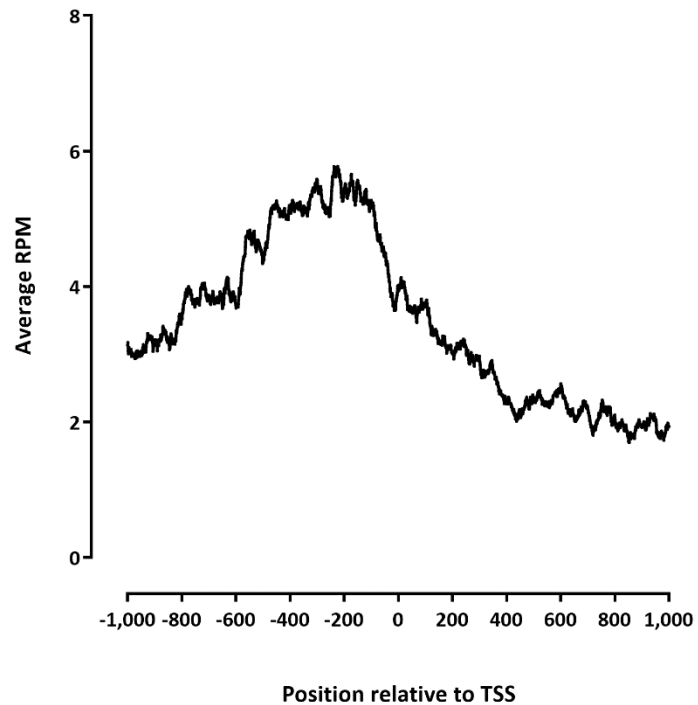


TSS profiles

L1910-GM16_Haematodinium_DVNP_rep1 sacCer3

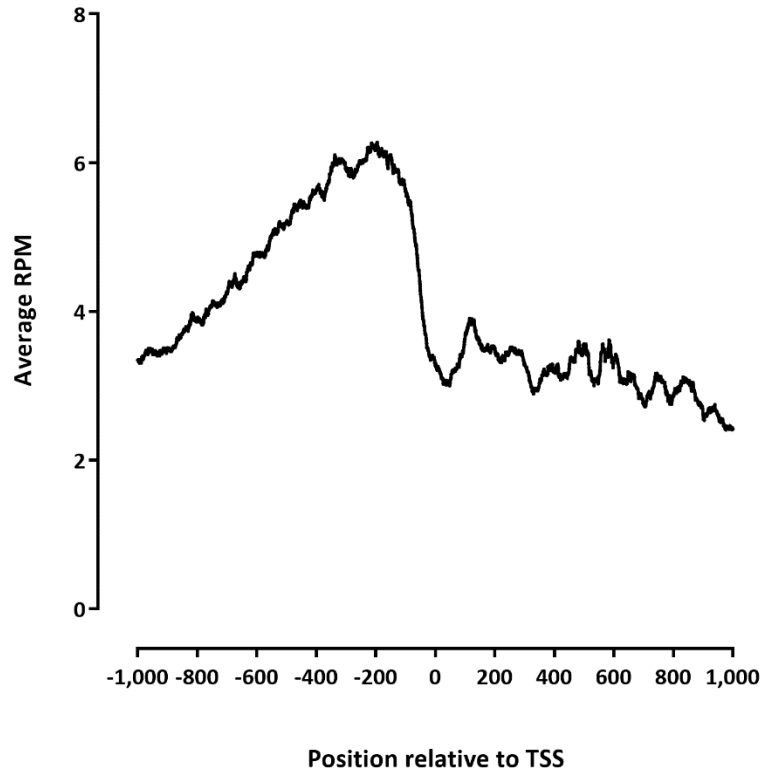


L1910-GM16_Haematodinium_DVNP_rep1 Candida

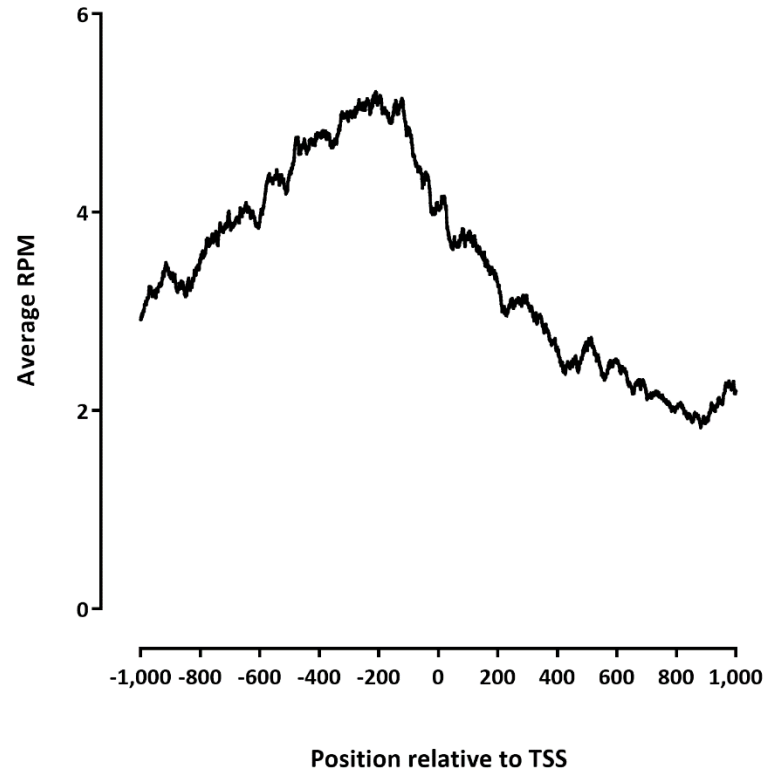


TSS profiles

L1911-GM16_Haematodinium_DVNP_rep2 sacCer3



L1911-GM16_Haematodinium_DVNP_rep2 Candida



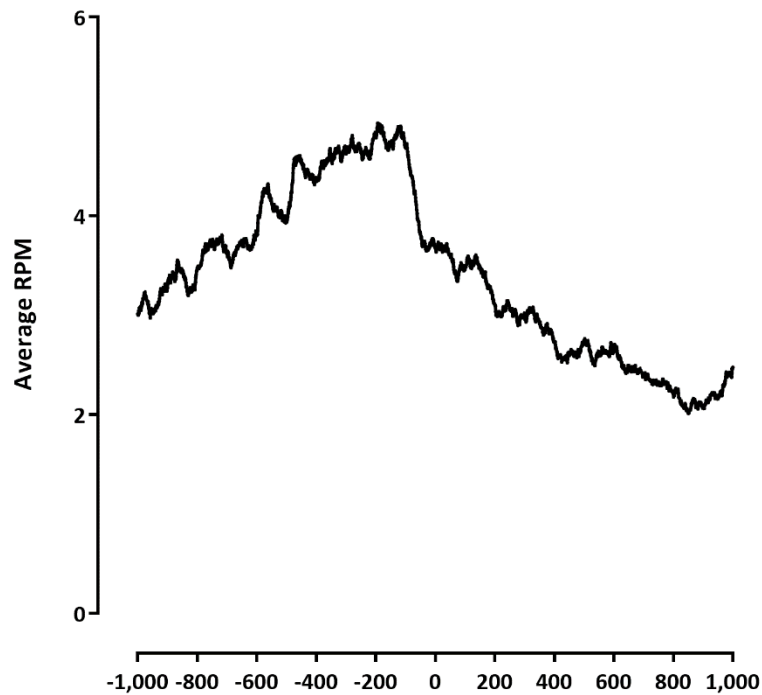
TSS profiles

L1912-GM16_Haematodinium_DVNP_ON_rep1 sacCer3



Position relative to TSS

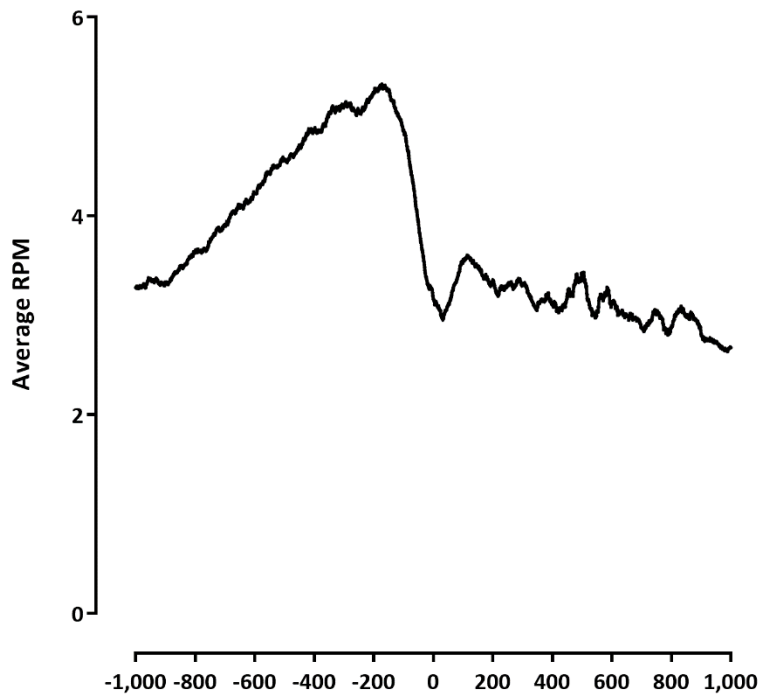
L1912-GM16_Haematodinium_DVNP_ON_rep1 Candida



Position relative to TSS

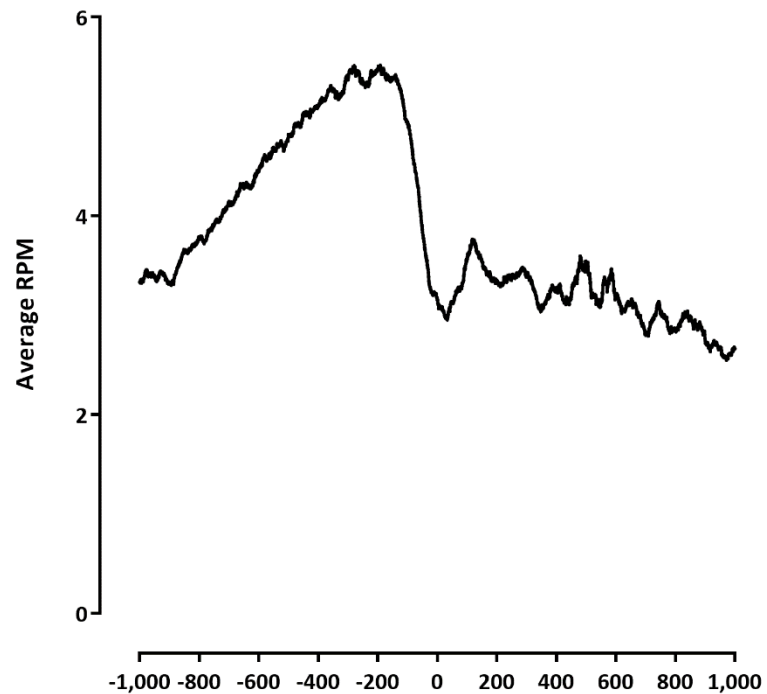
TSS profiles

L1913-GM16_Haematodinium_DVNP_ON_rep2 sacCer3



Position relative to TSS

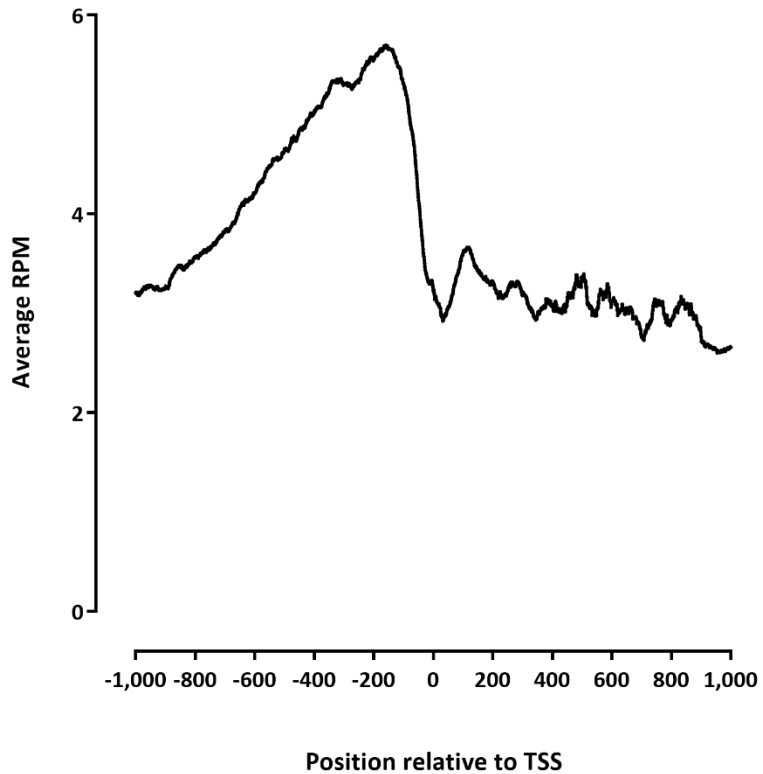
L1912-GM16_Haematodinium_DVNP_ON_rep1 sacCer3



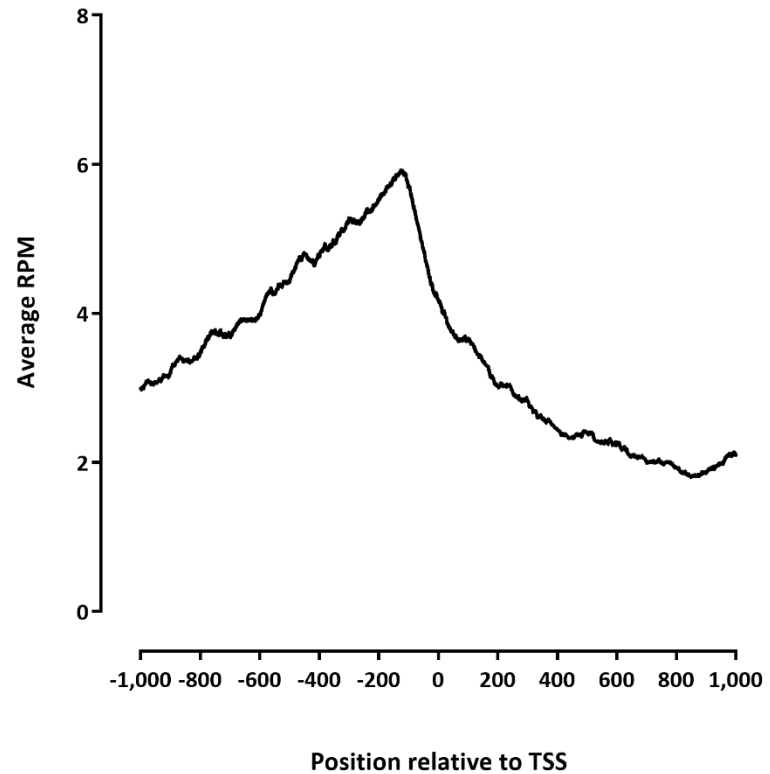
Position relative to TSS

TSS profiles

L1914-GM17_vehcile_DVNP_rep1 sacCer3

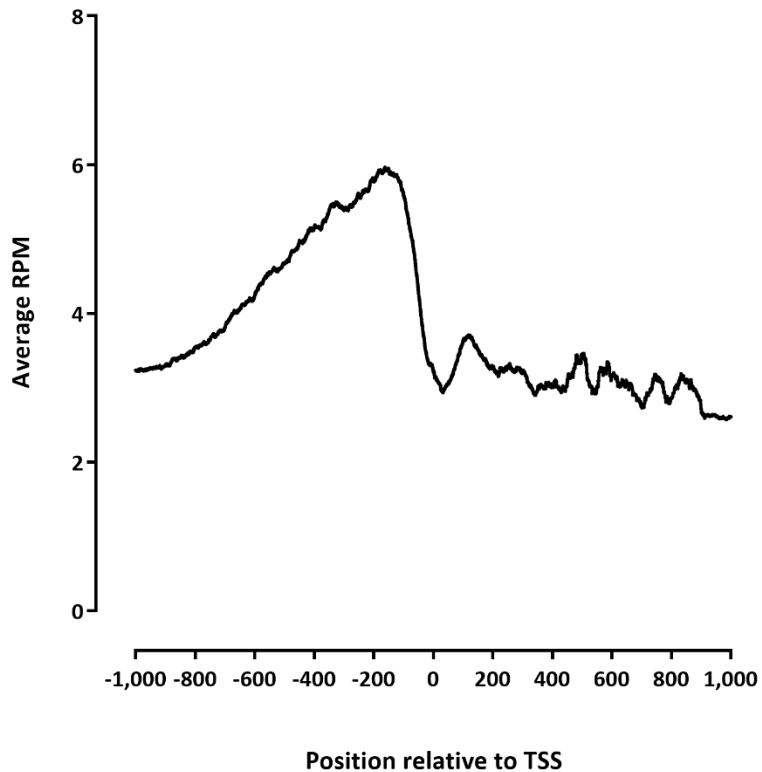


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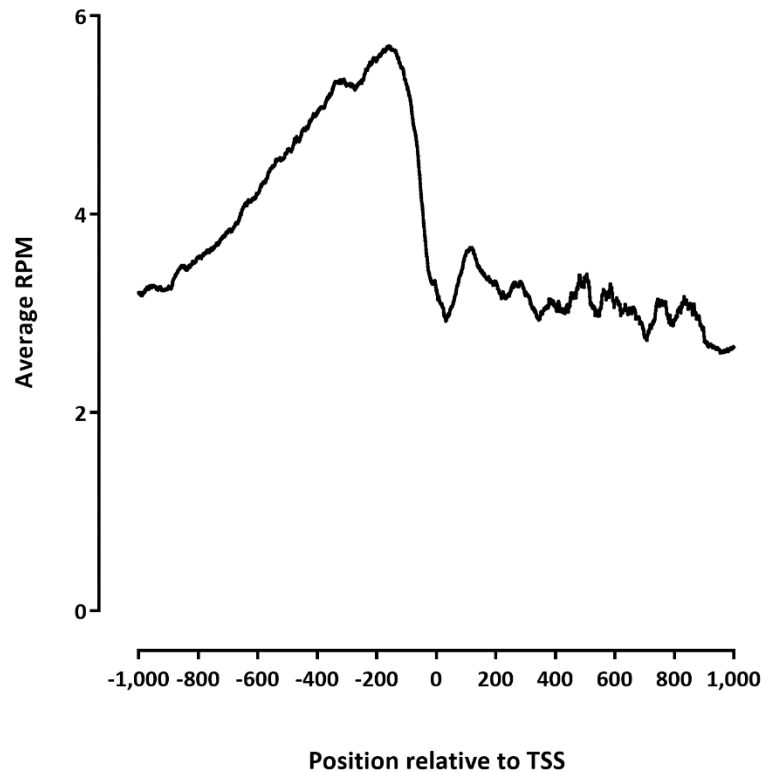


TSS profiles

L1915-GM17_vehcile_DVNP_rep2 sacCer3

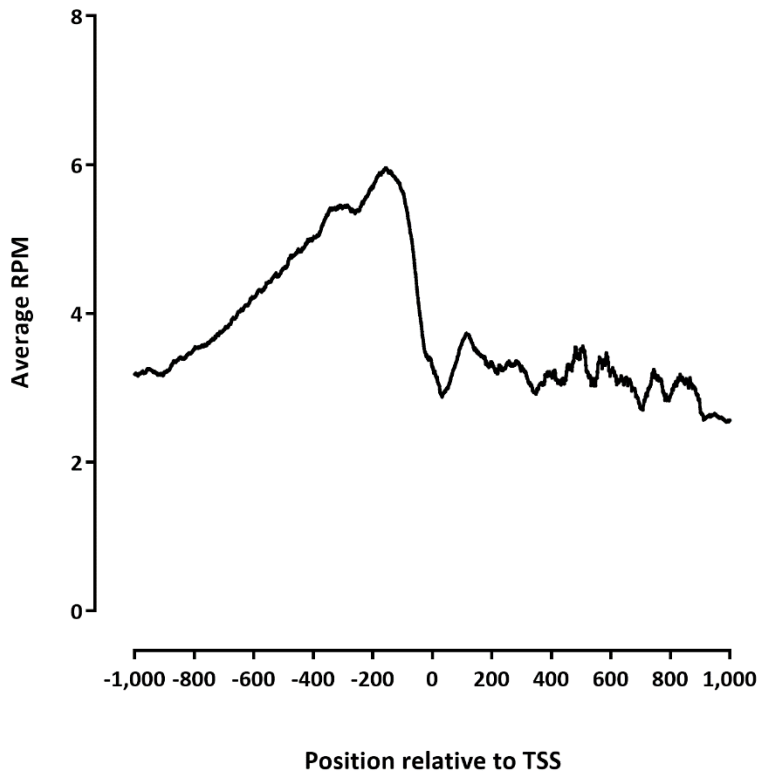


L1914-GM17_vehcile_DVNP_rep1 sacCer3

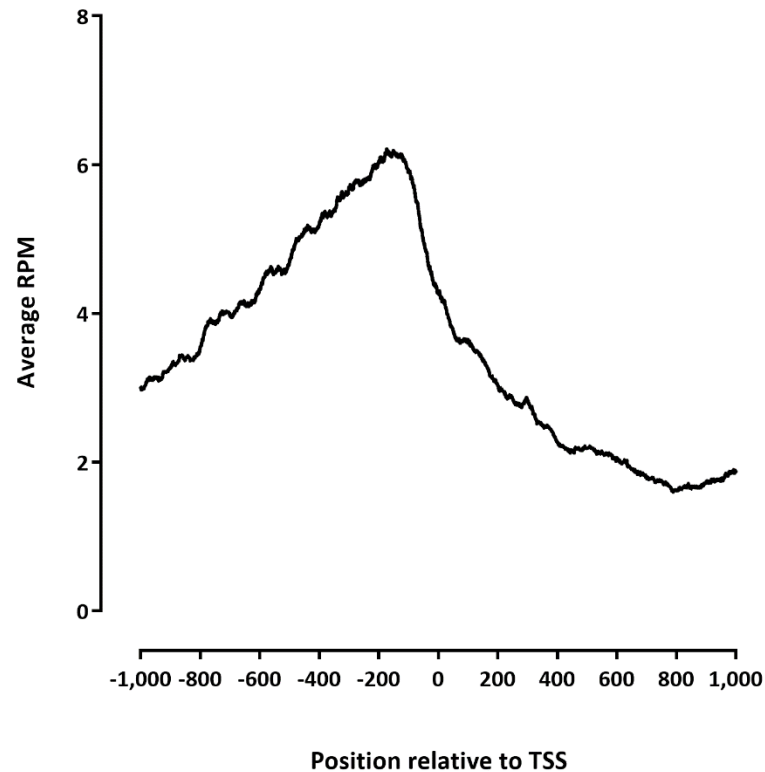


TSS profiles

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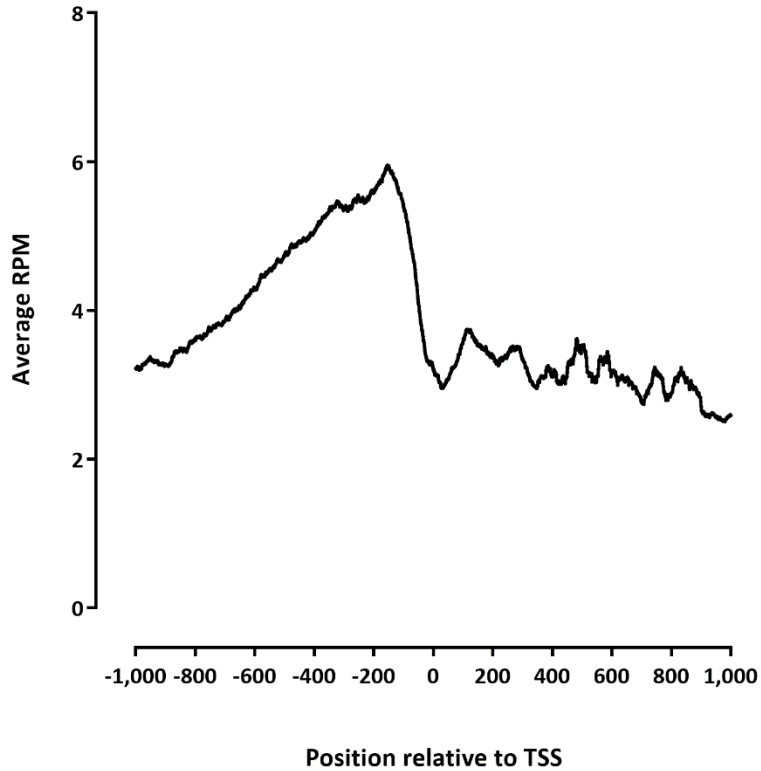


L1916-GM17_vehcile_DVNP_ON_rep1 Candida

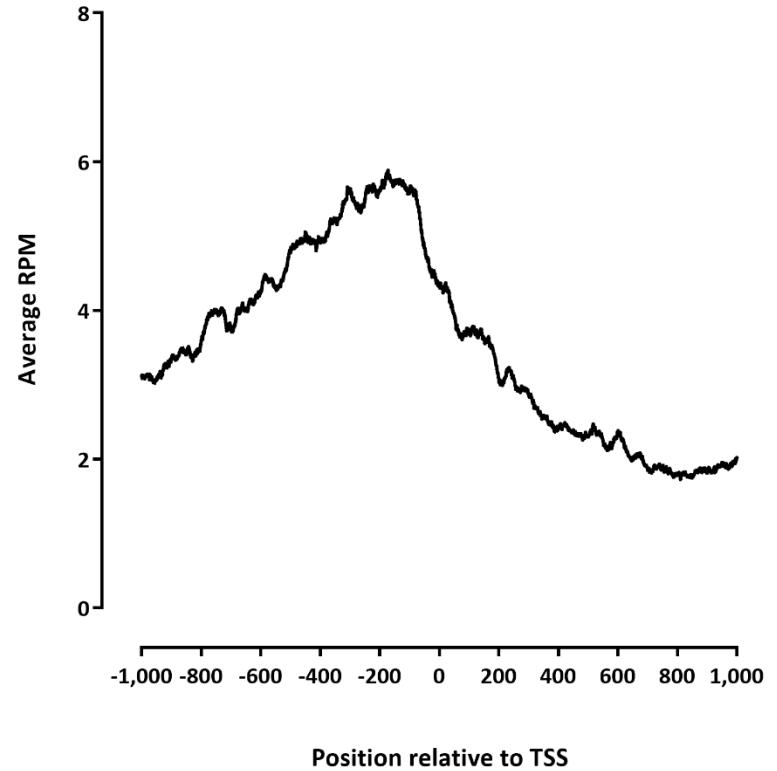


TSS profiles

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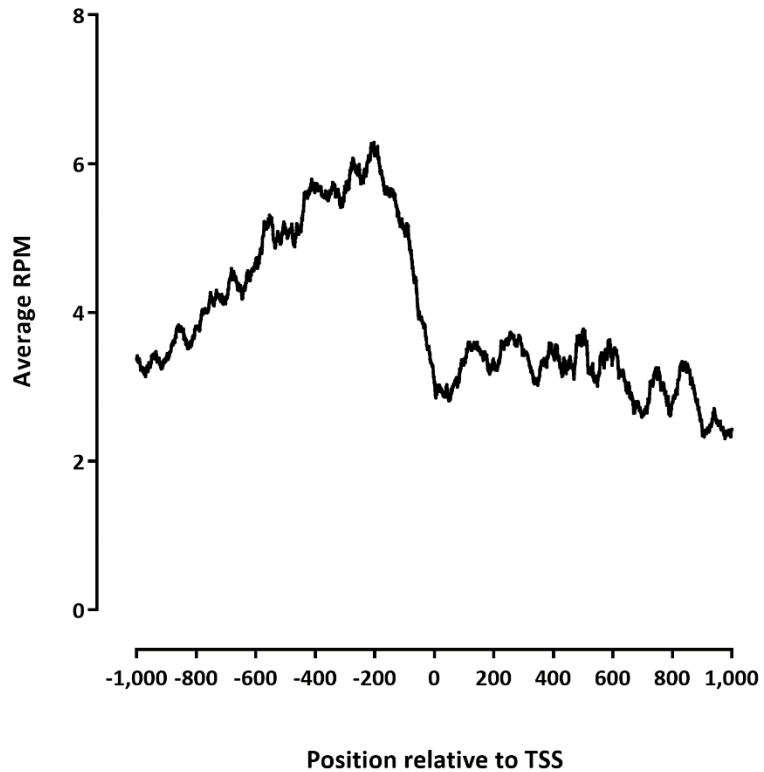


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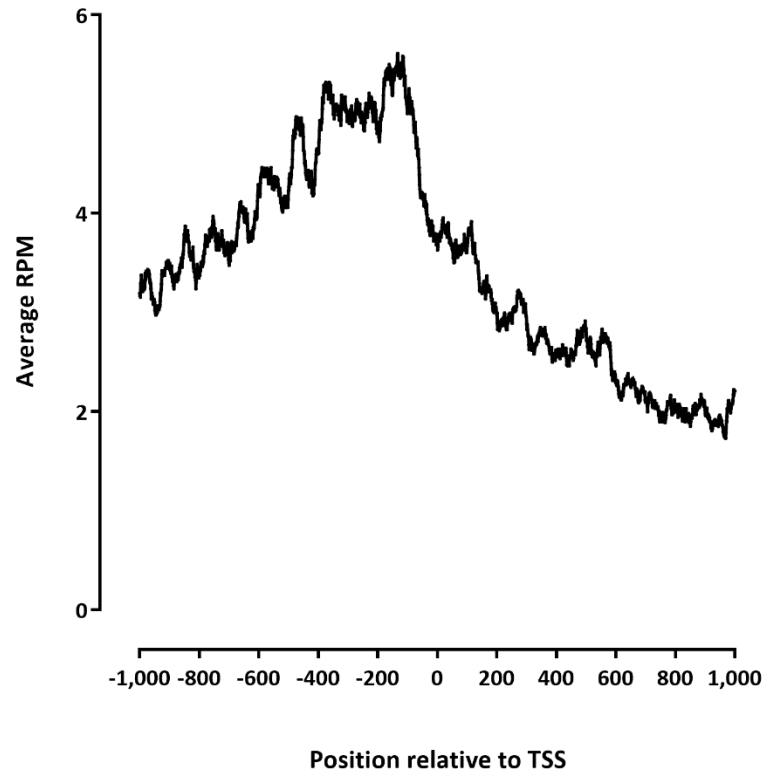


TSS profiles

L1918-GM18_rep1 sacCer3

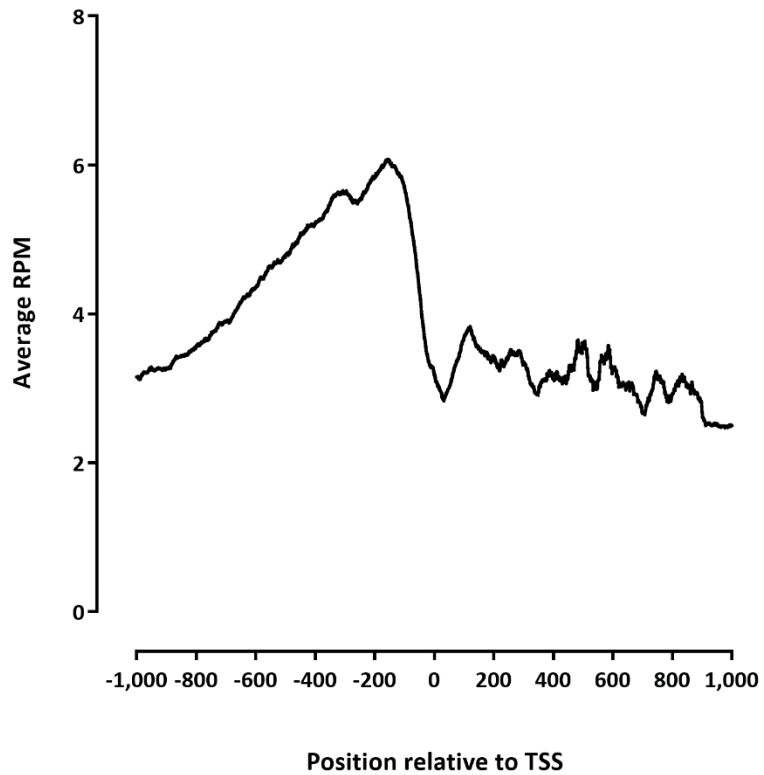


L1918-GM18_rep1 Candida

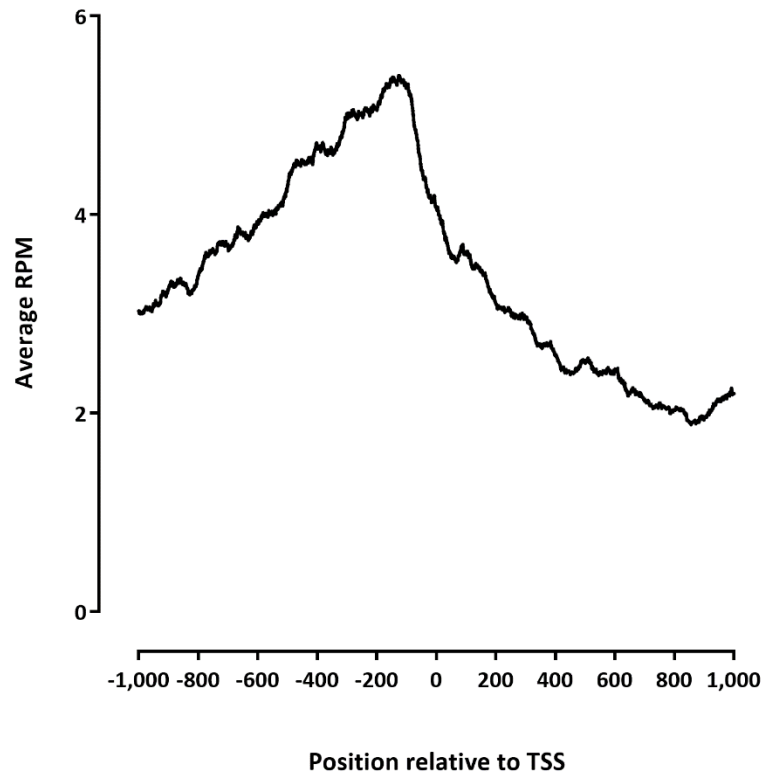


TSS profiles

L1919-GM18_rep2 sacCer3

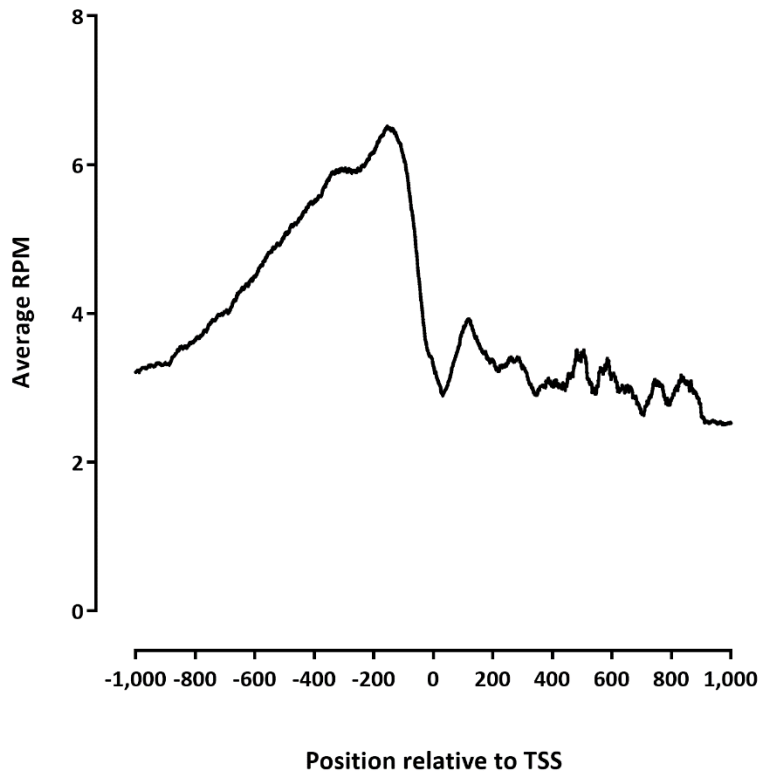


L1919-GM18_rep2 Candida

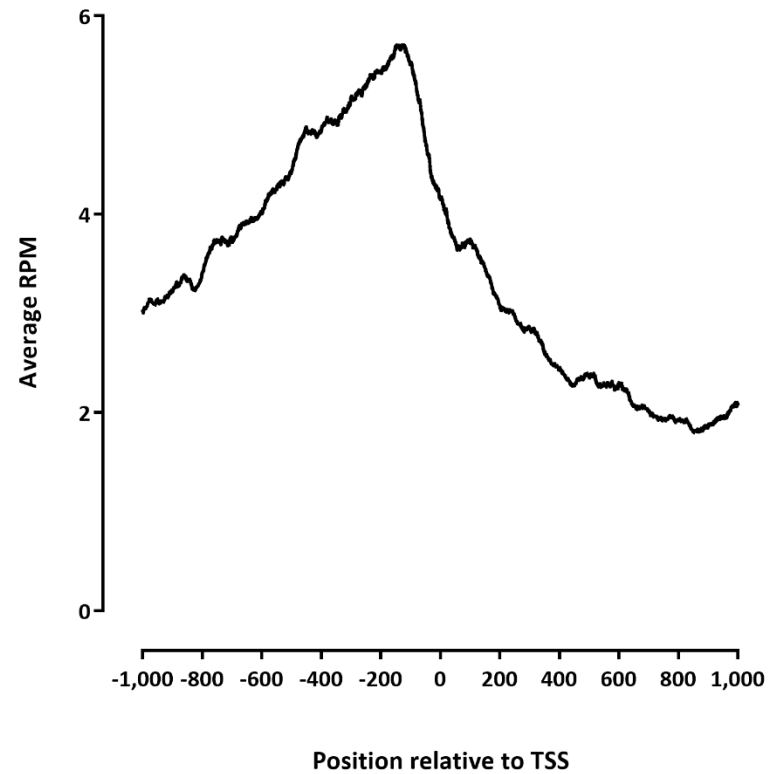


TSS profiles

L1920-GM19_rep1 sacCer3

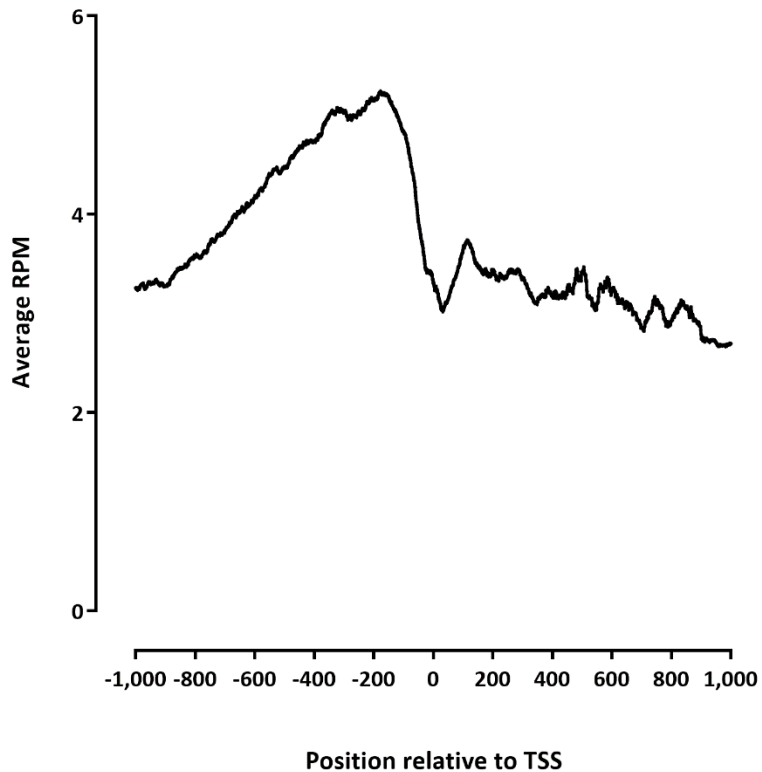


L1920-GM19_rep1 Candida

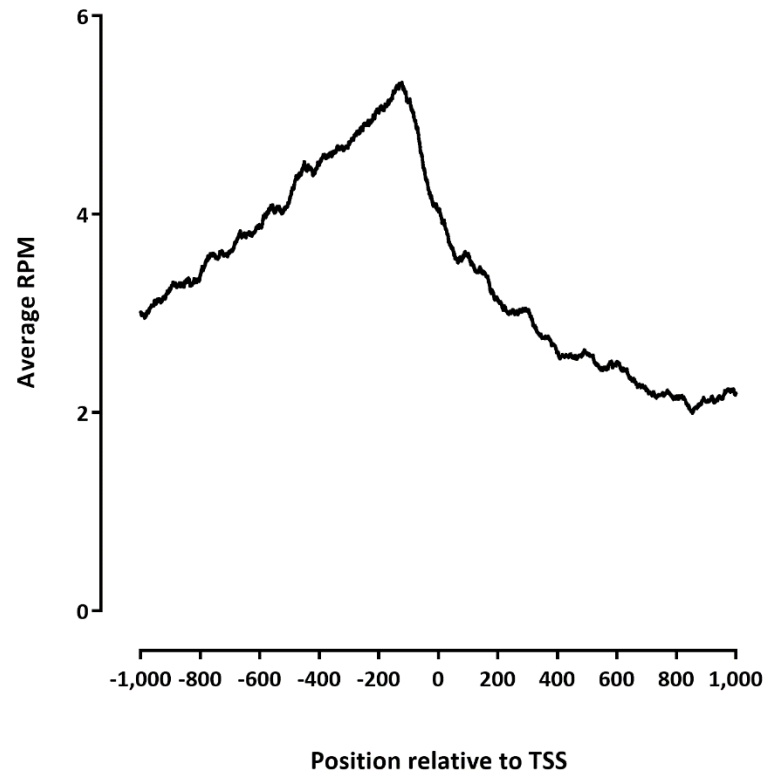


TSS profiles

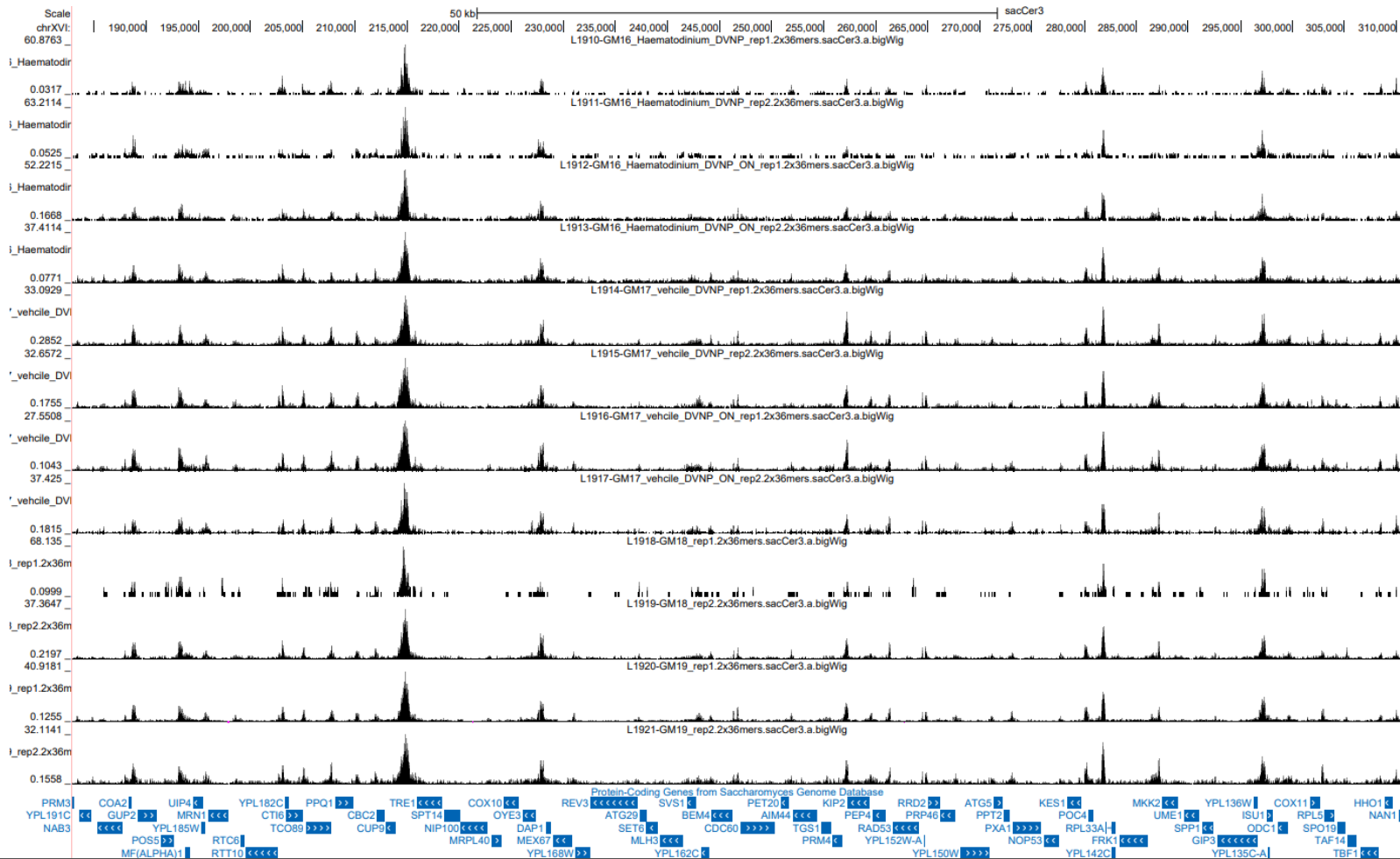
L1921-GM19_rep2 sacCer3



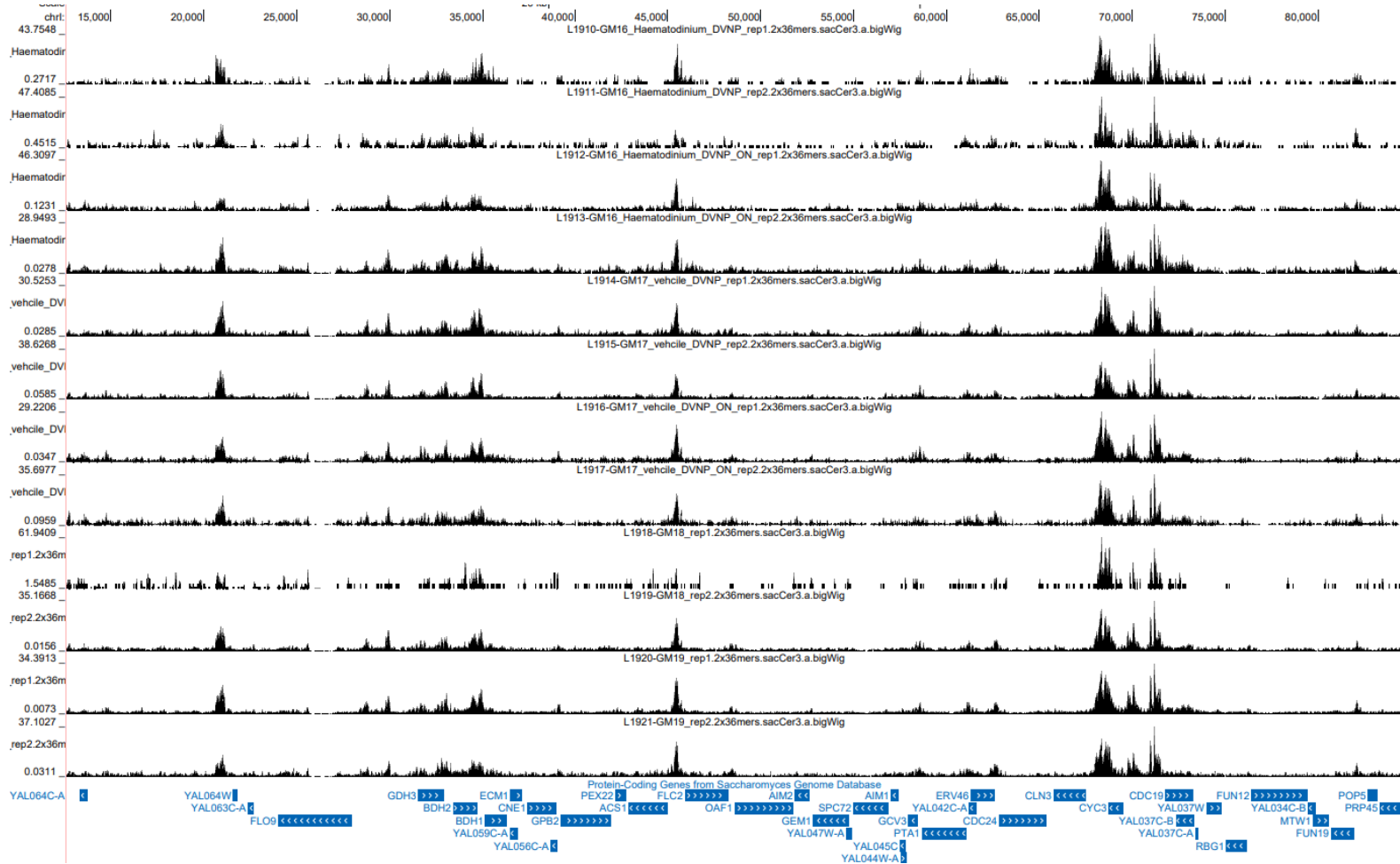
L1921-GM19_rep2 Candida



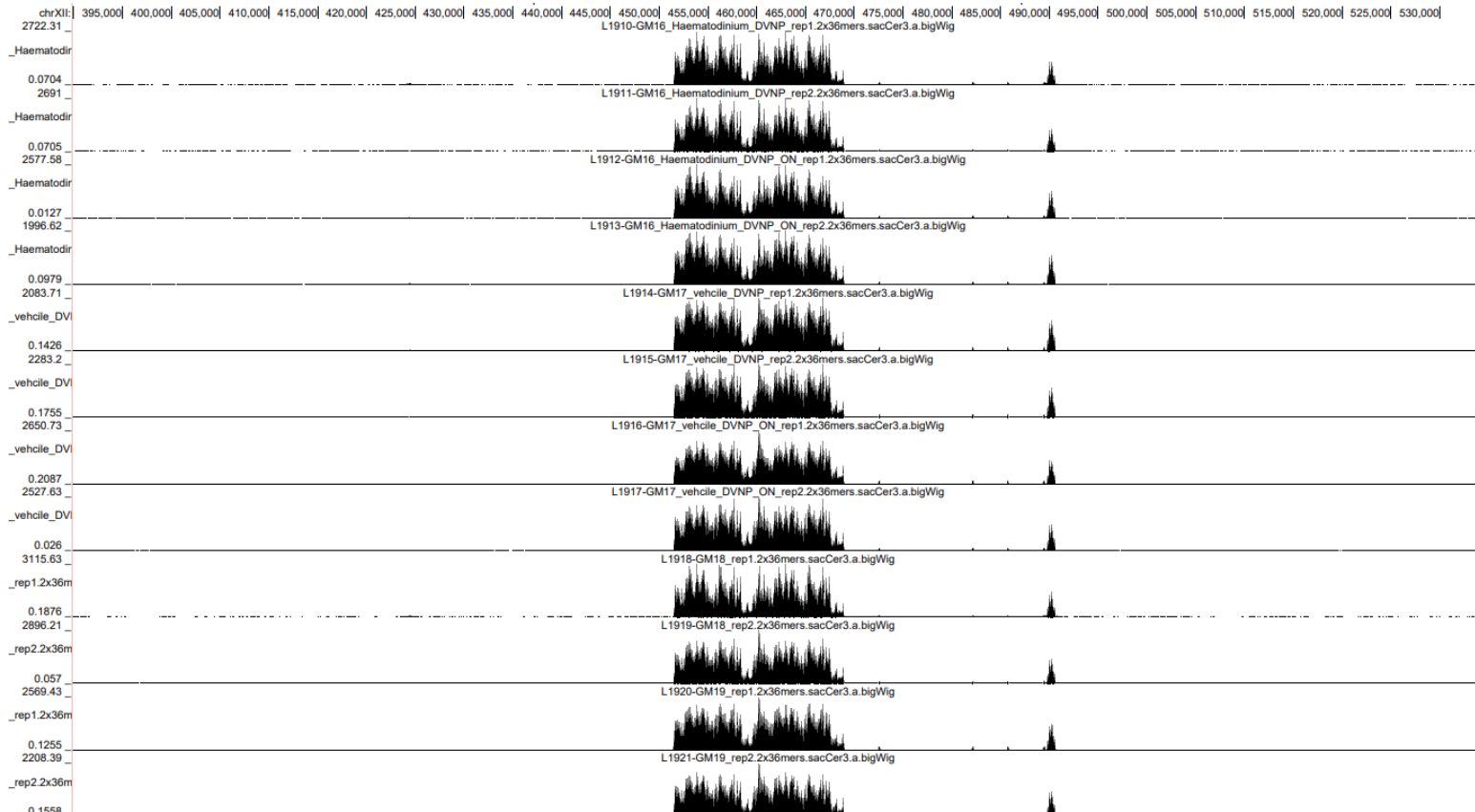
SNAPSHOTS



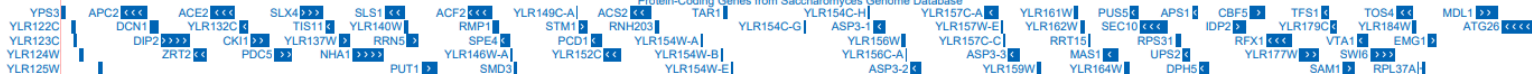
SNAPSHOTS



SNAPSHOTS



Protein-Coding Genes from Saccharomyces Genome Database



SNAPSHOTS

