

# Dataset stats

Library	Raw fragments	Species	2x36mers		1x36mers		Mappability Corrected RPKM	S.cerevisiae/Candida ratio	Correction factor
			Unique non-chrM reads	Complexity	Unique non-chrM reads	Complexity			
180718-L1-MS1-input	3,214,876	Candida	2,006,636	0.94	1,025,556	0.94	32.41	1.626	1.00
	3,214,876	sacCer3	3,083,456	0.93	1,577,836	0.93	52.72		
180718-L2-MS1-input-2	3,092,211	Candida	1,950,518	0.94	995,071	0.94	32.30	1.636	1.00
	3,092,211	sacCer3	3,014,266	0.93	1,539,465	0.93	52.83		
180718-L3-MS1-AB5	3,008,645	Candida	2,116,448	0.93	1,076,373	0.92	34.76	1.445	0.89
	3,008,645	sacCer3	2,887,178	0.93	1,471,672	0.92	50.24		
180718-L4-MS1-AB20	2,604,018	Candida	1,752,552	0.94	894,395	0.93	34.44	1.468	0.90
	2,604,018	sacCer3	2,433,246	0.94	1,242,384	0.93	50.58		
180718-L5-MS1-AB21	3,074,910	Candida	1,759,148	0.94	898,380	0.93	28.54	1.991	1.22
	3,074,910	sacCer3	3,310,860	0.93	1,691,998	0.92	56.82		
180718-L6-MS80-input	3,484,862	Candida	2,350,444	0.94	1,200,047	0.93	34.29	1.480	1.00
	3,484,862	sacCer3	3,285,458	0.93	1,679,597	0.92	50.74		
180718-L7-MS80-input-2	3,071,116	Candida	2,076,694	0.94	1,060,569	0.94	34.09	1.494	1.00
	3,071,116	sacCer3	2,931,312	0.93	1,498,980	0.93	50.94		
180718-L8-MS80-AB5	2,945,037	Candida	2,937,614	0.86	1,492,213	0.83	46.30	0.822	0.55
	2,945,037	sacCer3	2,273,602	0.91	1,159,960	0.90	38.04		
180718-L9-MS80-AB20	2,953,273	Candida	2,075,126	0.94	1,060,125	0.93	35.22	1.413	0.95
	2,953,273	sacCer3	2,773,896	0.94	1,416,776	0.93	49.76		
180718-L10-MS80-AB21	3,045,385	Candida	1,798,472	0.94	917,760	0.93	29.49	1.892	1.27
	3,045,385	sacCer3	3,215,782	0.93	1,642,782	0.92	55.81		
180718-L11-MS81-input	3,385,521	Candida	1,879,578	0.94	958,906	0.94	30.24	1.820	1.00
	3,385,521	sacCer3	3,233,396	0.93	1,650,467	0.92	55.02		
180718-L12-MS81-input-2	3,090,283	Candida	1,725,940	0.95	883,940	0.94	30.22	1.821	1.00
	3,090,283	sacCer3	2,969,806	0.93	1,522,474	0.93	55.03		
180718-L13-MS81-AB5	2,528,554	Candida	1,791,894	0.90	911,245	0.89	34.07	1.496	0.82
	2,528,554	sacCer3	2,526,524	0.90	1,289,403	0.89	50.97		
180718-L14-MS81-AB20	2,970,858	Candida	1,963,452	0.94	1,001,161	0.92	34.61	1.456	0.80
	2,970,858	sacCer3	2,705,322	0.93	1,379,258	0.93	50.40		
180718-L15-MS81-AB21	3,015,619	Candida	1,599,940	0.94	817,644	0.94	26.97	2.169	1.19
	3,015,619	sacCer3	3,281,438	0.93	1,677,249	0.92	58.48		
180718-L16-MS63-0nM-input	3,219,057	Candida	1,970,208	0.94	1,005,283	0.94	31.28	1.724	0.99
	3,219,057	sacCer3	3,208,120	0.93	1,638,965	0.92	53.92		
180718-L17-MS63-0nM-input-2	2,970,048	Candida	1,817,766	0.94	927,775	0.94	31.02	1.747	1.01
	2,970,048	sacCer3	2,999,584	0.93	1,533,413	0.92	54.20		
180718-L18-MS63-0nM-AB5	2,062,545	Candida	1,935,996	0.90	984,206	0.89	44.97	0.877	0.51
	2,062,545	sacCer3	1,601,960	0.93	816,475	0.92	39.44		
180718-L19-MS63-0nM-AB20	2,867,443	Candida	1,896,728	0.94	968,704	0.93	34.22	1.484	0.86
	2,867,443	sacCer3	2,665,092	0.93	1,360,234	0.93	50.80		

180718-L20-MS63-0nM-AB21	3,140,027	Candida	1,673,058	0.94	854,787	0.94	26.95	2.170	1.25
	3,140,027	sacCer3	3,433,350	0.93	1,755,006	0.92	58.49		
180718-L21-MS63-10nM-input	3,469,209	Candida	1,816,198	0.94	927,465	0.94	28.08	2.041	0.99
	3,469,209	sacCer3	3,503,064	0.93	1,790,392	0.92	57.30		
180718-L22-MS63-10nM-input-2	3,045,791	Candida	1,602,434	0.95	822,838	0.94	27.88	2.063	1.01
	3,045,791	sacCer3	3,120,900	0.93	1,605,640	0.92	57.51		
180718-L23-MS63-10nM-AB5	2,628,594	Candida	1,765,838	0.91	897,919	0.91	34.07	1.496	0.73
	2,628,594	sacCer3	2,490,418	0.91	1,270,395	0.91	50.96		
180718-L24-MS63-10nM-AB20	2,951,695	Candida	2,020,840	0.94	1,032,459	0.93	35.45	1.396	0.68
	2,951,695	sacCer3	2,669,872	0.93	1,363,711	0.92	49.51		
180718-L25-MS63-10nM-AB21	3,068,165	Candida	1,535,736	0.95	784,586	0.94	25.06	2.414	1.18
	3,068,165	sacCer3	3,504,956	0.92	1,791,915	0.92	60.50		
180718-L26-MS63-30nM-input	3,170,544	Candida	1,709,772	0.95	871,468	0.94	29.69	1.873	1.00
	3,170,544	sacCer3	3,024,464	0.93	1,543,975	0.93	55.60		
180718-L27-MS63-30nM-input-2	3,081,742	Candida	1,694,146	0.95	864,049	0.94	29.78	1.863	1.00
	3,081,742	sacCer3	2,981,146	0.93	1,523,065	0.93	55.50		
180718-L28-MS63-30nM-AB5	2,727,247	Candida	1,672,386	0.94	850,105	0.93	31.35	1.717	0.92
	2,727,247	sacCer3	2,711,936	0.93	1,380,821	0.93	53.84		
180718-L29-MS63-30nM-AB20	2,742,472	Candida	1,954,986	0.94	998,156	0.93	35.34	1.404	0.75
	2,742,472	sacCer3	2,596,786	0.93	1,326,011	0.93	49.63		
180718-L30-MS63-30nM-AB21	2,228,798	Candida	1,231,456	0.95	628,836	0.95	27.02	2.162	1.16
	2,228,798	sacCer3	2,515,582	0.94	1,285,803	0.93	58.42		
180723-L1-MS63-0nM-input1	5,643,003	Candida	2,540,404	0.94	1,293,243	0.93	23.24	2.685	1.00
	5,643,003	sacCer3	6,447,414	0.90	3,285,048	0.89	62.41		
180723-L2-MS63-0nM-AB1	5,839,477	Candida	2,304,434	0.91	1,173,357	0.90	19.17	3.481	1.30
	5,839,477	sacCer3	7,577,116	0.84	3,863,278	0.80	66.72		
180723-L3-MS63-0nM-AB4	5,704,272	Candida	3,036,070	0.89	1,545,916	0.88	25.62	2.338	0.87
	5,704,272	sacCer3	6,703,410	0.85	3,418,522	0.82	59.90		
180723-L4-MS63-0nM-AB5	6,368,849	Candida	5,076,622	0.84	2,577,872	0.80	38.29	1.215	0.45
	6,368,849	sacCer3	5,814,208	0.85	2,961,707	0.82	46.51		
180723-L5-MS63-0nM-AB20	4,943,749	Candida	2,271,024	0.94	1,171,061	0.93	23.75	2.605	0.97
	4,943,749	sacCer3	5,593,354	0.91	2,885,885	0.90	61.88		
180723-L6-MS63-0nM-AB21	5,519,273	Candida	2,051,766	0.94	1,045,336	0.93	19.02	3.515	1.31
	5,519,273	sacCer3	6,818,036	0.89	3,475,842	0.88	66.87		
180723-L7-MS63-10nM-input2	5,569,688	Candida	3,480,192	0.93	1,771,196	0.92	32.33	1.633	1.00
	5,569,688	sacCer3	5,368,582	0.91	2,736,703	0.90	52.81		
180723-L8-MS63-10nM-AB1	5,672,812	Candida	2,181,252	0.91	1,109,060	0.91	18.75	3.583	2.19
	5,672,812	sacCer3	7,380,158	0.84	3,758,586	0.81	67.17		
180723-L9-MS63-10nM-AB4	5,485,145	Candida	2,809,850	0.90	1,429,105	0.88	24.75	2.458	1.51
	5,485,145	sacCer3	6,520,318	0.85	3,322,970	0.82	60.83		

180723-L10-MS63-10nM-AB5	6,184,088	Candida	4,802,148	0.86	2,441,776	0.84	37.44	1.266	0.78
	6,184,088	sacCer3	5,735,278	0.87	2,924,491	0.84	47.40		
180723-L11-MS63-10nM-AB20	5,810,617	Candida	4,151,052	0.92	2,124,283	0.90	35.71	1.379	0.84
	5,810,617	sacCer3	5,414,028	0.90	2,770,212	0.90	49.23		
180723-L12-MS63-10nM-AB21	6,964,535	Candida	3,989,980	0.92	2,033,611	0.91	28.32	2.014	1.23
	6,964,535	sacCer3	7,594,566	0.88	3,874,167	0.87	57.04		
180723-L13-MS63-30nM-input2	6,236,689	Candida	4,465,564	0.92	2,272,581	0.90	37.45	1.266	1.00
	6,236,689	sacCer3	5,338,456	0.91	2,720,669	0.90	47.40		
180723-L14-MS63-30nM-AB1	4,902,169	Candida	1,874,464	0.92	954,291	0.91	18.46	3.655	2.89
	4,902,169	sacCer3	6,468,000	0.85	3,298,945	0.83	67.47		
180723-L15-MS63-30nM-AB4	5,393,853	Candida	2,612,572	0.90	1,328,839	0.89	23.19	2.693	2.13
	5,393,853	sacCer3	6,644,000	0.85	3,385,718	0.82	62.47		
180723-L16-MS63-30nM-AB5	5,044,284	Candida	3,841,552	0.86	1,954,818	0.83	36.78	1.308	1.03
	5,044,284	sacCer3	4,737,512	0.86	2,418,086	0.84	48.10		
180723-L17-MS63-30nM-AB20	5,025,505	Candida	4,166,708	0.91	2,125,639	0.90	40.33	1.100	0.87
	5,025,505	sacCer3	4,331,356	0.92	2,211,031	0.91	44.35		
180723-L18-MS63-30nM-AB21	4,805,780	Candida	3,470,076	0.92	1,770,945	0.91	35.00	1.428	1.13
	4,805,780	sacCer3	4,680,760	0.91	2,391,977	0.90	49.98		
180731-L1-MS1-input	6,793,256	Candida	3,454,074	0.93	1,764,614	0.92	26.71	2.199	1.00
	6,793,256	sacCer3	7,177,916	0.89	3,670,746	0.88	58.75		
180731-L2-MS1-AB1	7,166,795	Candida	2,596,028	0.90	1,320,898	0.89	17.02	4.053	1.84
	7,166,795	sacCer3	9,928,710	0.81	5,064,367	0.76	68.99		
180731-L3-MS1-AB4	6,755,125	Candida	3,517,668	0.88	1,788,847	0.86	24.33	2.518	1.15
	6,755,125	sacCer3	8,360,608	0.83	4,261,048	0.78	61.27		
180731-L4-MS1-AB5	7,194,784	Candida	4,900,754	0.87	2,494,202	0.85	32.91	1.586	0.72
	7,194,784	sacCer3	7,334,642	0.87	3,742,071	0.85	52.20		
180731-L5-MS1-AB20	6,100,053	Candida	3,347,722	0.93	1,708,400	0.92	27.62	2.092	0.95
	6,100,053	sacCer3	6,622,332	0.90	3,380,792	0.89	57.78		
180731-L6-MS1-AB21	6,286,510	Candida	2,770,430	0.93	1,416,761	0.92	22.10	2.879	1.31
	6,286,510	sacCer3	7,540,756	0.89	3,857,923	0.88	63.62		
180731-L7-MS80-input	11,401,236	Candida	15,874,400	0.83	8,095,392	0.80	67.91	0.224	1.00
	11,401,236	sacCer3	3,354,218	0.93	1,713,284	0.92	15.19		
180731-L8-MS80-AB1	10,575,754	Candida	9,982,258	0.79	5,080,180	0.73	43.45	0.945	4.22
	10,575,754	sacCer3	8,893,738	0.82	4,539,953	0.78	41.05		
180731-L9-MS80-AB4	12,628,595	Candida	13,664,556	0.75	6,960,422	0.66	49.54	0.699	3.12
	12,628,595	sacCer3	9,006,710	0.82	4,599,542	0.77	34.61		
180731-L10-MS80-AB5	5,701,581	Candida	4,400,326	0.85	2,235,090	0.82	36.36	1.335	5.96
	5,701,581	sacCer3	5,538,956	0.86	2,822,477	0.83	48.54		
180731-L11-MS80-AB20	12,542,415	Candida	19,693,706	0.79	10,055,258	0.75	71.90	0.153	0.68
	12,542,415	sacCer3	2,841,092	0.93	1,452,048	0.92	10.98		

180731-L12-MS80-AB21	11,799,303	Candida	16,419,182	0.82	8,385,308	0.78	63.96	0.303	1.35
	11,799,303	sacCer3	4,698,258	0.91	2,401,775	0.90	19.37		
180731-L13-MS81-input	5,723,145	Candida	2,571,502	0.94	1,313,427	0.93	23.54	2.639	1.00
	5,723,145	sacCer3	6,412,870	0.90	3,278,357	0.89	62.10		
180731-L14-MS81-AB1	5,852,367	Candida	2,175,116	0.91	1,108,336	0.90	17.65	3.872	1.47
	5,852,367	sacCer3	7,952,128	0.83	4,059,729	0.79	68.33		
180731-L15-MS81-AB4	6,814,027	Candida	3,650,470	0.88	1,860,717	0.86	25.27	2.385	0.90
	6,814,027	sacCer3	8,219,232	0.83	4,198,144	0.79	60.27		
180731-L16-MS81-AB5	6,278,111	Candida	5,028,684	0.85	2,562,385	0.81	37.98	1.233	0.47
	6,278,111	sacCer3	5,846,642	0.85	2,989,615	0.82	46.84		
180731-L17-MS81-AB20	6,976,868	Candida	3,212,680	0.93	1,638,663	0.92	23.40	2.660	1.01
	6,976,868	sacCer3	8,083,596	0.88	4,123,946	0.87	62.25		
180731-L18-MS81-AB21	6,349,351	Candida	2,521,902	0.94	1,285,575	0.93	20.00	3.291	1.25
	6,349,351	sacCer3	7,844,820	0.88	4,002,686	0.87	65.84		
180731-L19-MS55-1-input	14,470,672	Candida	21,638,396	0.79	11,024,127	0.75	72.60	0.141	1.00
	14,470,672	sacCer3	2,883,244	0.93	1,470,938	0.93	10.24		
180731-L20-MS55-1-AB1	12,551,795	Candida	15,786,380	0.77	8,032,511	0.69	56.82	0.474	3.36
	12,551,795	sacCer3	7,054,730	0.85	3,599,172	0.82	26.92		
180731-L21-MS55-1-AB4	12,138,333	Candida	15,823,254	0.76	8,084,307	0.68	58.97	0.418	2.96
	12,138,333	sacCer3	6,238,832	0.86	3,195,296	0.83	24.64		
180731-L22-MS55-1-AB5	10,929,840	Candida	13,645,678	0.76	6,933,185	0.67	57.38	0.459	3.25
	10,929,840	sacCer3	5,896,218	0.85	3,008,730	0.82	26.32		
180731-L23-MS55-1-AB20	11,969,771	Candida	19,994,346	0.80	10,221,316	0.76	75.68	0.092	0.65
	11,969,771	sacCer3	1,744,028	0.94	891,973	0.94	6.98		
180731-L24-MS55-1-AB21	11,665,732	Candida	17,912,346	0.81	9,160,706	0.78	69.71	0.191	1.35
	11,665,732	sacCer3	3,225,670	0.93	1,651,785	0.93	13.29		
180731-L25-MS55-2-input	6,088,789	Candida	3,740,794	0.92	1,908,273	0.91	33.65	1.528	1.00
	6,088,789	sacCer3	5,401,854	0.91	2,758,193	0.90	51.41		
180731-L26-MS55-2-AB1	6,263,004	Candida	2,517,968	0.90	1,280,646	0.89	19.25	3.461	2.27
	6,263,004	sacCer3	8,224,560	0.83	4,193,302	0.79	66.63		
180731-L27-MS55-2-AB4	6,675,444	Candida	3,455,478	0.88	1,762,239	0.87	24.71	2.463	1.61
	6,675,444	sacCer3	8,035,264	0.83	4,106,328	0.79	60.86		
180731-L28-MS55-2-AB5	12,427,697	Candida	17,028,036	0.75	8,655,369	0.66	63.57	0.311	0.20
	12,427,697	sacCer3	4,995,072	0.87	2,547,751	0.85	19.78		
180731-L29-MS55-2-AB20	6,444,511	Candida	4,898,064	0.91	2,498,467	0.90	38.33	1.212	0.79
	6,444,511	sacCer3	5,618,344	0.90	2,865,536	0.89	46.47		
180731-L30-MS55-2-AB21	6,862,932	Candida	3,947,628	0.92	2,017,408	0.91	28.70	1.974	1.29
	6,862,932	sacCer3	7,371,908	0.89	3,766,716	0.88	56.65		
180731-L31-MS55-3-input	10,680,482	Candida	15,897,406	0.83	8,105,455	0.80	73.82	0.121	1.00
	10,680,482	sacCer3	1,821,114	0.94	929,838	0.94	8.95		

180731-L32-MS55-3-AB1	11,260,849	Candida	14,463,024	0.77	7,361,203	0.69	57.98	0.443	3.66
	11,260,849	sacCer3	6,042,640	0.86	3,086,409	0.83	25.70		
180731-L33-MS55-3-AB4	11,613,170	Candida	15,551,524	0.75	7,917,711	0.67	60.22	0.387	3.20
	11,613,170	sacCer3	5,678,554	0.86	2,900,460	0.83	23.32		
180731-L34-MS55-3-AB5	10,637,016	Candida	16,389,150	0.79	8,332,263	0.73	70.39	0.179	1.48
	10,637,016	sacCer3	2,763,274	0.92	1,407,774	0.92	12.57		
180731-L35-MS55-3-AB20	12,163,790	Candida	20,217,640	0.79	10,321,590	0.75	75.68	0.092	0.76
	12,163,790	sacCer3	1,763,914	0.94	900,686	0.94	6.98		
180731-L36-MS55-3-AB21	11,925,606	Candida	18,949,858	0.80	9,669,870	0.77	71.92	0.152	1.26
	11,925,606	sacCer3	2,726,844	0.94	1,393,286	0.93	10.95		
180731-L37-MS55-4-input	12,475,605	Candida	14,944,796	0.83	7,622,179	0.81	60.08	0.391	1.00
	12,475,605	sacCer3	5,514,154	0.91	2,816,702	0.90	23.47		
180731-L38-MS55-4-AB1	12,062,149	Candida	10,019,638	0.81	5,098,983	0.75	38.86	1.181	3.02
	12,062,149	sacCer3	11,160,740	0.80	5,698,751	0.75	45.91		
180731-L39-MS55-4-AB4	12,767,212	Candida	11,949,662	0.76	6,086,622	0.67	43.15	0.959	2.45
	12,767,212	sacCer3	10,803,636	0.80	5,519,576	0.74	41.37		
180731-L40-MS55-4-AB5	11,742,617	Candida	18,408,912	0.74	9,348,792	0.65	70.70	0.173	0.44
	11,742,617	sacCer3	3,005,200	0.90	1,531,851	0.89	12.25		
180731-L41-MS55-4-AB20	11,121,145	Candida	15,409,482	0.82	7,865,114	0.79	64.62	0.289	0.74
	11,121,145	sacCer3	4,209,972	0.92	2,149,169	0.91	18.67		
180731-L42-MS55-4-AB21	11,023,445	Candida	12,786,052	0.84	6,526,649	0.82	54.41	0.542	1.39
	11,023,445	sacCer3	6,544,902	0.89	3,343,899	0.88	29.47		