

Mapping stats:

Aligned against	Read Length	Library	Raw fragments	Unique non-chrM reads	Multi reads	Complexity	chrM reads	chrM fraction	Unique non-chrM reads after dedup	TSS ratio
Homo sapiens hg38-male	2x36	L4358-M1-UT	32,207,277	40,873,920		0.86	432,482	0.01	38,909,630	14.85
Homo sapiens hg38-male	2x36	L4359-M1-UT	18,736,413	23,820,730		0.88	690,944	0.03	22,769,786	16.01
Homo sapiens hg38-male	2x36	L4360-M1-UT	17,845,581	22,793,890		0.88	483,668	0.02	21,853,216	15.72
Homo sapiens hg38-male	2x36	L4361-M1-AS	33,739,059	42,243,842		0.83	685,310	0.02	40,130,164	17.22
Homo sapiens hg38-male	2x36	L4362-M1-AS	18,206,004	23,089,864		0.87	637,442	0.03	22,064,134	18.13
Homo sapiens hg38-male	2x36	L4363-M1-AS	20,534,539	25,927,748		0.86	766,680	0.03	24,723,480	17.84
Homo sapiens hg38-male	2x36	L4364-M1-HS	31,368,396	40,043,556		0.83	503,096	0.01	38,104,440	16.44
Homo sapiens hg38-male	2x36	L4365-M1-HS	19,506,111	24,747,342		0.85	992,686	0.04	23,696,248	17.51
Homo sapiens hg38-male	2x36	L4366-M1-HS	29,003,020	36,743,380		0.83	1,489,832	0.04	34,896,714	17.63
Homo sapiens hg38-male	2x36	L4367-M2-UT	46,553,912	60,239,316		0.81	499,102	0.01	56,866,288	15.00
Homo sapiens hg38-male	2x36	L4368-M2-UT	29,267,471	37,746,166		0.85	851,182	0.02	36,060,732	14.60
Homo sapiens hg38-male	2x36	L4369-M2-UT	28,599,280	36,806,228		0.85	925,434	0.02	35,202,064	14.70
Homo sapiens hg38-male	2x36	L4370-M2-AS	48,167,381	62,743,170		0.78	343,952	0.01	58,799,112	15.40
Homo sapiens hg38-male	2x36	L4371-M2-AS	39,946,364	52,110,606		0.82	511,688	0.01	49,255,382	14.94
Homo sapiens hg38-male	2x36	L4372-M2-AS	35,636,343	46,204,670		0.83	508,336	0.01	43,875,180	15.00
Homo sapiens hg38-male	2x36	L4373-M2-HS	20,569,329	25,773,164		0.86	1,464,124	0.05	24,685,726	15.73
Homo sapiens hg38-male	2x36	L4374-M2-HS	40,890,859	52,003,066		0.81	2,255,388	0.04	49,015,154	15.25
Homo sapiens hg38-male	2x36	L4375-M2-HS	59,484,917	76,803,936		0.76	579,230	0.01	70,788,456	14.79