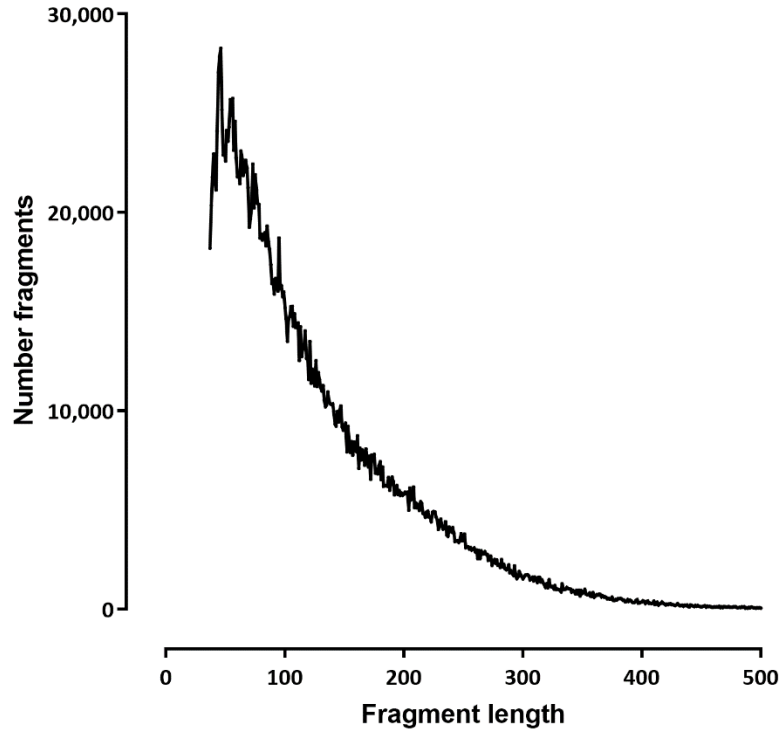


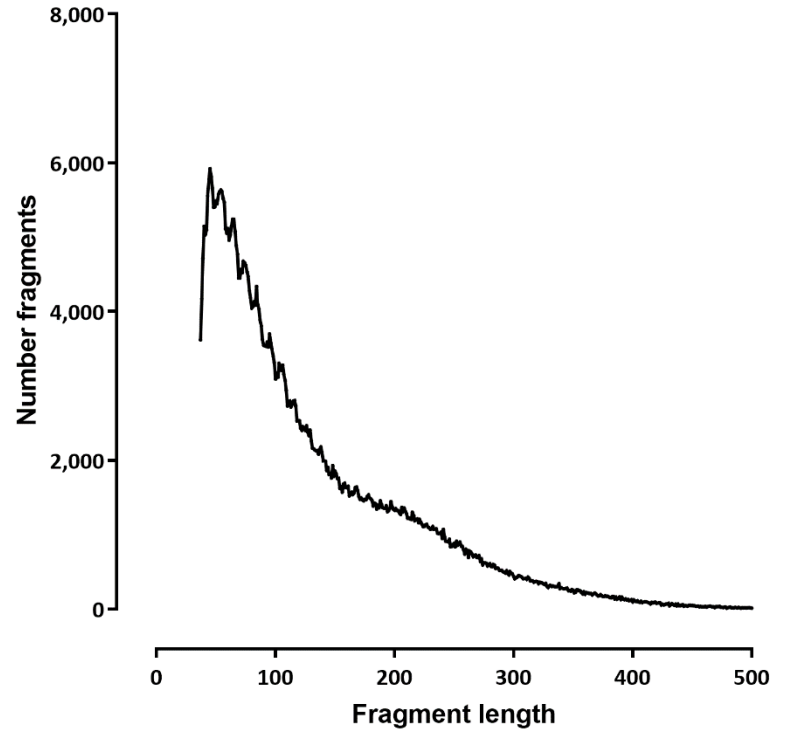
Mapping stats

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	L524-rDNA-146	1,343,984	1,308,468	945,318	0.79			1,225,238	1.49
<i>Saccharomyces cerevisiae</i>	2x36	L525-rDNA-147	2,816,430	2,458,692	2,562,146	0.76			2,235,744	1.47

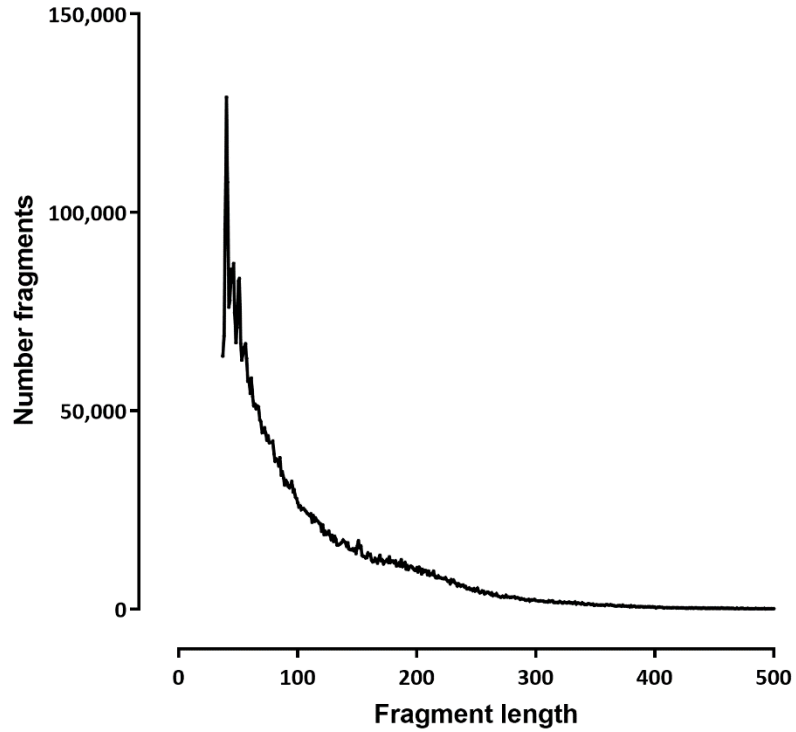
L524-rDNA-146.2x36mers.a



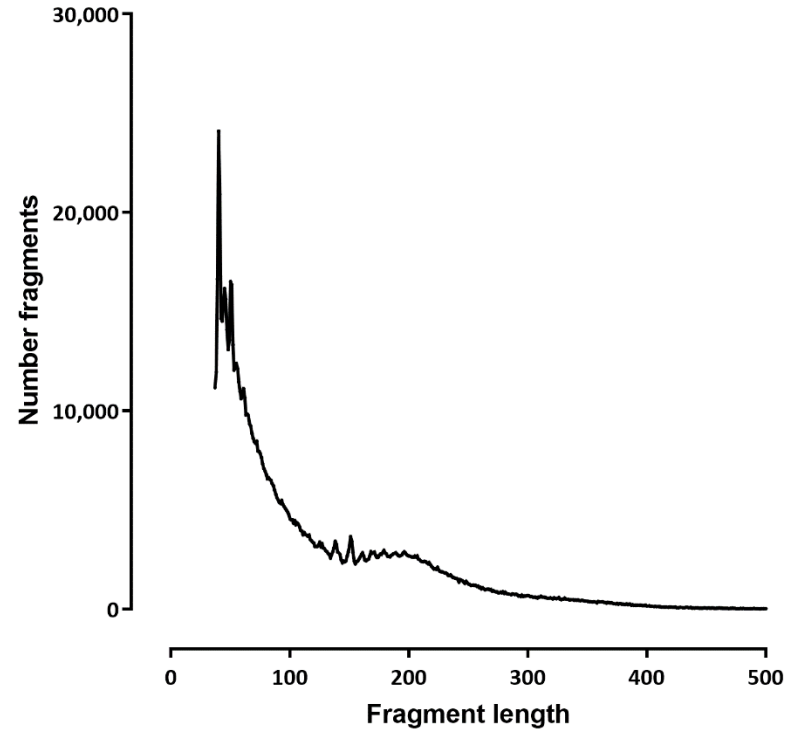
L524-rDNA-146.2x36mers.unique.dedup



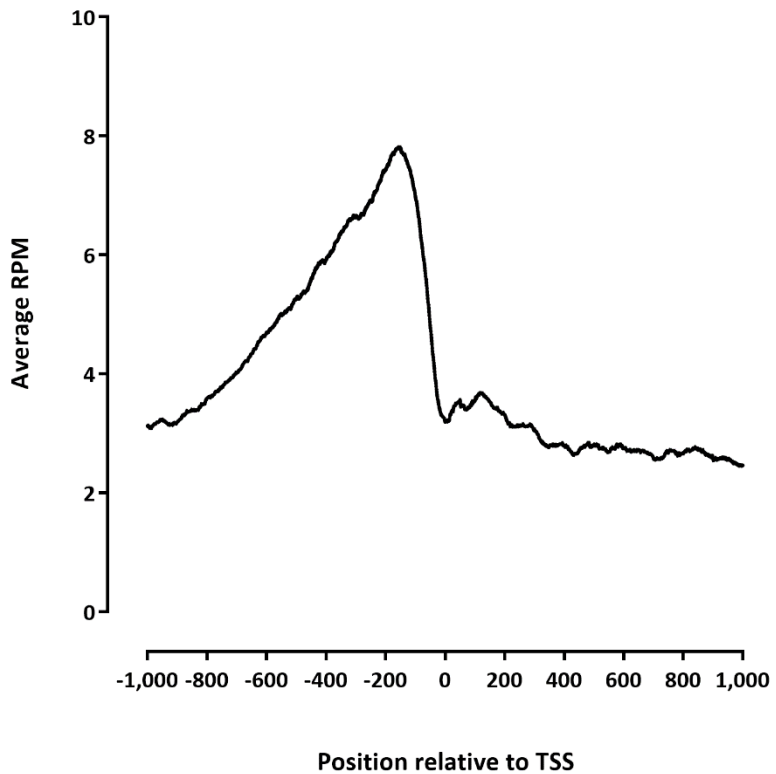
L525-rDNA-147.2x36mers.a



L525-rDNA-147.2x36mers.unique.dedup



L524-rDNA-146



L525-rDNA-147

