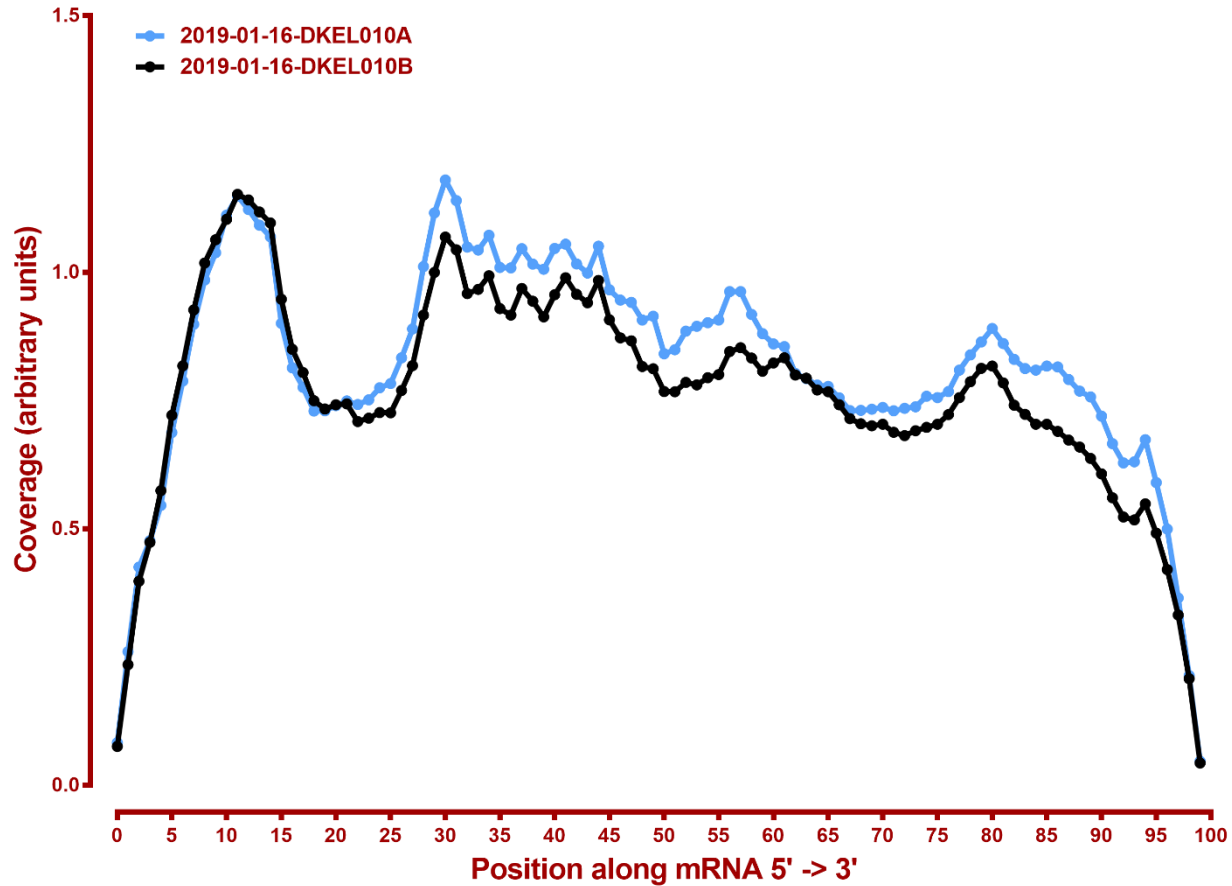


Read mapping stats

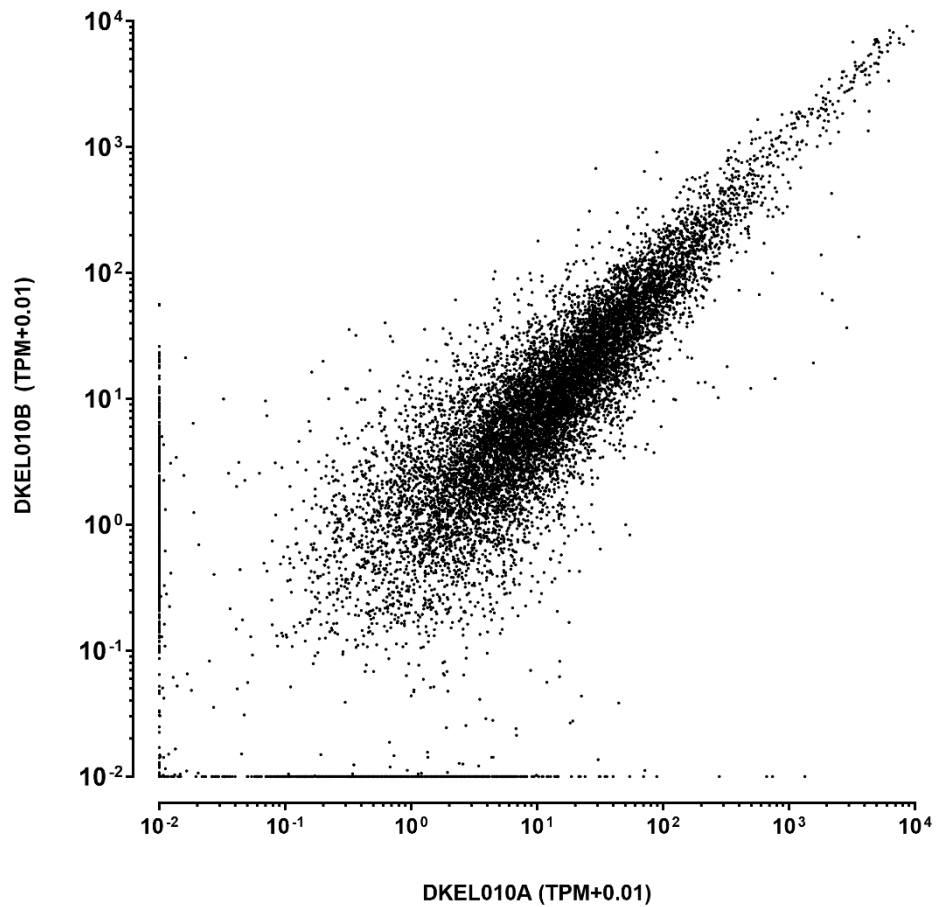
Species	Read Length	Mapping	Library	Raw fragments	Complexity	Unique	Unique Splices	Multi	Multi Splices	Fraction mapped
hg20	2x75	TopHat-2.0.8 GENCODE V25	2019-01-16-DKEL010A	2,795,238	0.32	3,411,147	1,323,926	200,240	30,227	0.89
hg20	2x75	TopHat-2.0.8 GENCODE V25	2019-01-16-DKEL010B	2,412,158	0.20	2,702,359	1,066,474	192,366	23,215	0.83

#	Exonic:	Intergenic:	Intronic:
2019-01-16-DKEL010A	0.96	0.01	0.03
2019-01-16-DKEL010B	0.95	0.02	0.03

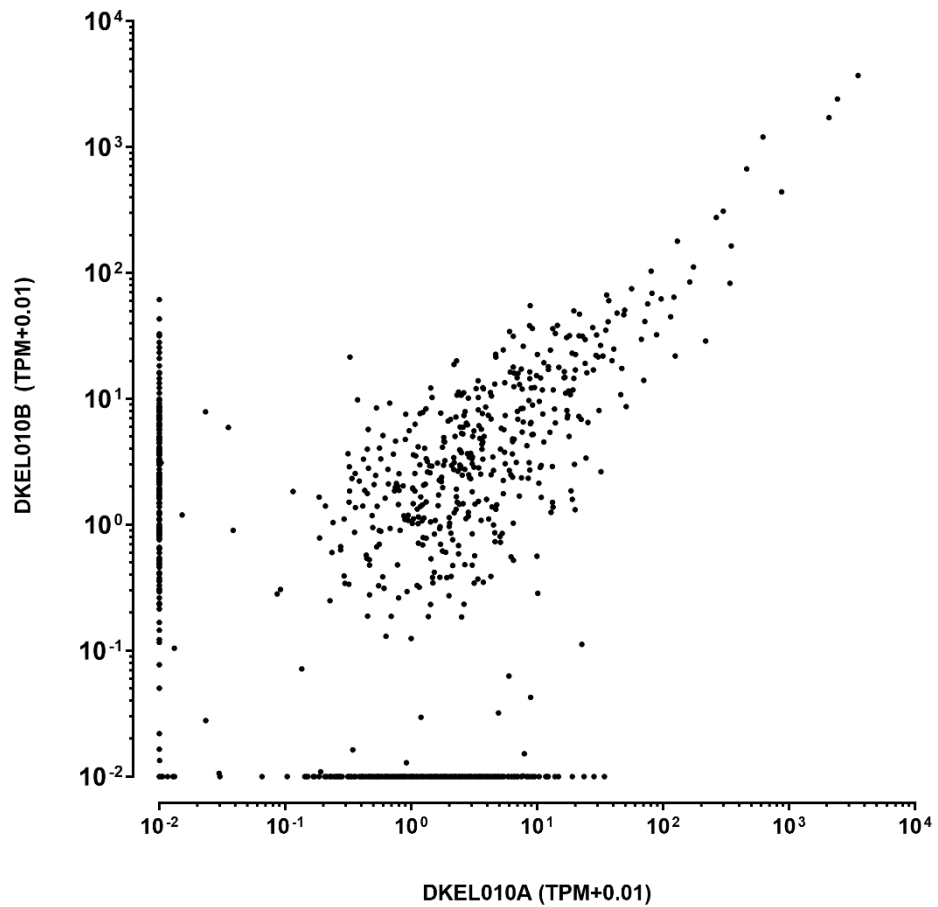
Coverage of genes; >1000bp



protein coding



lincRNAs



humrep

