fraction aligned reads (normalised by # input reads)

Stage	MKL−1 scr DMSO EV			MKL−1 scr Dox EV			MKL−1 sT DMSO EV			MKL−1 sT Dox EV			MKL−1 wt cells			MKL−1 wt EV	
	2608_MKL1_scr_DMSO	2701_MKL1_scr_DMSO	2802_MKL1_scr_DMSO	2608_MKL1_scr_Dox	2701_MKL1_scr_Dox	2802_MKL1_scr_Dox	2608_MKL1_sT_DMSO	2701_MKL1_sT_DMSO	2802_MKL1_sT_DMSO	2608_MKL1_sT_Dox	2701_MKL1_sT_Dox	2802_MKL1_sT_Dox	nf780	nf796	nf797	2404_MKL1_wt_EVs	2608_MKL1_wt_EVs
input	100.0%	100.0%	100.0%	100.0%	100.0%	100.0%	100.0%	100.0%	100.0%	100.0%	100.0%	100.0%	100.0%	100.0%	100.0%	100.0%	100.0%
successfully_clipped	100.0%	100.0%	100.0%	100.0%	100.0%	100.0%	100.0%	100.0%	100.0%	100.0%	100.0%	100.0%	100.0%	100.0%	100.0%	100.0%	100.0%
failed_quality_filter	1.3%	1.3%	1.0%	1.1%	1.3%	1.2%	1.6%	0.9%	1.2%	1.6%	1.6%	1.2%	7.7%	5.5%	5.5%	1.0%	1.1%
failed_homopolymer_filter	0.6%	0.5%	0.3%	0.5%	0.4%	0.4%	0.5%	0.4%	0.5%	0.4%	0.5%	0.4%	2.6%	2.0%	2.0%	0.5%	0.4%
calibrator	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
UniVec_contaminants	0.0%	0.0%	0.0%	0.1%	0.0%	0.0%	0.0%	0.1%	0.0%	0.1%	0.2%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
rRNA	8.0%	6.5%	7.2%	7.0%	6.4%	7.7%	7.1%	7.5%	8.6%	5.5%	9.9%	7.4%	3.4%	11.6%	16.0%	7.9%	7.6%
reads_used_for_alignment	90.1%	91.7%	91.4%	91.3%	91.9%	90.7%	90.8%	91.1%	89.7%	92.5%	87.8%	91.1%	86.3%	81.0%	76.5%	90.5%	90.8%
genome	49.5%	39.0%	64.9%	50.3%	40.9%	60.7%	40.6%	42.9%	58.6%	39.8%	47.8%	60.6%	84.4%	76.4%	71.9%	53.9%	51.4%
miRNA_sense	36.1%	26.6%	49.7%	37.4%	27.6%	47.3%	27.7%	31.2%	43.8%	25.6%	34.7%	47.8%	1.7%	1.3%	0.8%	39.7%	38.9%
miRNA_antisense	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
miRNAprecursor_sense	0.0%	0.1%	0.0%	0.0%	0.0%	0.1%	0.0%	0.1%	0.1%	0.1%	0.1%	0.0%	0.1%	0.0%	0.0%	0.0%	0.1%
miRNAprecursor_antisense	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
tRNA_sense	2.4%	1.6%	2.1%	2.4%	1.8%	1.6%	1.7%	1.6%	2.1%	1.8%	1.7%	1.9%	0.2%	9.7%	8.2%	2.9%	2.4%
tRNA_antisense	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
piRNA_sense	0.6%	0.3%	0.4%	0.3%	0.3%	0.4%	0.4%	0.3%	0.5%	0.2%	0.4%	0.4%	0.0%	0.0%	0.0%	0.6%	0.4%
piRNA_antisense	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
gencode_sense	5.9%	5.5%	9.8%	6.0%	6.0%	8.2%	6.0%	5.5%	8.9%	6.1%	6.5%	7.5%	76.3%	58.2%	51.6%	7.0%	5.8%
gencode_antisense	0.2%	0.2%	0.2%	0.2%	0.3%	0.2%	0.3%	0.2%	0.2%	0.2%	0.2%	0.2%	0.1%	0.2%	0.3%	0.3%	0.2%
circularRNA_sense	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
circularRNA_antisense	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
not_mapped_to_genome_or_libs	40.6%	52.7%	26.5%	41.0%	51.0%	30.0%	50.2%	48.2%	31.1%	52.6%	40.0%	30.5%	1.8%	4.5%	4.6%	36.6%	39.4%
repetitiveElements	1.0%	1.3%	0.4%	1.0%	1.2%	0.6%	1.0%	1.0%	0.5%	1.4%	1.0%	0.5%	0.0%	0.0%	0.0%	0.6%	1.2%
endogenous_gapped	1.1%	1.3%	0.5%	0.9%	1.4%	0.8%	1.1%	1.0%	0.5%	1.3%	1.0%	0.7%	0.0%	0.2%	0.2%	0.8%	1.1%
input_to_exogenous_miRNA	38.6%	50.1%	25.6%	39.1%	48.4%	28.6%	48.1%	46.2%	30.1%	49.9%	38.0%	29.3%	1.8%	4.4%	4.4%	35.2%	37.1%
exogenous_miRNA	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.1%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%	0.0%
input_to_exogenous_rRNA	38.6%	50.1%	25.6%	39.1%	48.4%	28.5%	48.1%	46.1%	30.1%	49.9%	38.0%	29.3%	1.8%	4.4%	4.4%	35.1%	37.1%
exogenous_rRNA	0.7%	1.4%	0.8%	0.6%	0.9%	0.6%	1.2%	0.8%	0.5%	1.1%	0.7%	0.5%	0.0%	0.0%	0.0%	0.7%	0.7%
input_to_exogenous_genomes	37.9%	48.6%	24.8%	38.4%	47.5%	27.9%	46.9%	45.3%	29.5%	48.9%	37.3%	28.8%	1.8%	4.4%	4.4%	34.4%	36.4%
exogenous_genomes	13.1%	16.2%	5.9%	12.9%	16.9%	8.1%	13.9%	15.5%	8.1%	19.7%	13.0%	7.7%	0.1%	0.1%	0.2%	11.0%	12.1%

Sample

ReadFraction
1.00
0.75
0.50
0.25
0.00