

fraction aligned reads (normalised by # input reads)

Stage	MKL-1 wt cells			MKL-1 wt EV	WaGa scr DMSO EV			WaGa scr Dox EV			WaGa sT DMSO EV			WaGa sT Dox EV			WaGa wt cells			WaGa wt EV		
	nf780	nf796	nf797	nf655	nf933	nf938	nf973	nf934	nf939	nf974	nf931	nf936	nf971	nf932	nf937	nf972	nf774	nf961	nf962	nf657	nf930	nf935
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%	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%	100.0%	100.0%	100.0%	100.0%	100.0%
failed_quality_filter	7.7%	5.5%	5.5%	6.8%	5.2%	6.6%	2.8%	13.9%	5.1%	2.4%	7.4%	4.9%	4.5%	6.8%	5.3%	3.0%	8.1%	0.5%	0.5%	7.0%	6.0%	5.0%
failed_homopolymer_filter	2.6%	2.0%	2.0%	0.6%	1.0%	1.0%	0.2%	1.2%	1.2%	0.3%	1.1%	1.5%	0.3%	0.9%	1.1%	0.4%	1.4%	0.1%	0.1%	1.8%	1.8%	0.5%
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%	0.0%	0.0%	0.0%	0.0%	0.0%
UniVec_contaminants	0.0%	0.0%	0.0%	0.0%	0.1%	0.1%	0.4%	0.1%	0.2%	0.2%	0.1%	0.1%	0.3%	0.0%	0.0%	1.1%	0.0%	0.0%	0.0%	0.1%	4.0%	0.1%
rRNA	3.4%	11.6%	16.0%	17.2%	8.3%	9.3%	3.1%	6.4%	8.7%	2.6%	9.3%	9.8%	3.0%	10.9%	9.0%	3.2%	2.7%	2.7%	3.3%	16.3%	8.1%	2.5%
reads_used_for_alignment	86.3%	81.0%	76.5%	75.4%	85.4%	83.1%	93.5%	78.5%	84.9%	94.6%	82.1%	83.8%	91.9%	81.4%	84.6%	92.4%	87.8%	96.7%	96.1%	74.8%	80.1%	92.0%
genome	84.4%	76.4%	71.9%	62.5%	62.6%	65.8%	84.7%	54.4%	66.4%	84.4%	53.9%	60.2%	84.4%	63.9%	68.8%	84.0%	85.5%	95.7%	95.2%	45.9%	36.0%	19.6%
miRNA_sense	1.7%	1.3%	0.8%	30.9%	6.4%	12.7%	72.9%	5.5%	12.5%	67.1%	10.6%	12.2%	74.5%	12.9%	13.7%	73.2%	3.2%	85.7%	83.2%	15.8%	5.7%	3.2%
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%	0.0%	0.0%	0.0%	0.0%	0.0%
miRNAprecursor_sense	0.1%	0.0%	0.0%	0.1%	0.0%	0.0%	0.0%	0.0%	0.1%	0.0%	0.0%	0.0%	0.0%	0.0%	0.1%	0.0%	0.1%	0.1%	0.1%	0.0%	0.0%	0.0%
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%	0.0%	0.0%	0.0%	0.0%	0.0%
tRNA_sense	0.2%	9.7%	8.2%	11.2%	44.5%	39.3%	1.2%	38.5%	35.1%	1.6%	33.9%	28.4%	1.2%	39.7%	38.3%	1.3%	0.4%	2.8%	2.9%	16.2%	20.3%	7.7%
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%	0.0%	0.0%	0.0%	0.0%	0.0%
piRNA_sense	0.0%	0.0%	0.0%	0.7%	0.4%	0.3%	1.0%	0.4%	0.3%	0.9%	0.3%	0.5%	1.5%	0.4%	0.2%	1.2%	0.0%	0.4%	0.4%	0.6%	0.3%	0.1%
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%	0.0%	0.0%	0.0%	0.0%	0.0%
gencode_sense	76.3%	58.2%	51.6%	17.8%	9.5%	10.7%	5.1%	8.5%	14.1%	7.5%	7.0%	15.3%	4.2%	8.8%	13.3%	4.5%	75.4%	5.2%	6.7%	11.3%	6.9%	6.2%
gencode_antisense	0.1%	0.2%	0.3%	0.1%	0.2%	0.2%	0.3%	0.1%	0.3%	0.5%	0.2%	0.3%	0.2%	0.2%	0.2%	0.2%	0.1%	0.2%	0.2%	0.1%	0.2%	0.7%
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%	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%	0.0%	0.0%	0.0%	0.0%	0.0%
not_mapped_to_genome_or_libs	1.8%	4.5%	4.6%	12.9%	22.8%	17.2%	8.8%	24.2%	18.4%	10.2%	28.2%	23.6%	7.5%	17.5%	15.8%	8.4%	2.3%	0.9%	0.9%	28.9%	44.1%	72.4%
repetitiveElements	0.0%	0.0%	0.0%	0.1%	0.2%	0.1%	0.1%	0.2%	0.1%	0.2%	0.2%	0.2%	0.1%	0.1%	0.1%	0.1%	0.0%	0.0%	0.0%	0.2%	0.2%	0.1%
endogenous_gapped	0.0%	0.2%	0.2%	0.2%	0.6%	0.2%	0.6%	0.7%	0.3%	0.7%	0.8%	0.5%	0.4%	0.4%	0.2%	0.5%	0.0%	0.1%	0.1%	0.4%	1.1%	0.2%
input_to_exogenous_miRNA	1.8%	4.4%	4.4%	12.7%	22.1%	16.9%	8.1%	23.3%	18.1%	9.3%	27.3%	22.9%	6.9%	17.0%	15.5%	7.7%	2.3%	0.9%	0.9%	28.3%	42.8%	72.0%
exogenous_miRNA	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%	0.0%	0.0%	0.0%	0.0%	0.0%
input_to_exogenous_rRNA	1.8%	4.4%	4.4%	12.7%	22.1%	16.9%	8.1%	23.3%	18.1%	9.3%	27.3%	22.9%	6.9%	17.0%	15.5%	7.7%	2.3%	0.9%	0.9%	28.3%	42.8%	72.0%
exogenous_rRNA	0.0%	0.0%	0.0%	0.3%	0.7%	0.2%	0.2%	0.6%	0.2%	0.3%	0.7%	0.6%	0.2%	0.4%	0.2%	0.2%	0.0%	0.0%	0.0%	0.8%	1.2%	0.1%
input_to_exogenous_genomes	1.8%	4.4%	4.4%	12.4%	21.4%	16.7%	7.9%	22.7%	17.8%	9.1%	26.6%	22.3%	6.8%	16.6%	15.3%	7.5%	2.3%	0.9%	0.9%	27.5%	41.7%	71.9%
exogenous_genomes	0.1%	0.1%	0.2%	2.8%	5.9%	2.7%	3.0%	7.4%	2.9%	2.6%	7.7%	4.7%	1.5%	3.6%	2.1%	2.0%	0.0%	0.0%	0.0%	4.7%	14.2%	3.2%

