Cards for bioinformatic resources

.footnote[11/18/2019, Huihan]

name: light layout: false class: center, top

Mapping efficiency & total reads

.pure-table.pure-table-bordered.pure-table-striped.smaller-font[

s am ple ID	readMiliion	multiMap%	uniMap%	noMap%
2xFlag_TetTKO_input	37.546244	0.5341571	0.4290482	0.0367948
2xFlag_WT_input	33.497314	0.3641530	0.6016375	0.0342095

]

???

Compare the proportion of multiple hits on repeats in DNAseq of tetTKO and tetWT

Figure2: 2018 March. TetTKO by ZHY (TetKl: 2xFlag)

class: center, middle

For this moment, we couldn't explain the difference.

???

Technical problem?