

Memory Usage Distribution by Rule

Total benchmarks: 126819

ppgb construct
samtools fasta unmapped
minimap2 align batches
meryl build alleles db meryl difference kfilter index
meryl build reference db
gfainject inject
samtools faidx batches
bwamem2 mem samtools sort
pangene prepare
kfilter unmapped
viz odgi
pansnspc target
gafpack coverage
bwamem2 index
pangene viz
odgi dissimilarity
bedtools getfasta
subset gtf
filter nodes
generate batches
samtools fasta mapped
panplexity filter
merge paf per region
pangene graph
odgi utils
make clusters
impg refine
pangene getaa
cosigt genotype
impg index
make bed refined
impg project batches
make reference bed
make alignment bed
combine mapped unmapped

