**BGI：**

**lh /fslgroup/cotton\_seq/compute/ReSeq\_Project/BGI/Map2Texas/\*.recal.bam |wc -l**

**580**

**BYUSC**

**lh /fslgroup/cotton\_seq/compute/ReSeq\_Project/BYUSC/Map2Texas/\*.recal.bam |wc -l**

**19**

**Mutant**

**lh /fslgroup/cotton\_seq/compute/ReSeq\_Project/Mutant/Map2Texas/\*gvcf.gz |wc -l**

**18**

**AD2\_NovoGene**

**lh /fslgroup/cotton\_seq/compute/ReSeq\_Project/AD2\_Novogene/Map2Texas/\*.recal.bam |wc -l**

**15**

**Peru**

**lh /fslgroup/cotton\_seq/compute/ReSeq\_Project/Peru/Map2Texas/\*.recal.bam |wc -l**

**16**

Justin\_PLoS\_Genetics

/fslgroup/fslg\_CG\_W2014/compute/Daojun/JustinPlosGenetics/Map2Tesxas/\*dedup.bam |wc -l

35

DavidFang

lh /fslhome/daojuny/compute/DavidFang/Map2Texas/\*bam |wc -l

16

NAU\_GenomeBiology

lh /fslhome/daojuny/compute/NAU/Map2Texas/\*bam |wc -l

146

NAU\_NatureGenetics

lh /fslgroup/cotton\_seq/compute/ReSeq\_Project/NAU/Map2Texas/\*.bam |wc -l

269

lh /fslgroup/cotton\_seq/compute/ReSeq\_Project/HAU/Map2Texas/\*.bam |wc -l

282