

### R-3.0.2-Packages-BlueCrystal Phase 3

```
lme4
caper
irr
mvtnorm
MCMCglmm
nlme
ggplot2
multicore
ape
ade4
mixtools
boot
plotrix
sna
bipartite
vegan
permute
tnet
survival
igraph
MASS
phangorn
sem
rms
Hmisc
kohonen
TreePar
speedglm
CpGassoc
forecast
SKAT
MetaSKAT
```

### BioConductor Packages

```
source("http://bioconductor.org/biocLite.R")
biocLite()
biocLite("globaltest")
biocLite("annotate")
biocLite("org.Hs.eg.db")

biocLite("affycoretools")
biocLite("watermelon")
```