

R 2.15.1 Packages BlueCrystal Phase 3

```
install.packages("caper", depend = TRUE)
```

```
caper  
irr  
lme4  
mvtnorm  
MCMCglmm  
multicore  
ape  
ade4  
mixtools  
plotrix  
sna  
bipartite  
vegan  
permute  
tnet  
survival  
igraph  
rgl  
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")
```