

Red Abalone Reference Transcriptome - Post-Diginorm Stats Lane 4 & Lane 5

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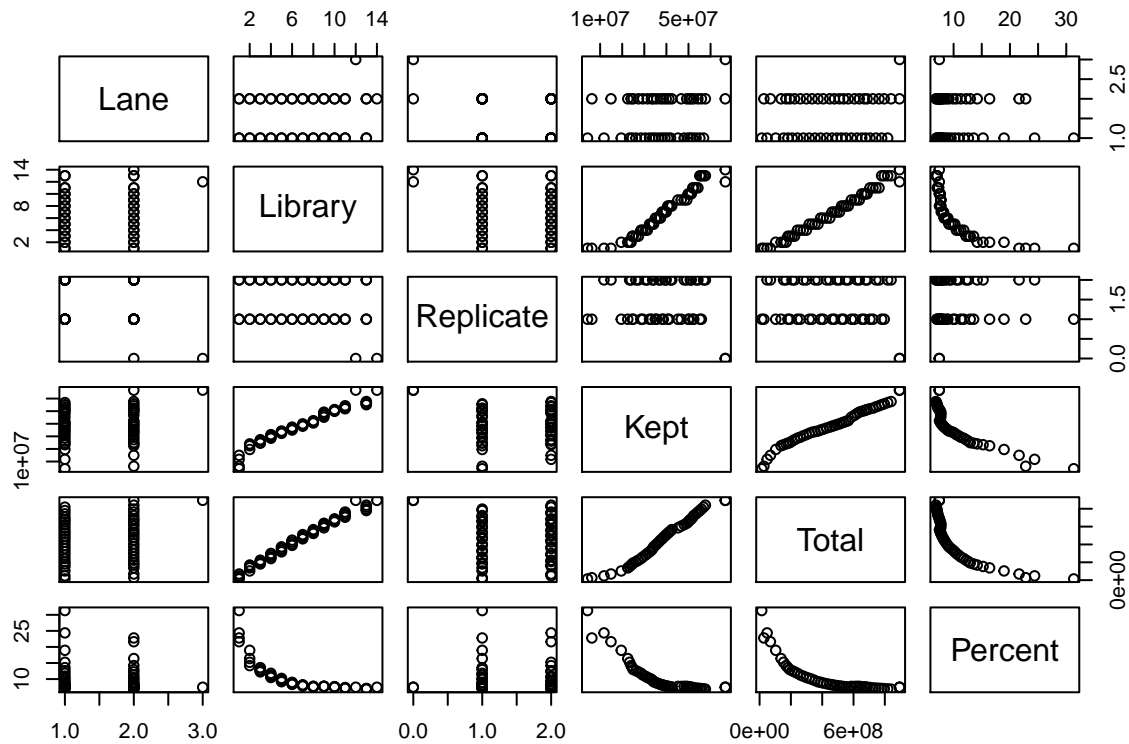
07/09/2017

**This is post-diginorm stats for lane 4 and lane 5 female tissues only.
Here are the summary statics.**

```
library(wesanderson)
library(ggplot2)
library(stats)
library(devtools)
library(jsonlite)
abalone<-read.csv("post-diginormlane4lane5female.csv")
summary(abalone)
```

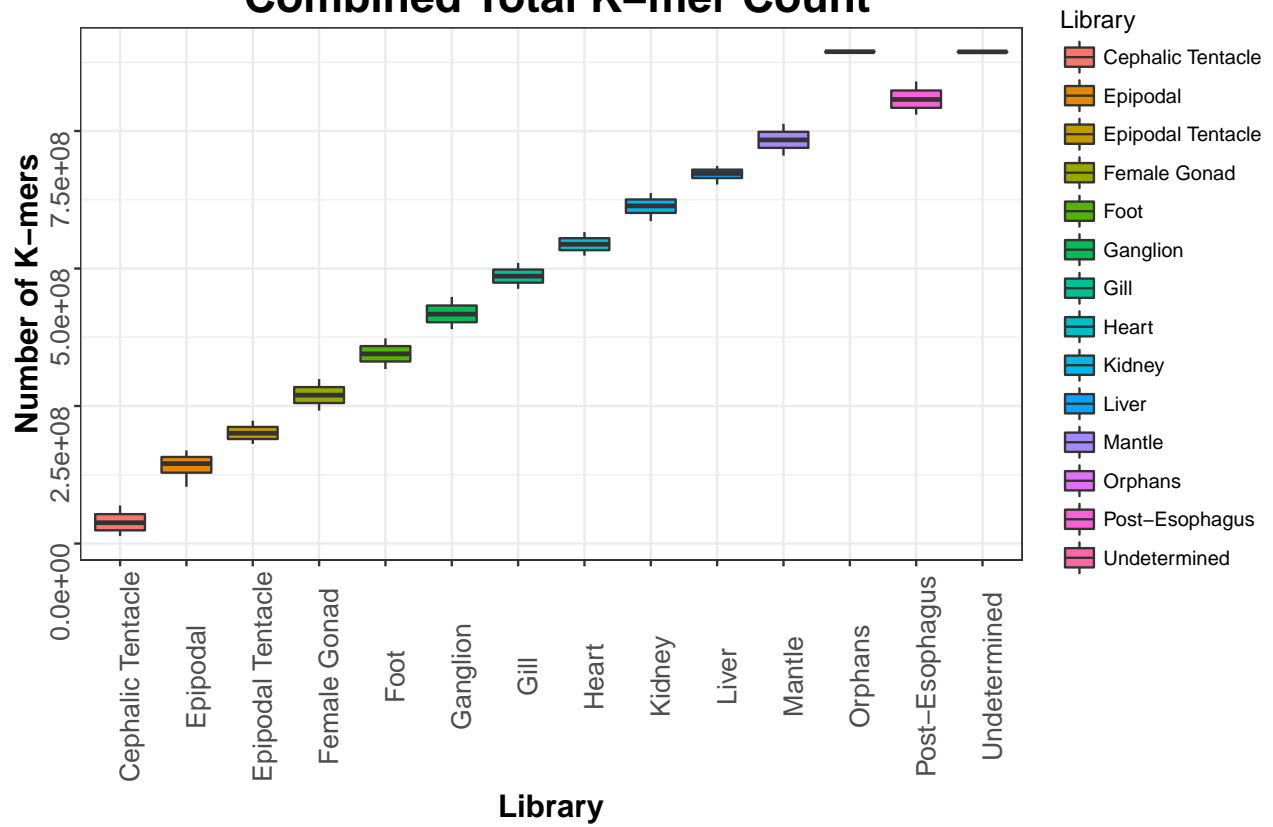
```
##              Lane              Library      Replicate
## Lane 4 - Female Tissues:24 Cephalic Tentacle: 4 Min.      :0.00
## Lane 5- Female Tissues :25 Epipodal           : 4 1st Qu.:1.00
## Orphans                  : 1 Epipodal Tentacle: 4 Median   :1.00
##                           Female Gonad      : 4 Mean     :1.44
##                           Foot              : 4 3rd Qu.:2.00
##                           Ganglion          : 4 Max.      :2.00
##                           (Other)           :26
##      Kept              Total              Percent
## Min.      : 4291354 Min.      : 13696626 Min.      : 6.864
## 1st Qu.:28717910 1st Qu.:246278790 1st Qu.: 7.514
## Median :38331852 Median :470568722 Median : 8.147
## Mean   :38243390 Mean   :457547601 Mean   :10.590
## 3rd Qu.:50238414 3rd Qu.:664455241 3rd Qu.:11.668
## Max.    :66697147 Max.    :894163071 Max.    :31.331
##
```

```
plot(abalone)
```



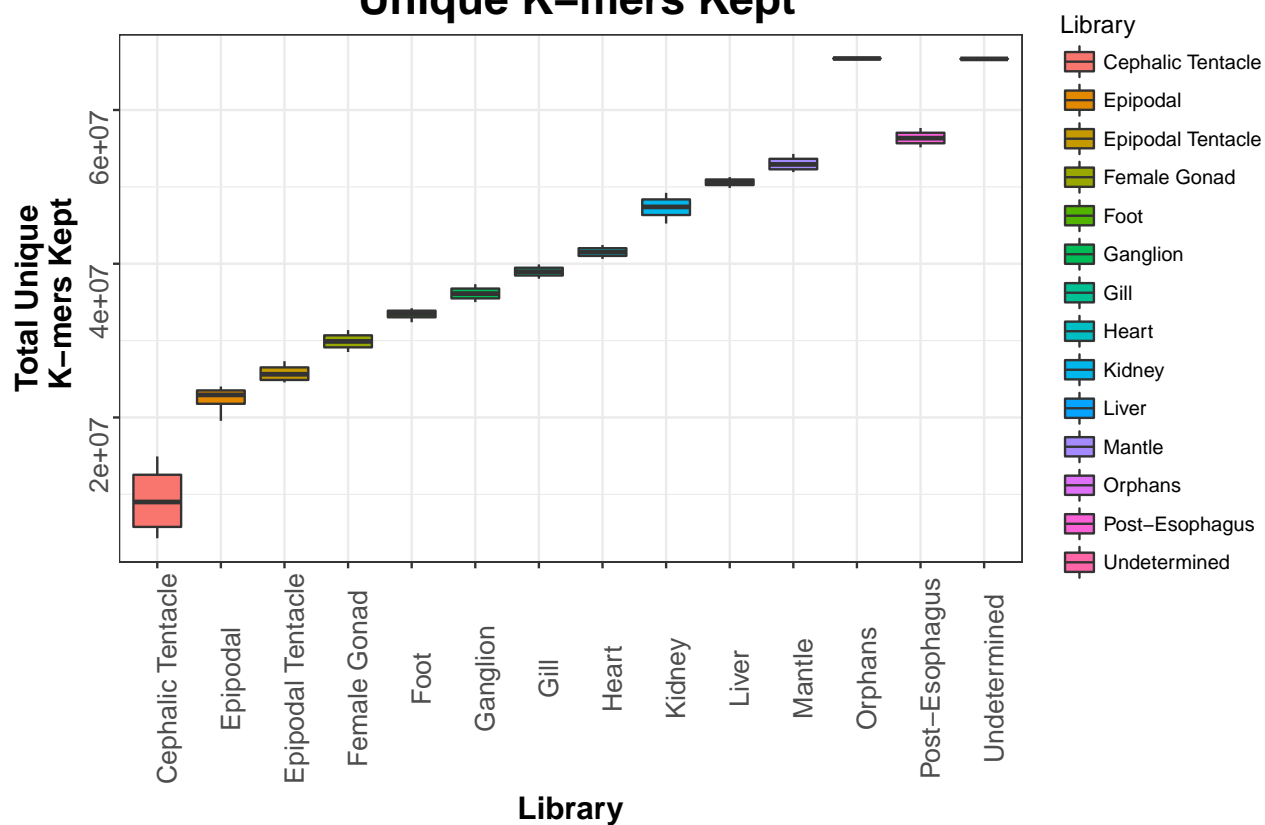
```
p=ggplot(abalone,aes(Library>Total,fill=Library))+geom_boxplot()
p+scale_y_continuous(name="Number of K-mers")+theme_bw()+ggtitle("Post-Diginorm Stats\n Lane 4 & Lane 5")
```

Post-Diginorm Stats Lane 4 & Lane 5 Female Tissues Combined Total K-mer Count

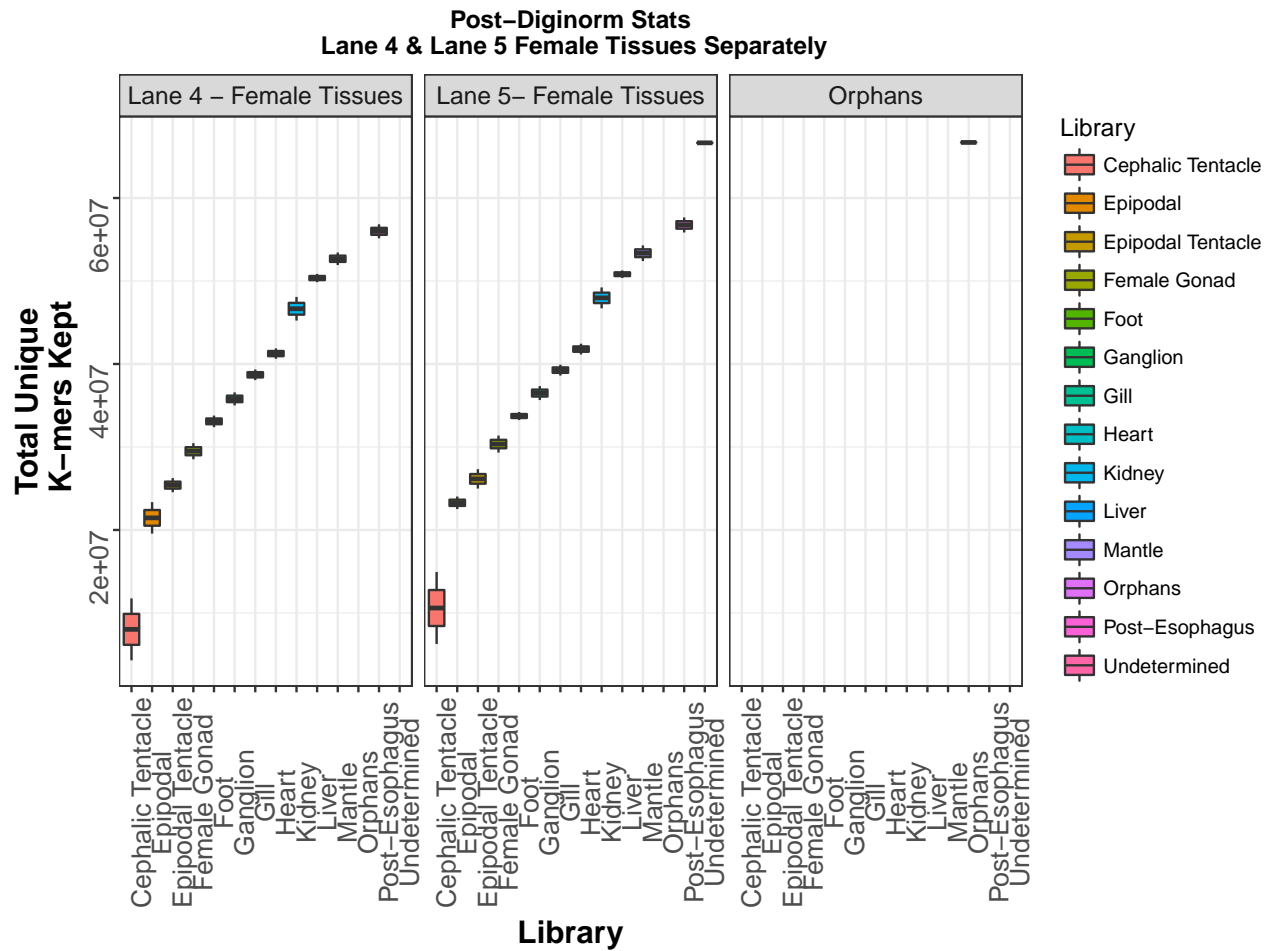


```
p=ggplot(abalone,aes(Library,Kept,fill=Library))+geom_boxplot()
p+scale_y_continuous(name="Total Unique\n K-mers Kept")+theme_bw()+ggtitle("Post-Diginorm Stats\n Lane 4 & Lane 5 Female Tissues\n Combined Total K-mer Count")
```

Post-Diginorm Stats Lane 4 & Lane 5 Female Tissues Combined: Unique K-mers Kept

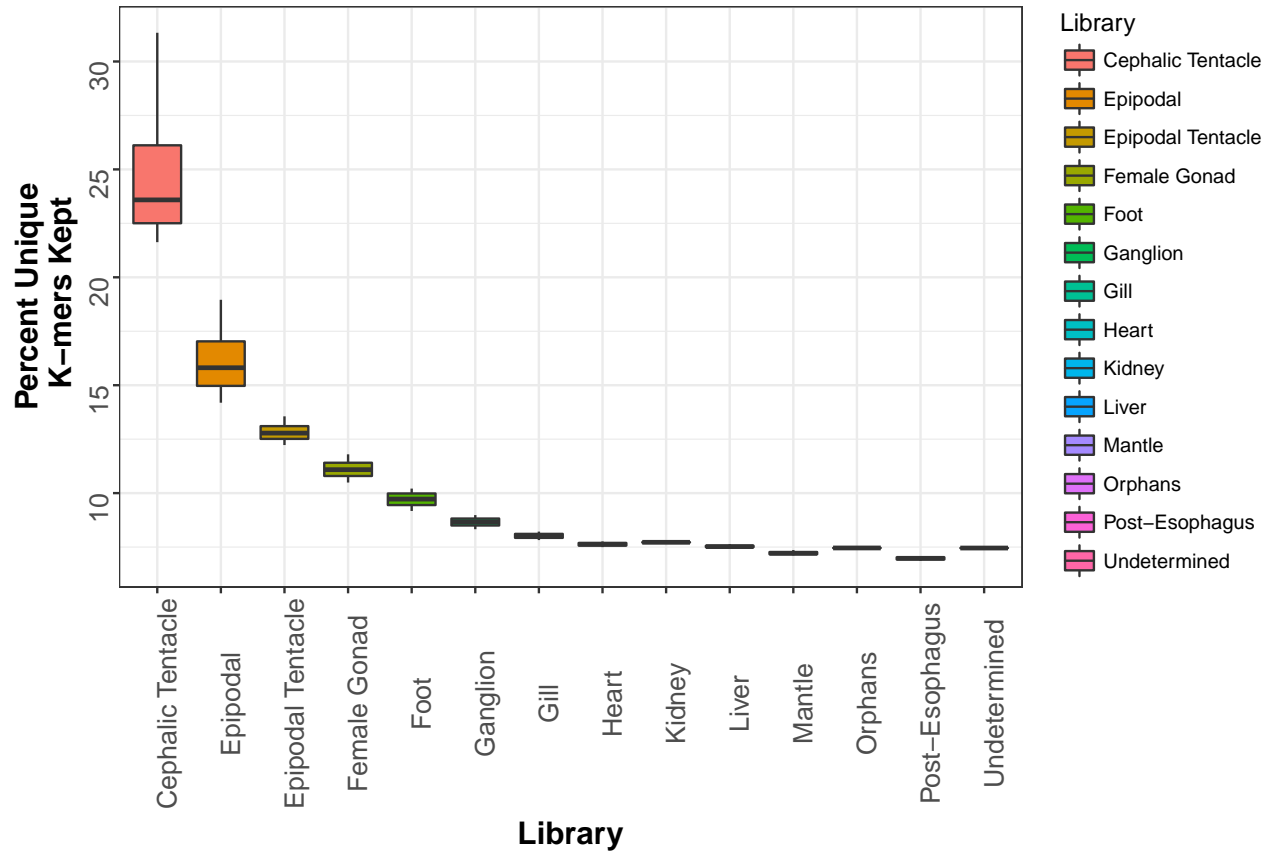


```
p=ggplot(abalone,aes(Library,Kept,fill=Library))+geom_boxplot()
p+scale_y_continuous(name="Total Unique\n K-mers Kept")+theme_bw()+ggtitle("Post-Diginorm Stats\n Lane 4 & Lane 5 Female Tissues Combined: Unique K-mers Kept")
```



```
p=ggplot(abalone,aes(Library,Percent,fill=Library))+geom_boxplot()
p+scale_y_continuous(name="Percent Unique\n K-mers Kept")+theme_bw()+ggtitle("Post-Diginorm Stats\n Lane 4 & Lane 5 Female Tissues Separately")
```

Post-Diginorm Stats Lane 4 & Lane 5 Female Tissues Combined



```
p=ggplot(abalone,aes(Library,Percent,fill=Library))+geom_boxplot()
p+scale_y_continuous(name="Percent Unique\n K-mers Kept")+theme_bw()+ggtitle("Post-Diginorm Stats\n Lane 4 & Lane 5 Female Tissues Combined")
```

Post-Diginorm Stats
Lane 4 & Lane 5 Female Tissues

