ReadCoverage

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```
readCovDat <- read.table("~/biohazard/data/oikopleura/last/embryos/readCovDist_lastsplitOKI2018_I69_1.0_
```
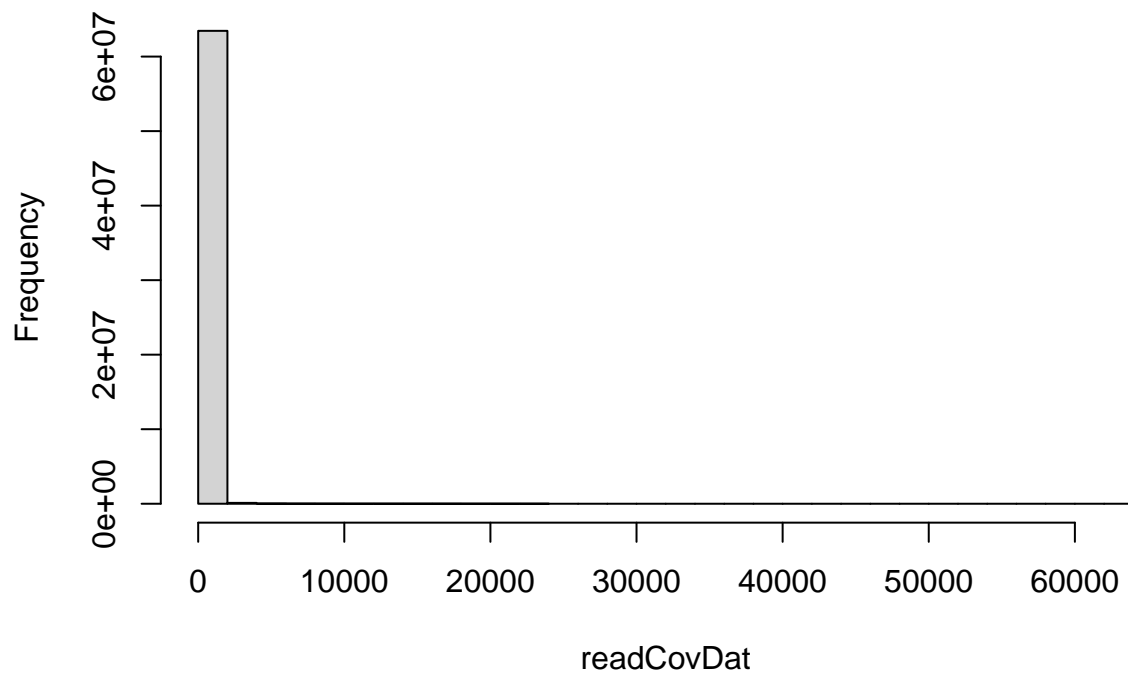
```
mean(readCovDat)
```

```
## [1] 85.33687
```

```
sd(readCovDat)
```

```
## [1] 626.3116
```

```
hist(readCovDat)
```

Histogram of readCovDat


```
hist(readCovDat, ylim=c(0, 10000))
```

Histogram of readCovDat