# Weighted expected binomial model

#### Anthony Aylward

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## Prepare the data.

```
library(chenimbalance)
total_reads <- rowSums(accb[, c("cA", "cC", "cG", "cT")])</pre>
data <- data.frame(</pre>
 total = total_reads,
 allelicRatio = sapply(
   1:nrow(accb),
   function(i) {
      accb[[paste("c", accb[["ref"]][[i]], sep = "")]][[i]] / total_reads[[i]]
 )
)
head(data)
#> total allelicRatio
#> 1 45 1.0000000
#> 2 59 0.5423729
#> 3 114 0.4736842
#> 4 53 0.5094340
#> 5 119 0.5042017
#> 6 21 0.0952381
```

#### Binomial distribution

```
# graded weights for SSE calculation
binSize <- 40
bins <- pretty(0:1, binSize)
w.grad <- graded_weights_for_sse_calculation(r_min = 0, r_max = 1, bins = bins)
# empirical allelic Ratio
minN <- 6
maxN <- min(2500, max(data[["total"]]))
apropor <- length(data[["total"]]][data[["total"]]] <= 2500]) / nrow(data)</pre>
```

```
empirical <- empirical_allelic_ratio(
  data,
  bins,
  maxN = maxN,
  minN = minN,
  plot = TRUE
)</pre>
```

## Histogram of data.match[["allelicRatio"]]

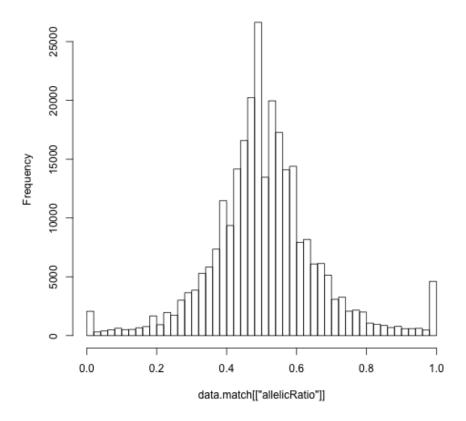


Figure 1: plot of chunk unnamed-chunk-2

Plot the empirical and weighted expected binomial distributions

```
d.combined.sorted.binned <- weighted_expected_binomial(
  data[["total"]],
  minN = minN,
  binSize = binSize
)</pre>
```

```
yuplimit <- 0.15</pre>
barplot(
  empirical,
  ylim = c(0, yuplimit),
  ylab = "density",
  xlab = "allelicRatio",
  names.arg = bins[2:length(bins)] - bins[[2]] / 2,
  main = paste("n=", minN, "-", maxN)
par(new = TRUE)
plot(
  d.combined.sorted.binned,
  ylim = c(0, yuplimit),
  pch=16,
  type='b',
  col='red',
 bty='n',
 ylab='',
 xlab='',
  yaxt='n',
  xaxt='n',
 yaxs="i"
Compute the sum of squared errors for the binomial distribution.
sse = sum((empirical - d.combined.sorted.binned[,2])^2)
sse
#> [1] 0.005153999
```

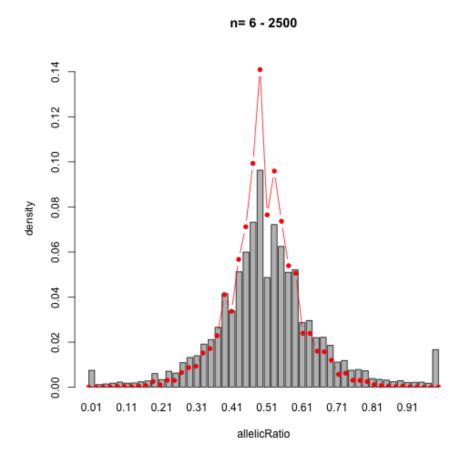


Figure 2: plot of chunk unnamed-chunk-3  $\,$