Transcription vs Metilation

Lucas Michel Todó

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Contents

```
library(ggplot2)
library(reshape2)
```

1 Taules Resum

1.1 Regió 5'

```
## [1] "10G"
##    Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##   11.28   30.53   42.98   40.40   51.08   73.44
## [1] "1.2B"
##    Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##   10.52   29.59   41.77   40.41   48.67   77.90
## [1] "A7"
##    Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##    5.56   21.81   30.12   29.55   38.11   58.04
## [1] "C2"
##    Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##   4.581  12.389  16.740  16.872  21.238  31.232
## [1] "E5"
##    Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##    5.08   17.66   24.89   24.97   33.15   51.76
```

1.2 ORF
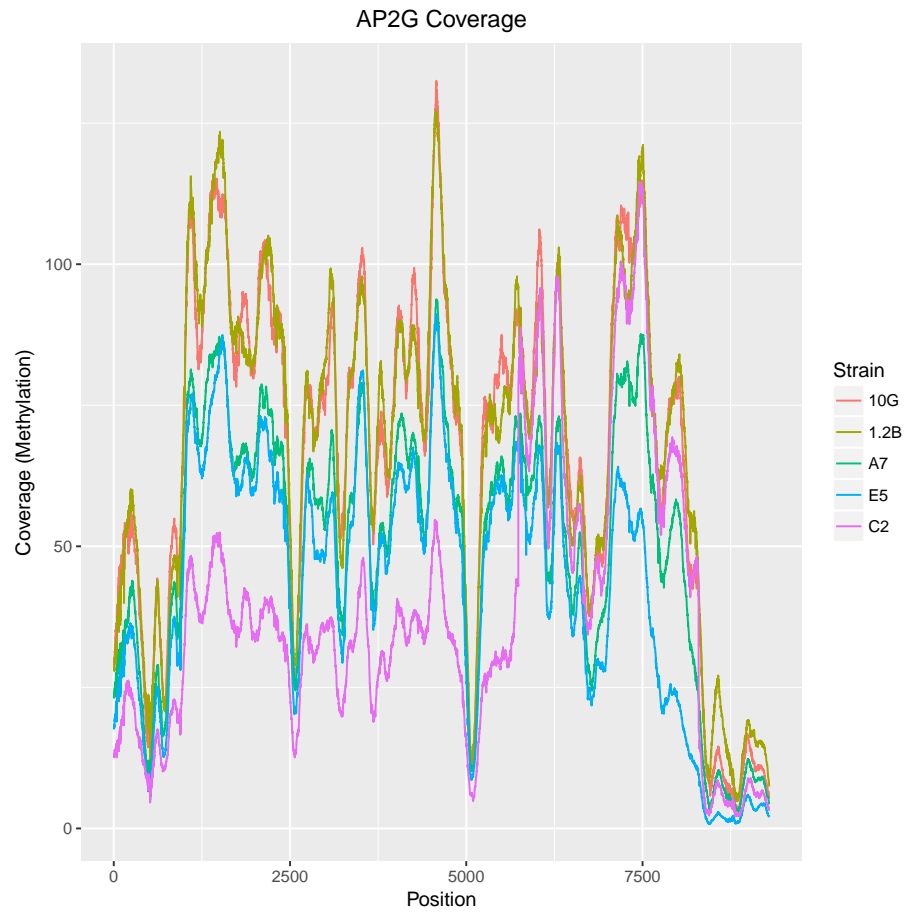
```
##    Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##   10.31   66.11   79.51   78.46   92.62  132.52
##    Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##   11.60   67.29   80.06   78.89   92.90  127.29
##    Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##   10.13   49.79   61.61   59.44   71.20   93.74
##    Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##   4.831  32.648  38.811  46.081  53.802 114.518
##    Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##   8.573  39.413  55.289  51.499  63.227  91.169
```

1.3 Regió 3'

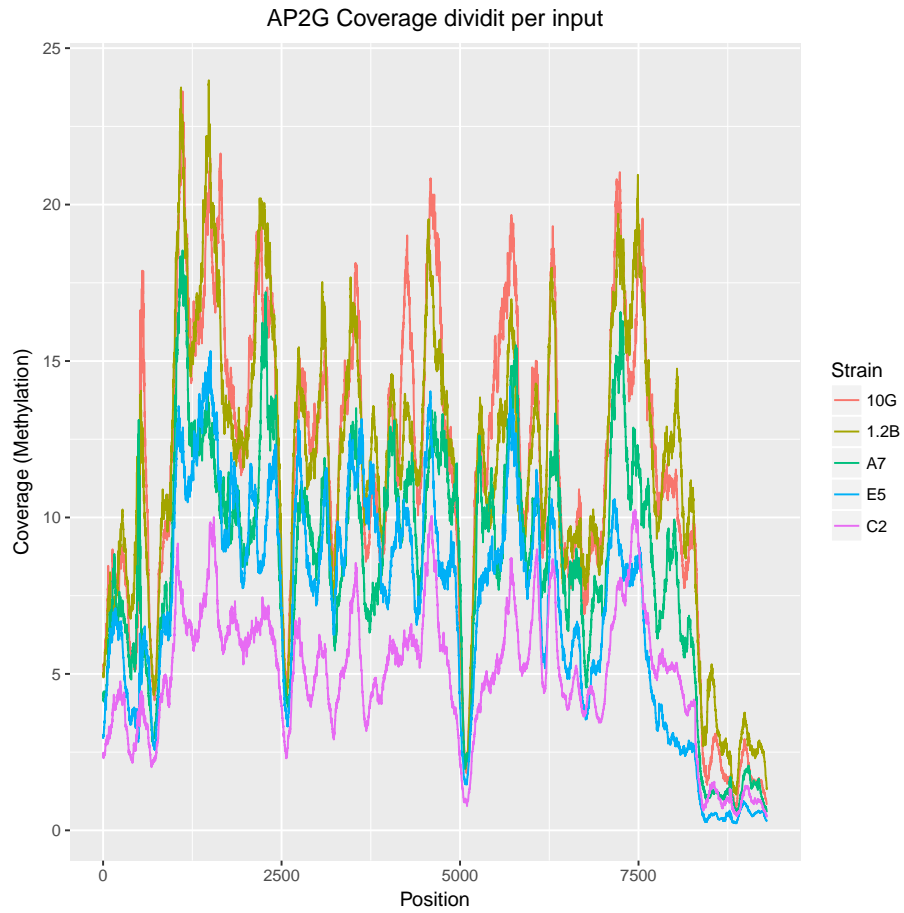```
Min. 1st Qu. Median   Mean 3rd Qu.   Max.
2.891   7.493 10.409 11.215 12.818 40.575
Min. 1st Qu. Median   Mean 3rd Qu.   Max.
4.807 11.271 15.083 15.651 18.253 42.265
Min. 1st Qu. Median   Mean 3rd Qu.   Max.
2.994   5.297   7.814   7.895   9.344 22.439
Min. 1st Qu. Median   Mean 3rd Qu.   Max.
2.165   4.060   5.580   5.707   6.934 17.490
Min. 1st Qu. Median   Mean 3rd Qu.   Max.
0.7938 1.5479 2.4806 2.9586 4.0980 9.4463
```
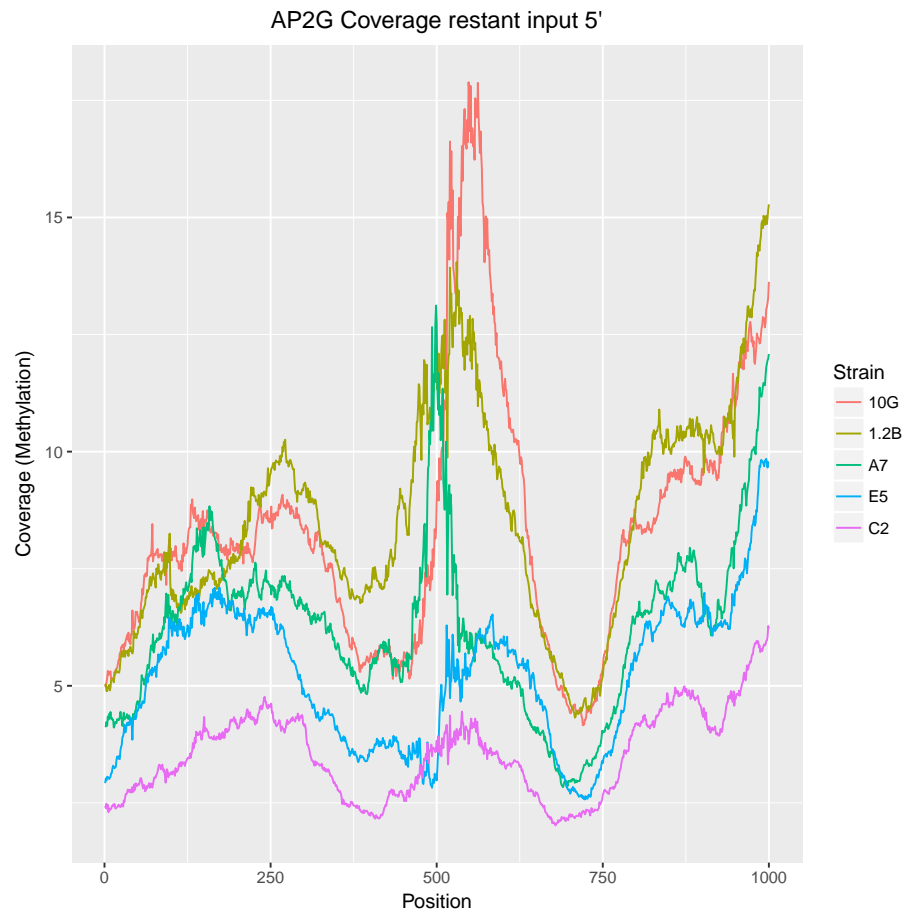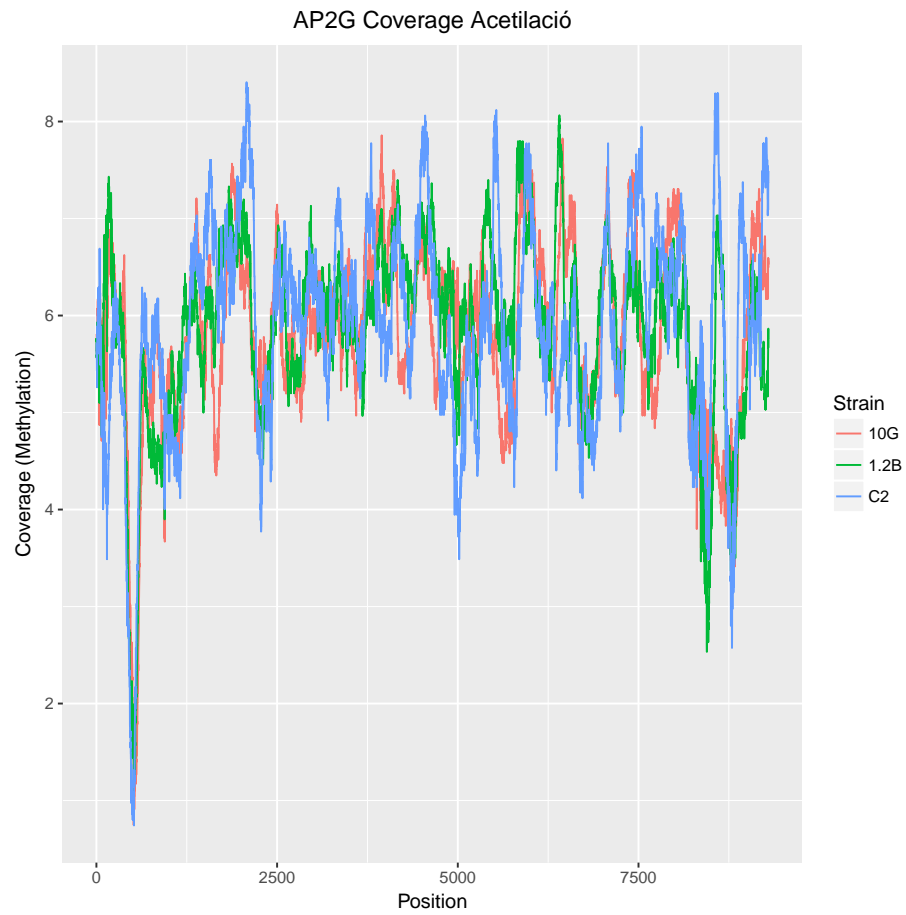
# 2 Gràfics

## 2.1 Coverage
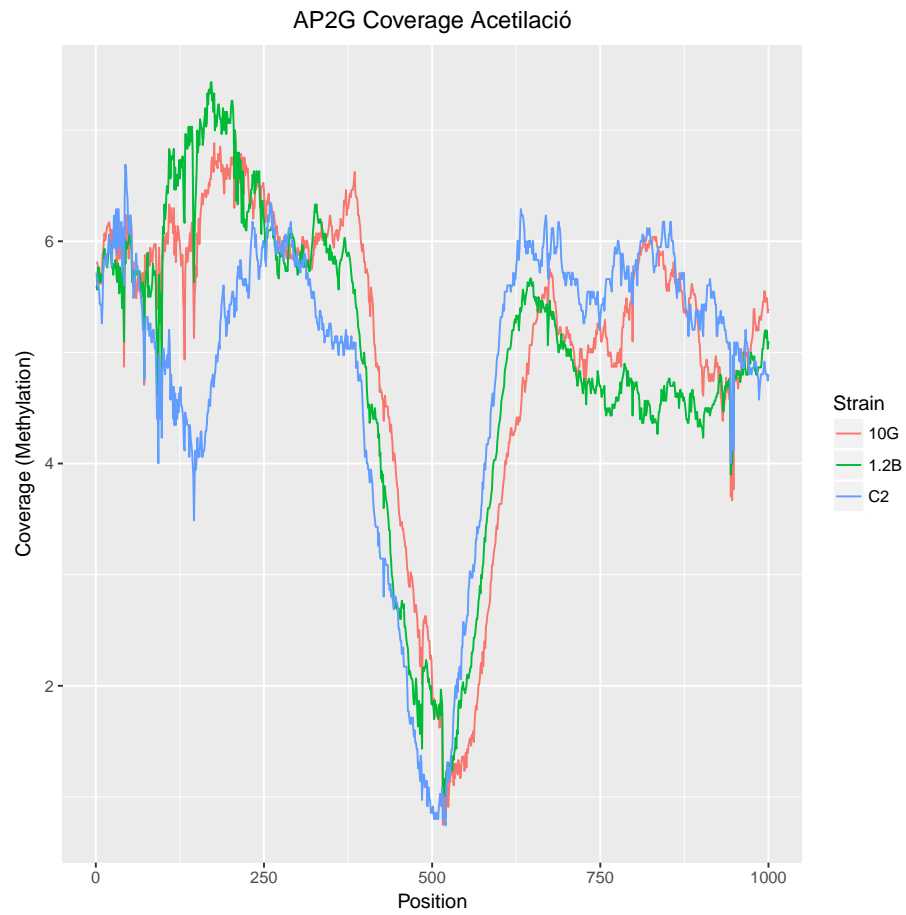


AP2G Coverage

## 2.2 Coverage Normalitzat



AP2G Coverage dividit per input

## 2.3 Coverage Normalitzat a regió 5'



AP2G Coverage restant input 5'

## 2.4   Coverage Acetilació



AP2G Coverage Acetilació

## 2.5 Coverage Acetilació a 5'

AP2G Coverage Acetilació

```
df_ac_met <- df_ac[,-4] / df[,c(1,2,5)]
df_ac_met["pos"] <- df$pos
df_ac_met_m <- melt(df_ac_met, id = "pos")

p <- ggplot(df_ac_met_m, aes(x = pos, y = df_ac_met_m$value, colour = df_ac_met_m$variable))
p + geom_line() +
    labs(title = "AP2G Coverage Acetilació", x = "Position", y = "Coverage (Methylation)",
    theme(plot.title = element_text(hjust = 0.5))
```


AP2G Coverage Acetilació

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