Untitled

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Study the effect of promoter, coding, and cg content on mutation rate. For promoters and coding windows, remove those that have "n/a" as gene names, for coding also remove utrs

```
load("../analysis/161119_mutation_rate_calibration_2.Rdata")
```

258 control

```
control_model_258_summary
```

```
##             Estimate Std. Error    z value      Pr(>|z|)
## (Intercept)  0.3420736  0.0987237   3.464959 5.303119e-04
## coding       0.1259754  0.1005675   1.252645 2.103348e-01
## promoter    -0.1951714  0.1306264  -1.494119 1.351445e-01
## cg          -2.2436740  0.2180893 -10.287868 7.992662e-25
```

manuscript cases

```
ASD_model_manuscript_summary
```

```
##             Estimate Std. Error    z value     Pr(>|z|)
## (Intercept)  0.05831605 0.07344410  0.7940196 0.4271840302
## coding       0.20751452 0.06544651  3.1707501 0.0015204588
## promoter    -0.21024447 0.08799983 -2.3891464 0.0168875717
## cg          -0.57395305 0.15224742 -3.7698705 0.0001633323
```

manusript control

```
control_model_693_summary
```

```
##             Estimate Std. Error    z value     Pr(>|z|)
## (Intercept) -0.281413299 0.05941043 -4.73676560 2.171561e-06
## coding       0.357831797 0.05098383  7.01853552 2.242058e-12
## promoter    -0.004112942 0.06663336 -0.06172496 9.507819e-01
## cg          -0.719811819 0.12361210 -5.82314996 5.774869e-09
```

scherer cases, partly overlapped with manuscript cases

ASD_model_Scherer_summary

```
##               Estimate Std. Error   z value      Pr(>|z|)
## (Intercept) -0.1152721 0.10297037 -1.119468 2.629403e-01
## coding       0.3855092 0.08650837  4.456323 8.337741e-06
## promoter    -0.3116285 0.13118010 -2.375577 1.752152e-02
## cg          -0.7237196 0.21452135 -3.373648 7.417903e-04
```