

**Title of talk**

**Authors**

Institution

Email

## Title of slide 1

Ordinary text, with maths inline:  $\beta_1 < 0$  or displayed:

$$\hat{p}_{\text{HM}}(y|u, X) = \left[ \frac{1}{N} \sum_{t=1}^N p(y|X, u, \beta^{(t)}, z^{(t)})^{-1} \right]^{-1}$$

And then some more text.

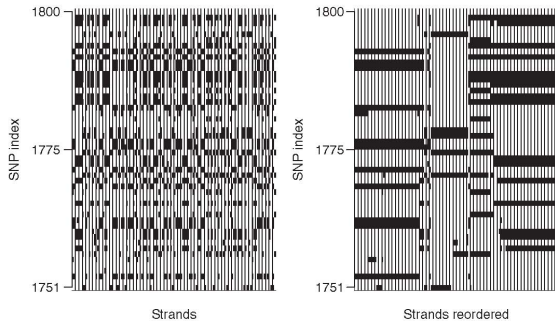
## Title of slide 2

Our complete model combines the strand and disease models, linked though the unobserved alleles  $\{z_{ij}\}$ .

## Heading in middle of slide

We observe the SNP data  $X$  and disease indicators  $y$ , and wish to make inference primarily about the location  $u$  of the causal gene, and also the regression parameters  $\beta$ .

## Slide 3



Text below picture.

# Slide 4

Text to left of picture

