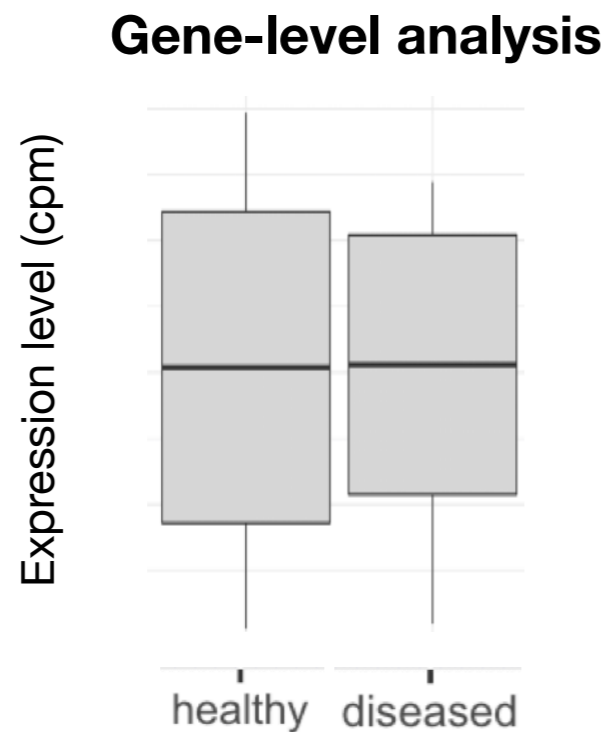
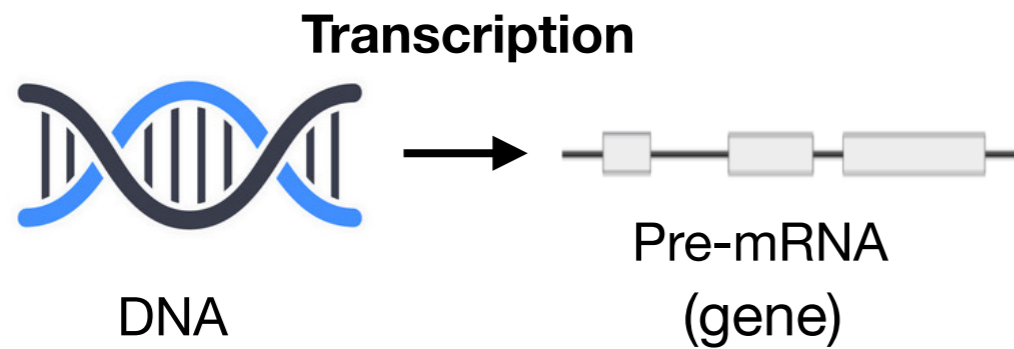
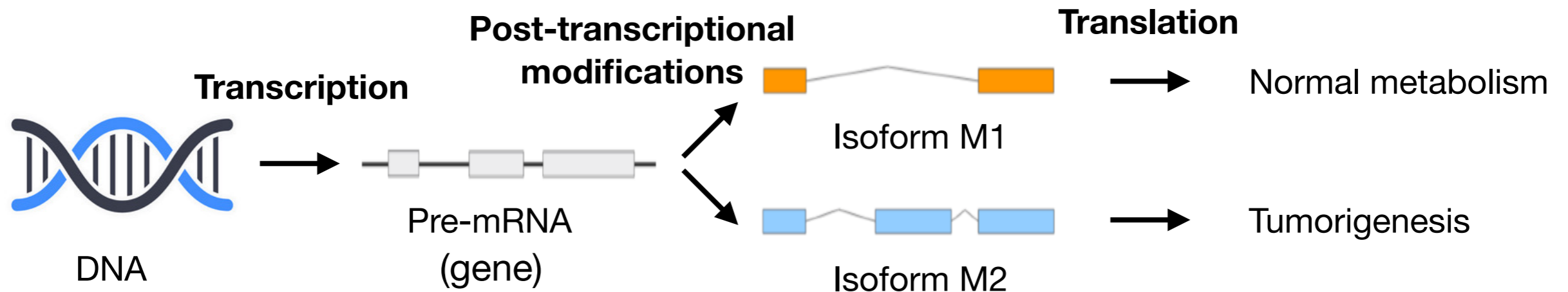


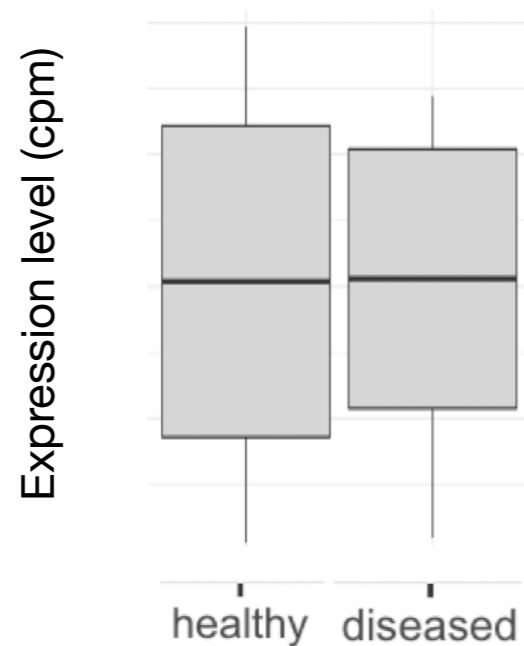
# Differential Transcript Usage (DTU)



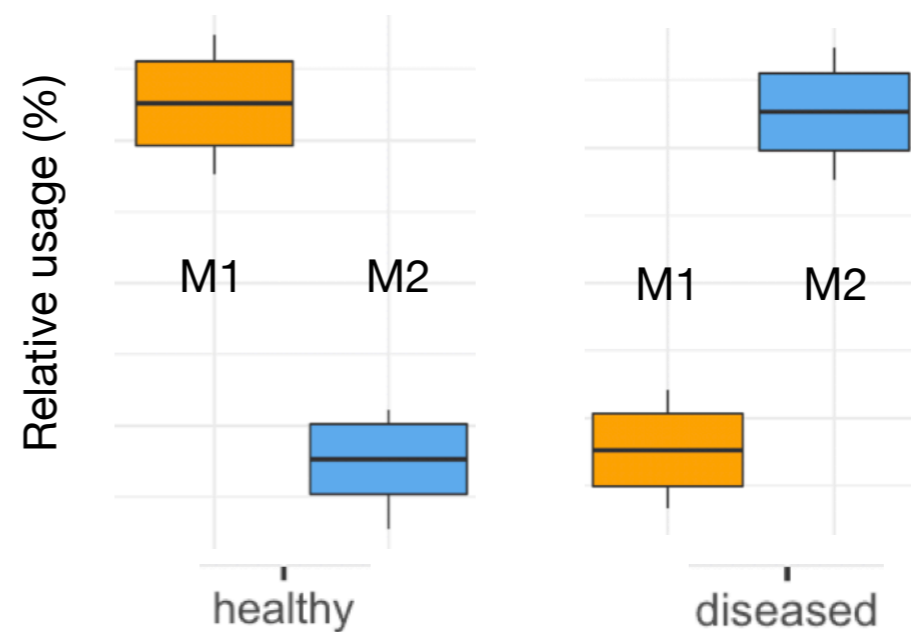
# Differential Transcript Usage (DTU)



### Gene-level analysis



### Transcript-level DTU analysis





# Software development

- Denote the expression of transcript  $t$  of gene  $g$  in sample  $i$  as  $Y_{gti}$
- Denote the usage of transcript  $t$  of gene  $g$  in sample  $i$  as:

$$U_{gti} = \frac{Y_{gti}}{Y_{g.i}}$$

- Describe the **quasi-binomial** GLM:

$$\left\{ \begin{array}{l} E[U_{gti} | \mathbf{X}_i, Y_{g.i}] = \pi_{gti} \\ \log\left(\frac{\pi_{gti}}{1 - \pi_{gti}}\right) = \eta_{gti} \\ \eta_{gti} = \mathbf{X}_i^T \boldsymbol{\beta}_{gt} \end{array} \right.$$

- With variance:

$$\text{Var}[U_{gti} | \mathbf{X}_i, Y_{g.i}] = \frac{\pi_{gti} * (1 - \pi_{gti})}{Y_{g.i}} * \phi_{gt}$$