

## **Methods for Computational Gene Prediction by William H. Majoros (2007)**

### **Errata:**

Page 55 (original print edition)

The text currently reads:

"If we set our rejection criterion for  $PD(x)$  at  $\alpha=5\%$ , then we can expect that of the null hypotheses which we reject, roughly 5% were in fact true and should not have been rejected, thereby constituting what are known as Type I errors."

This should read:

"If we set our rejection region for  $PD(x)$  at  $\alpha=5\%$ , then we can expect that of the true null hypotheses that we test, roughly 5% will be rejected, which would constitute Type I errors."