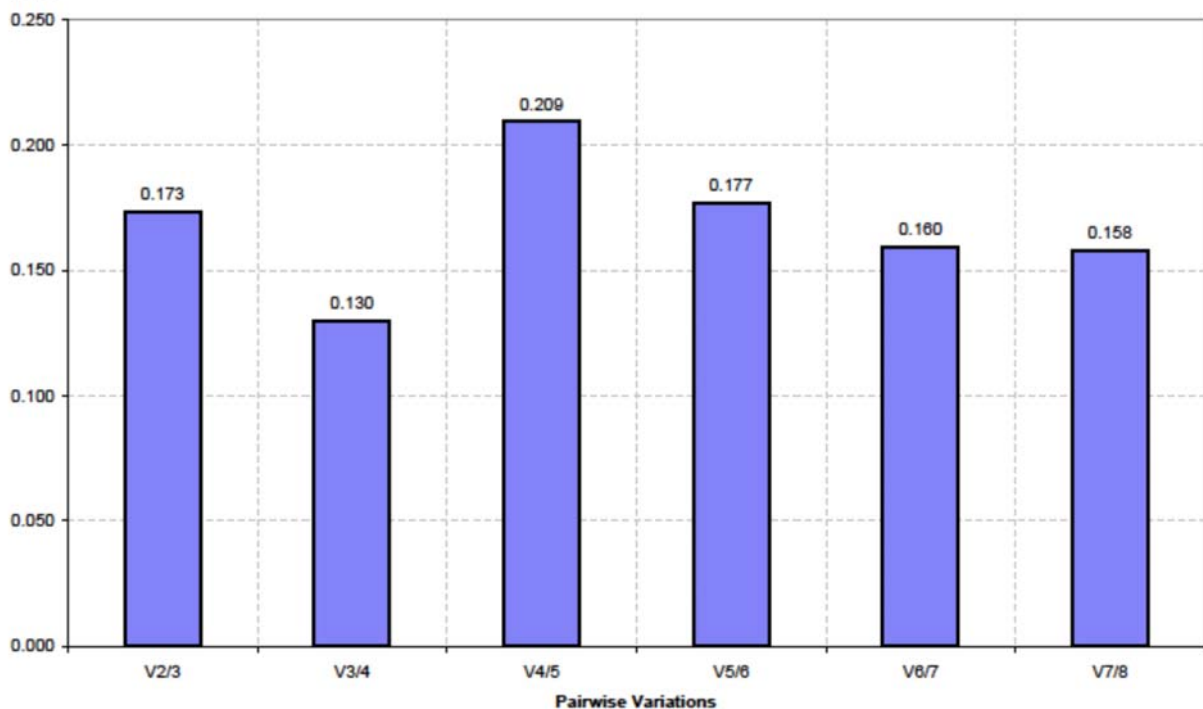


### Determination of the optimal number of control genes for normalization

**A**



**B**

### Average expression stability values of remaining control genes

