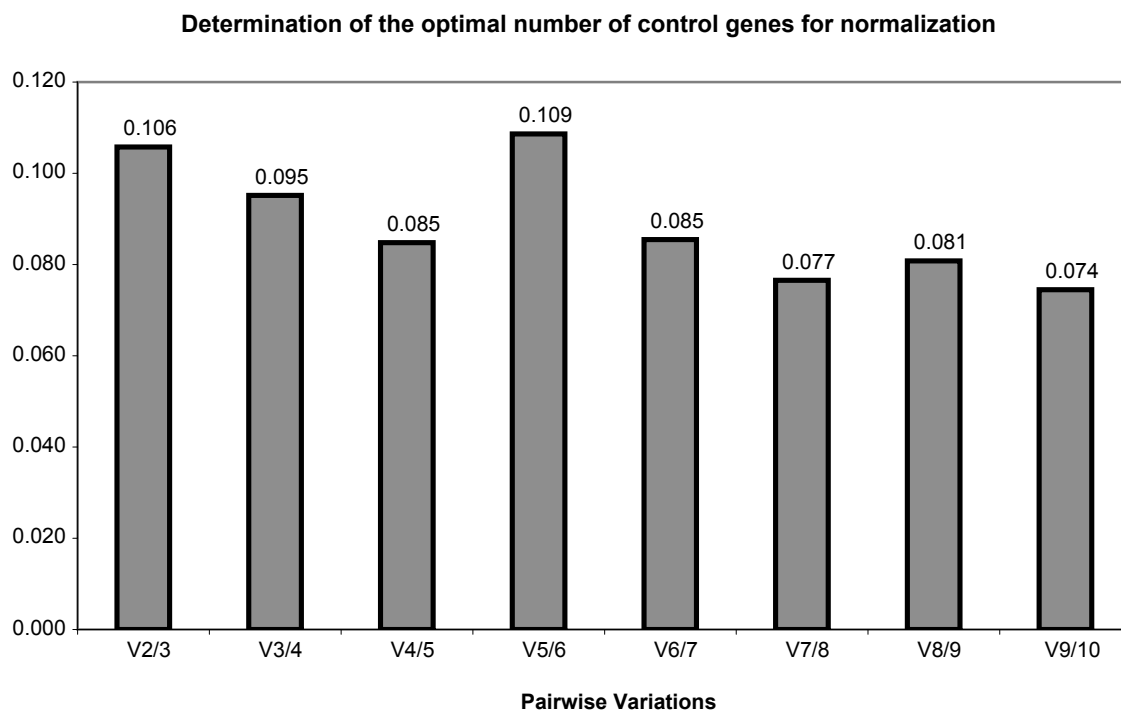
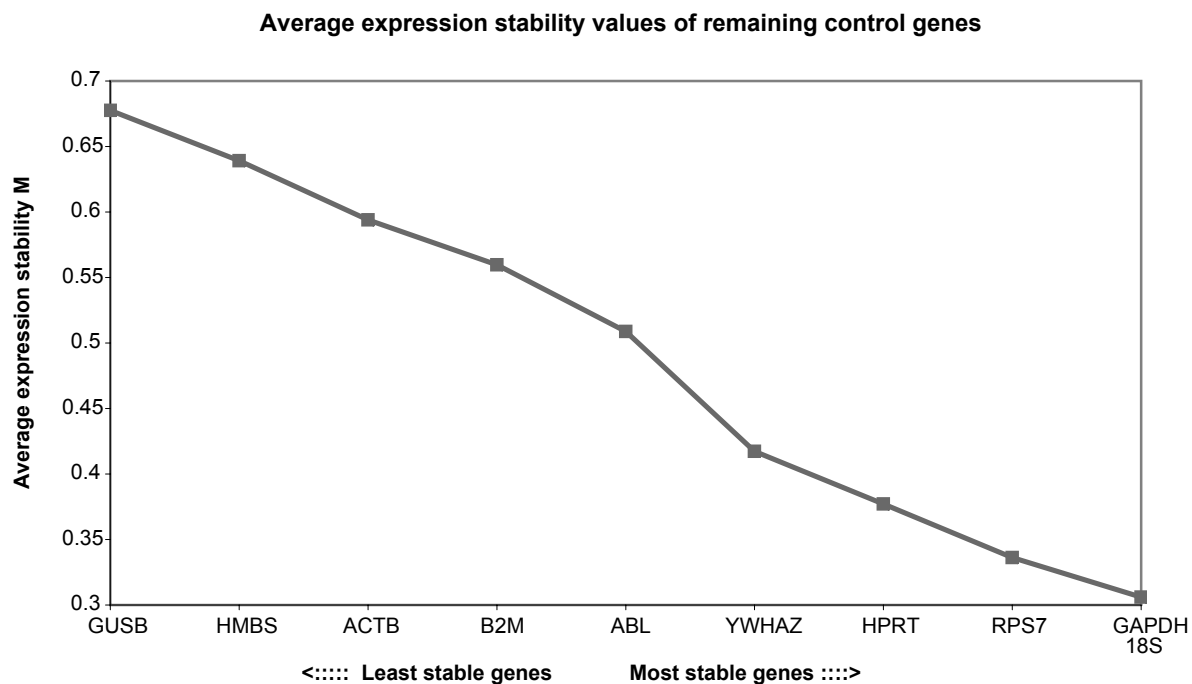
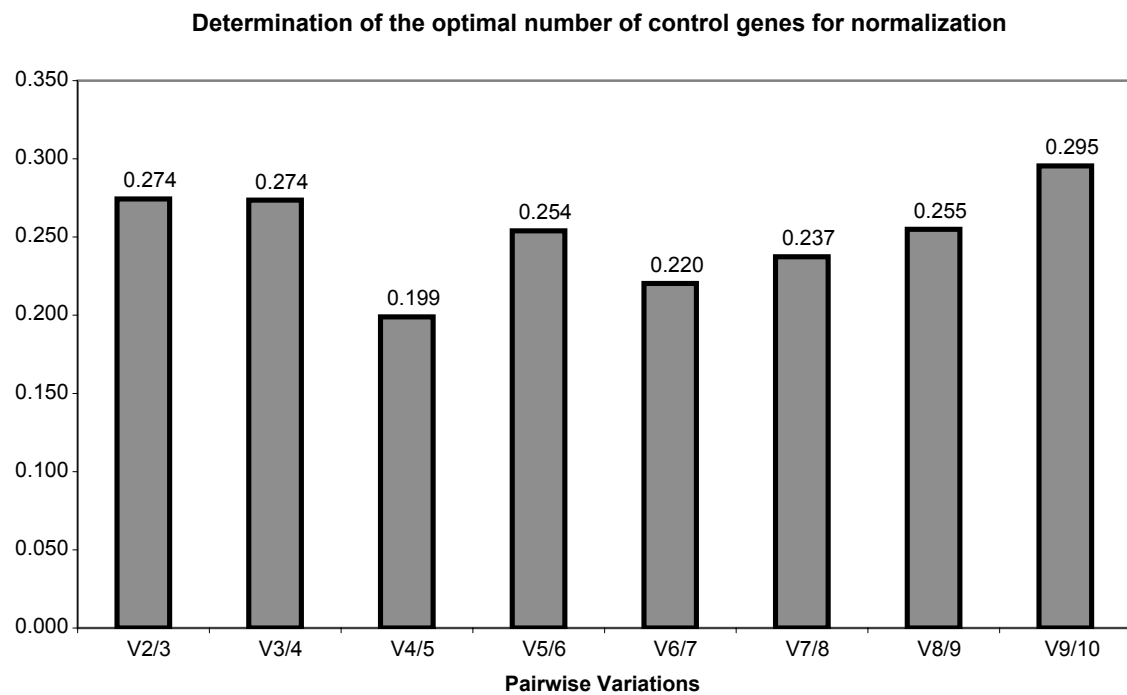
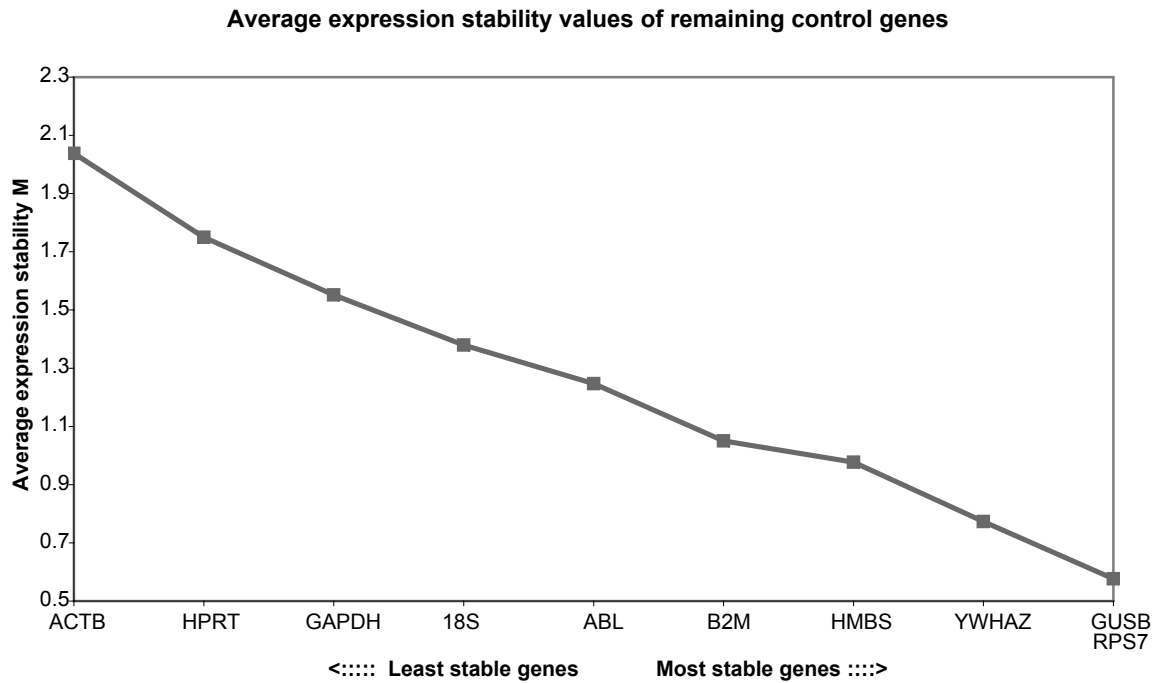


**Additional File 2**  
**a) geNorm output: Adrenal gland**

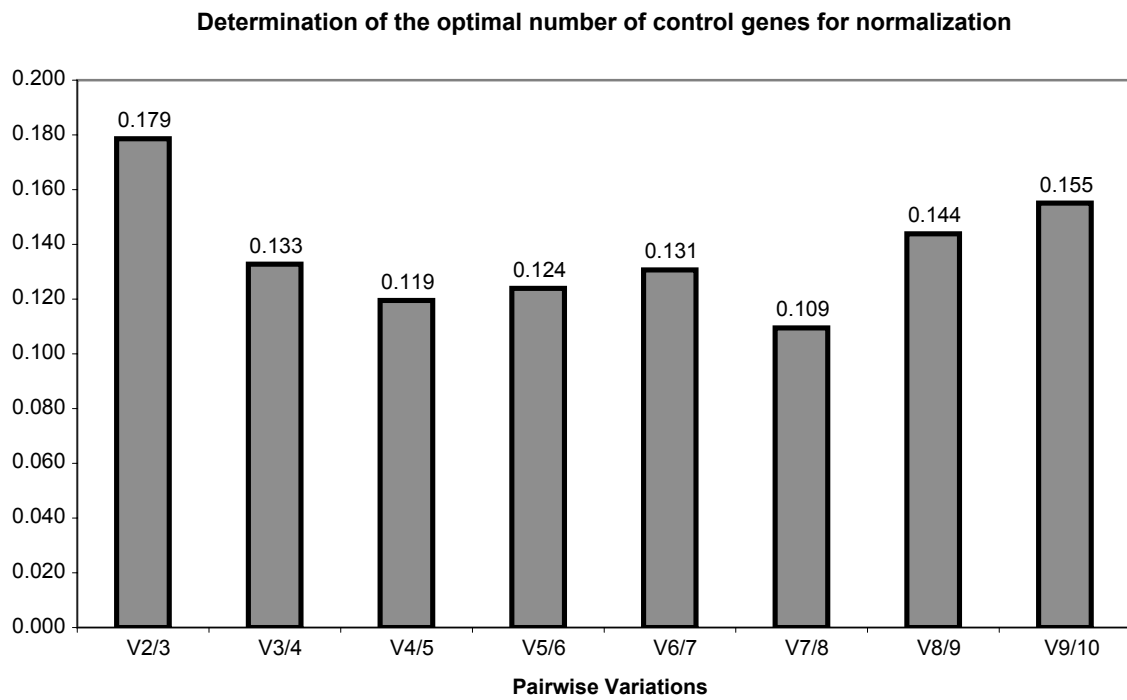
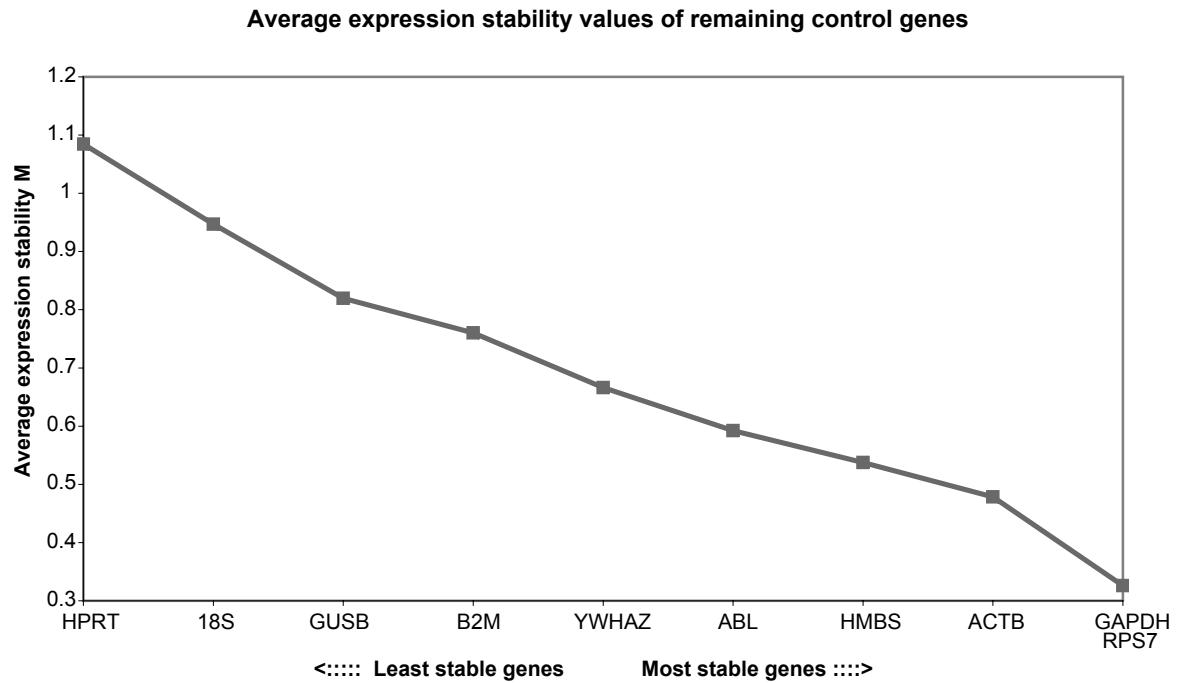


**Additional File 2**  
**b) geNorm output: Pancreas**

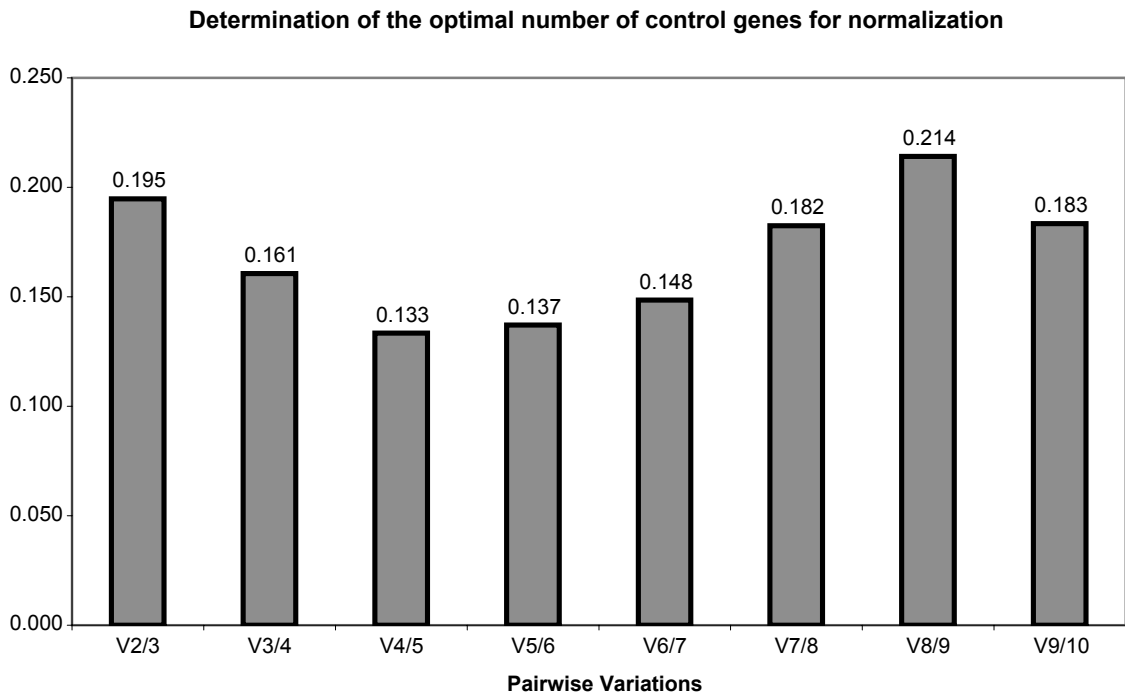
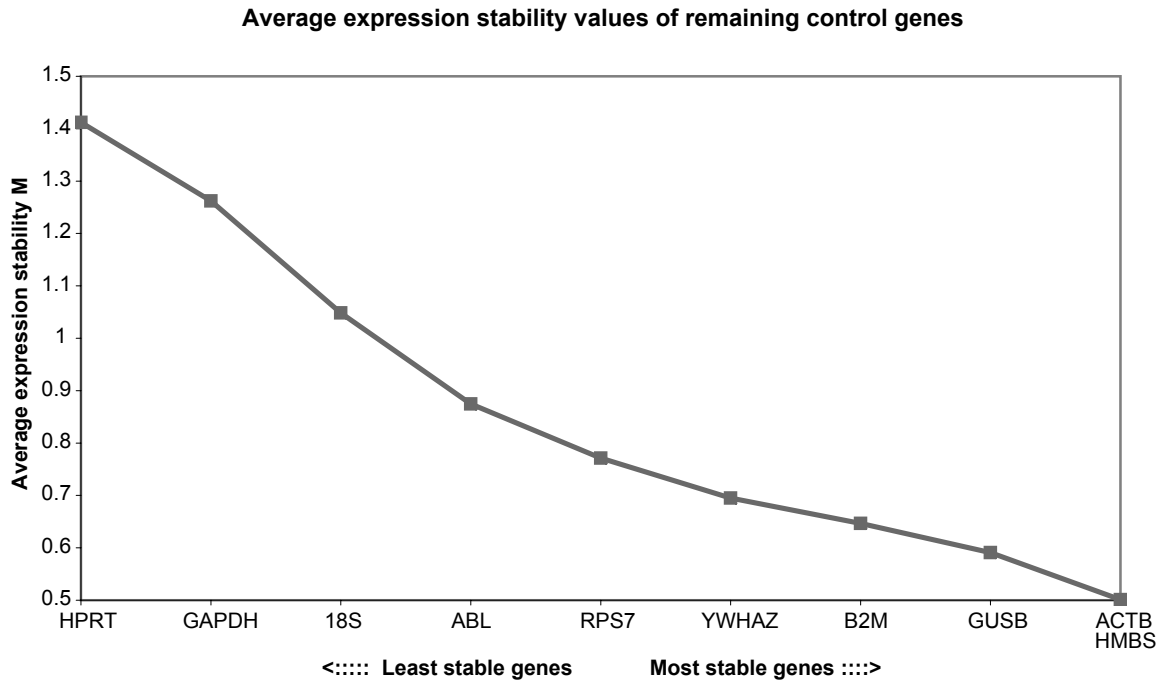


**Additional File 2**

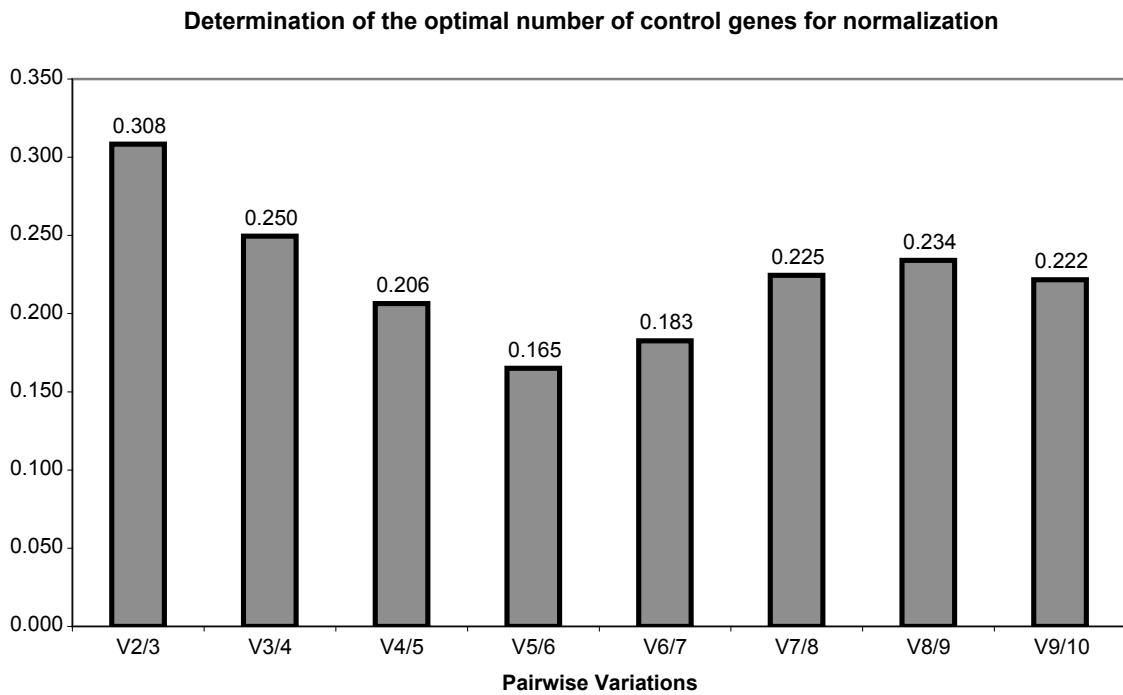
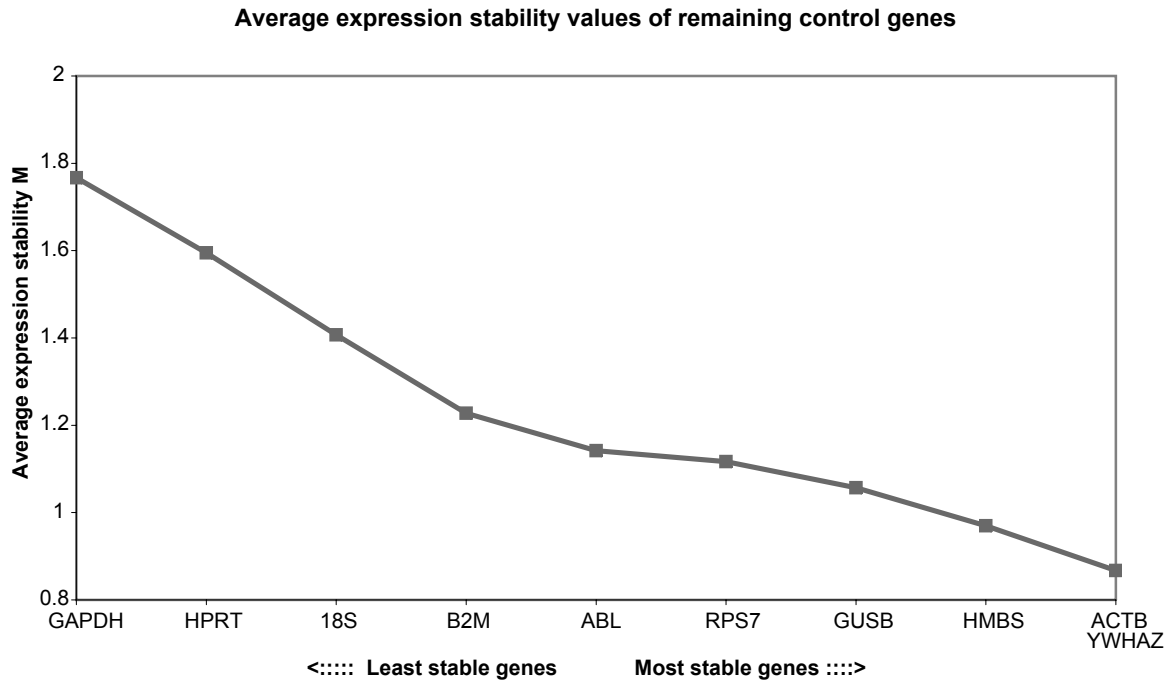
**c) geNorm output: Parathyroid**



**Additional File 2**  
**d) geNorm output: Thyroid**



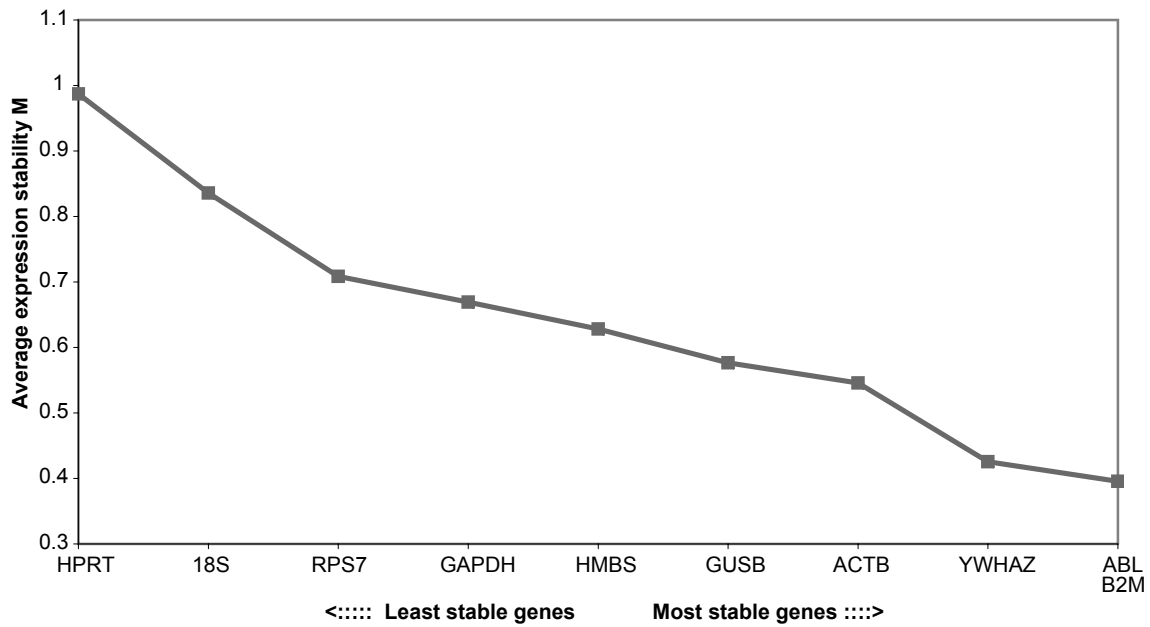
**Additional File 2**  
**e) geNorm output: Bone marrow**



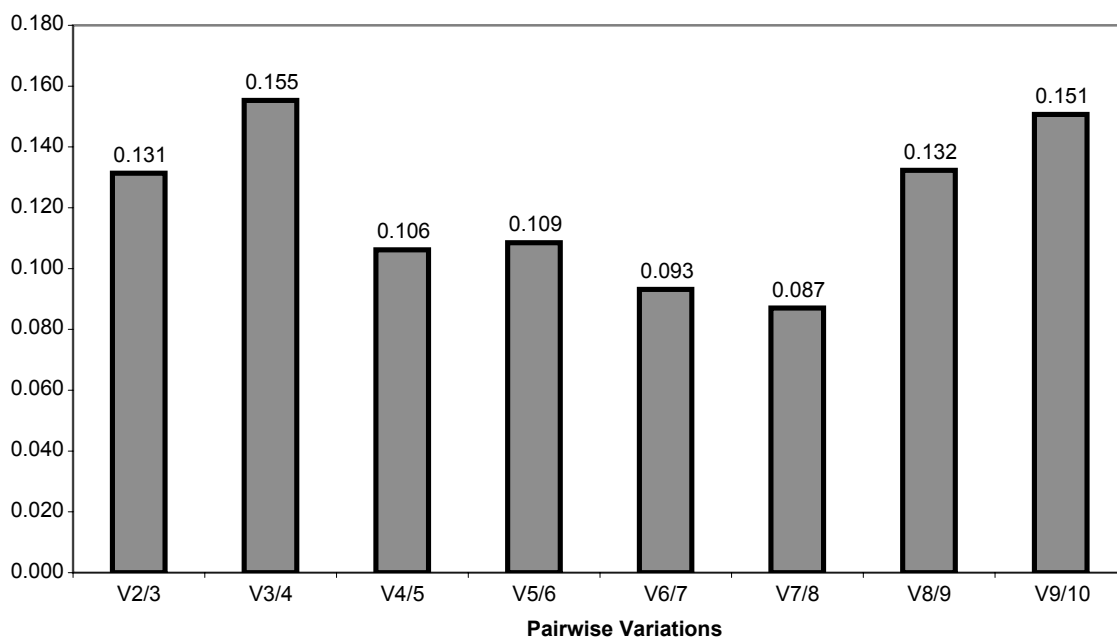
**Additional File 2**

**f) geNorm output: Mesenteric lymph node**

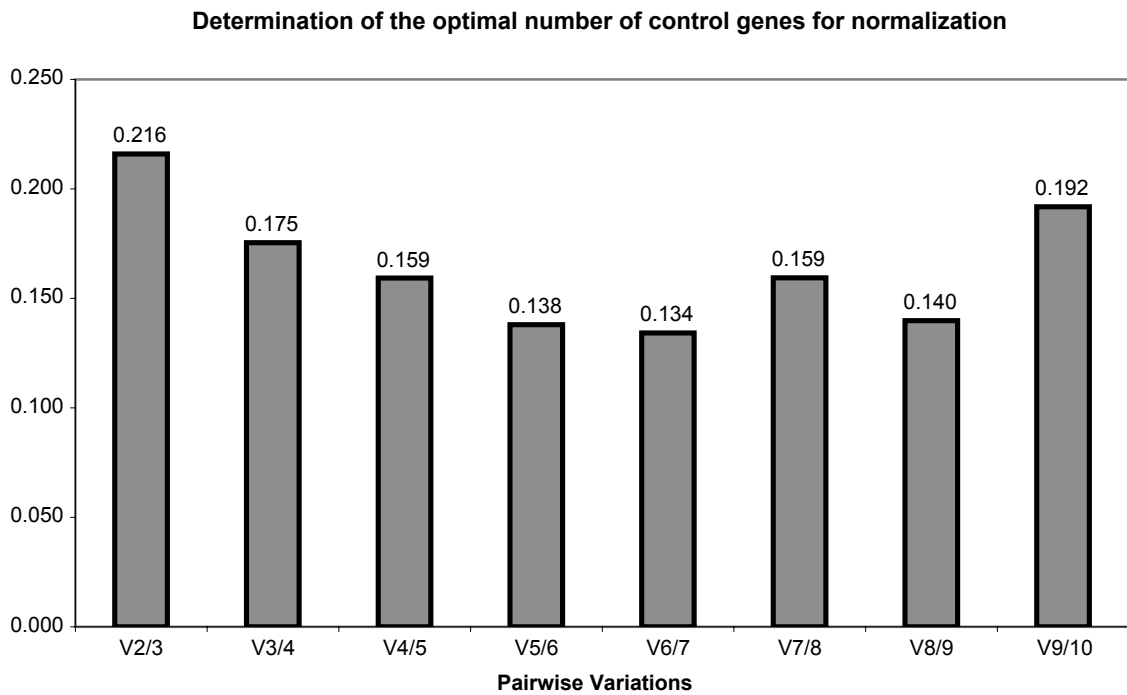
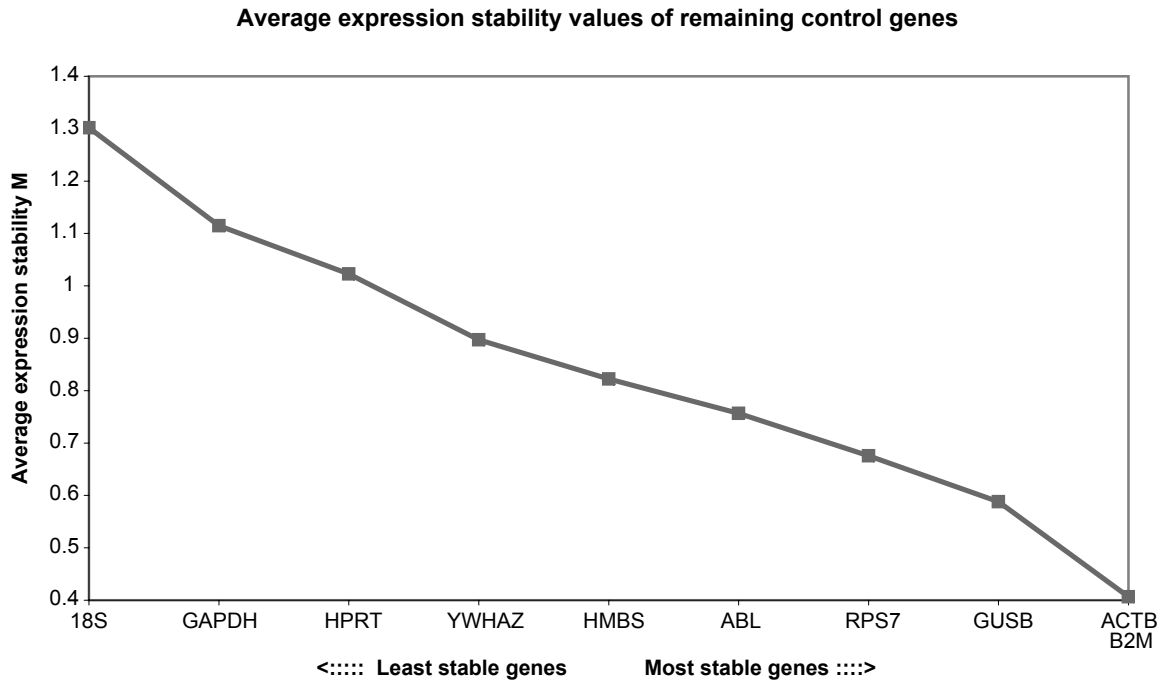
**Average expression stability values of remaining control genes**



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

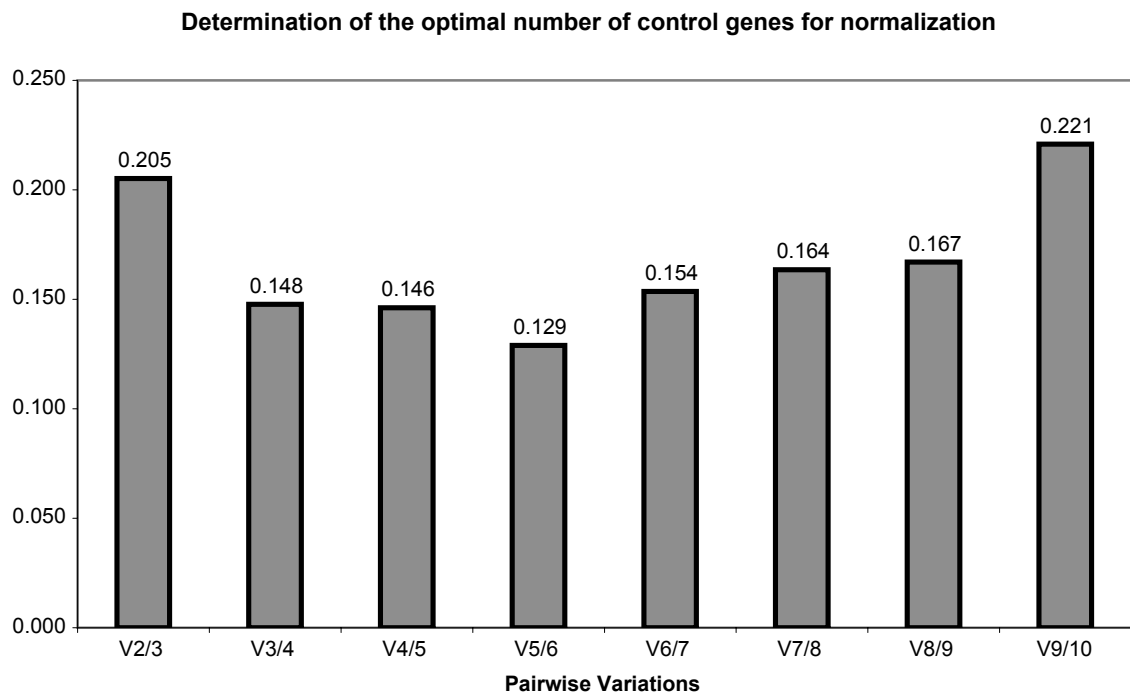
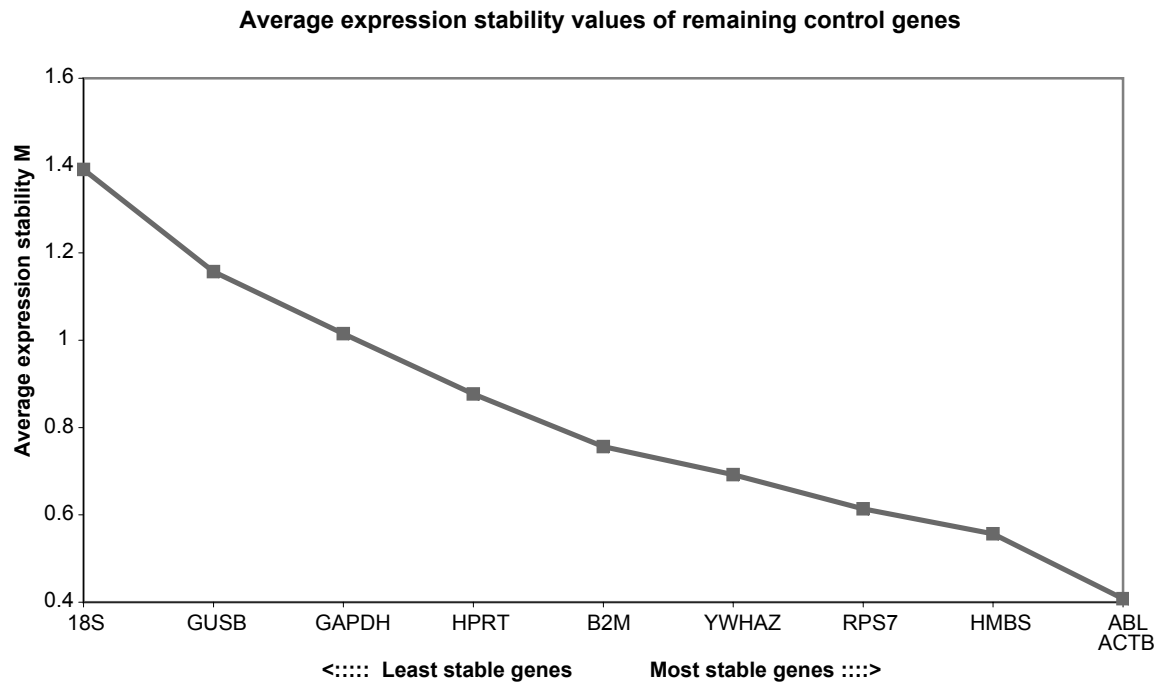


**Additional File 2**  
**g) geNorm output: Spleen**



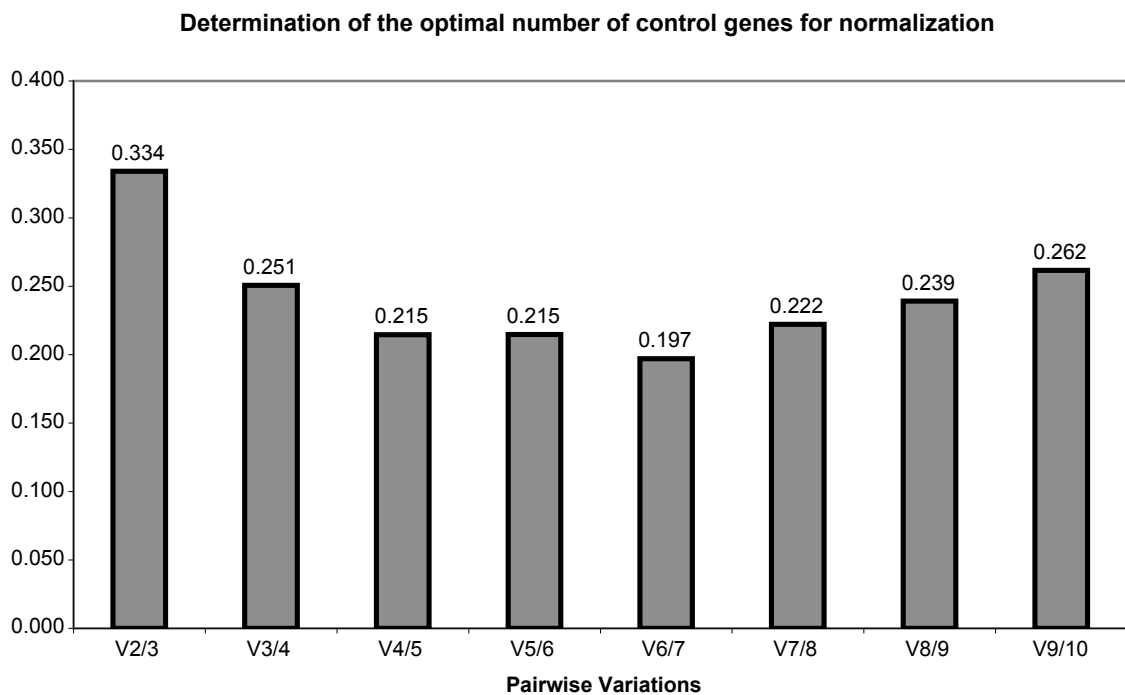
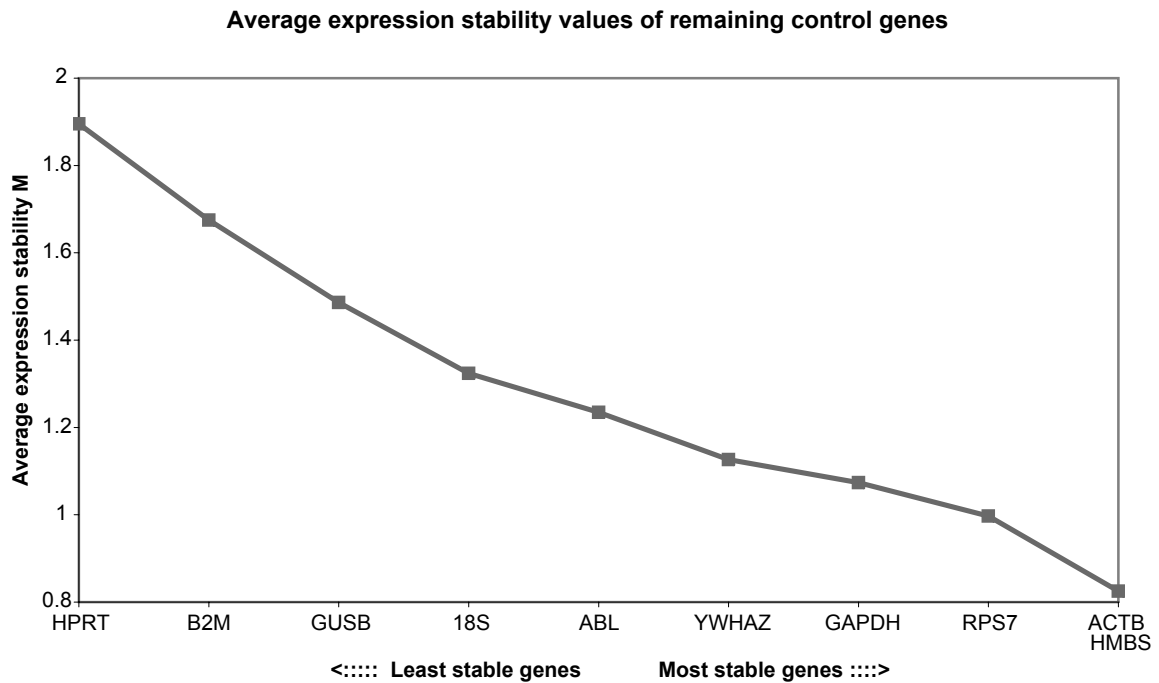
**Additional File 2**

**h) geNorm output: Parotid gland**

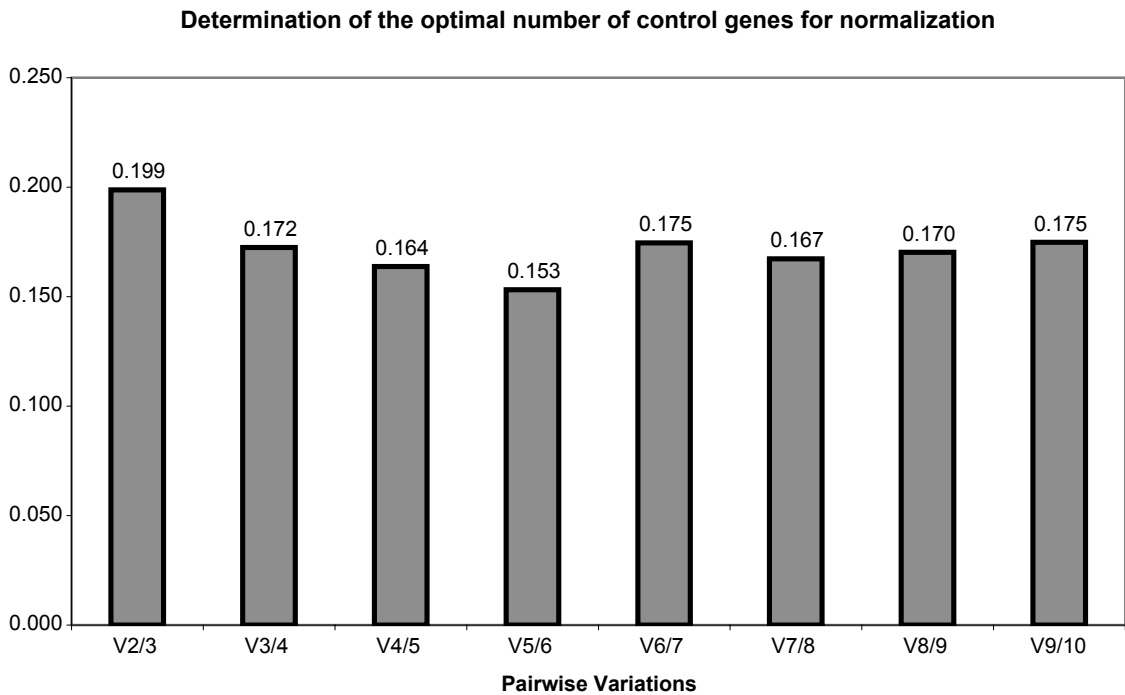
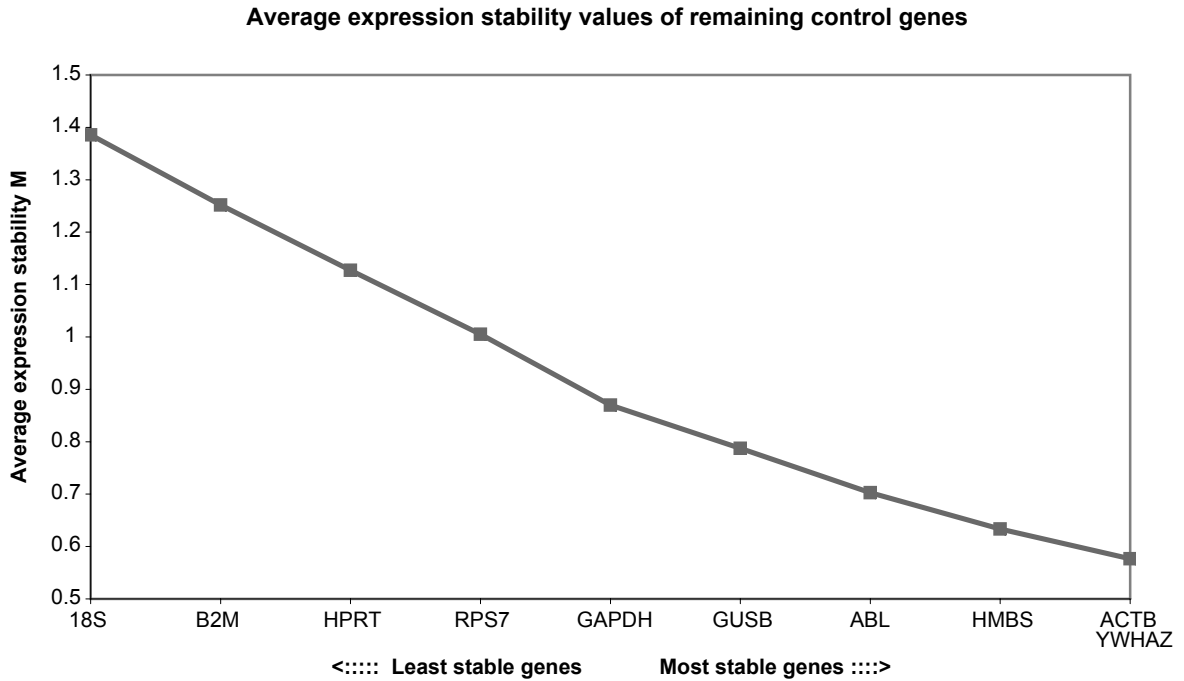


**Additional File 2**

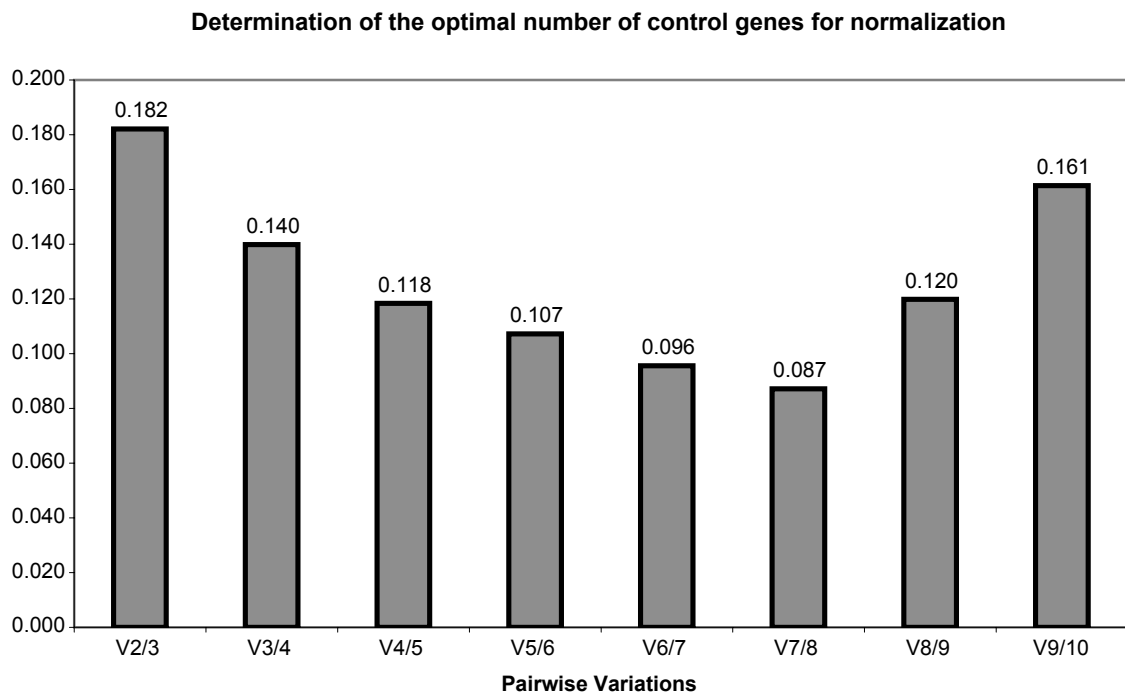
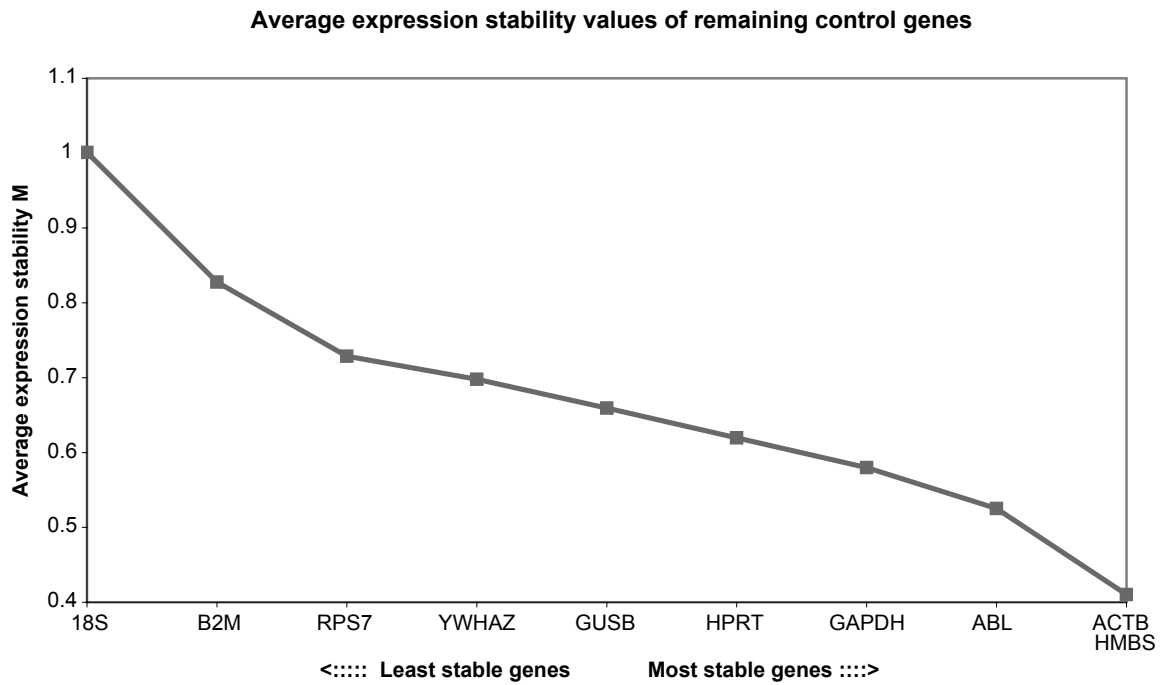
**i) geNorm output: Duodenum**



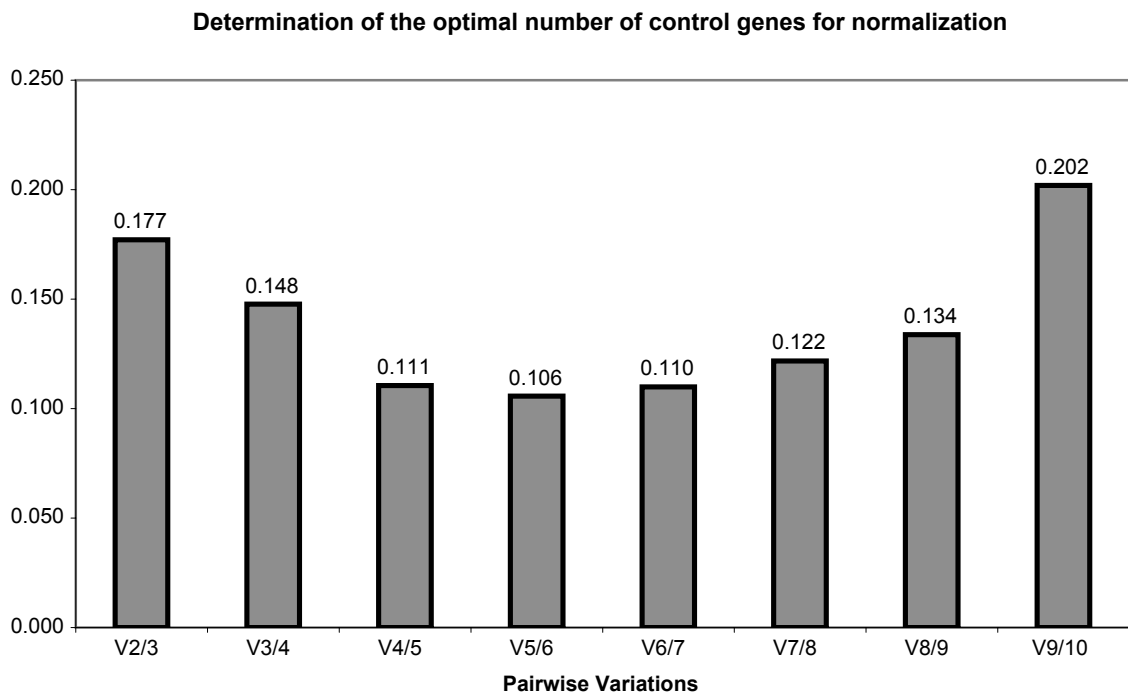
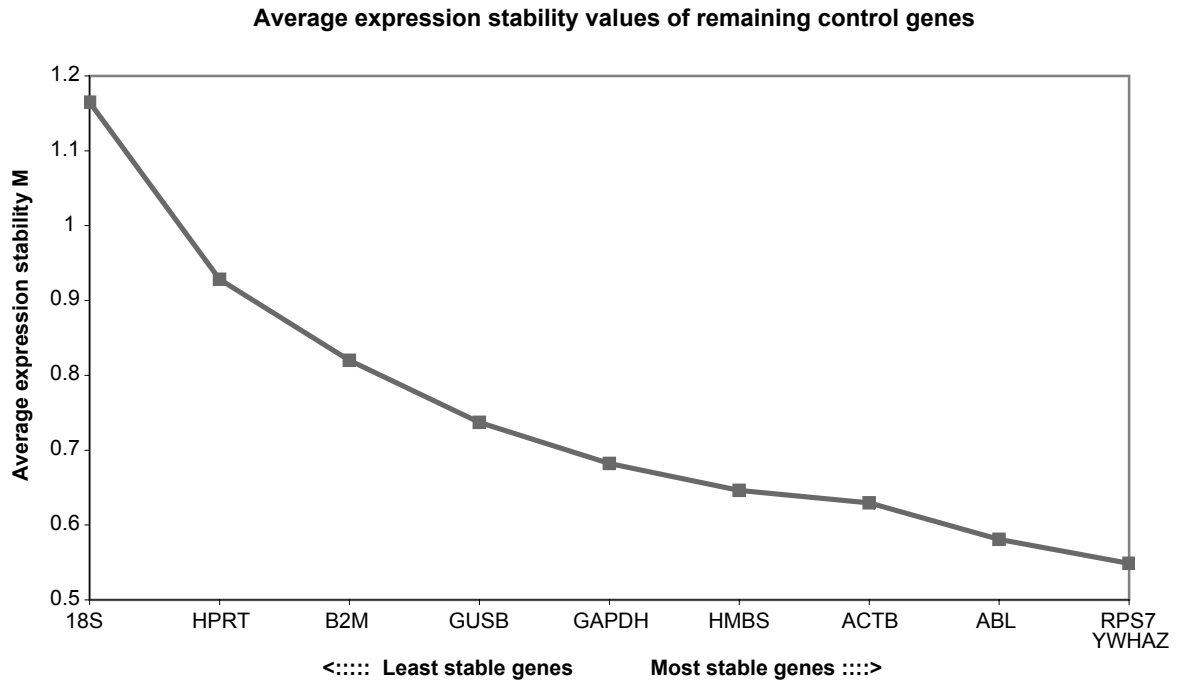
**Additional File 2**  
**j) geNorm output: Ileum**



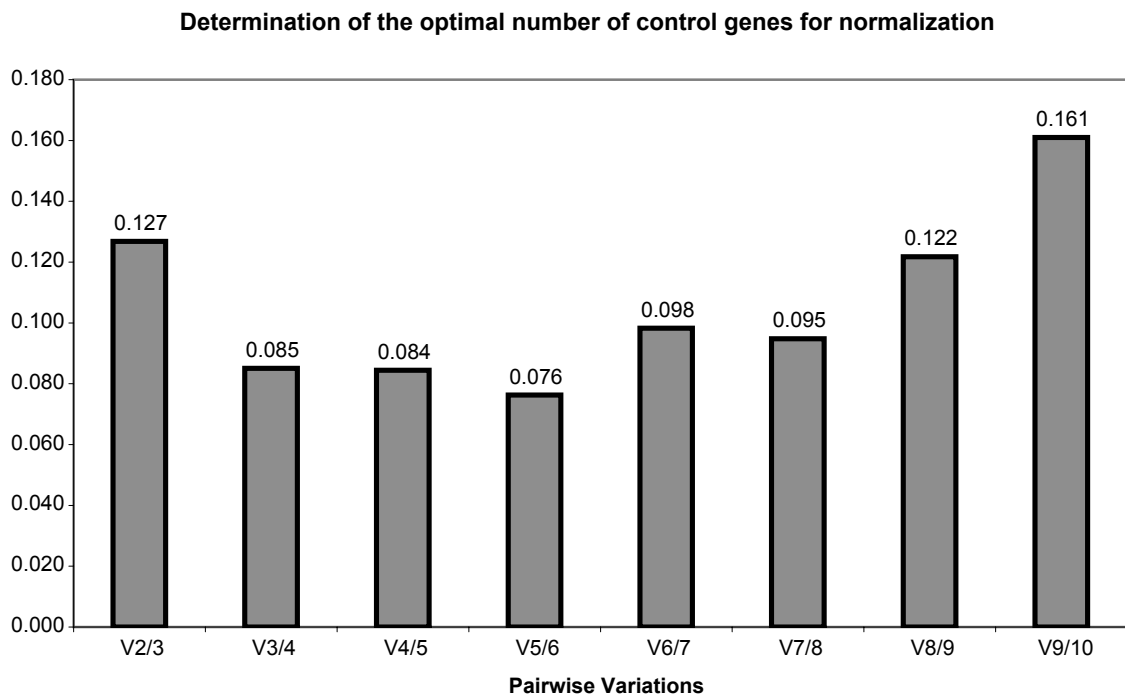
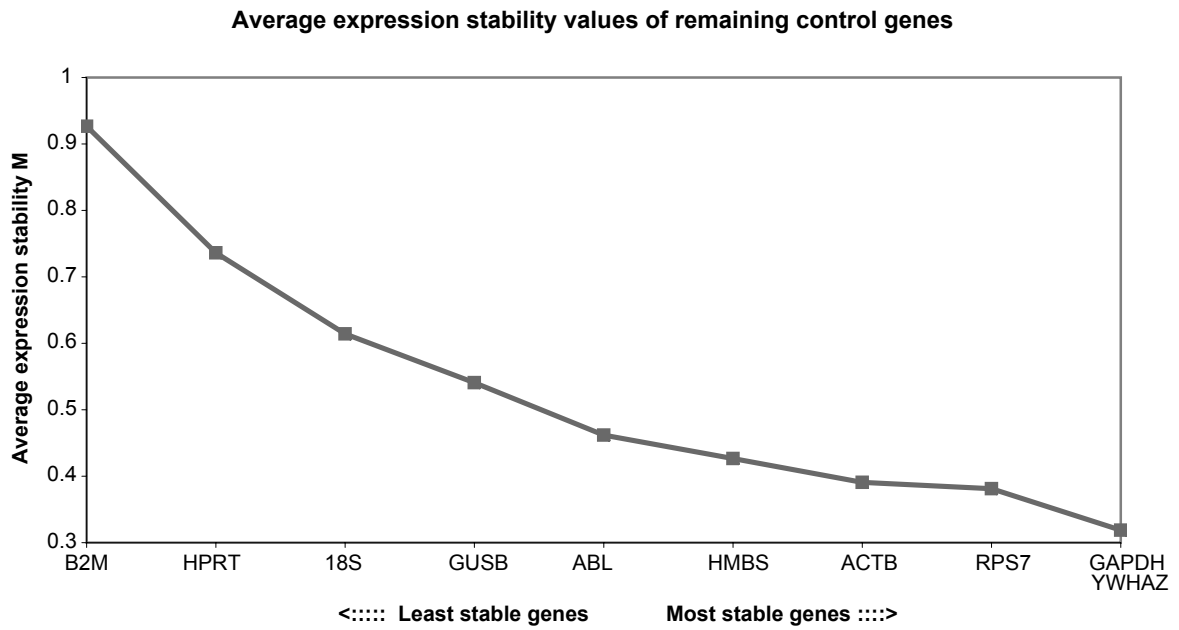
**Additional File 2**  
**k) geNorm output: Liver**



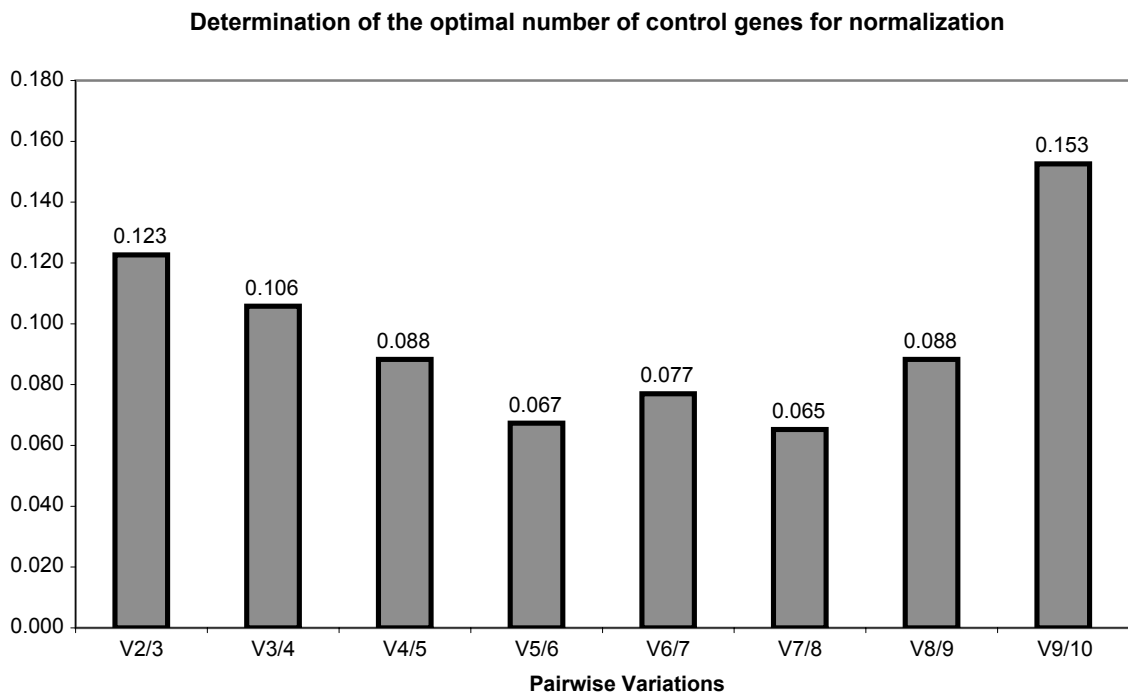
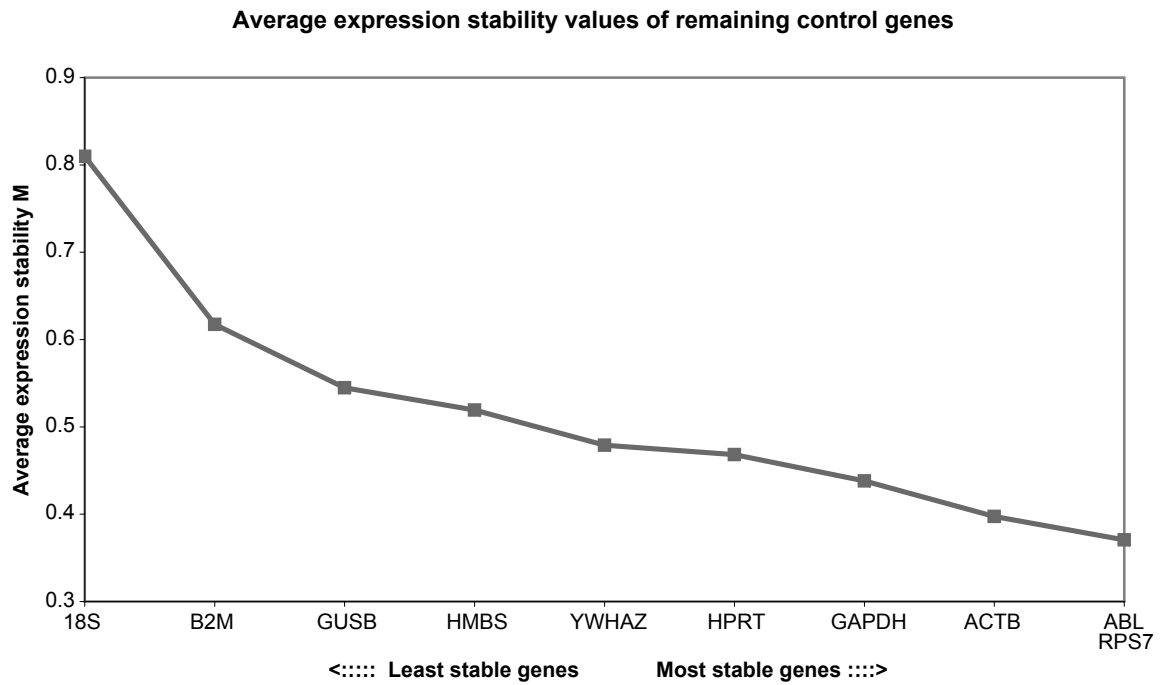
**Additional File 2**  
**l) geNorm output: Kidney**



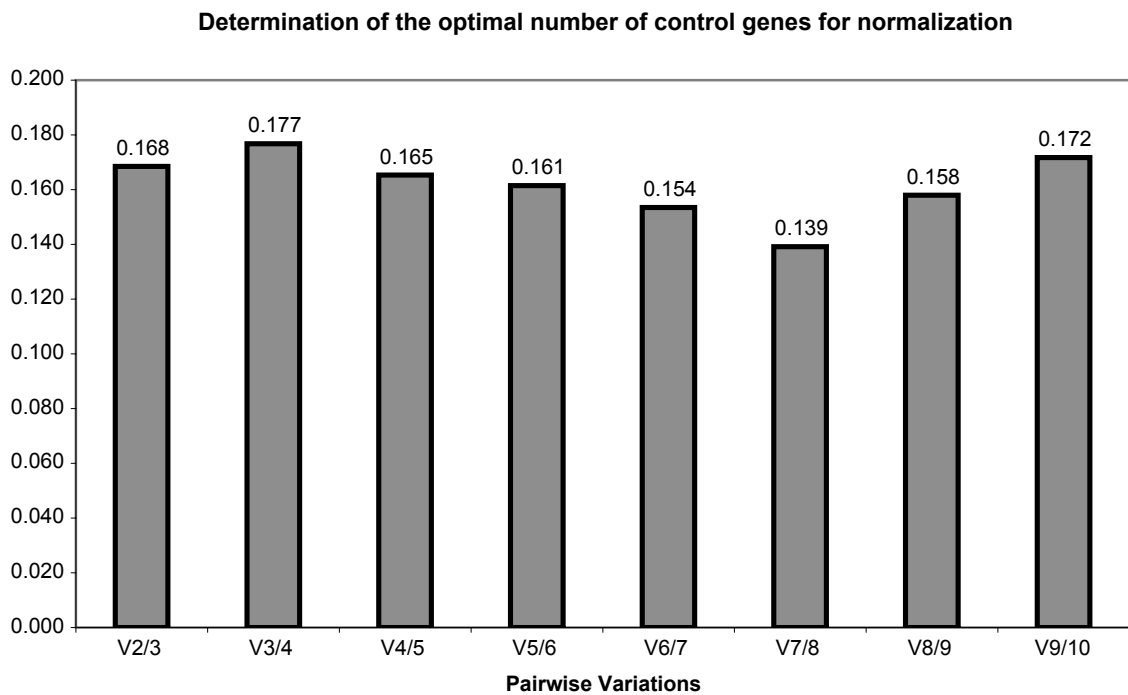
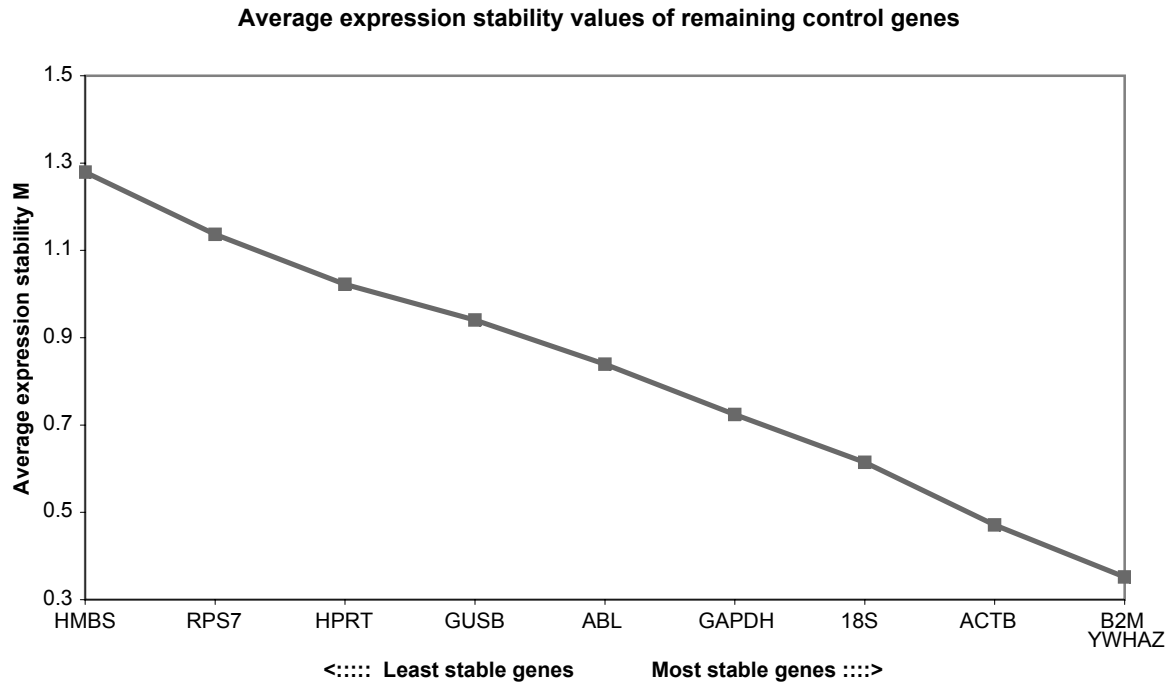
**Additional File 2**  
**m) geNorm output: Myocardium**



**Additional File 2**  
**n) geNorm output: Brain**

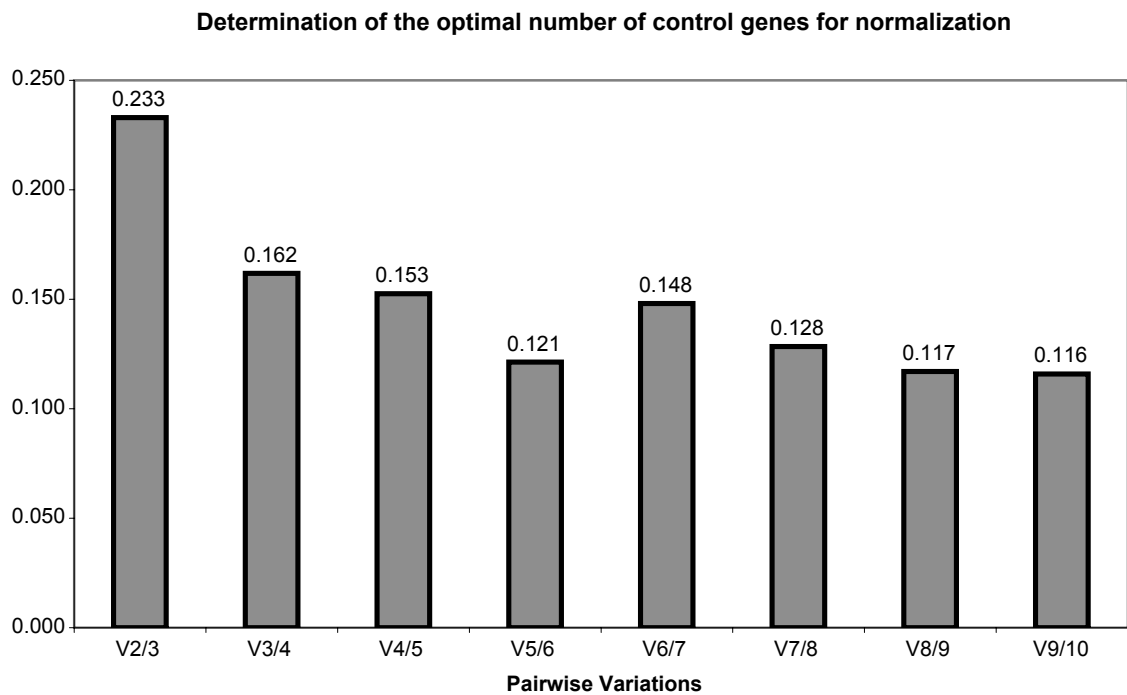
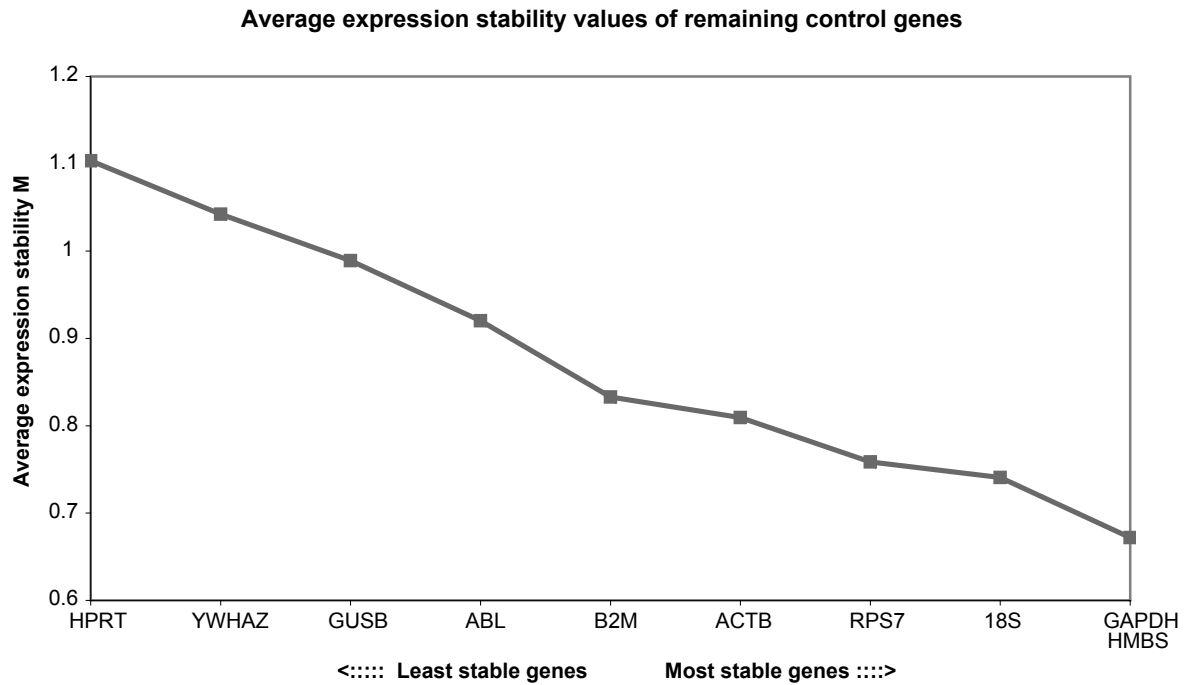


**Additional File 2**  
**o) geNorm output: Blood**



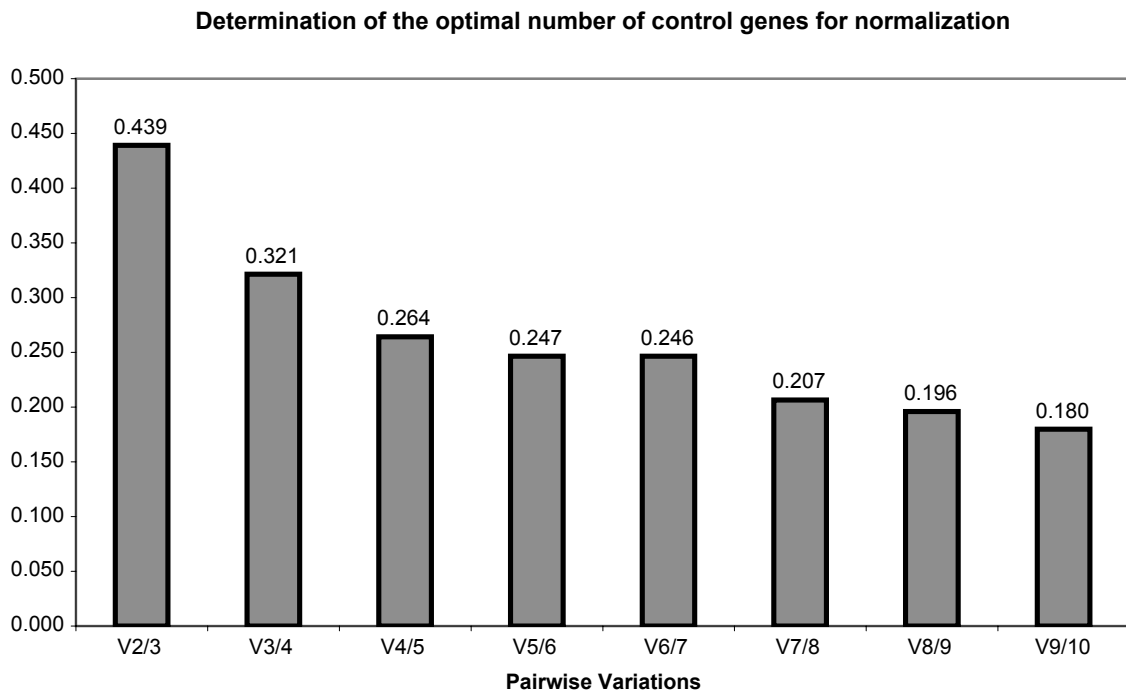
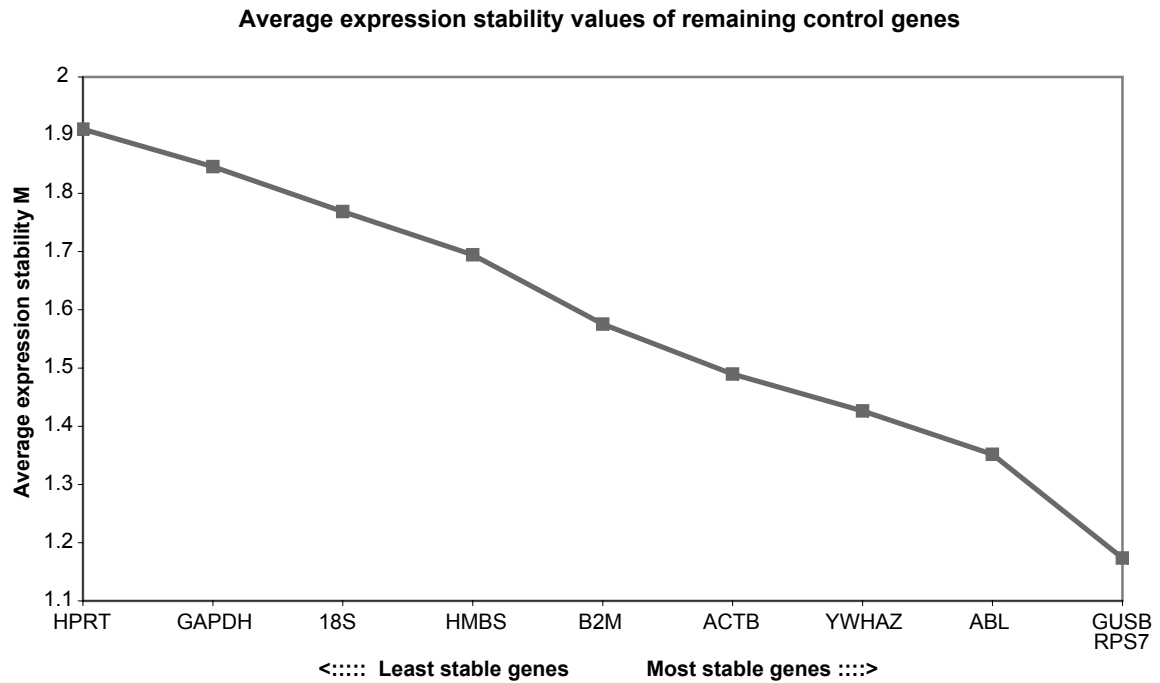
**Additional File 2**

**p) geNorm output: Neoplastic tissues**



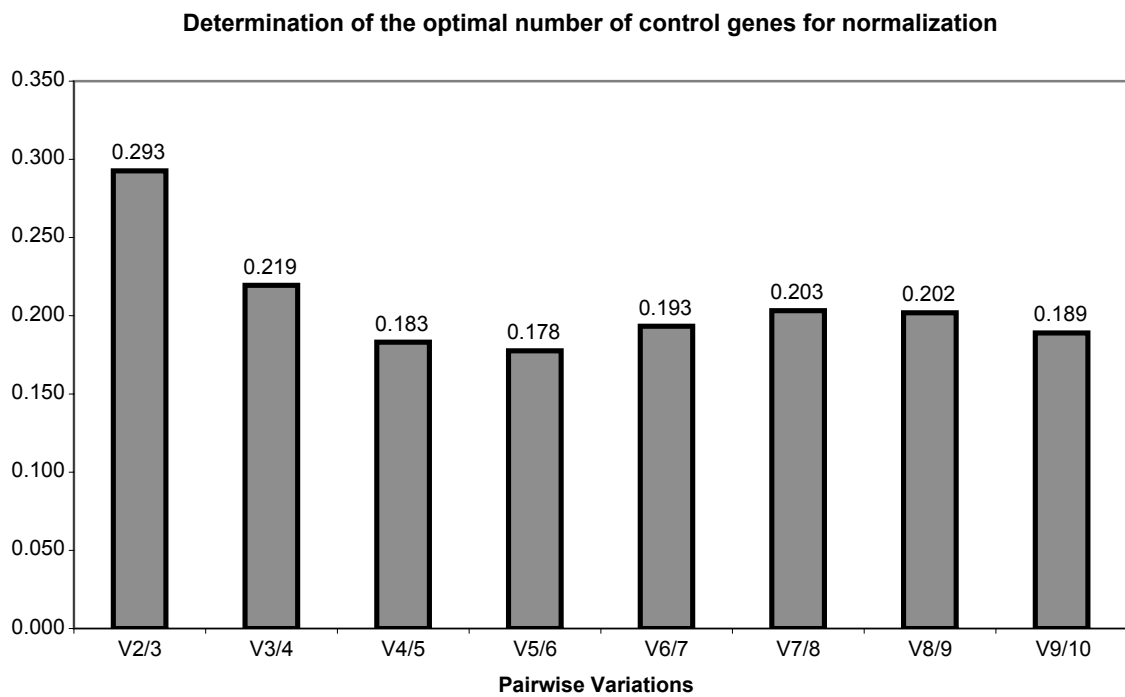
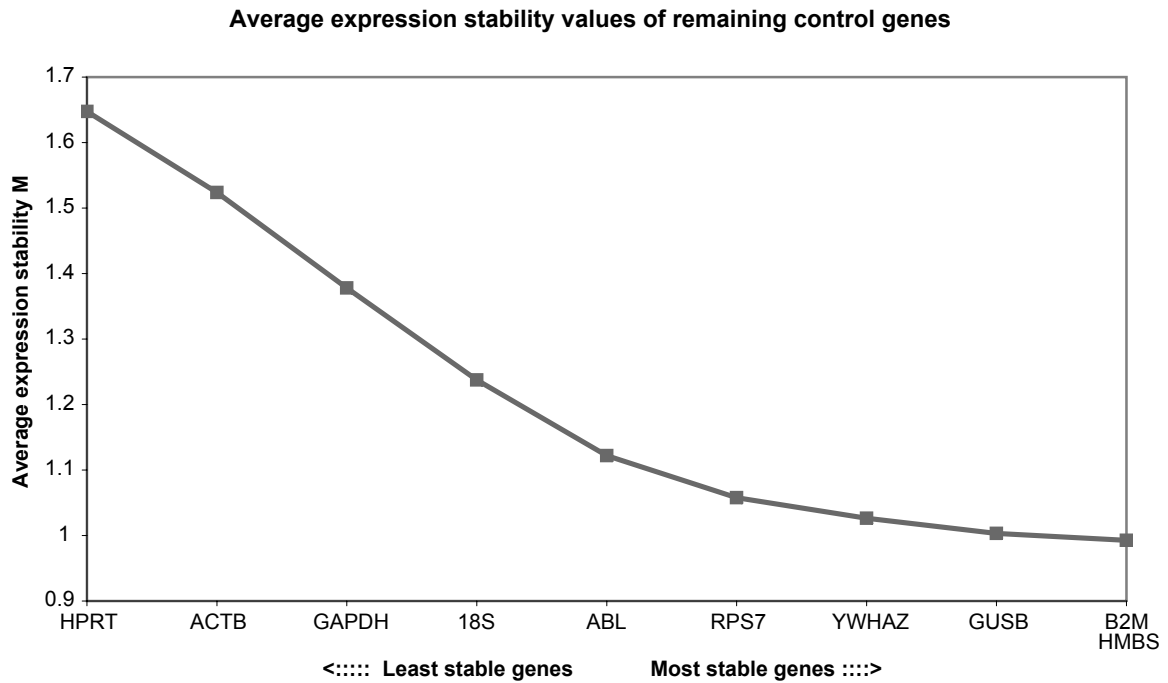
**Additional File 2**

**q) geNorm output: All healthy tissues**

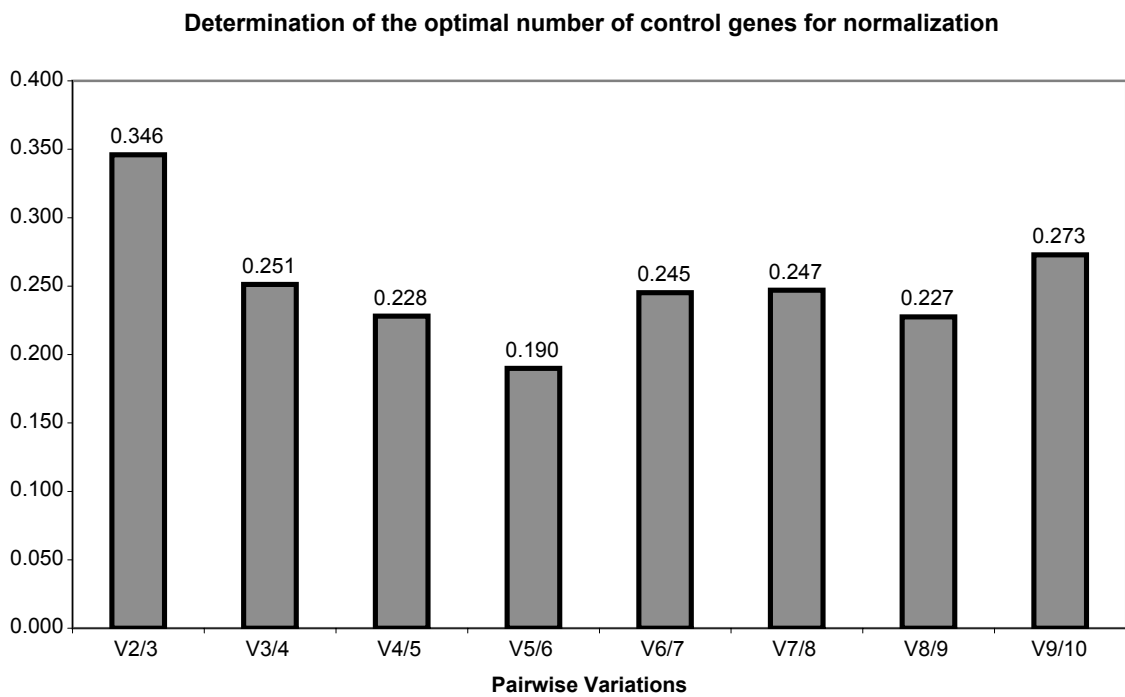
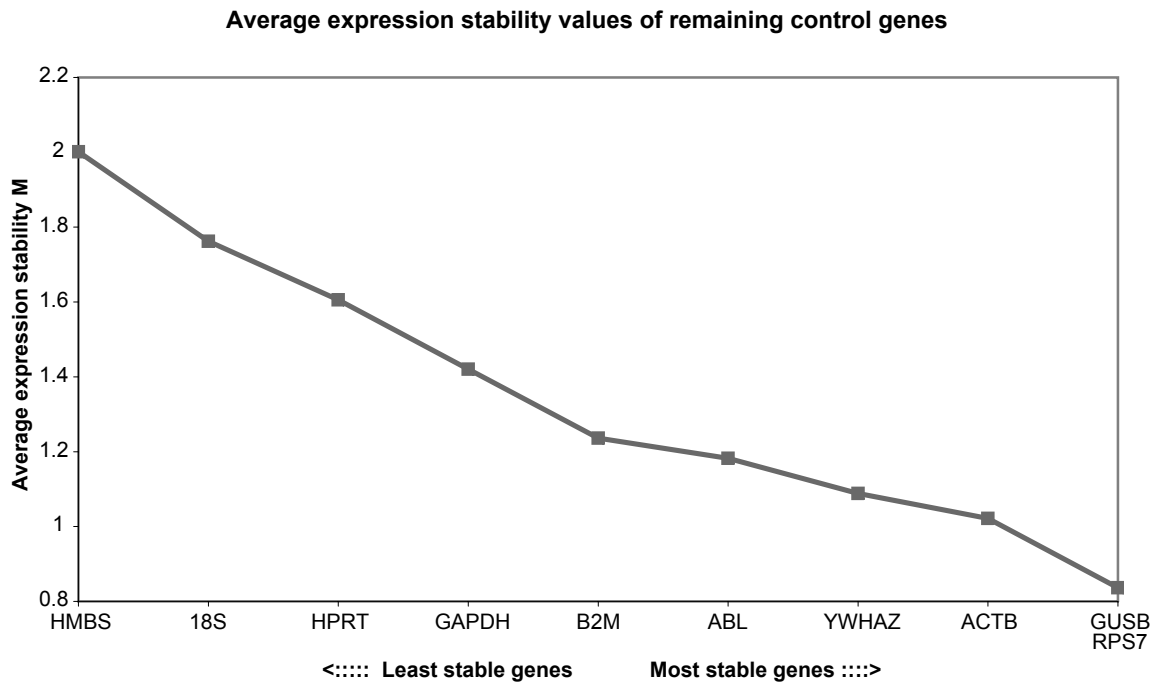


**Additional File 2**

**r) geNorm output: Endocrine tissues**



**Additional File 2**  
**s) geNorm output: Lymphoid tissues**



**Additional File 2**

**t) geNorm output: Gastrointestinal tissues**

