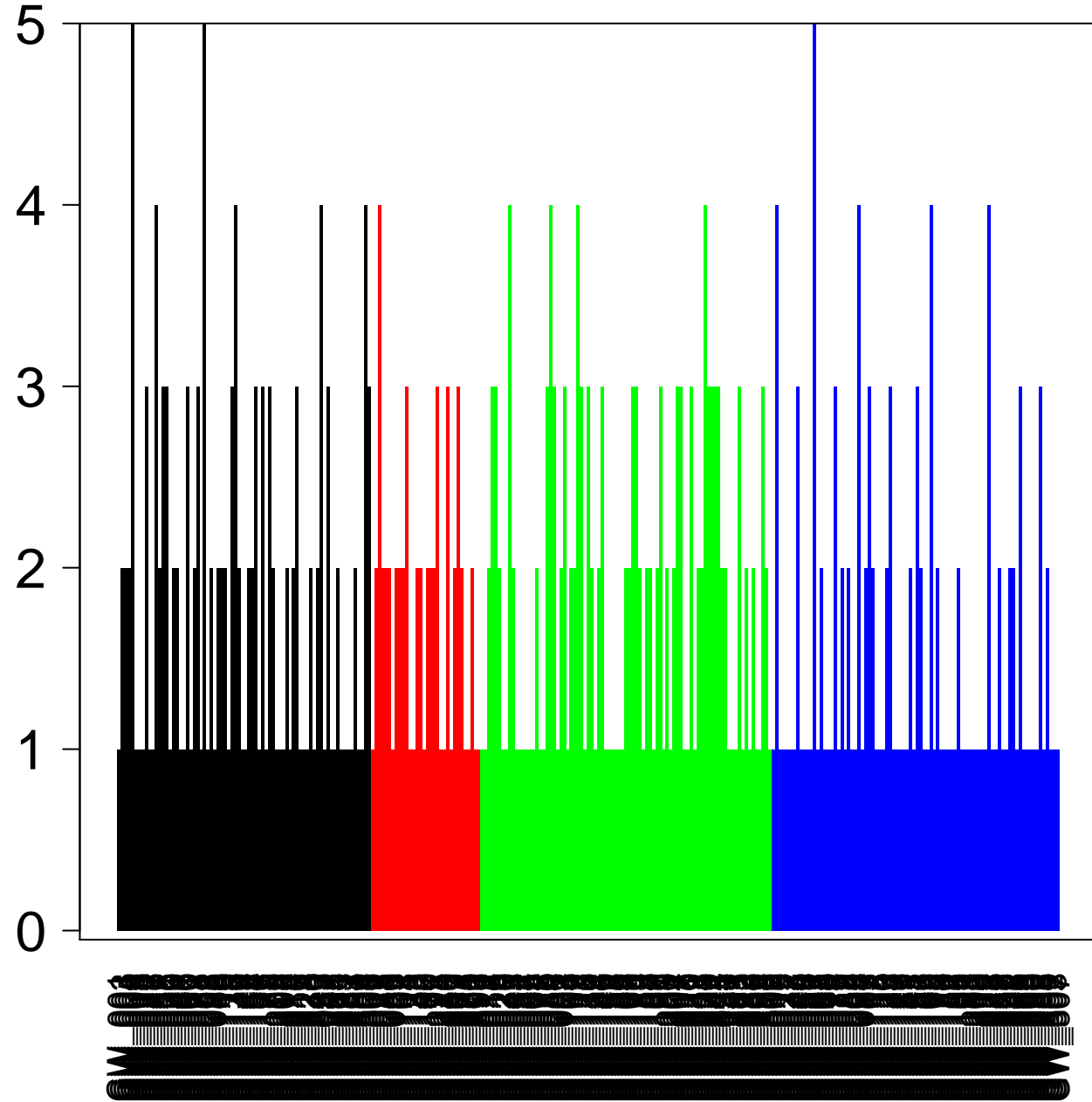
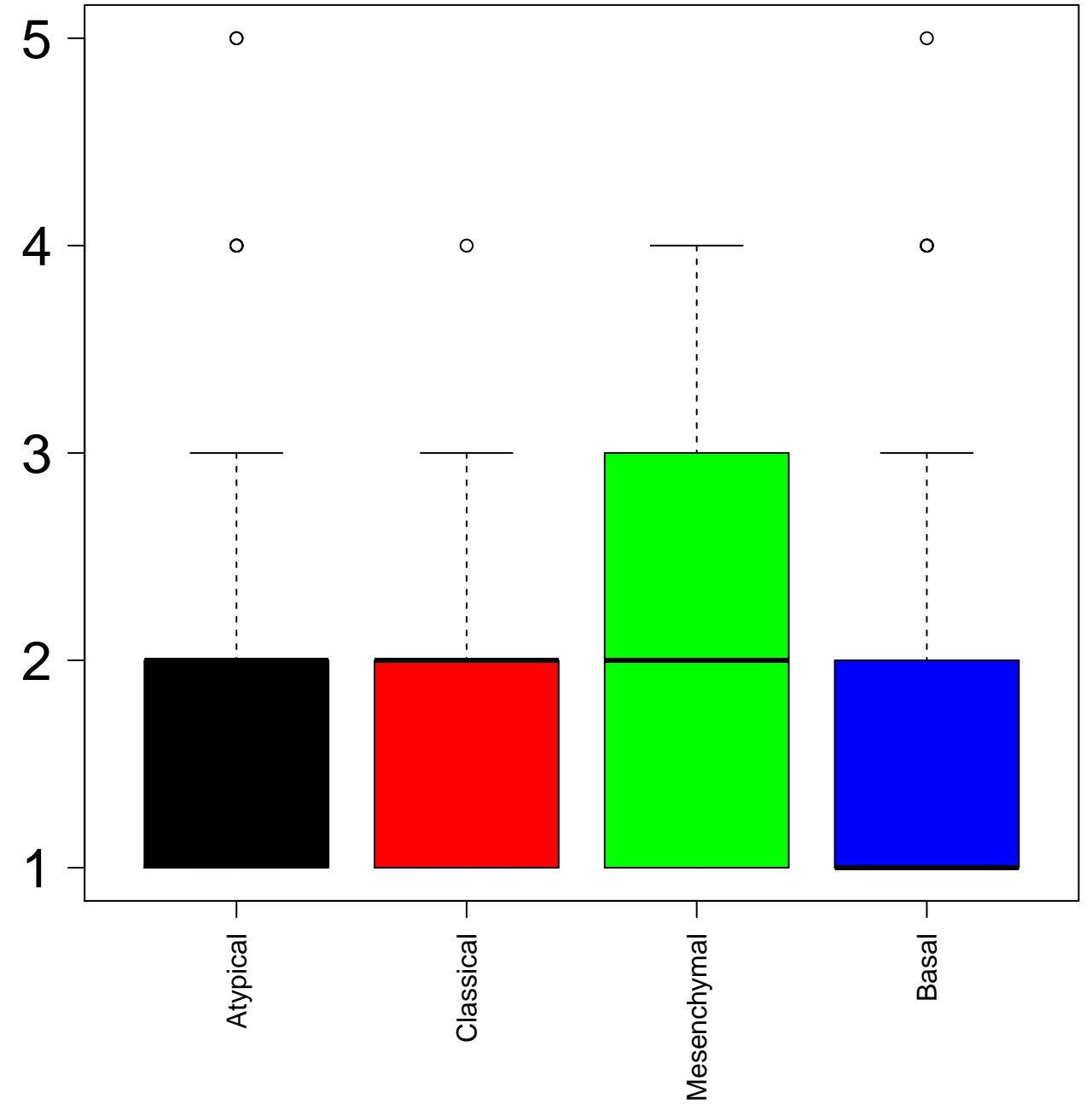


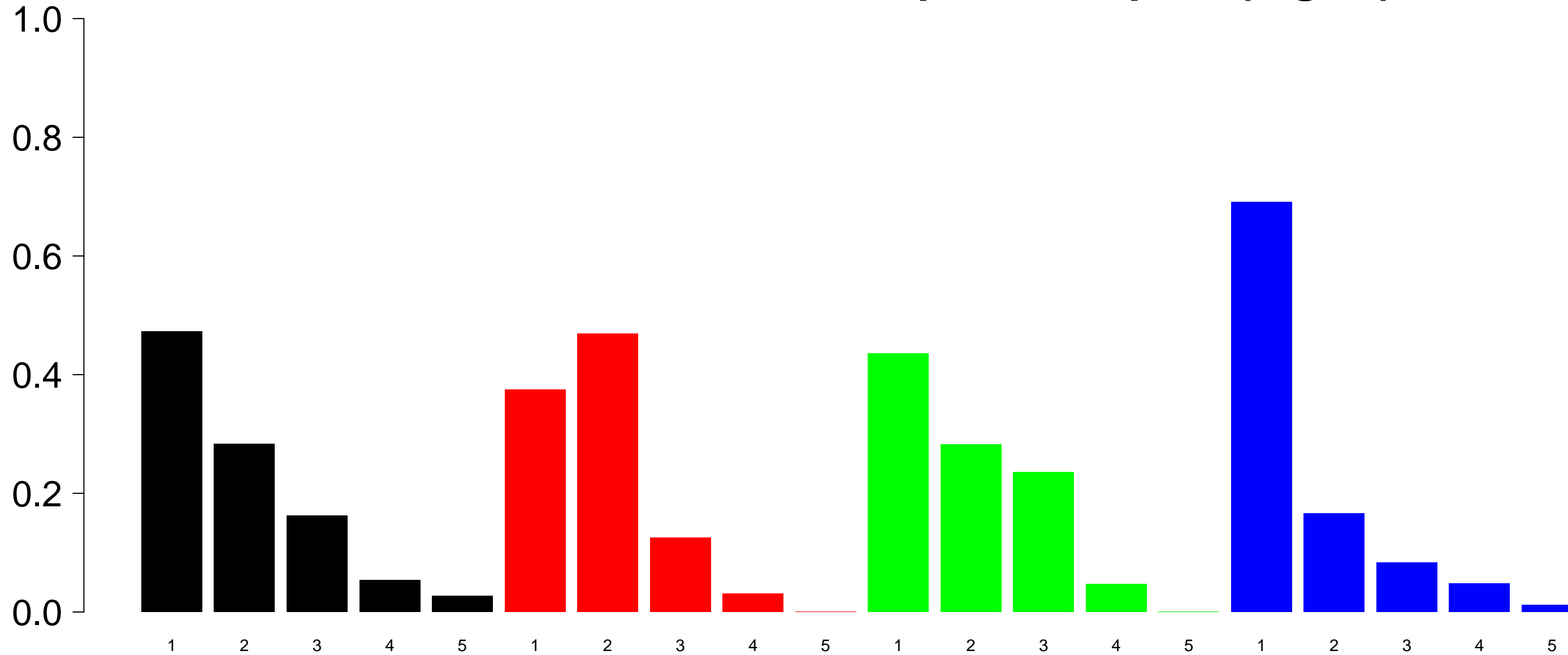
# Number of overexpressed spots (logFC)



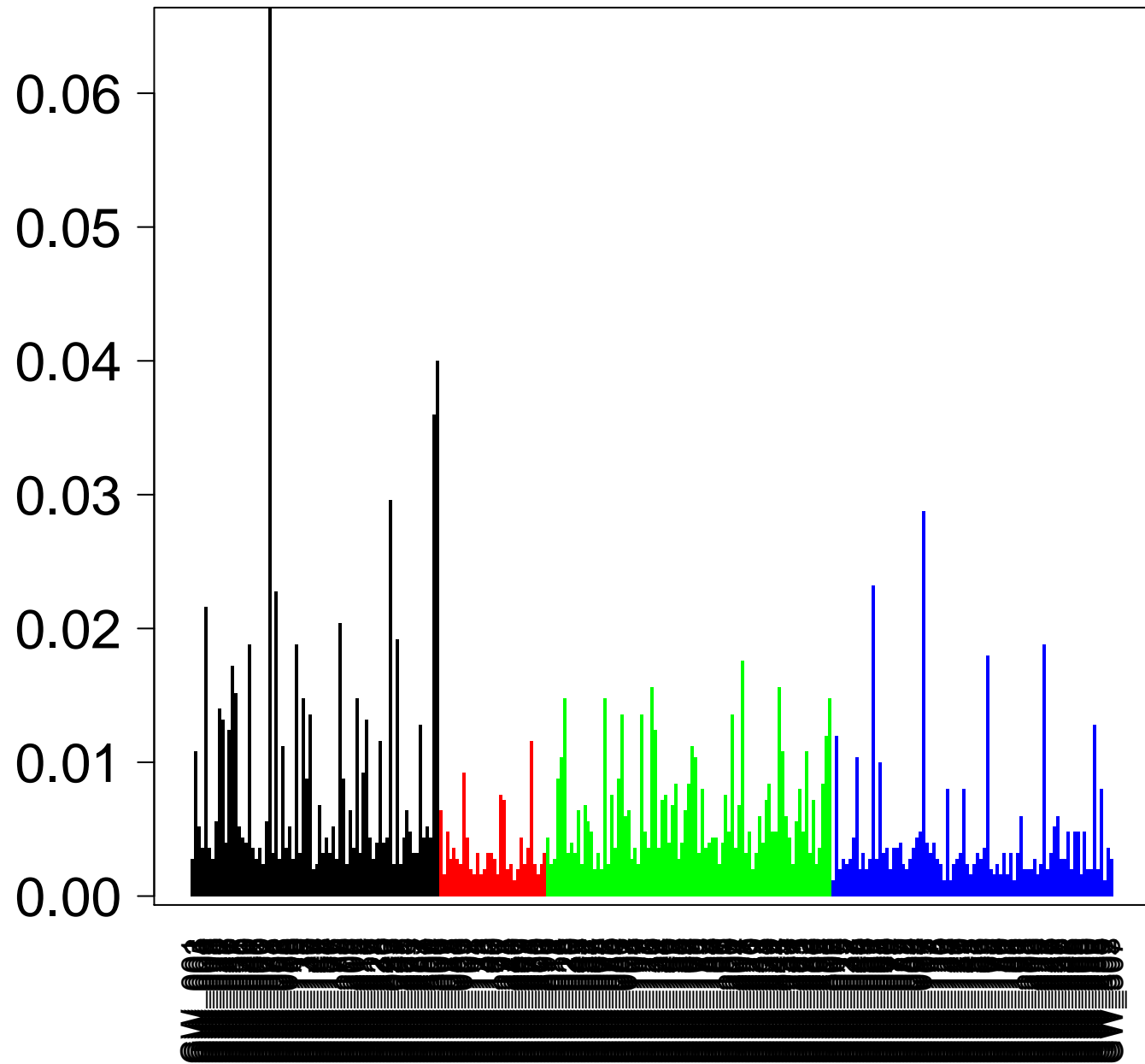
# Number of overexpressed spots (logFC)



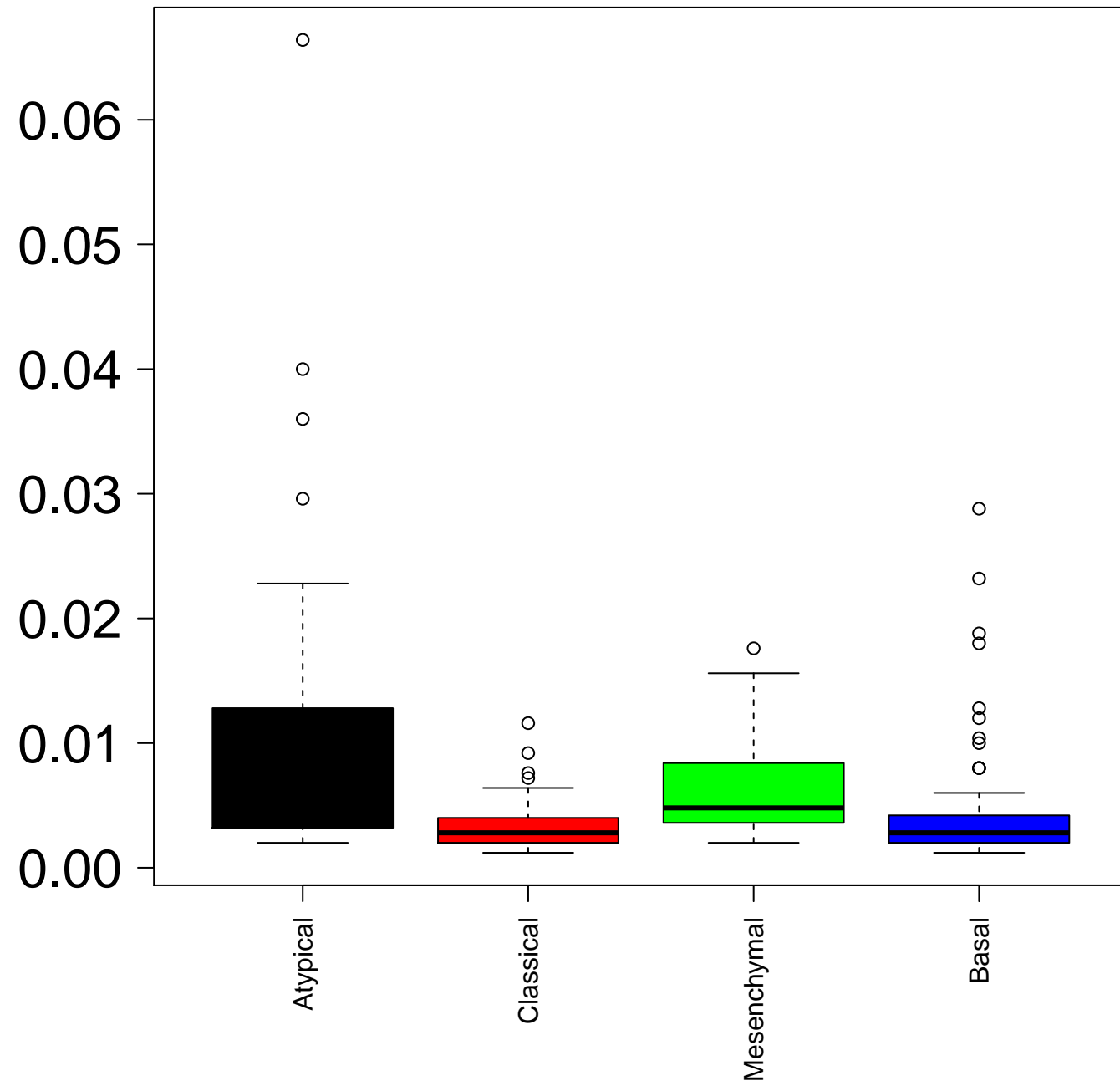
# Relative numbers of overexpressed spots (logFC)



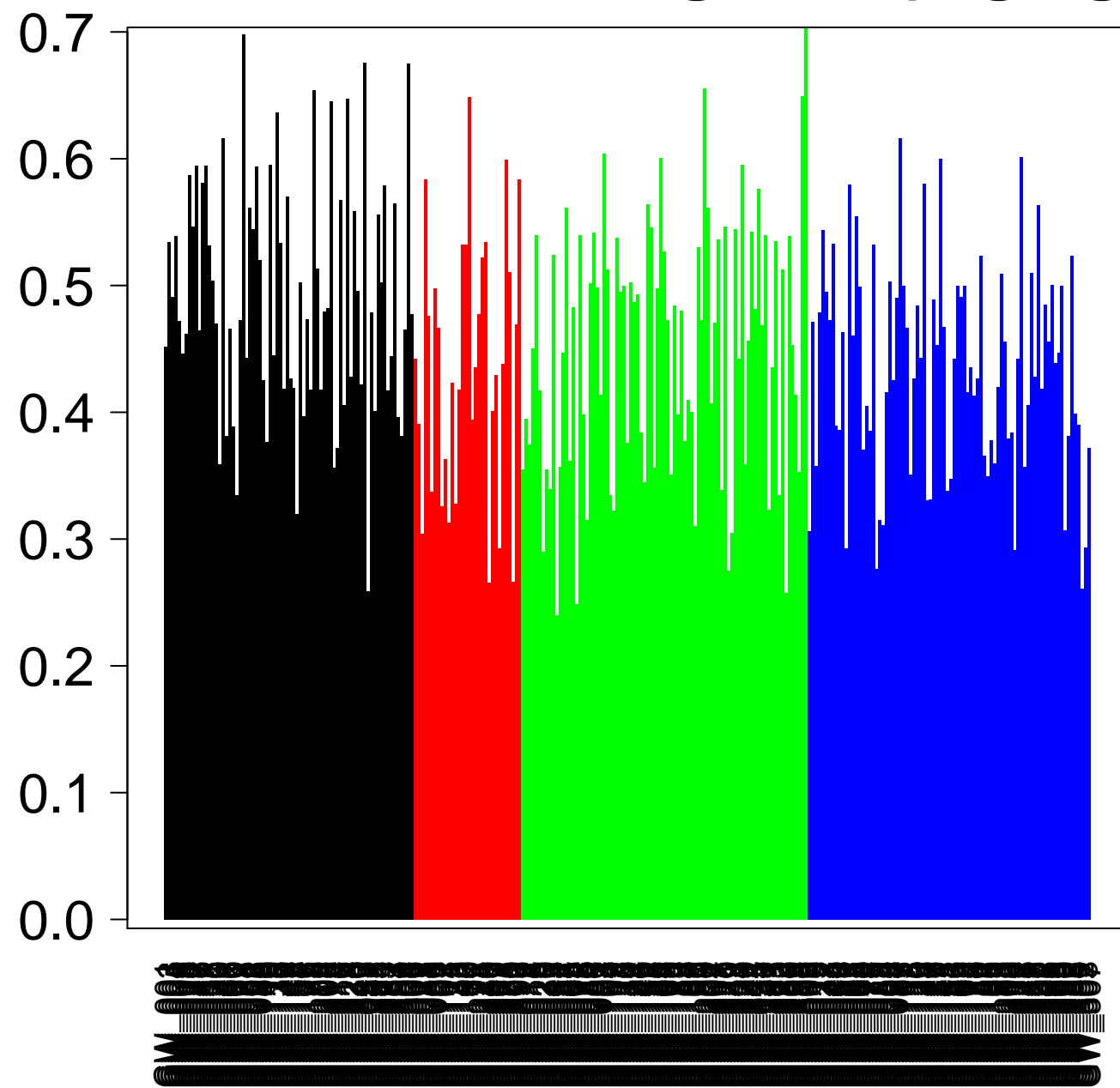
### Fraction of red metagenes (logFC)



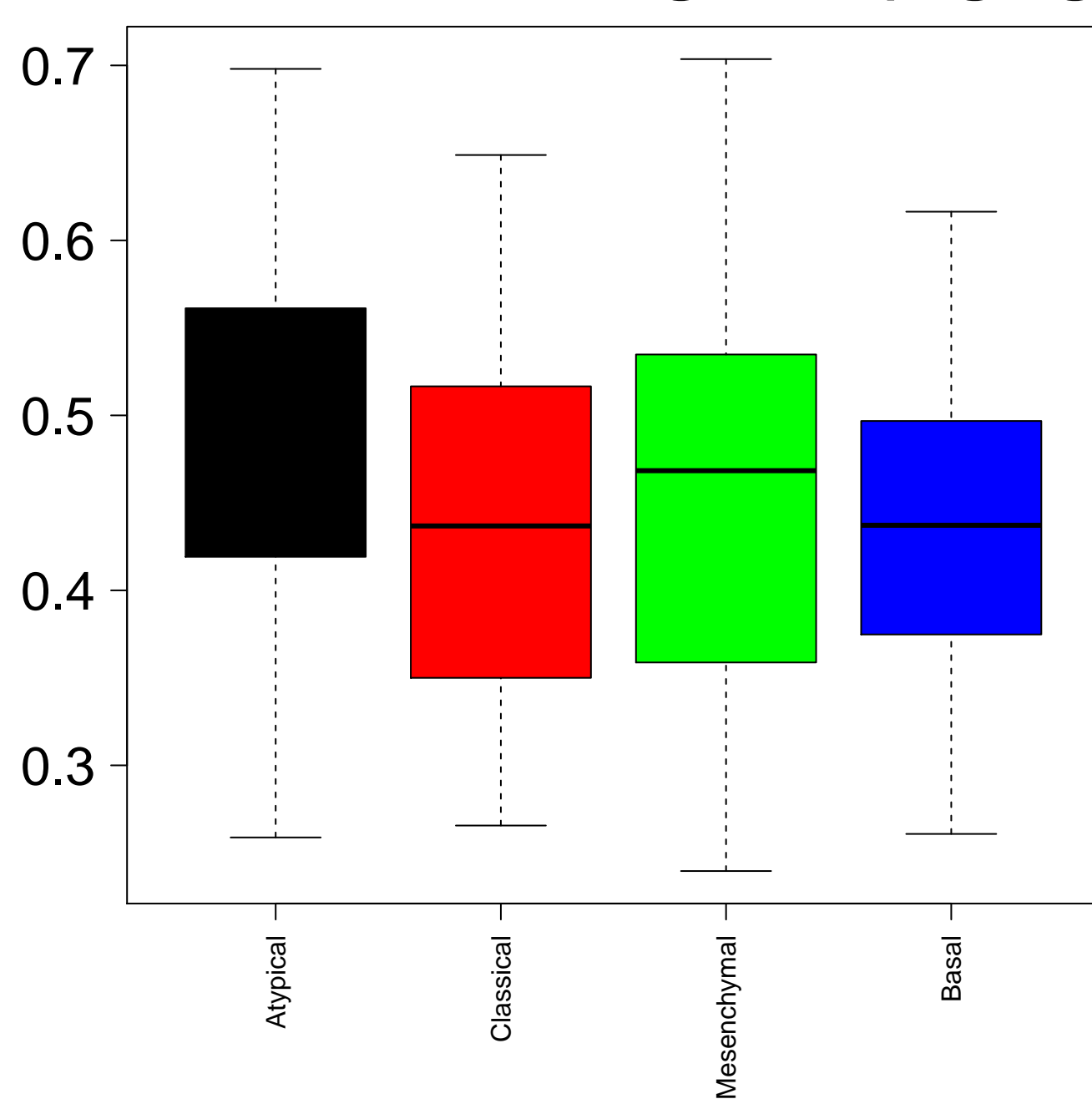
### Fraction of red metagenes (logFC)



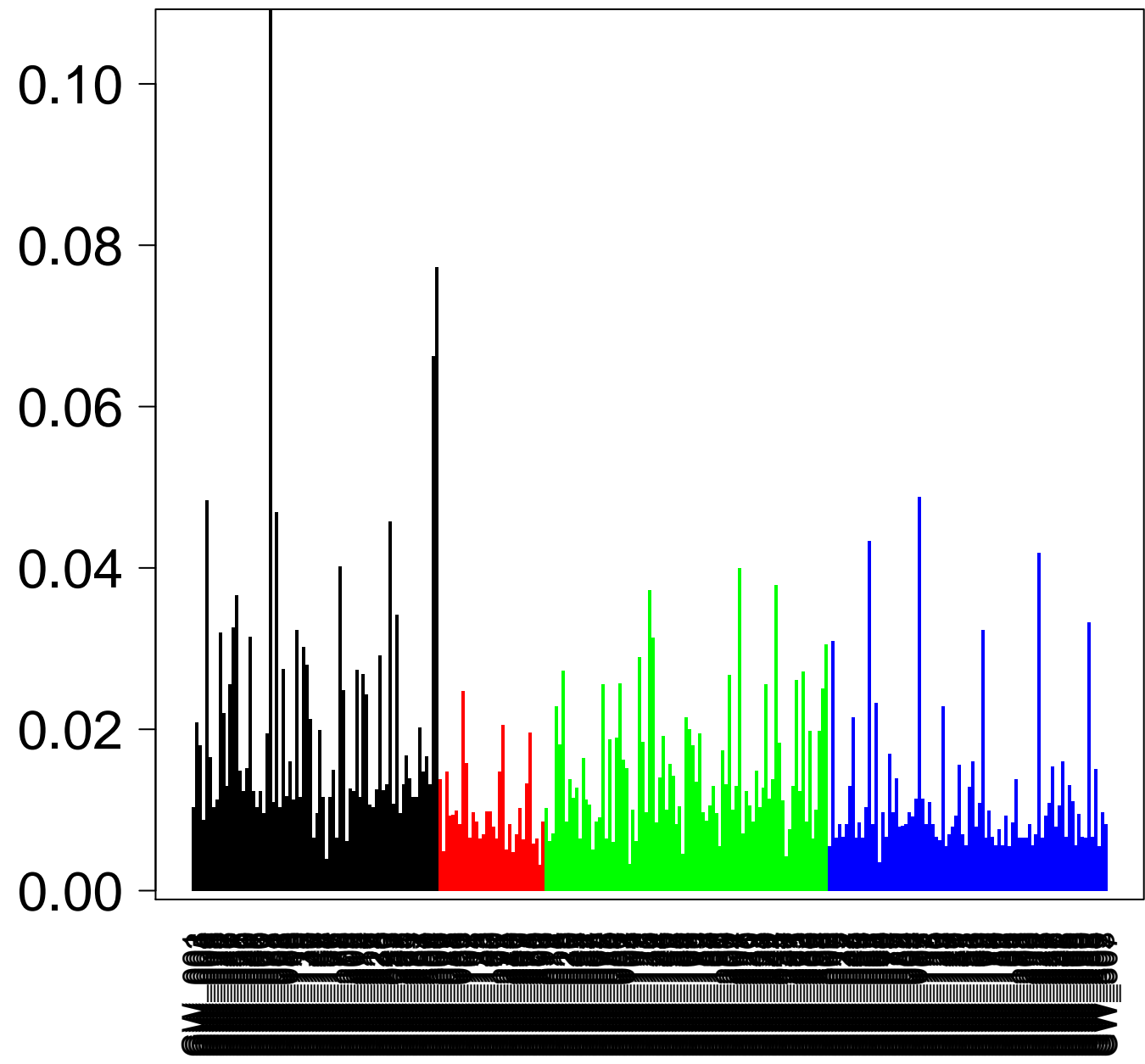
### Fraction of red metagenes (loglogFC)



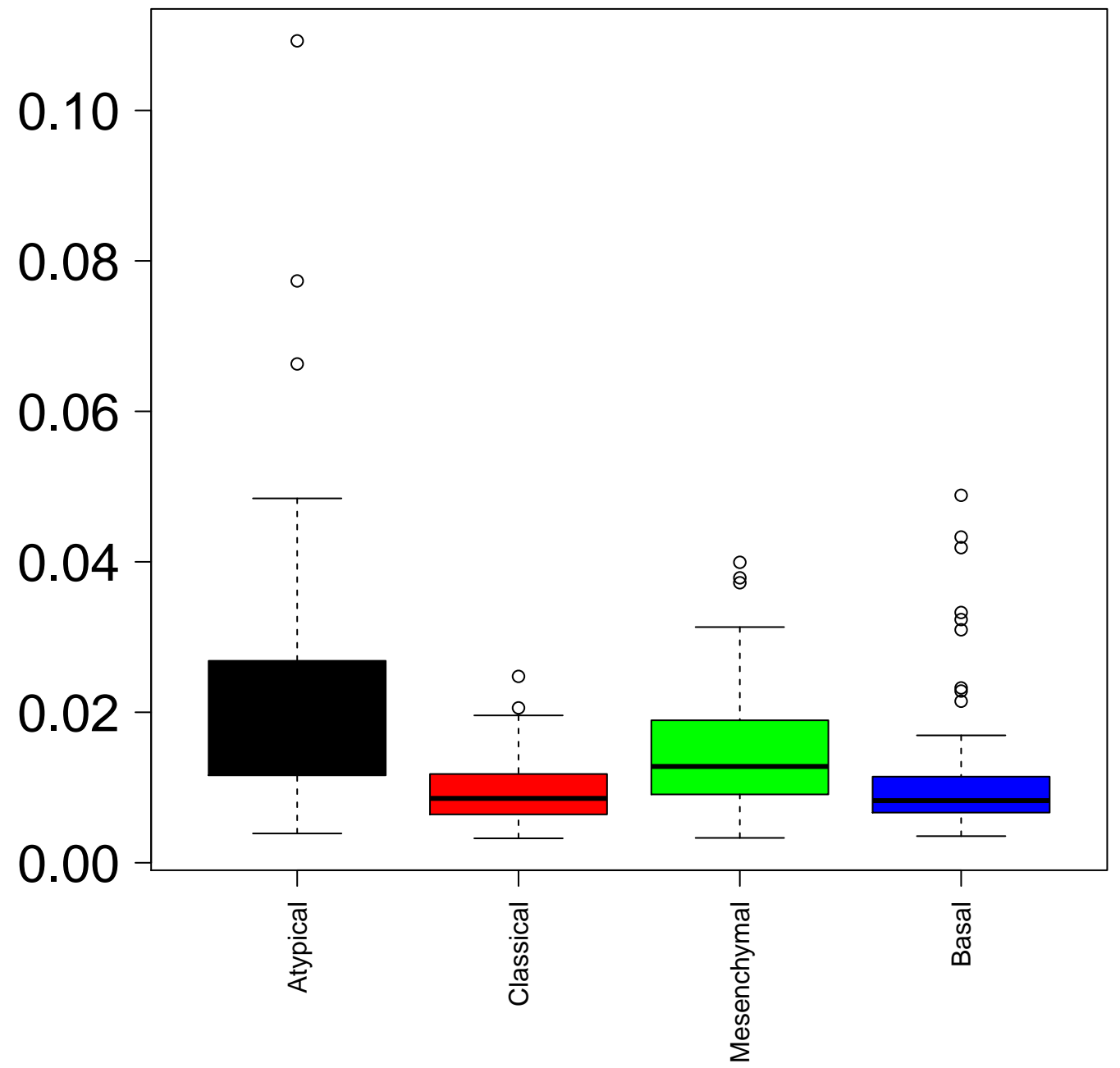
### Fraction of red metagenes (loglogFC)



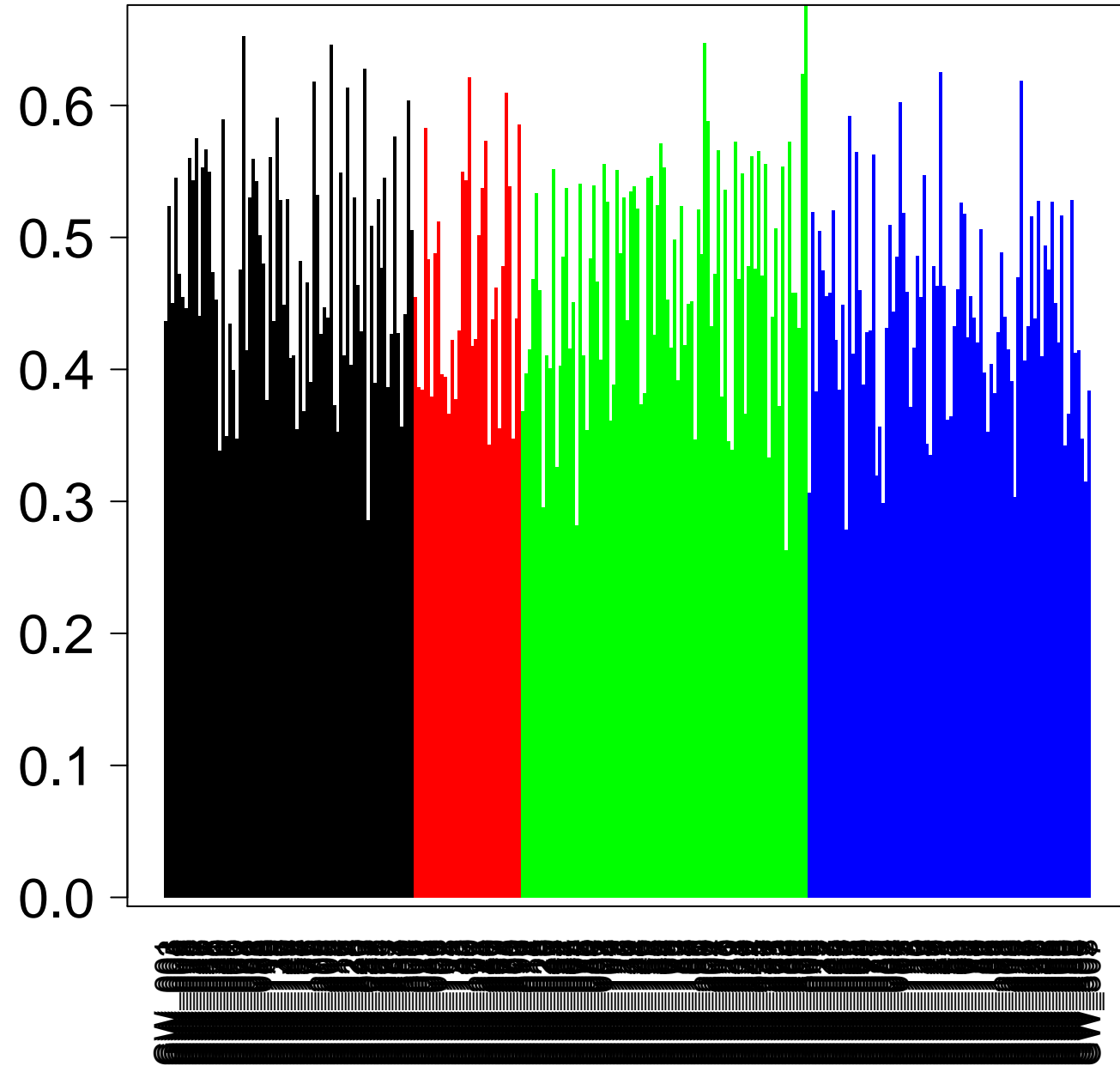
# Fraction of red genes (logFC)



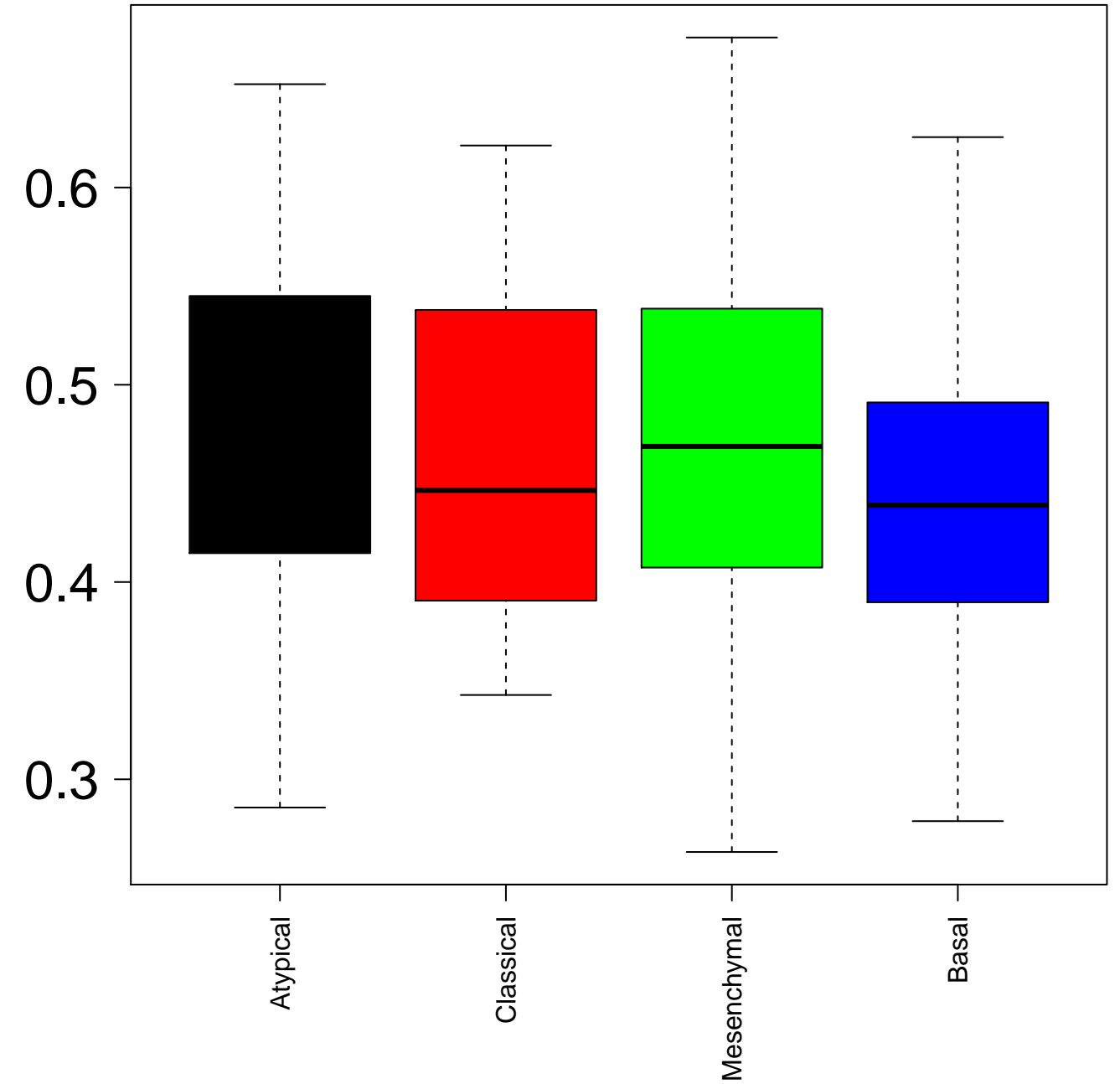
# Fraction of red genes (logFC)



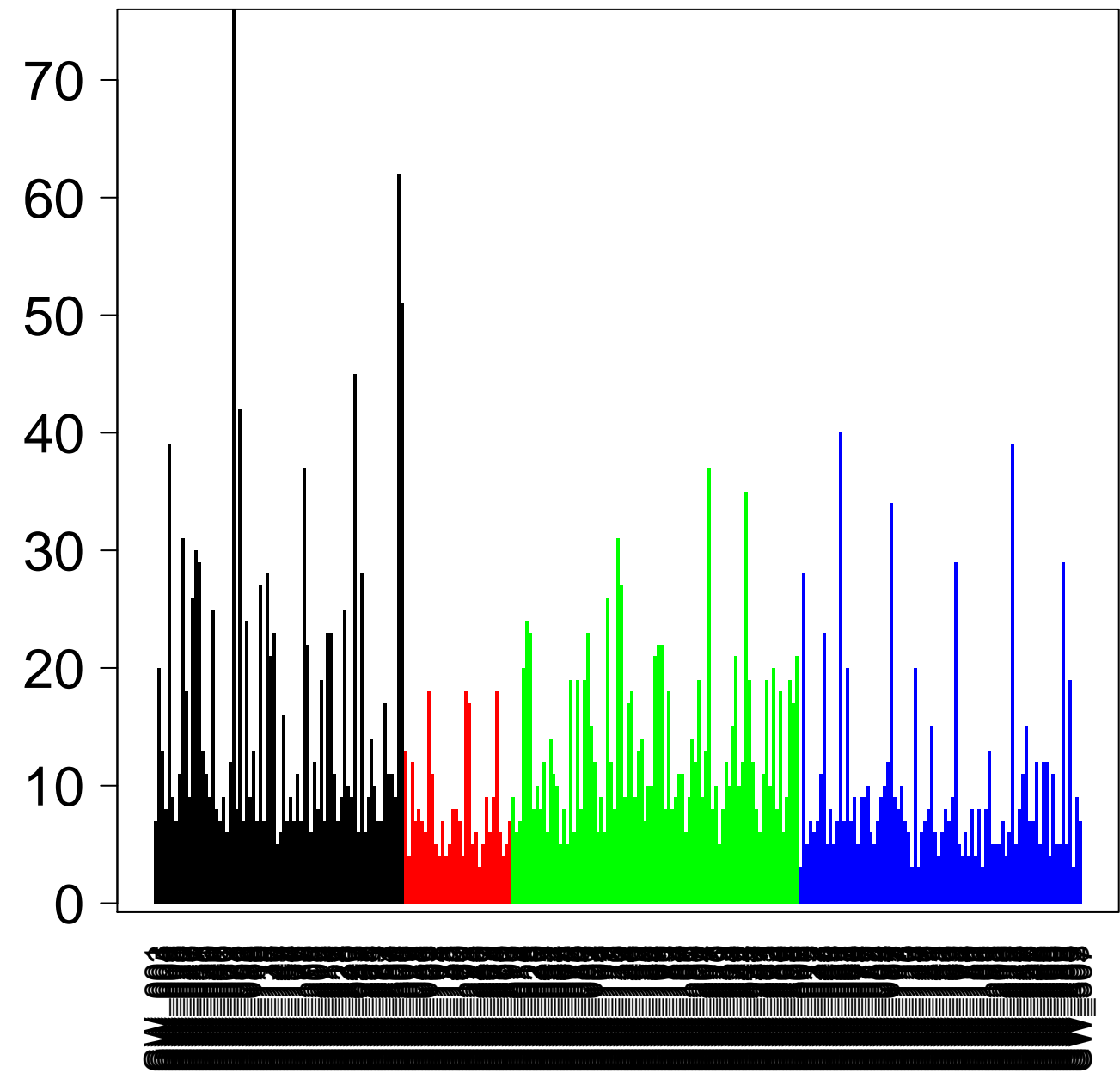
### Fraction of red genes (loglogFC)



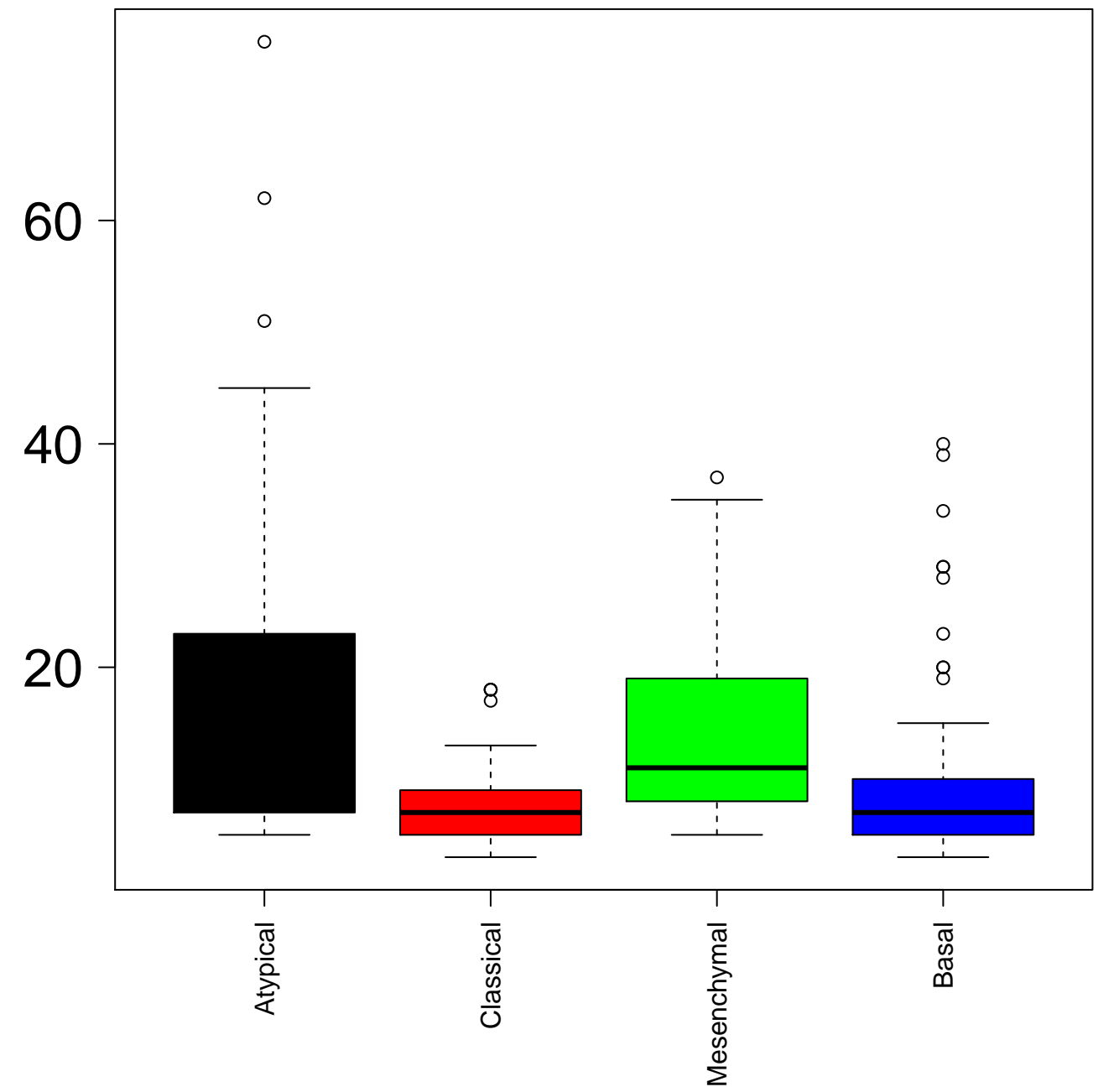
### Fraction of red genes (loglogFC)



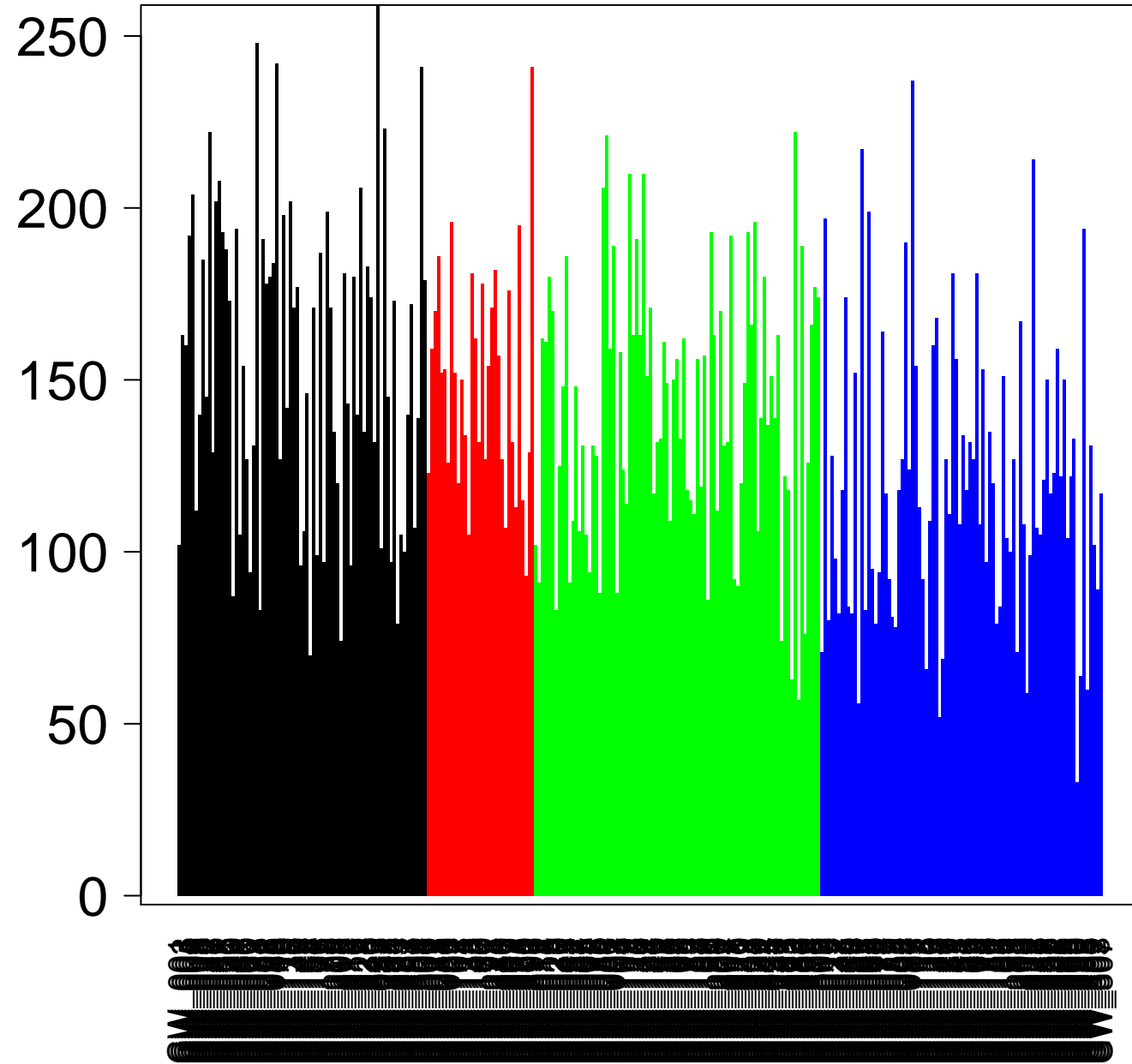
# Length of borderline (logFC)



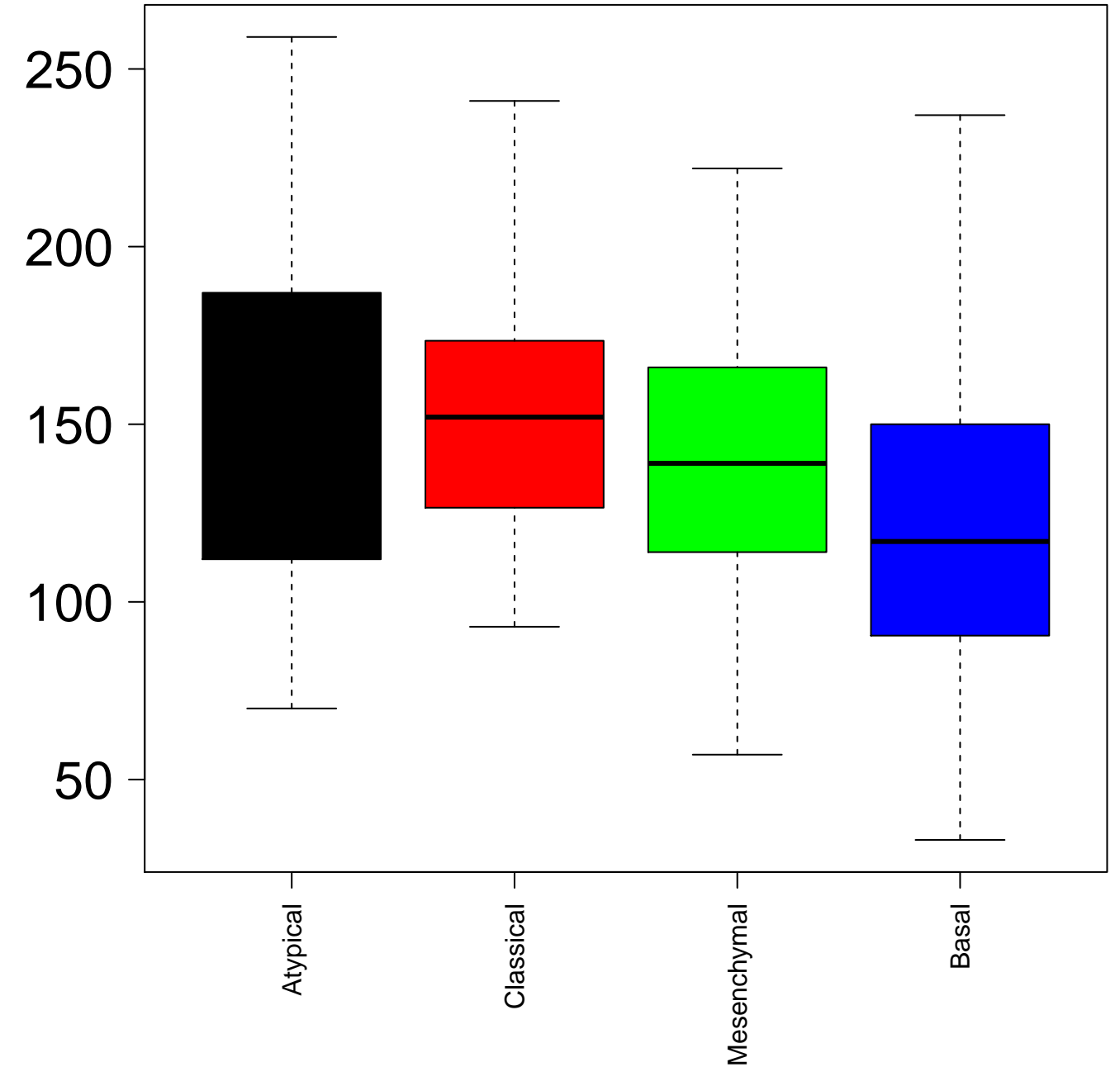
# Length of borderline (logFC)



# Length of borderline (loglogFC)

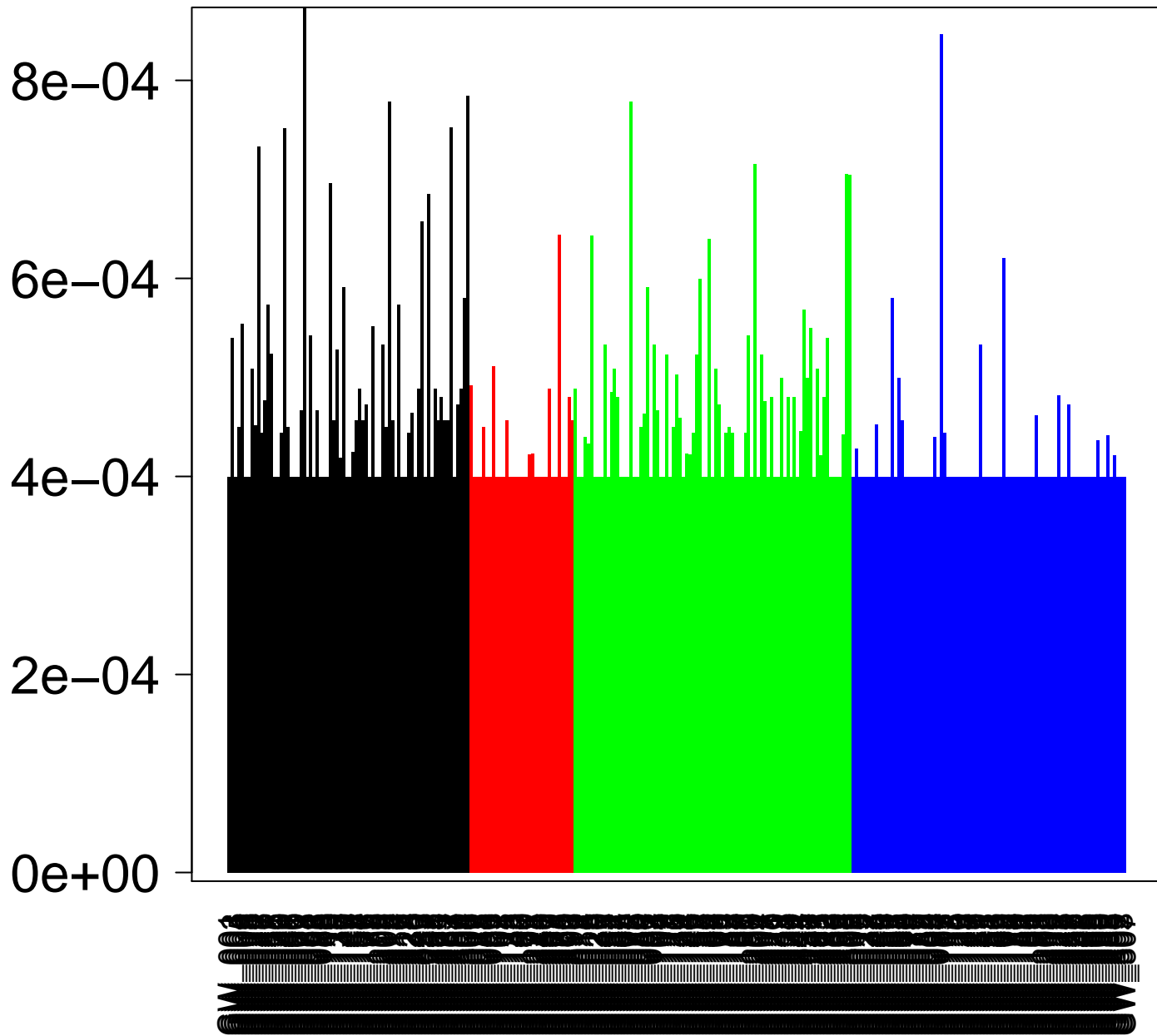


# Length of borderline (loglogFC)

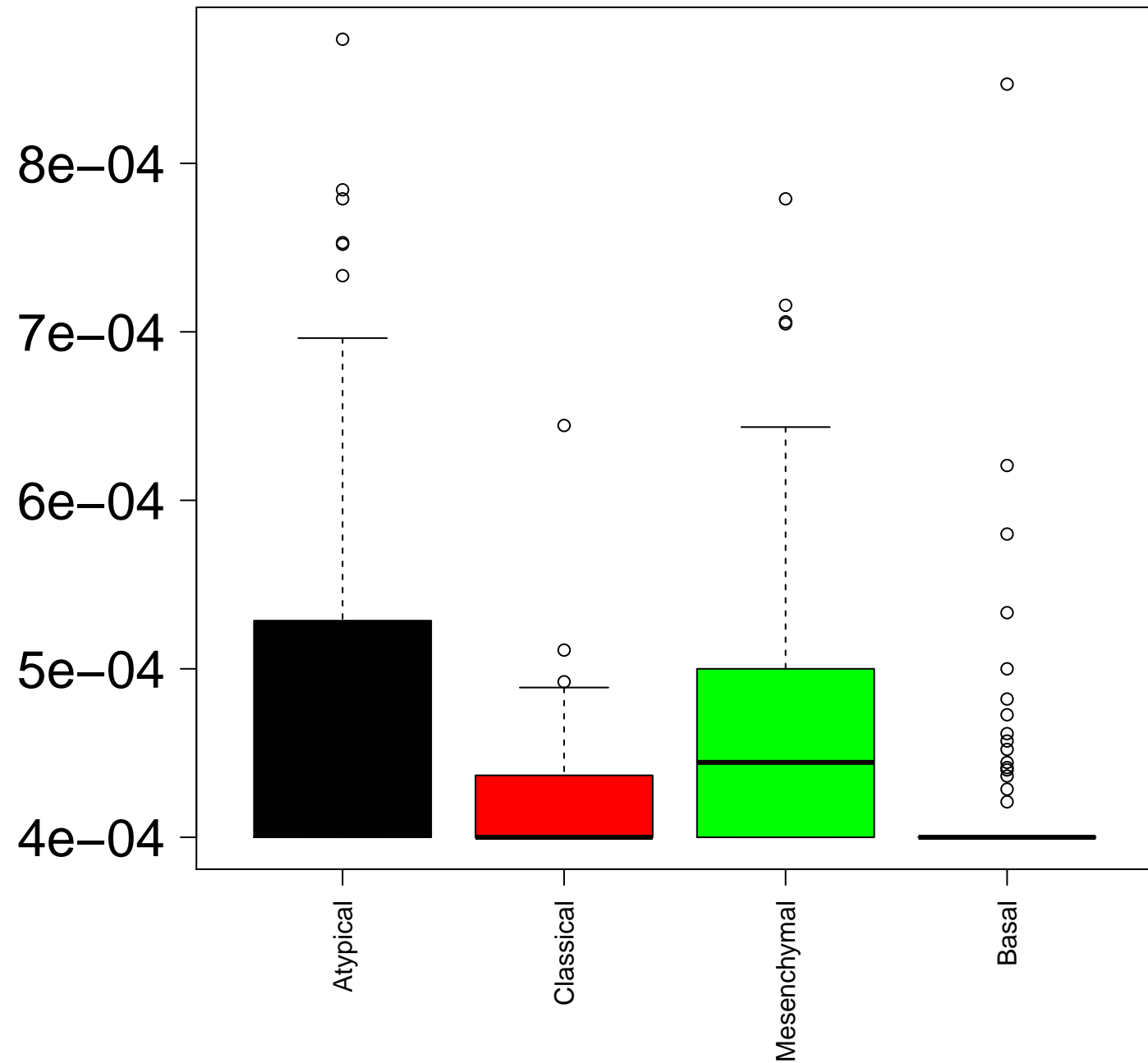




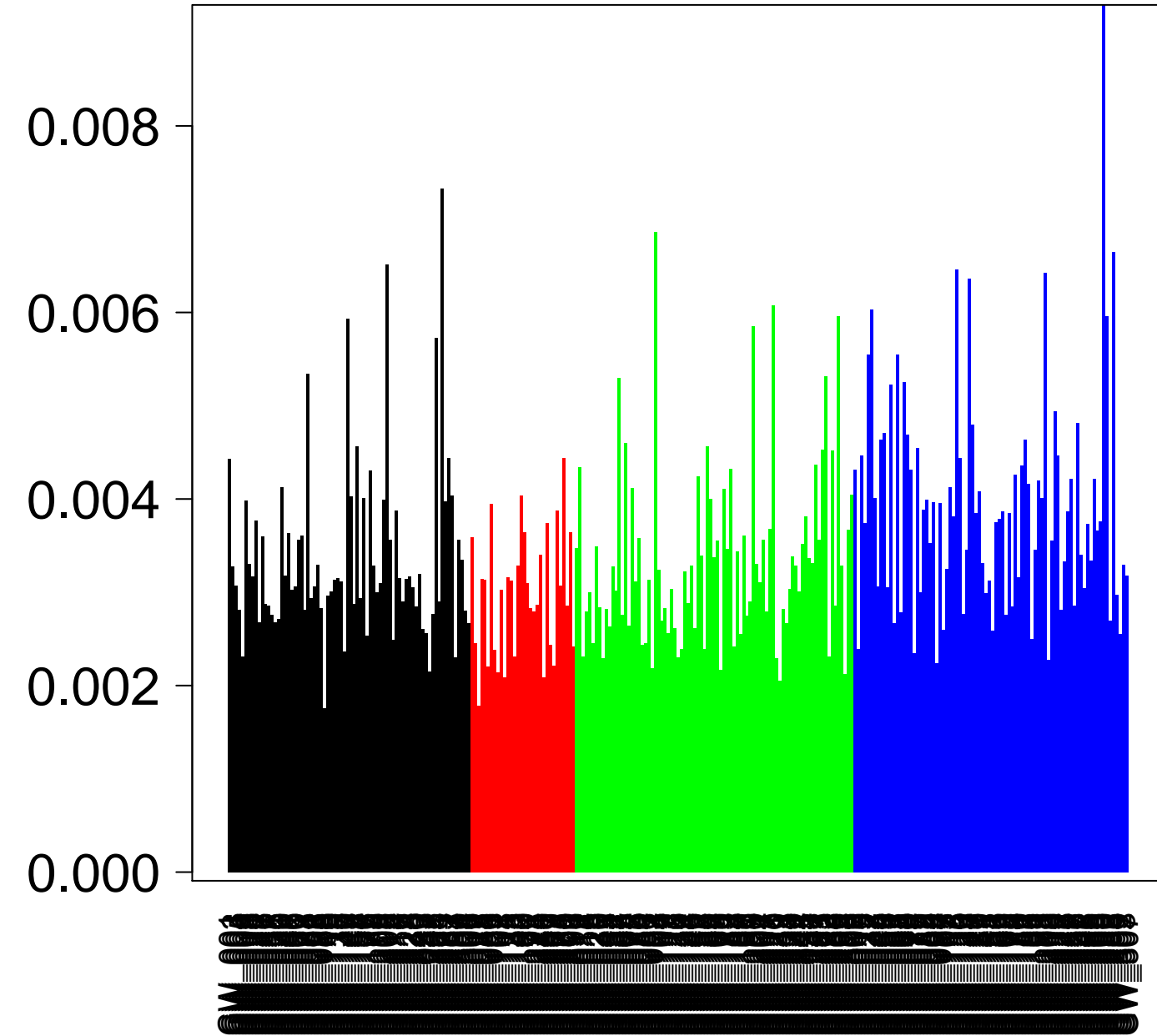
### Compactness of spots (logFC)



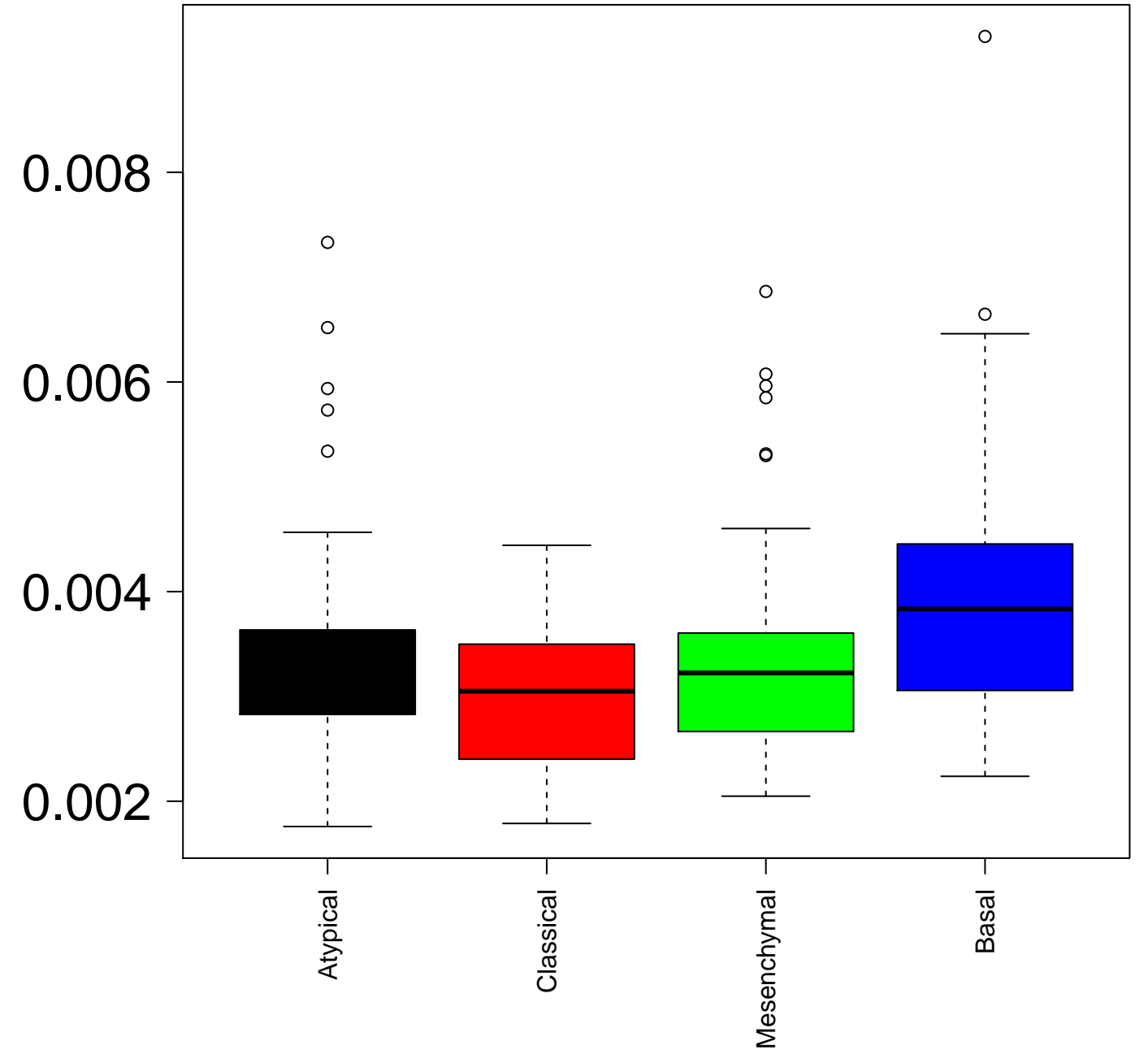
### Compactness of spots (logFC)



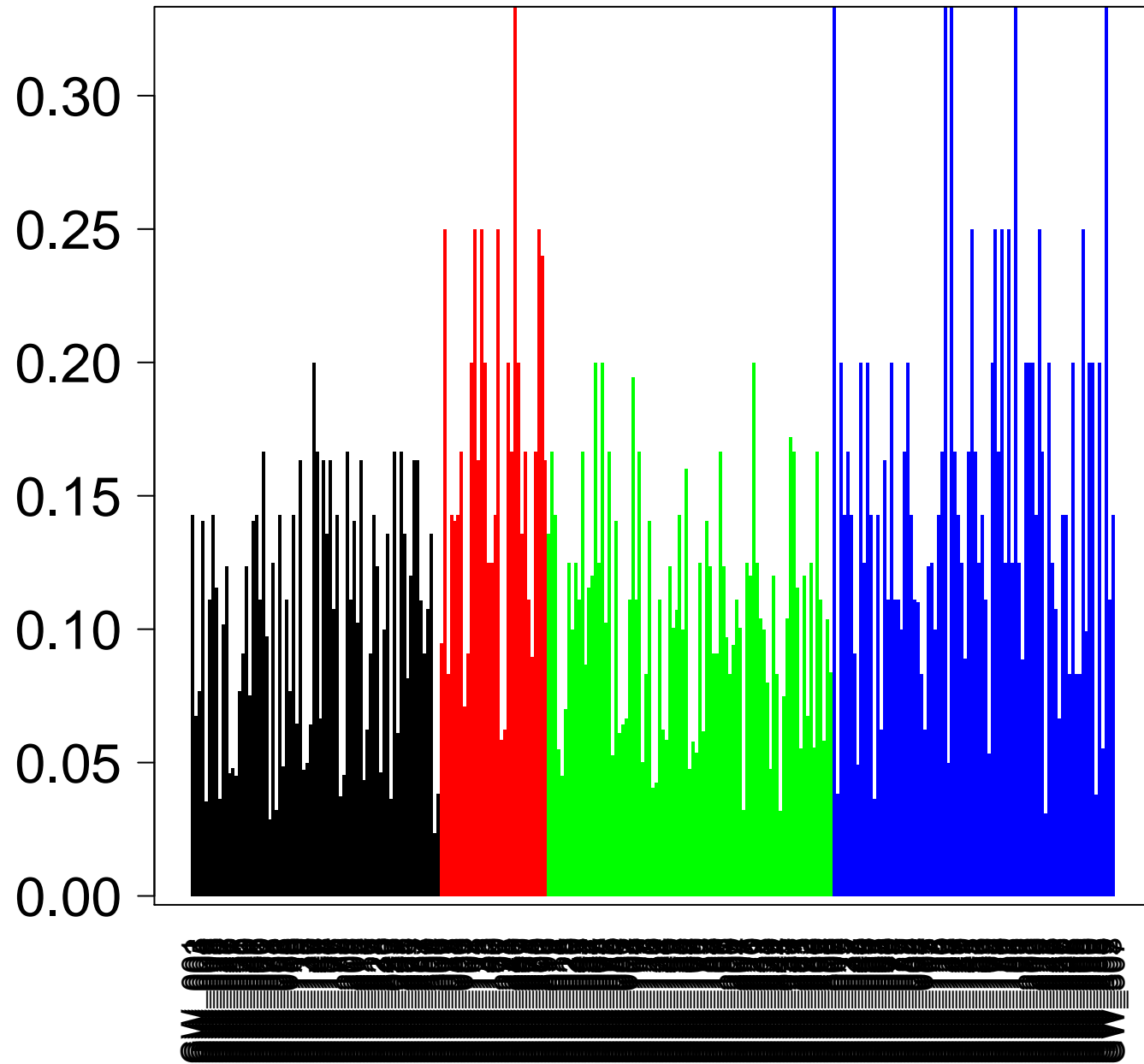
# Compactness of spots (loglogFC)



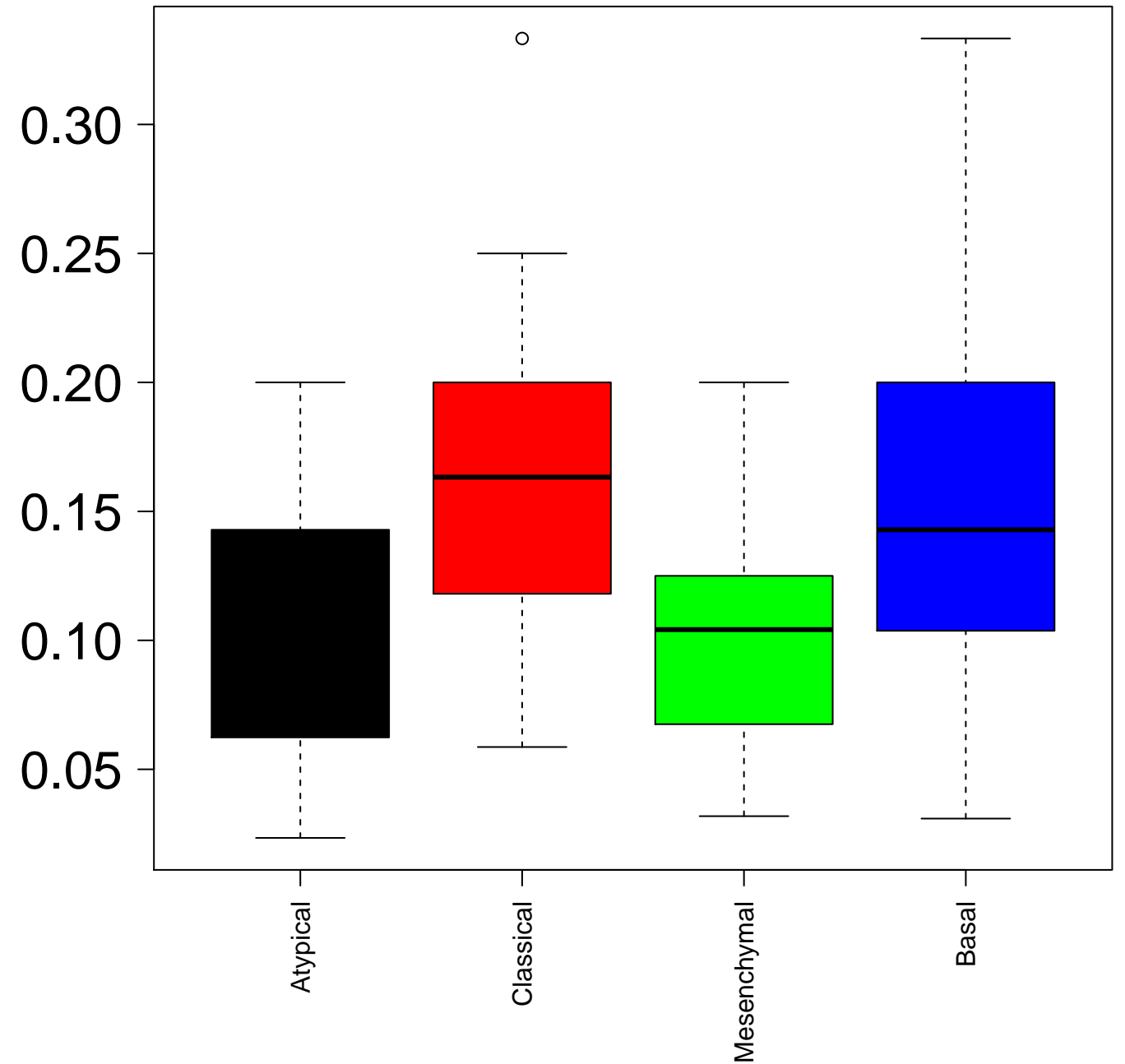
# Compactness of spots (loglogFC)



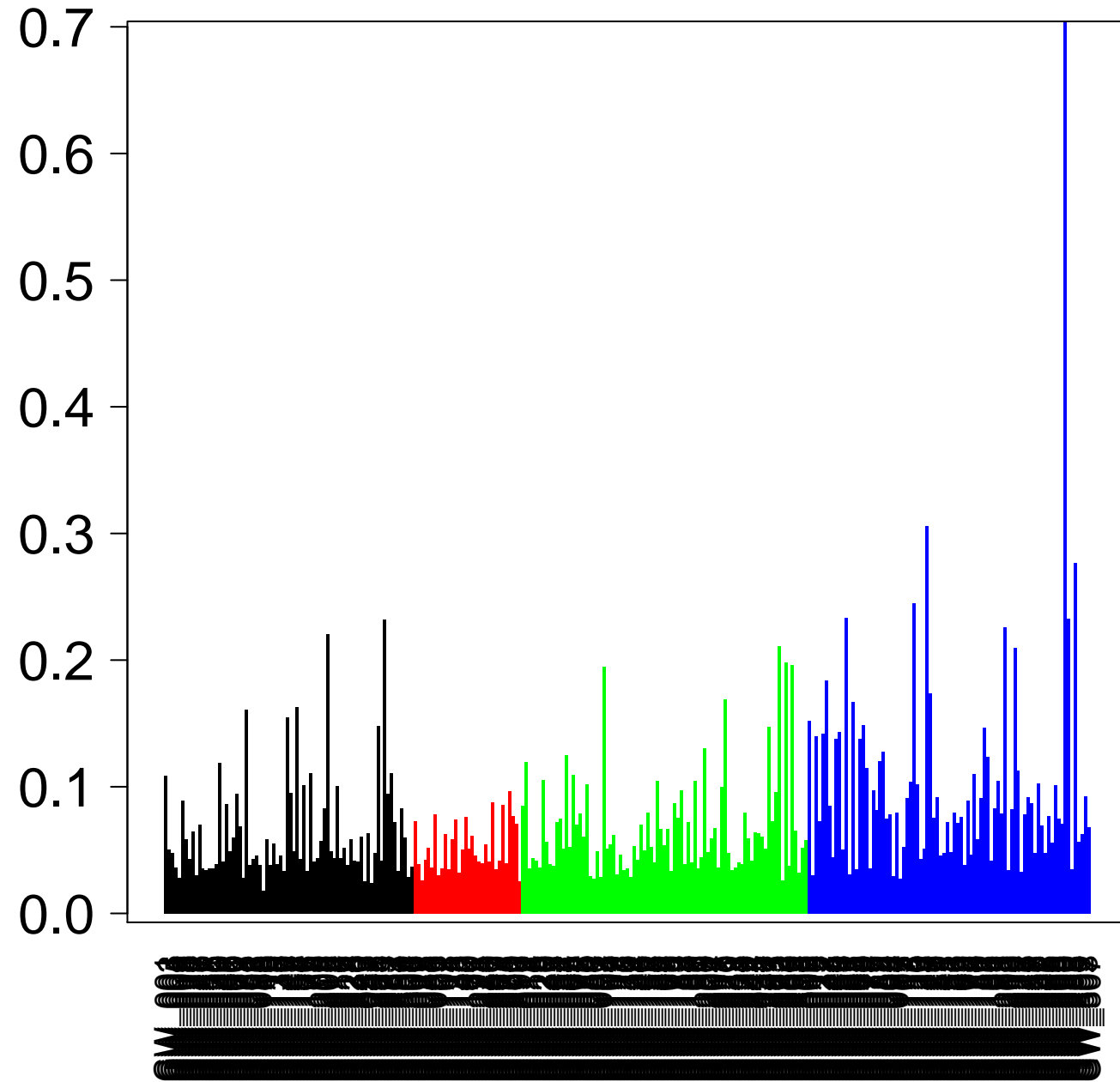
# Shape of spots (logFC)



# Shape of spots (logFC)



# Shape of spots (loglogFC)



# Shape of spots (loglogFC)

