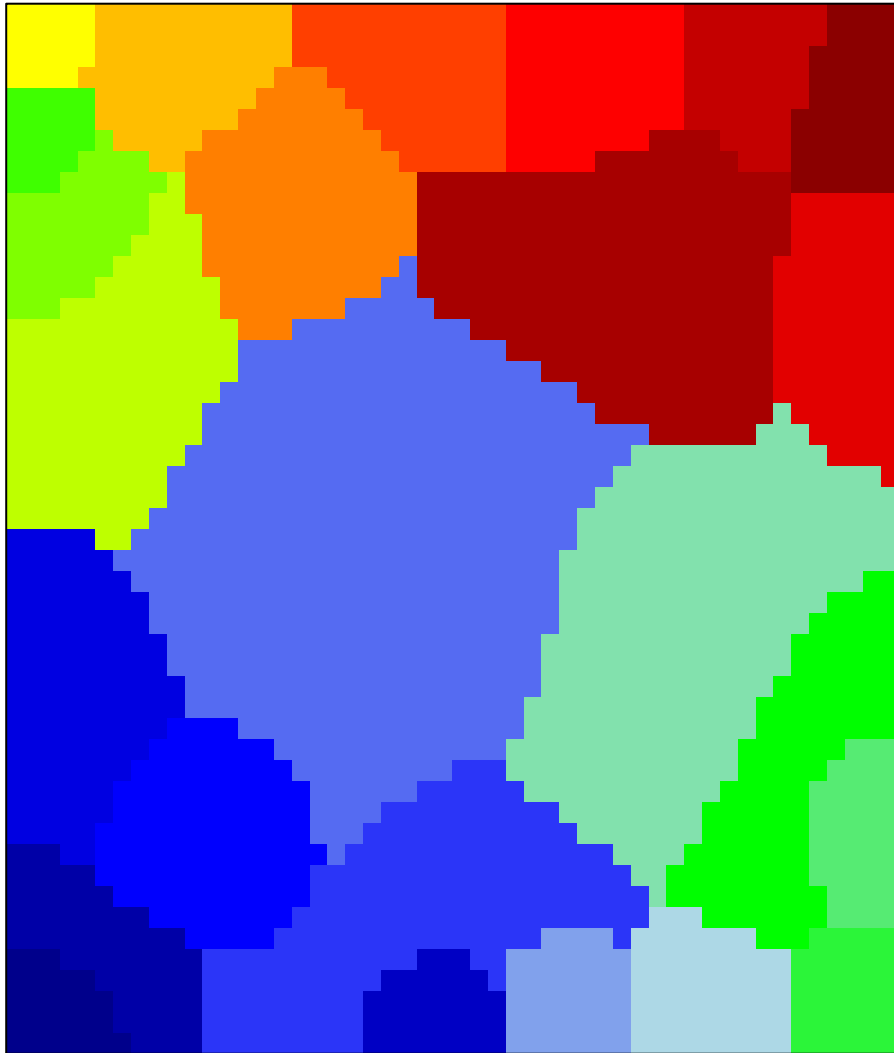
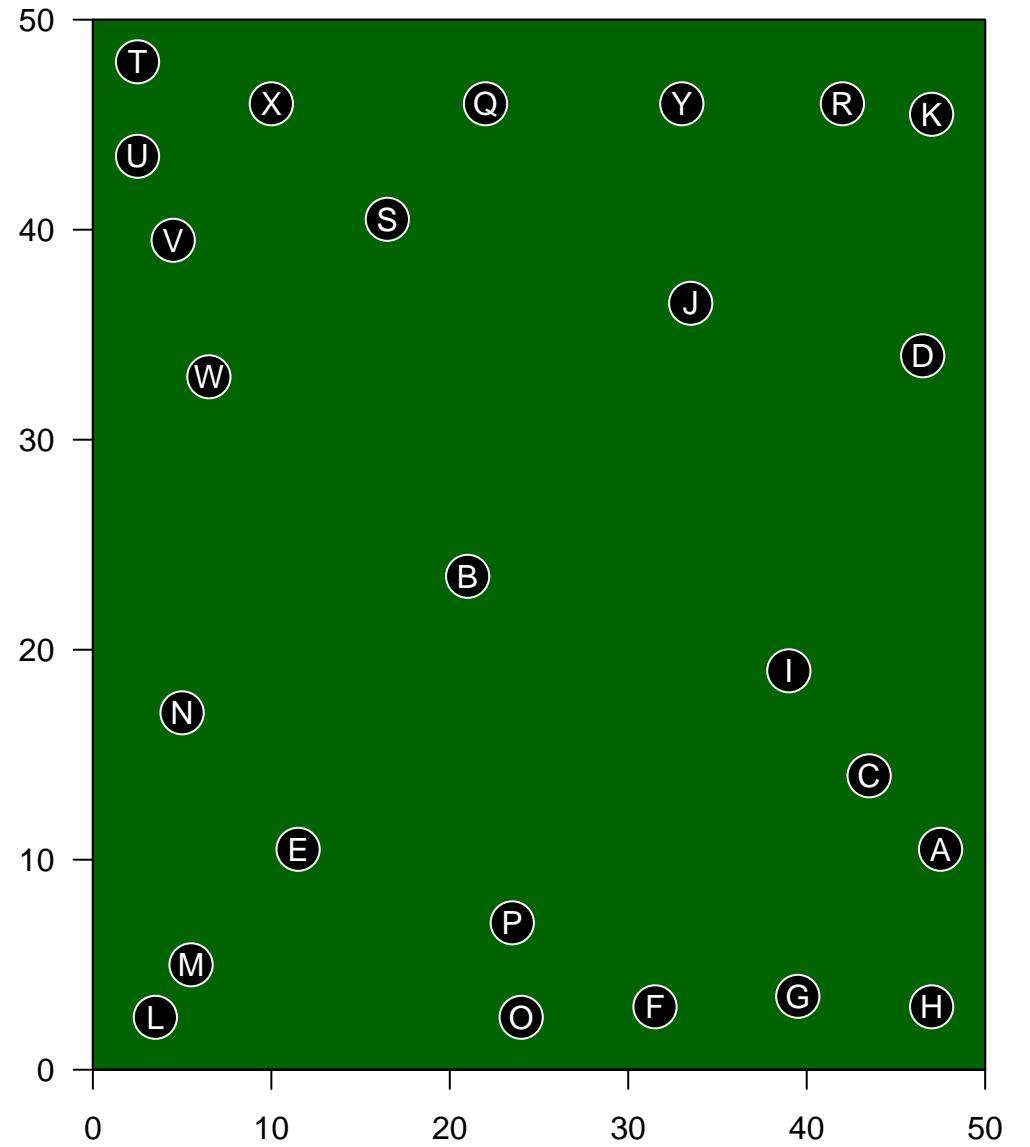
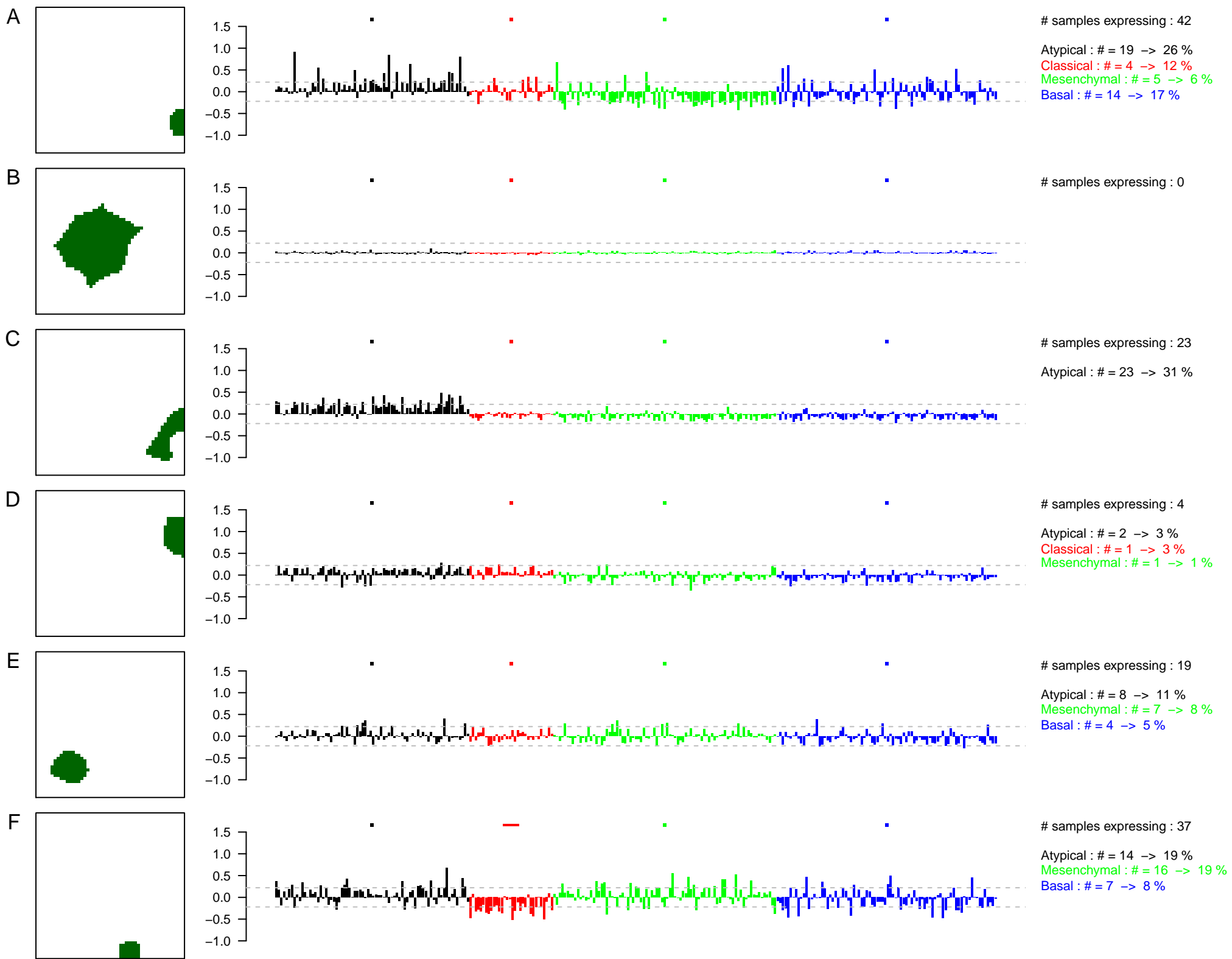


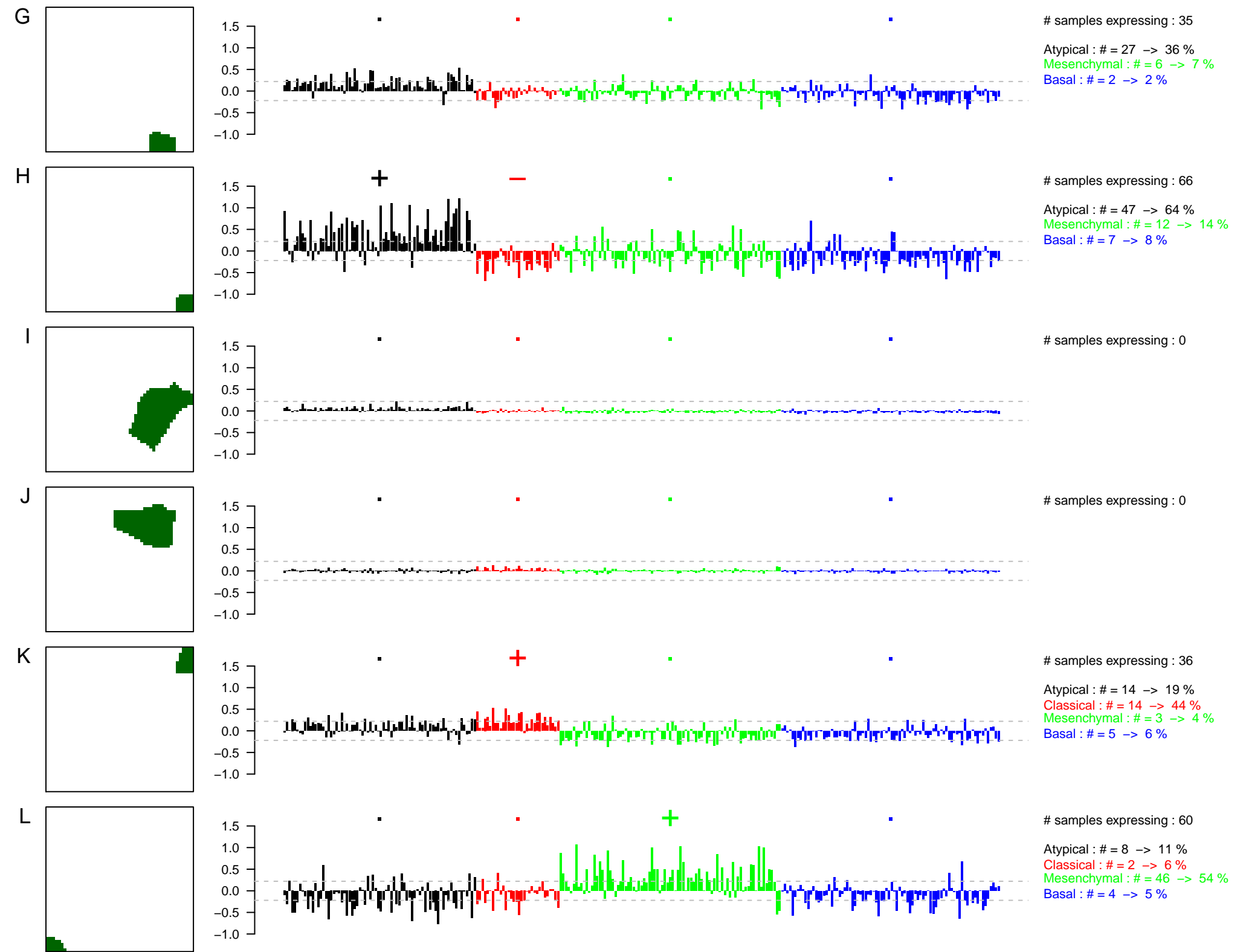
K-Means Cluster

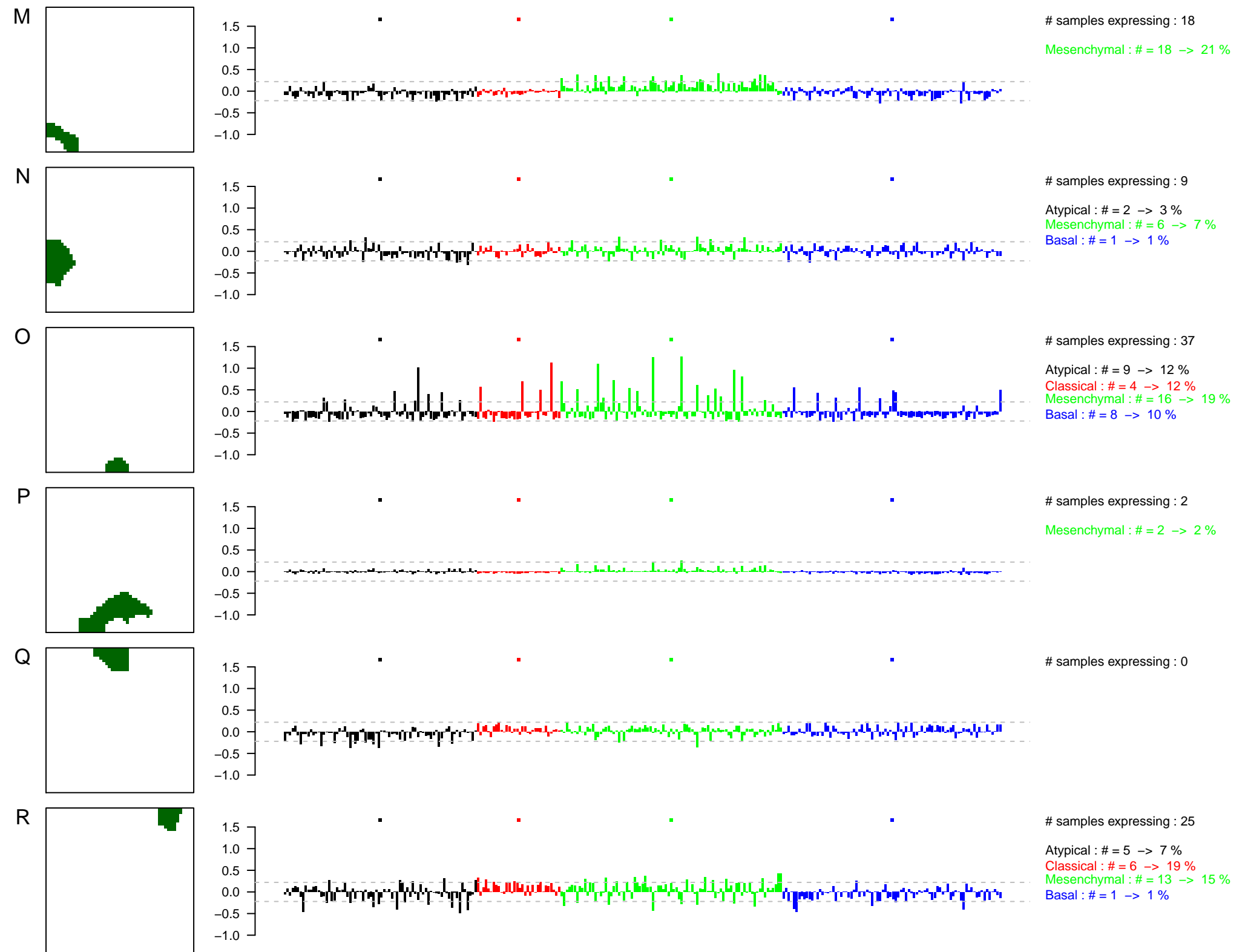


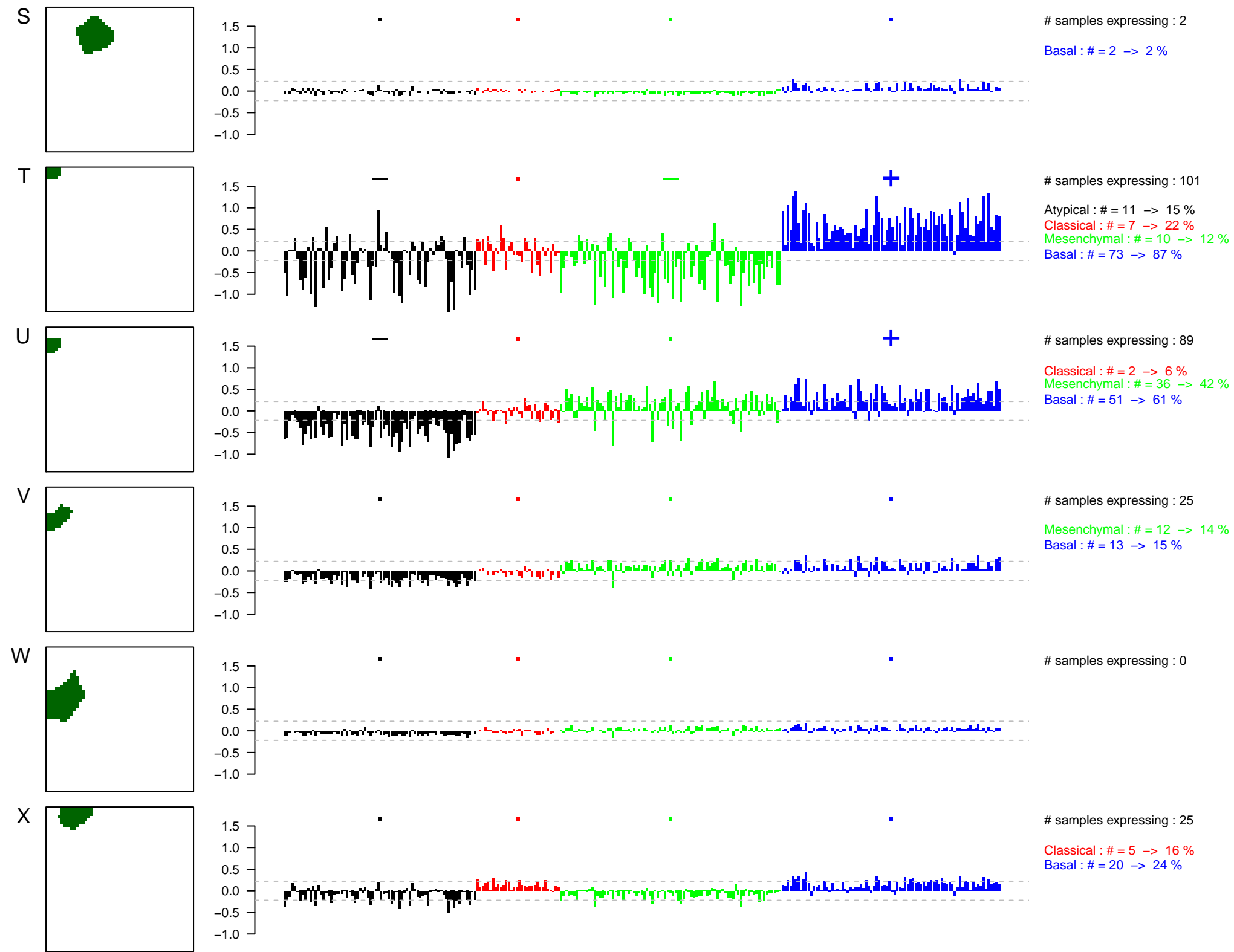
Spots



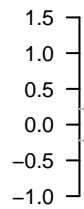
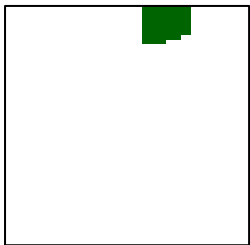








Y



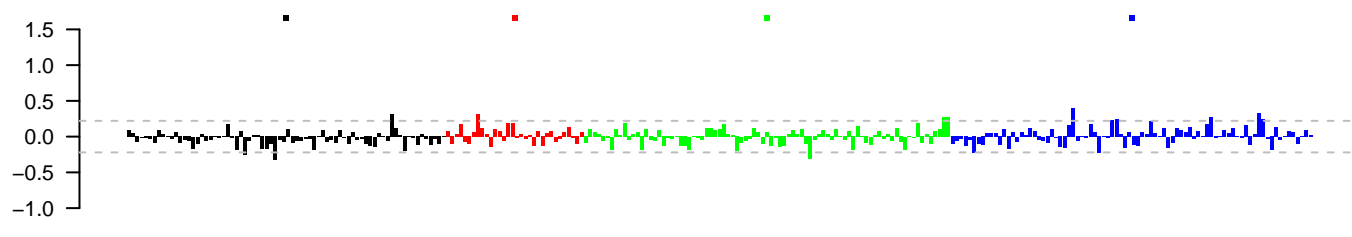
samples expressing : 10

Atypical : # = 1 -> 1 %

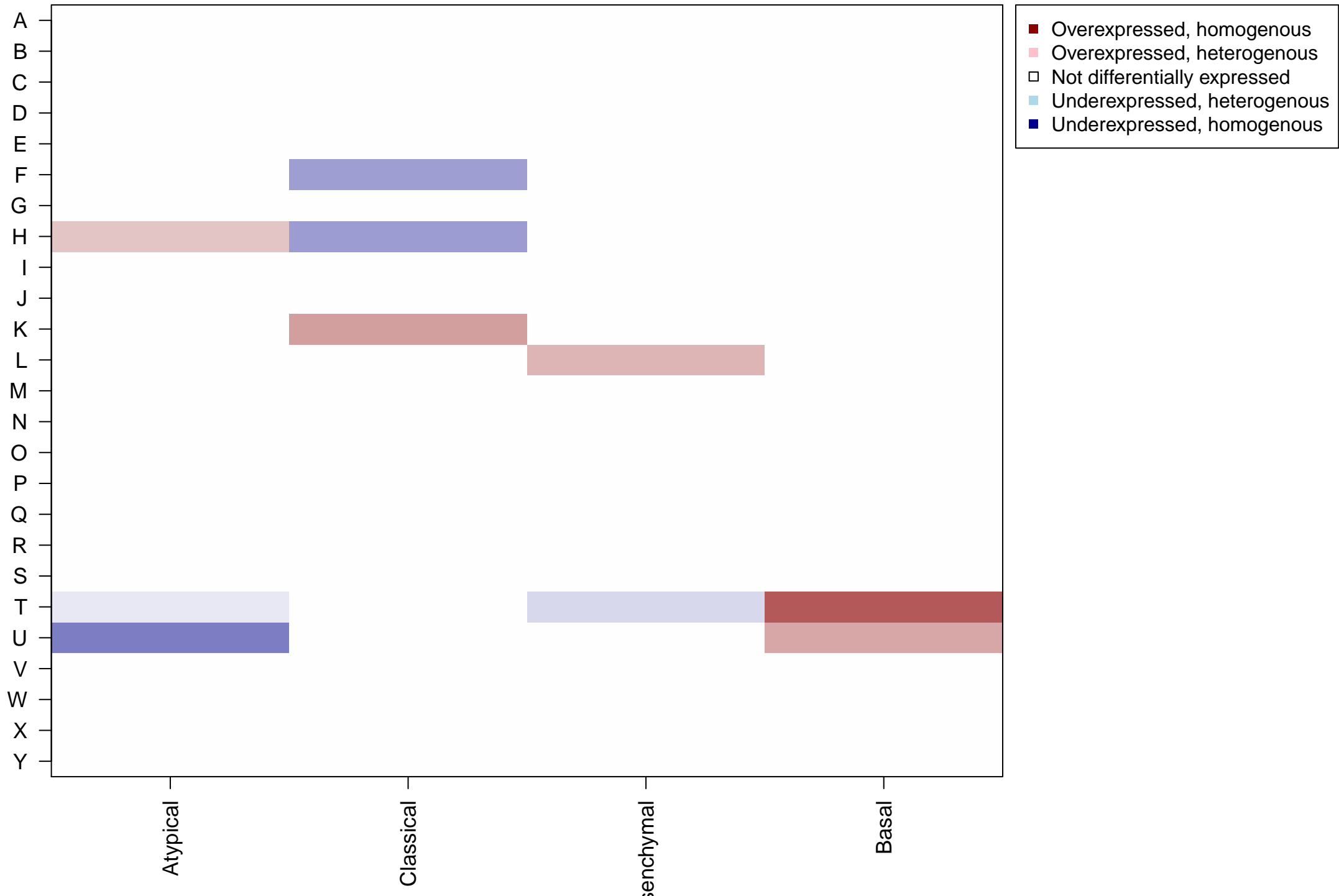
Classical : # = 1 -> 3 %

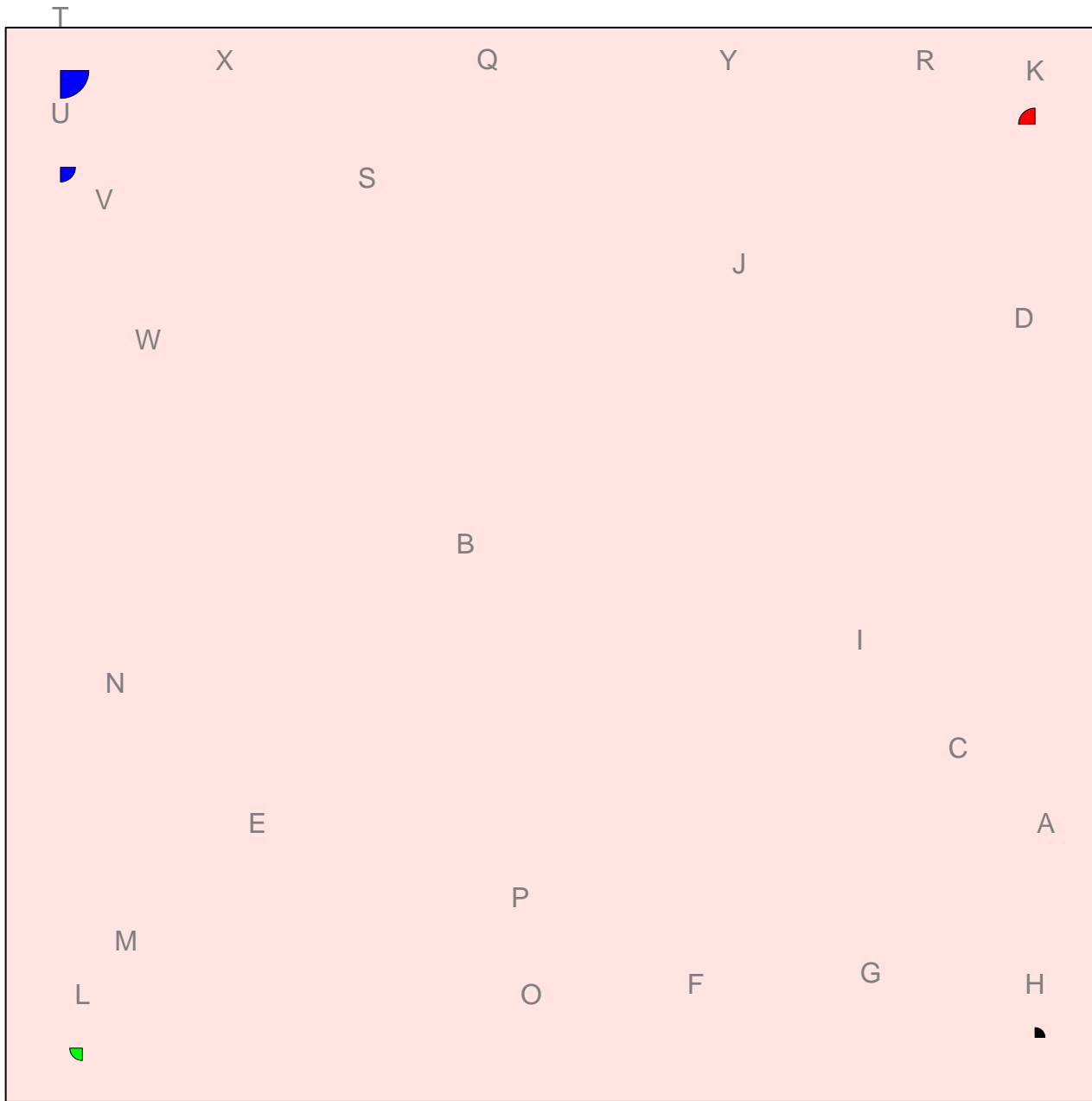
Mesenchymal : # = 2 -> 2 %

Basal : # = 6 -> 7 %

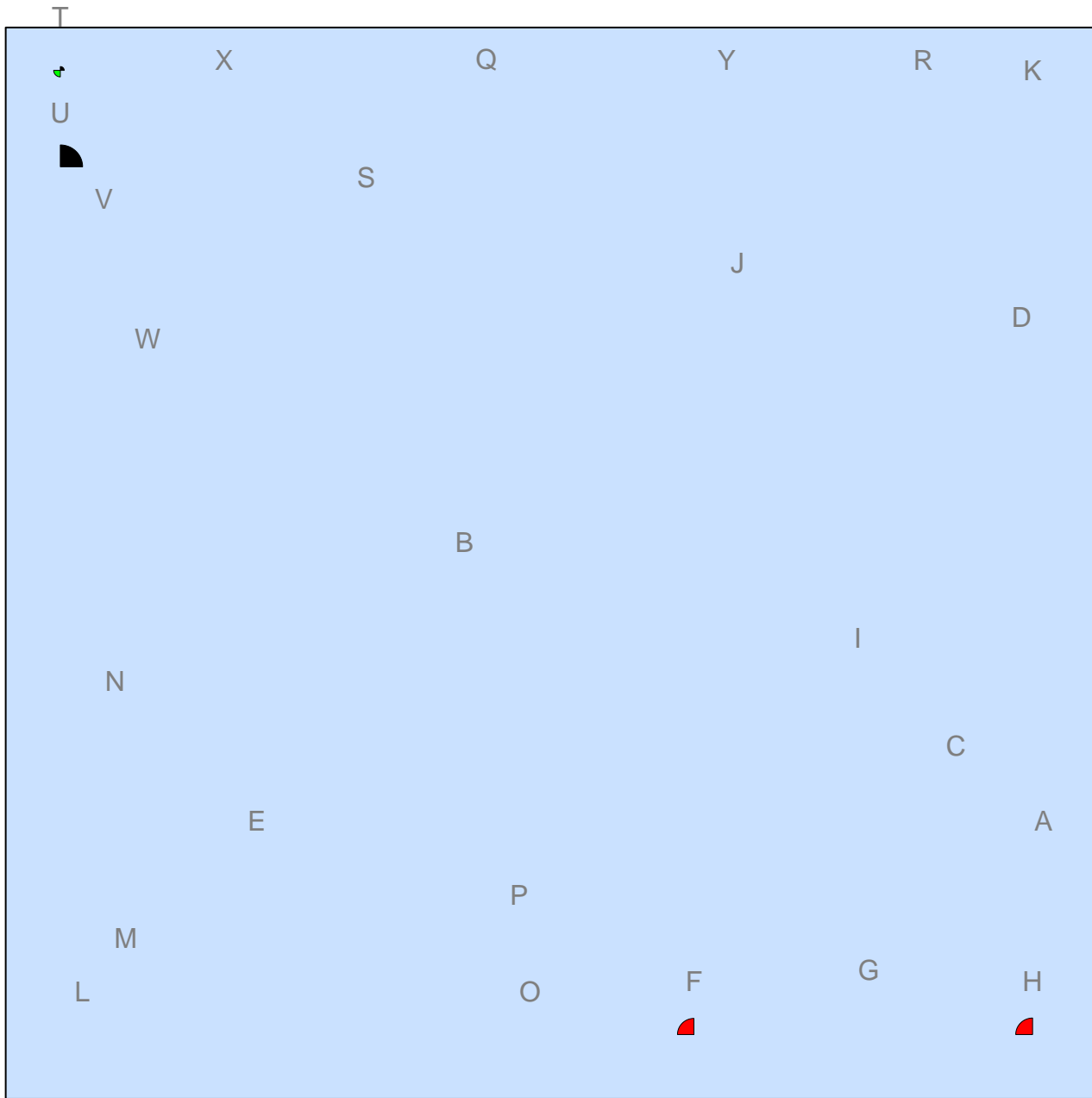


Spot homogeneity map



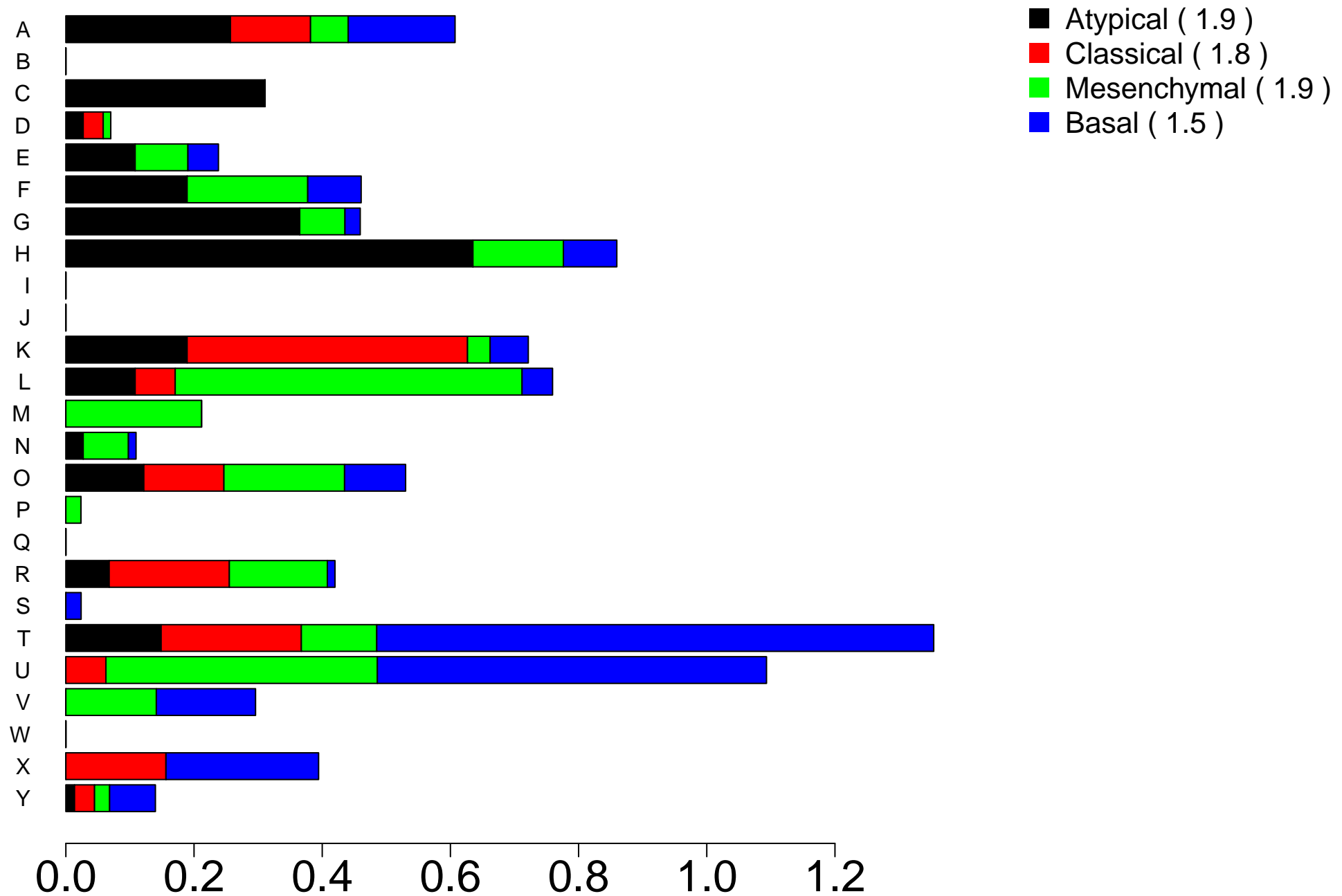


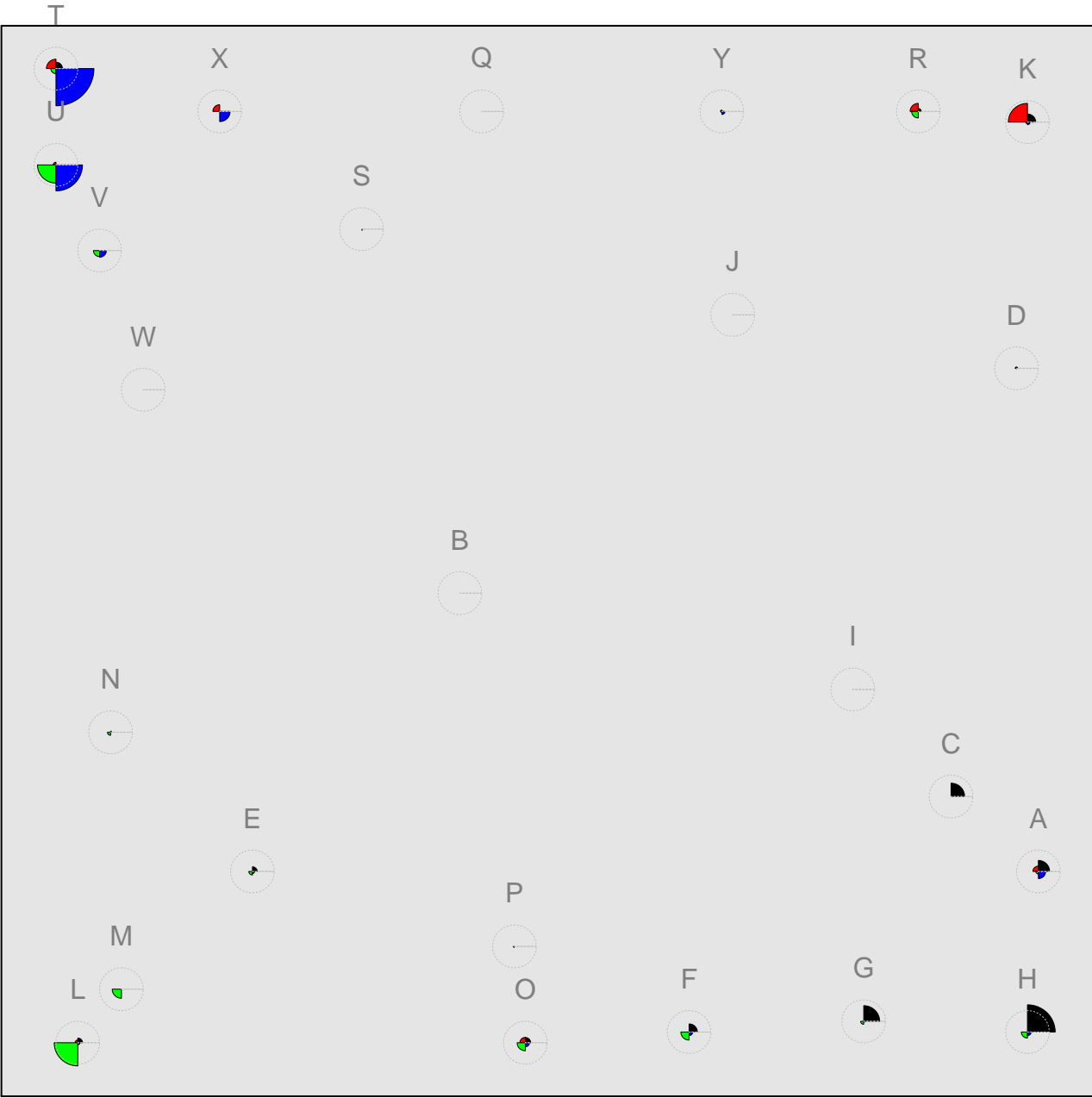
- Atypical
- Classical
- Mesenchymal
- Basal



Group Association

< #spots >

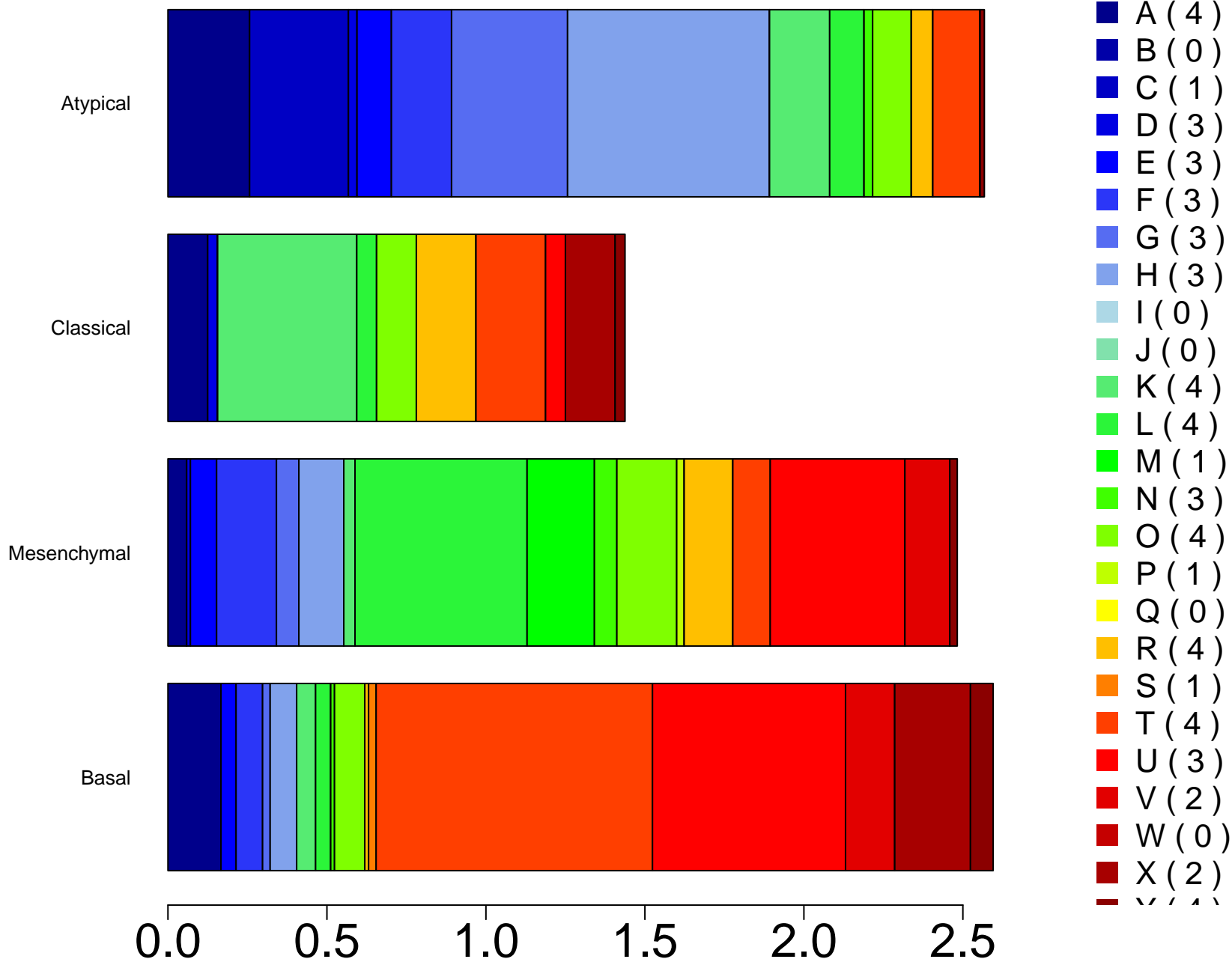


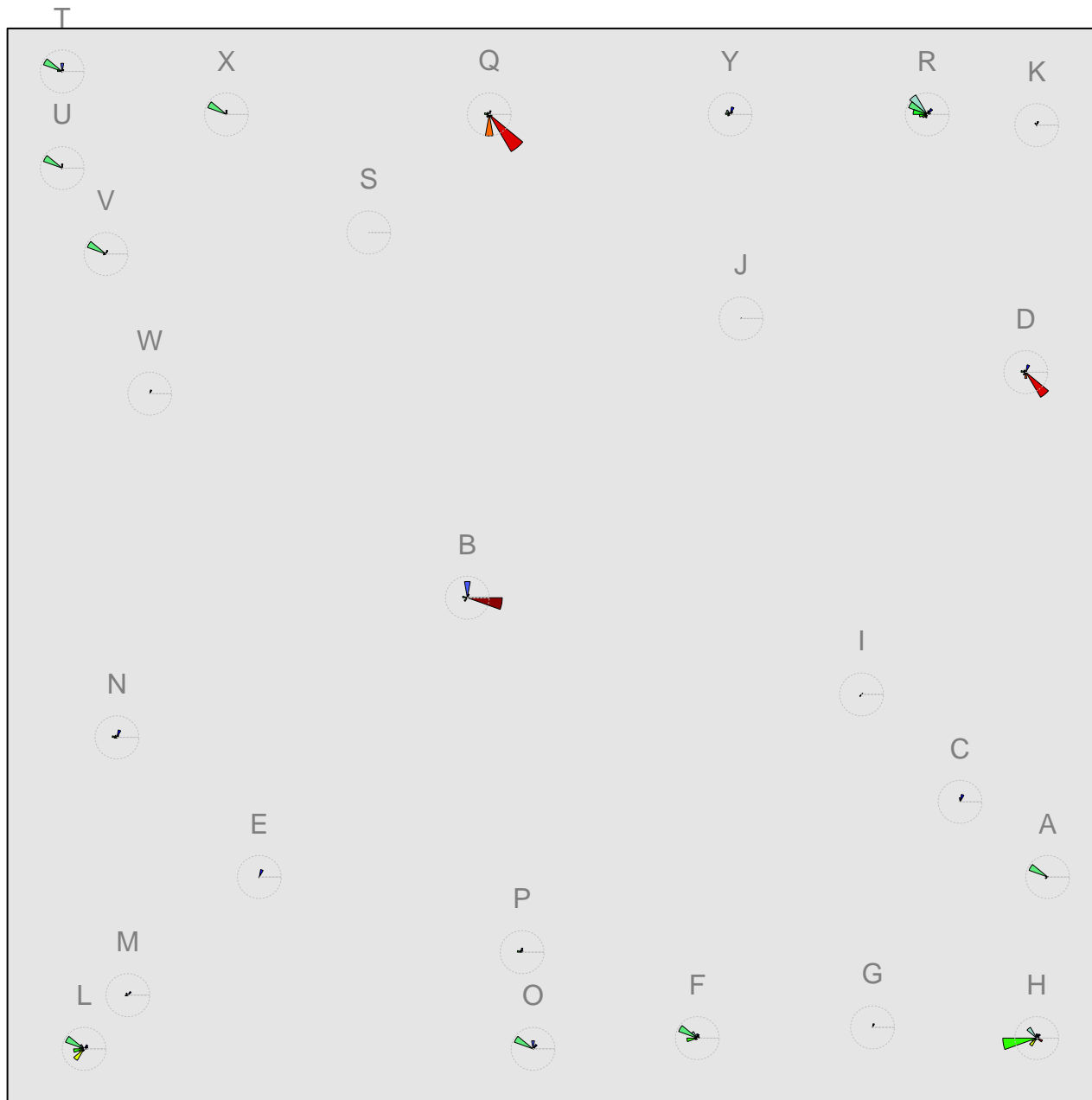


- Atypical
- Classical
- Mesenchymal
- Basal

Spot Association

#groups





- BP
- MF
- CC
- Chr
- H.Tiss
- Pathw Act
- Cancer
- Disease
- TF
- Lymphoma
- MMML CGS
- Glio
- GSEA C2
- miRNA 3UTR
- miRNA target
- miRNA target starBase
- TF Tissue
- miRNA Disease