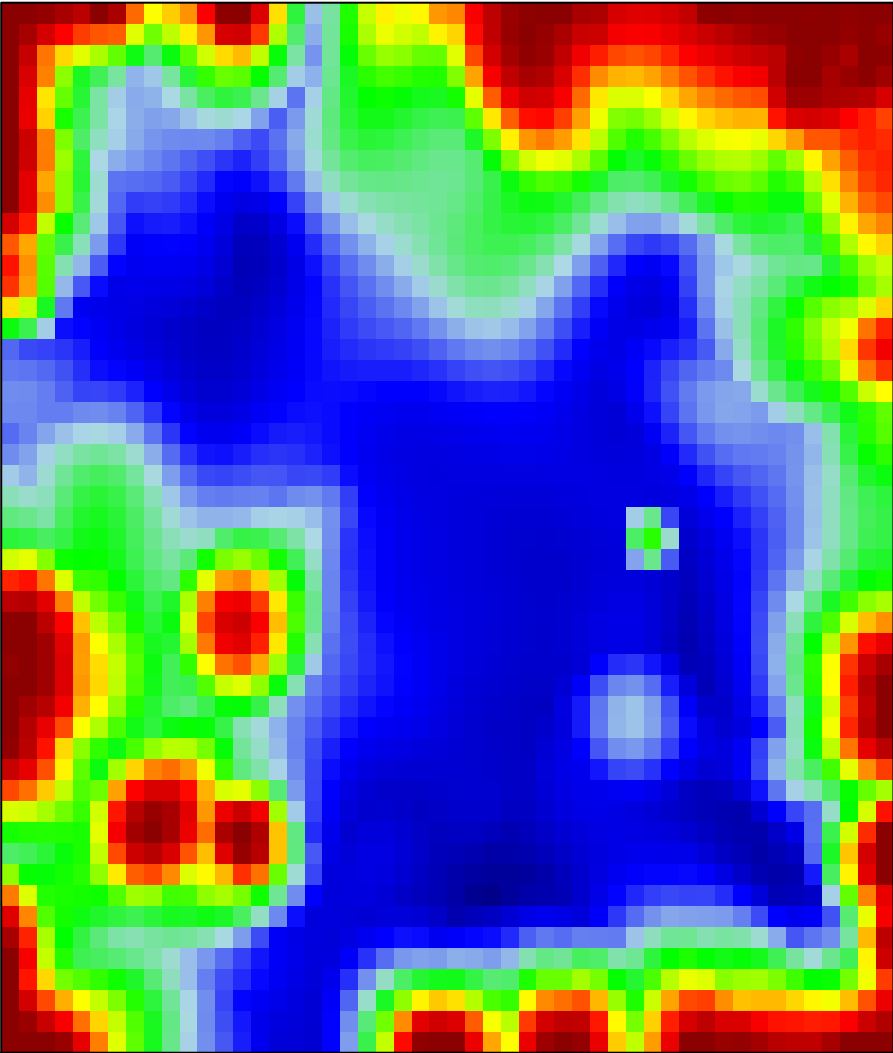
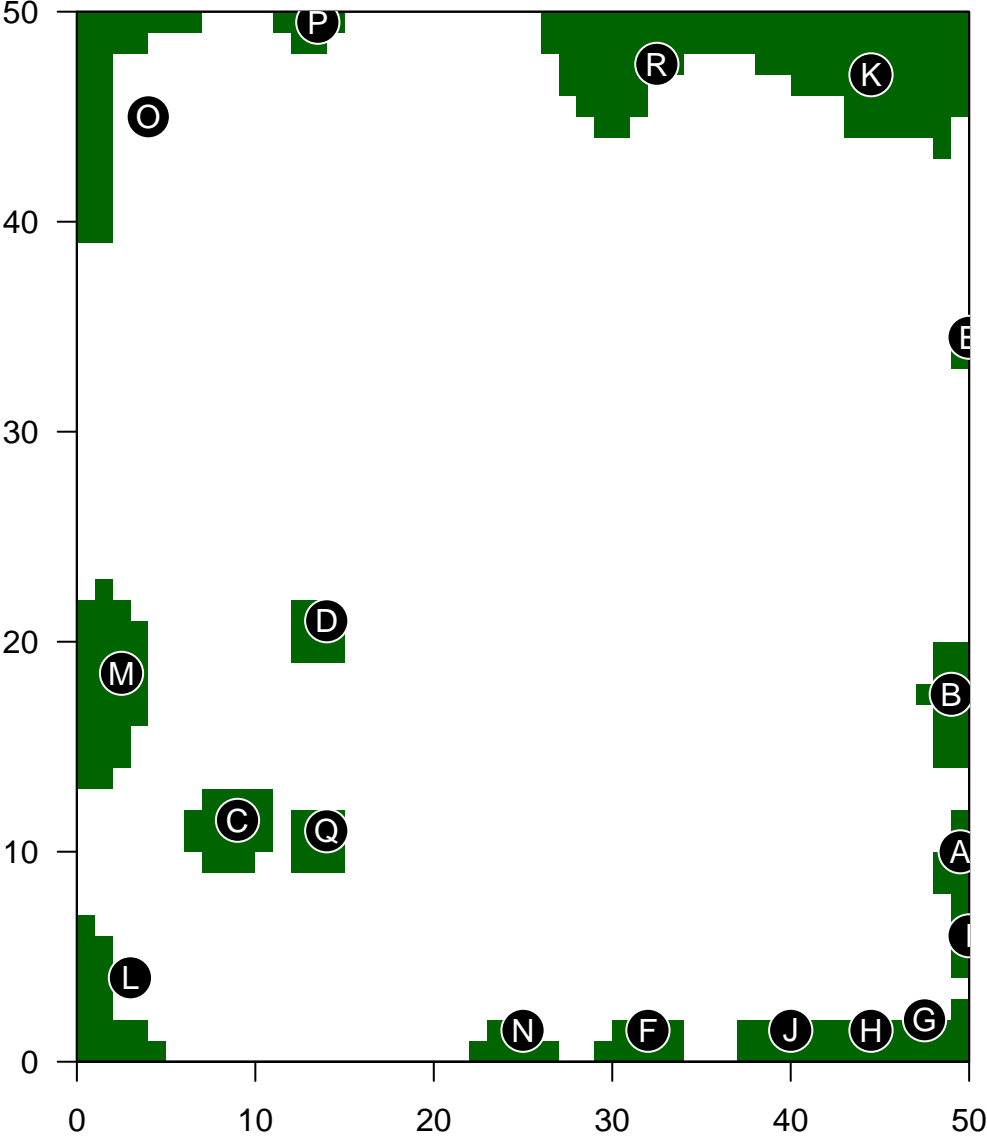
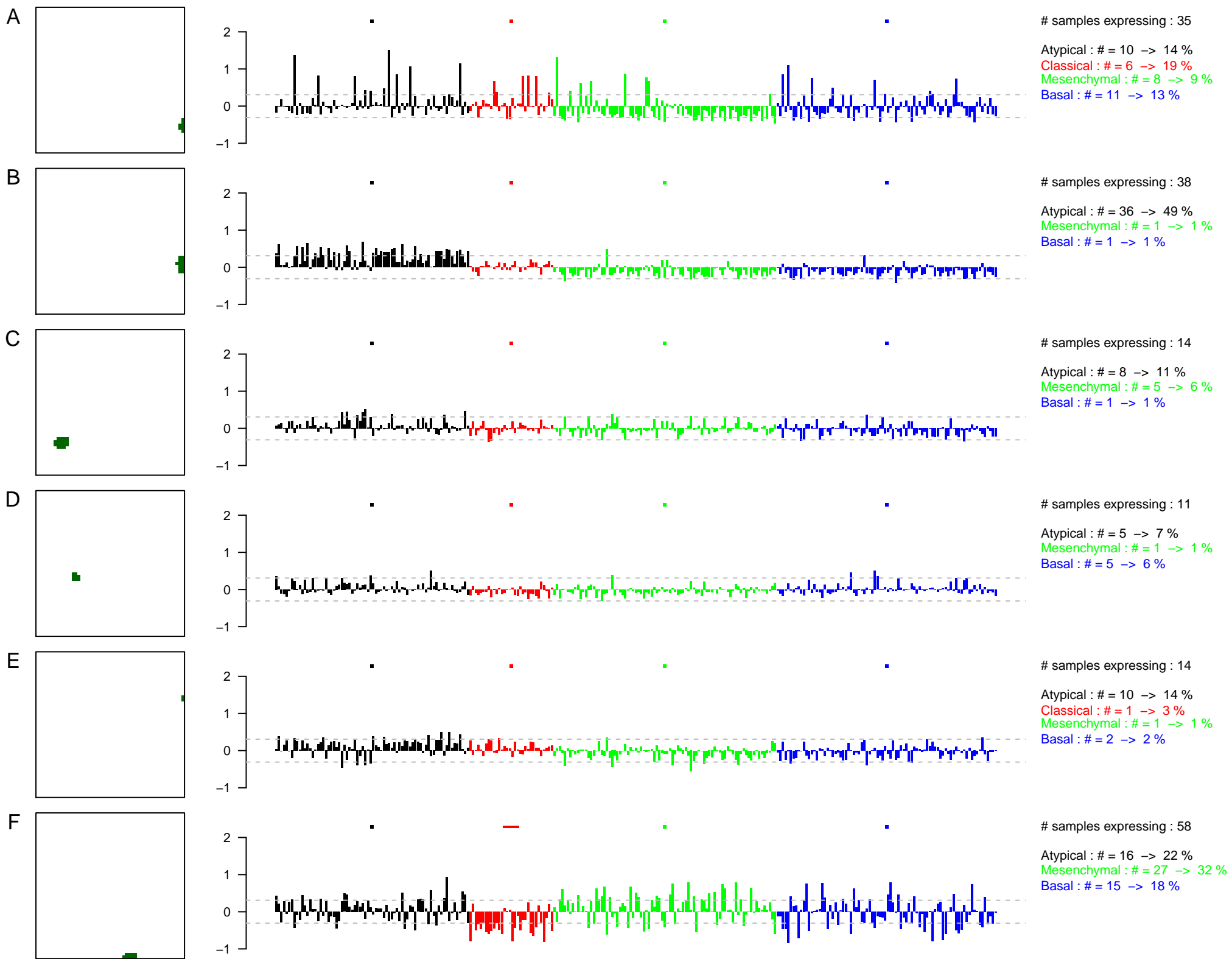


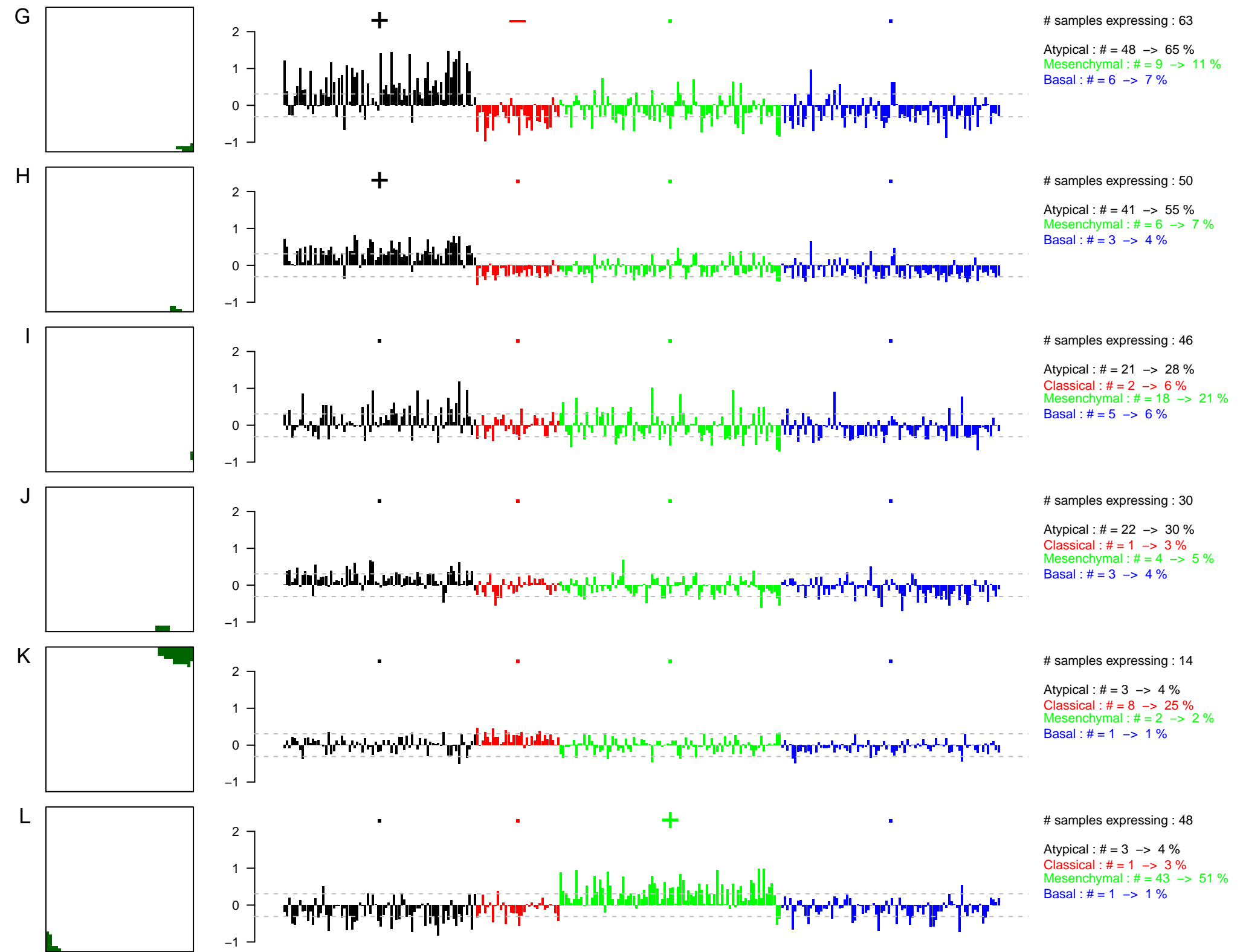
Sample-Overexpression

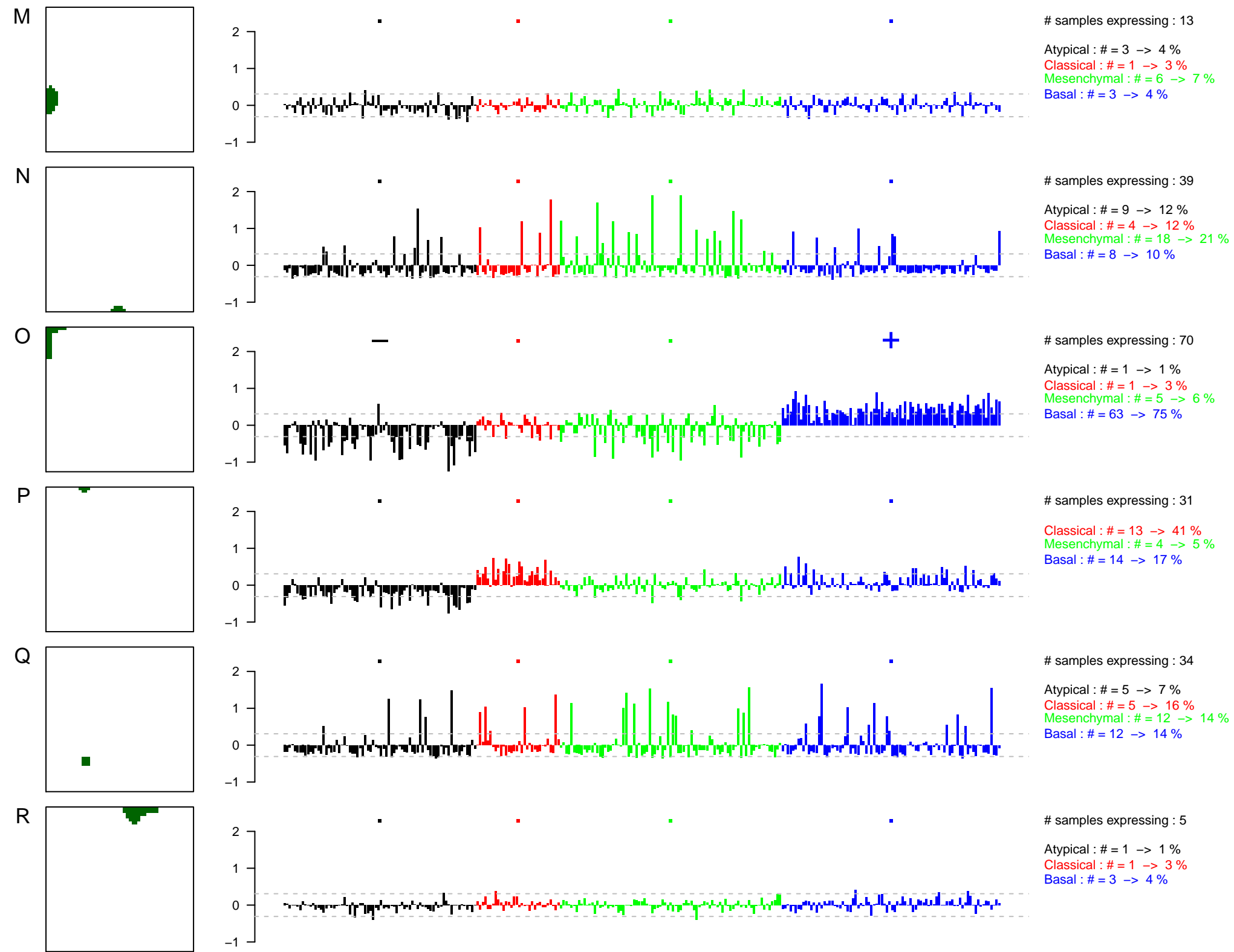


Spots

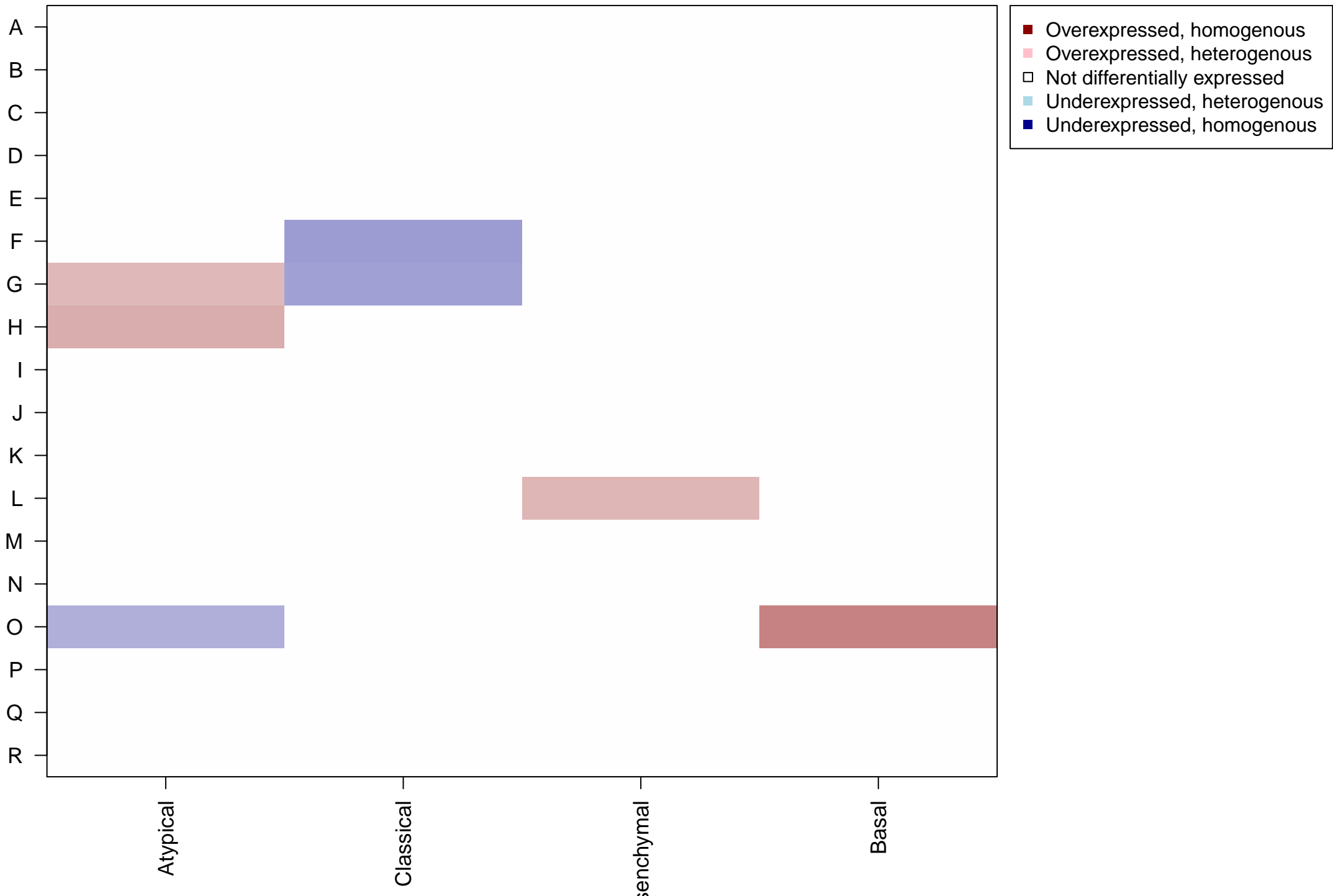


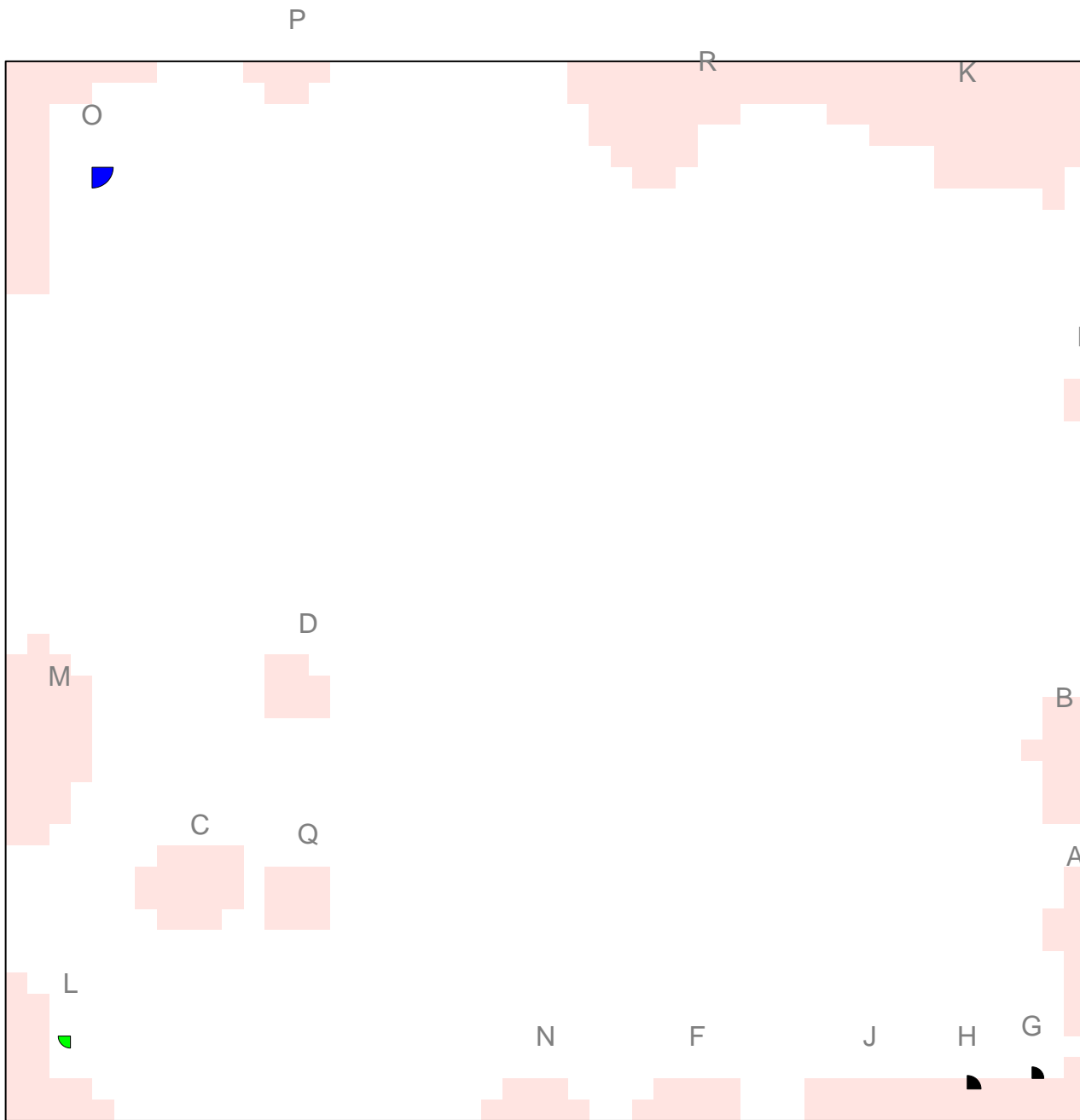




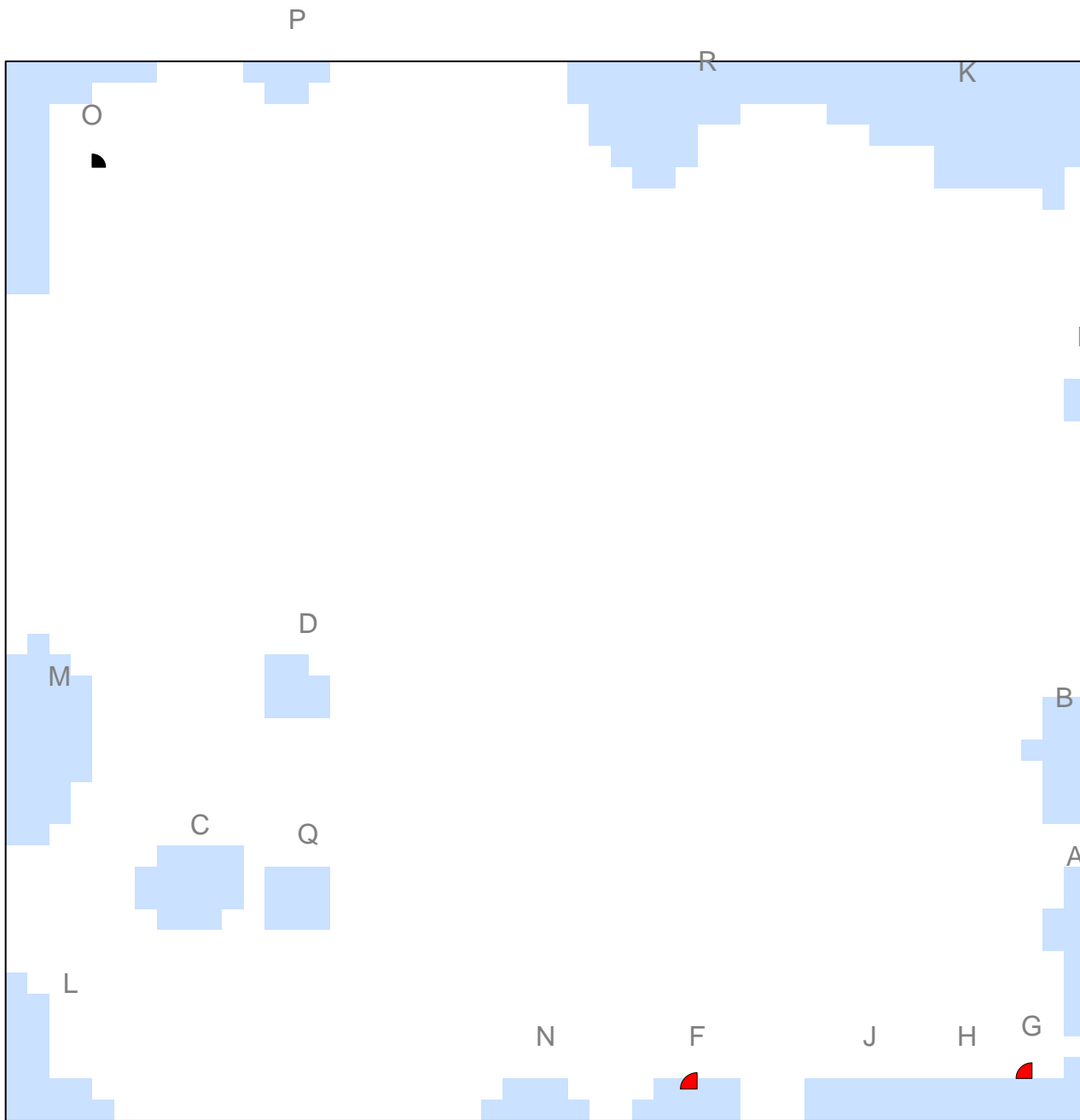


Spot homogeneity map





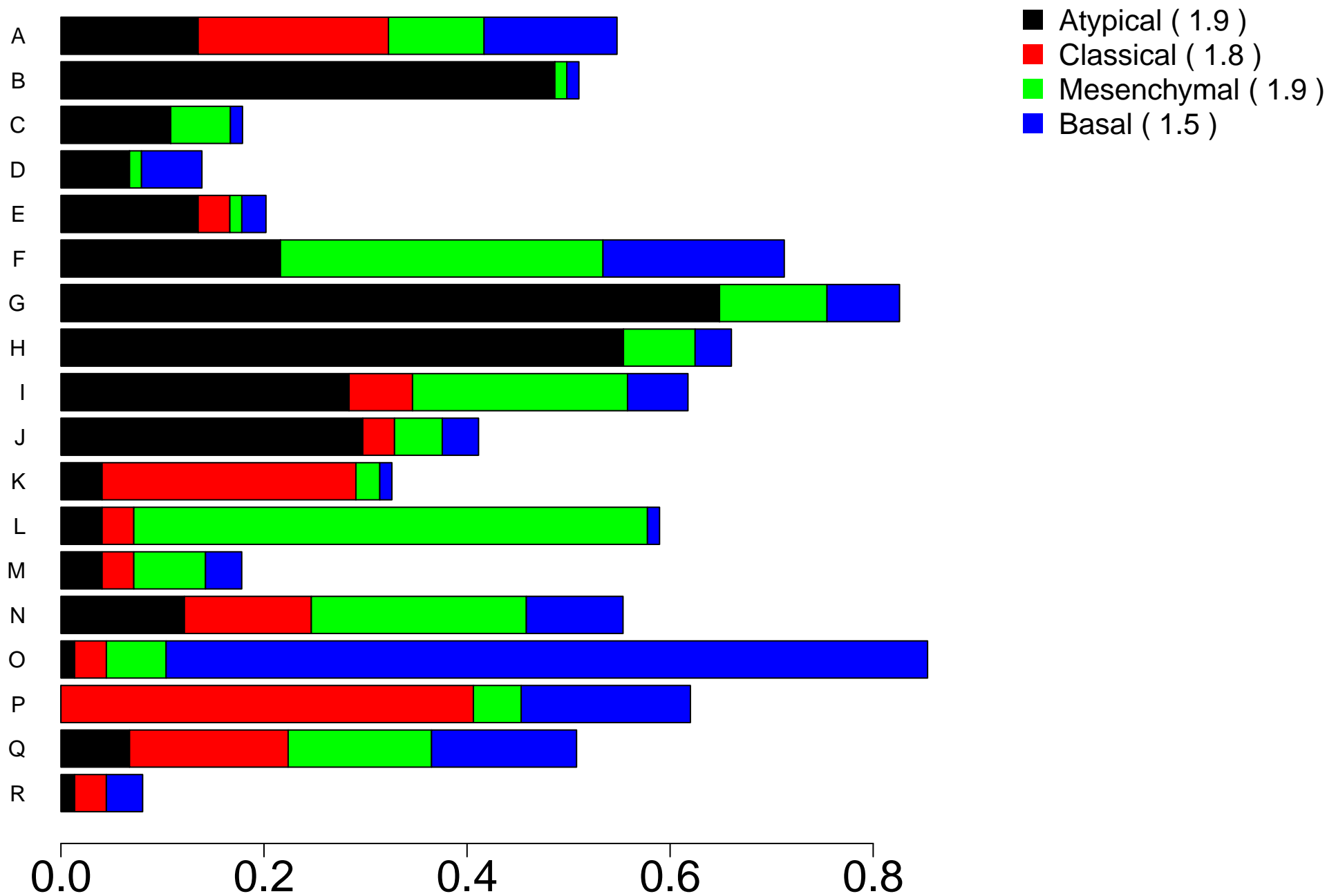
- Atypical
- Classical
- Mesenchymal
- Basal

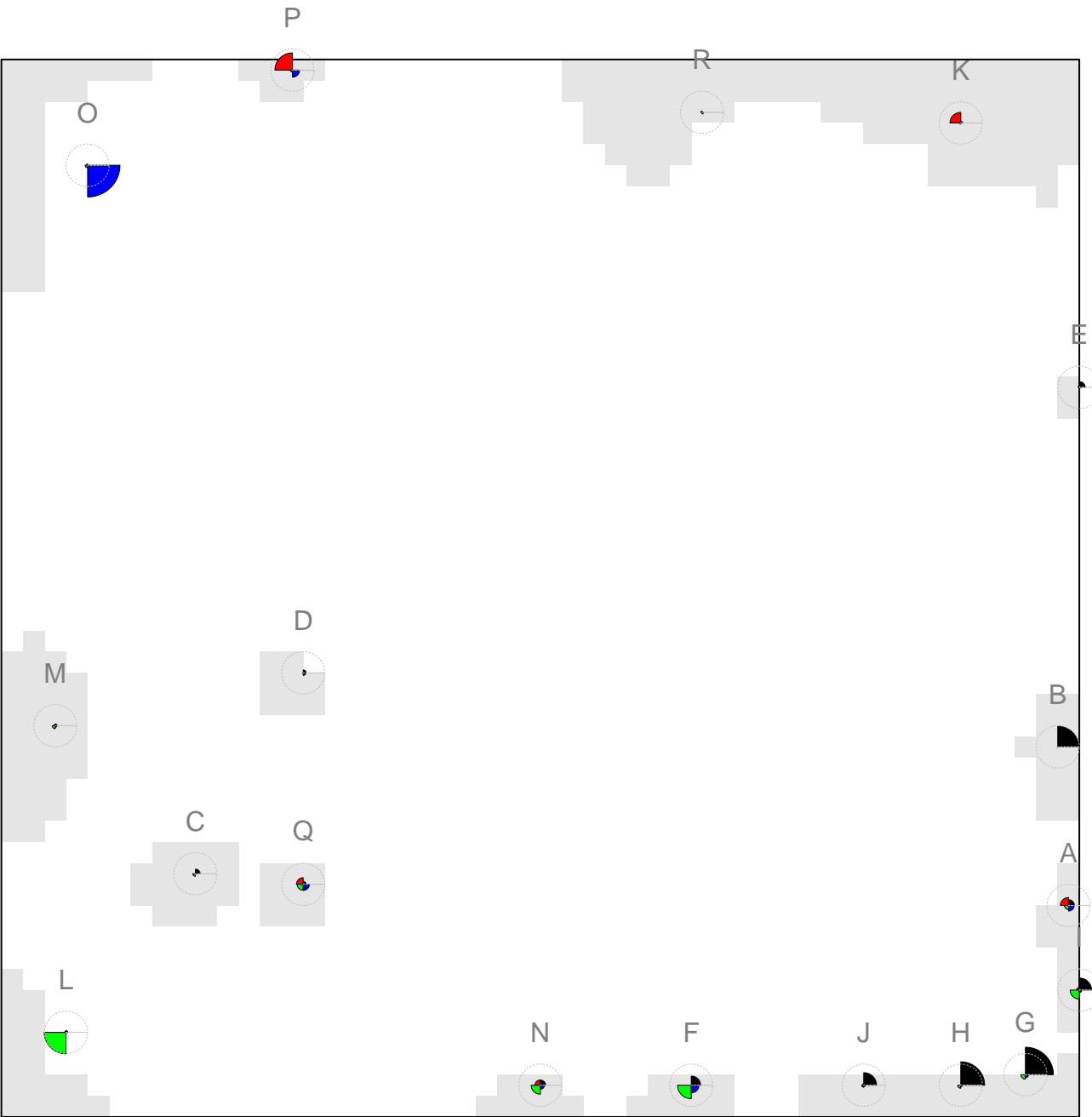


- Atypical
- Classical
- Mesenchymal
- Basal

Group Association

< #spots >

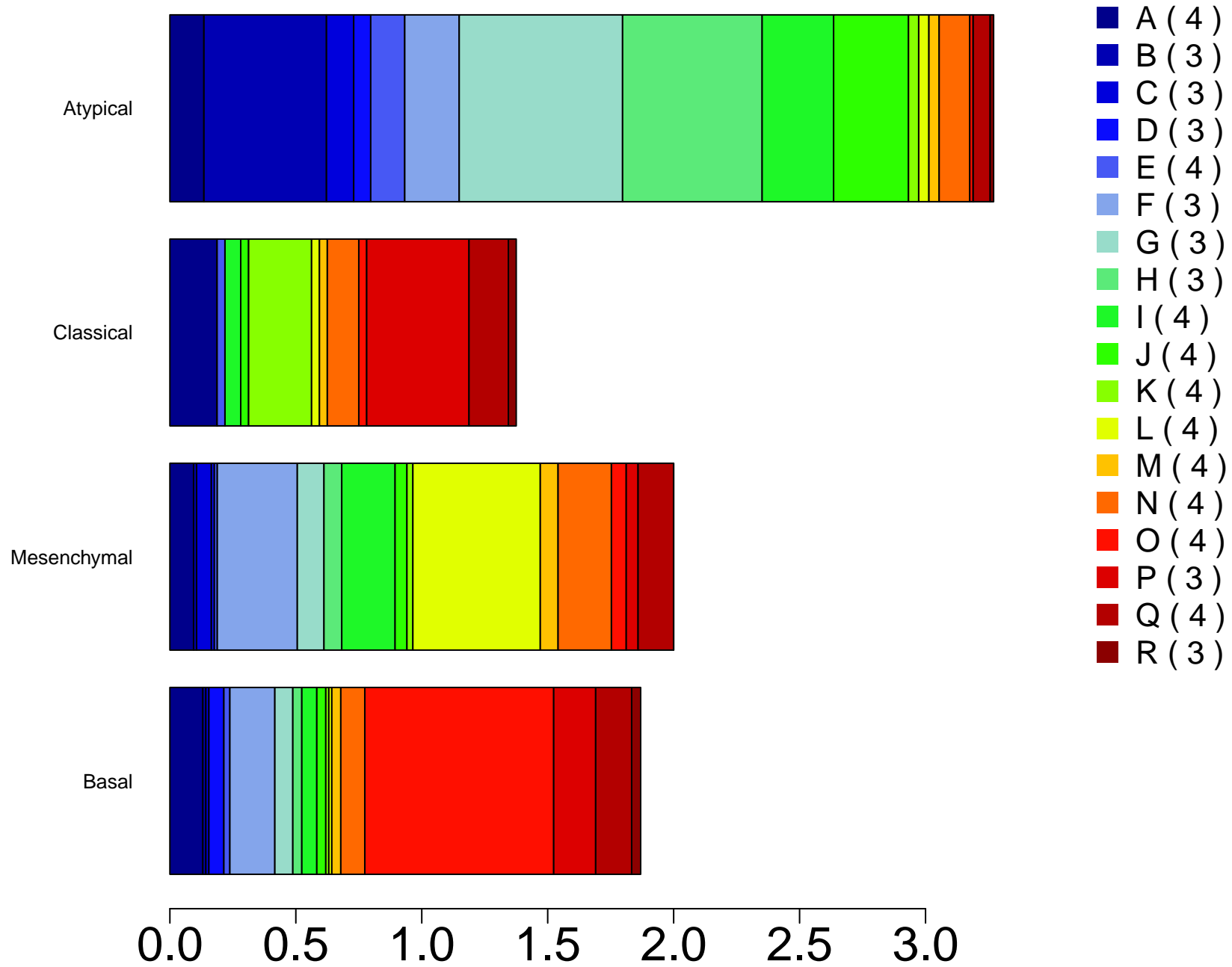




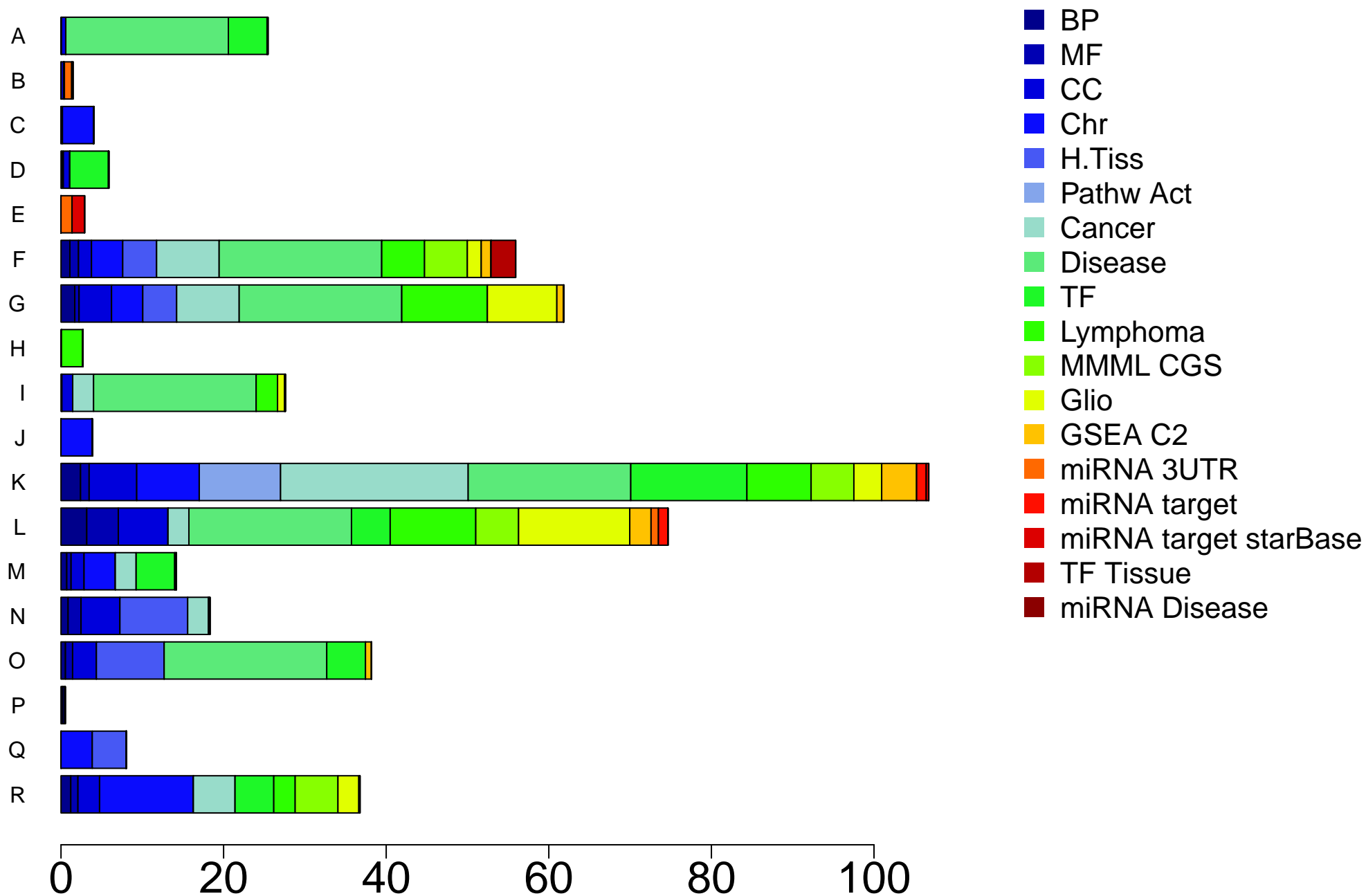
- Atypical
- Classical
- Mesenchymal
- Basal

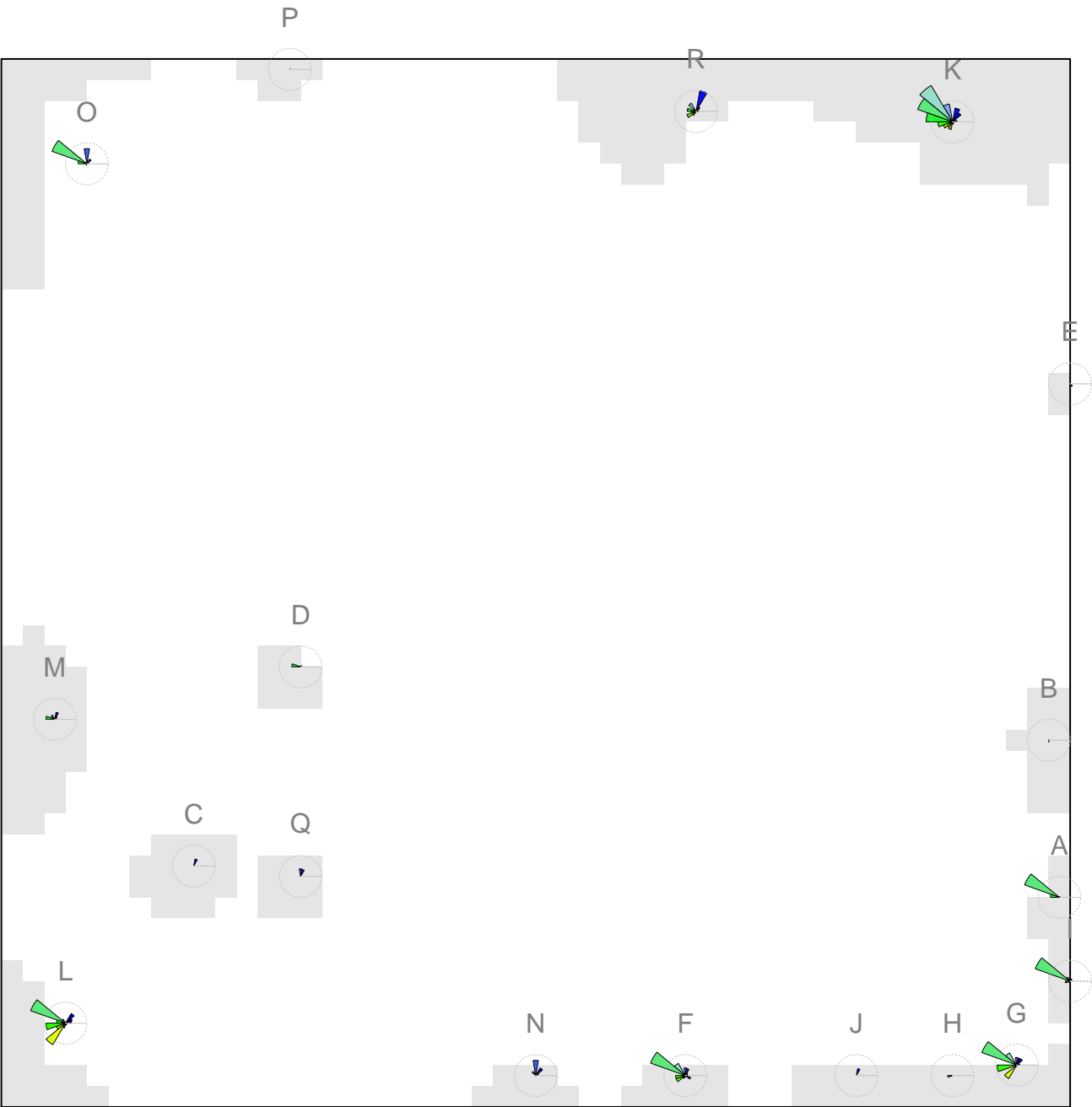
Spot Association

#groups



%Significant Genesets





- 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