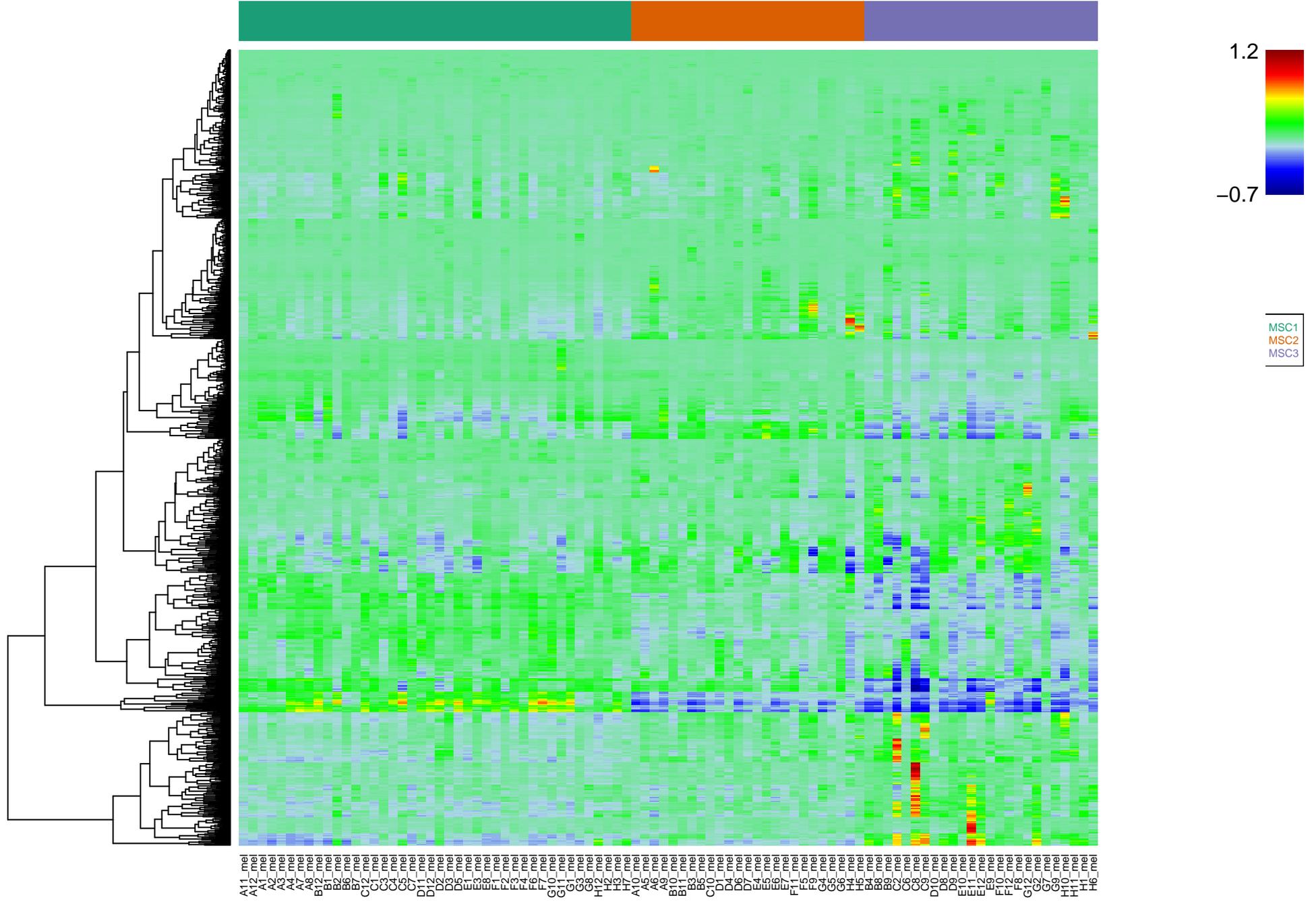
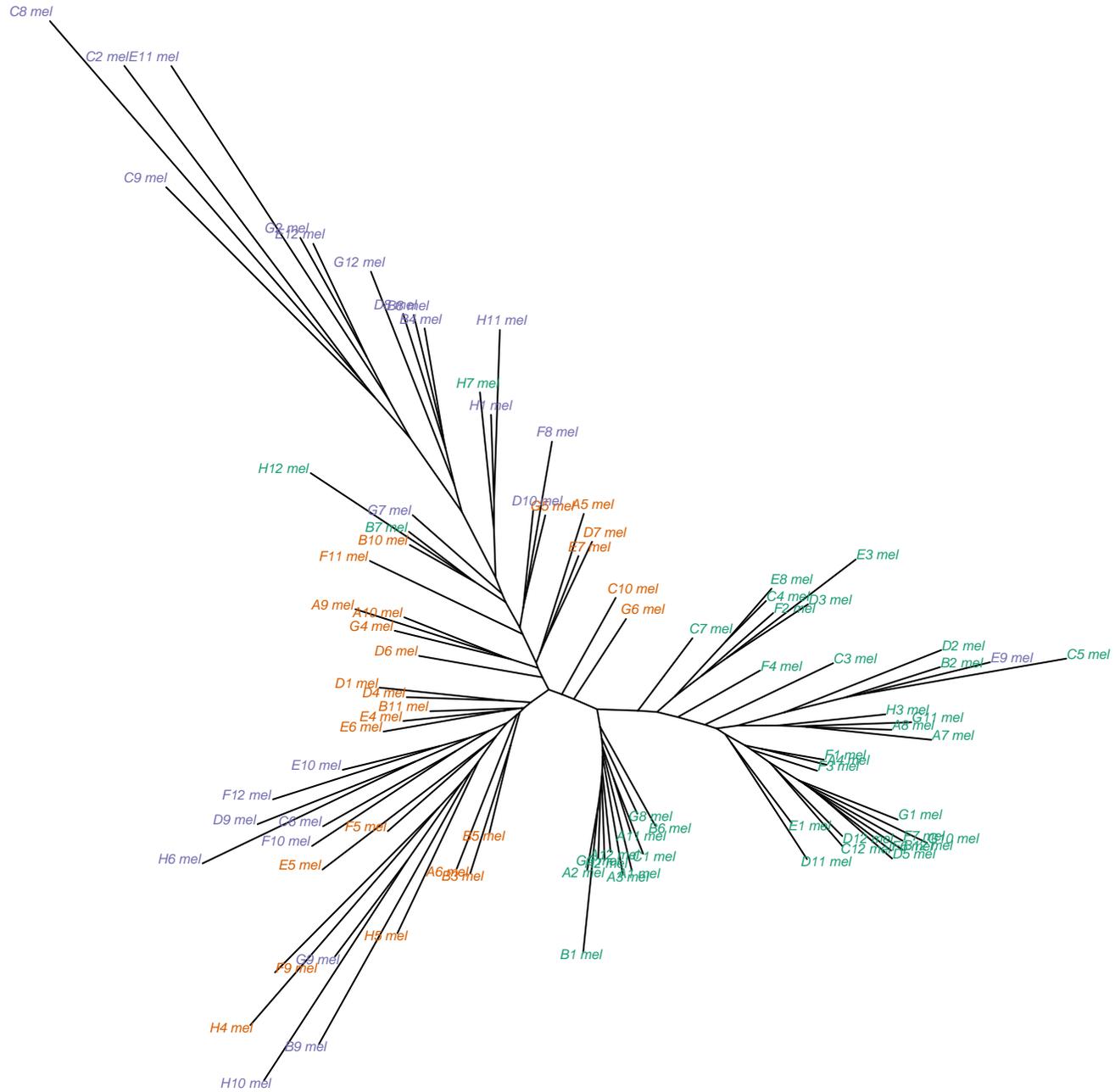


Clustering heatmap, all 2500 Metagenes

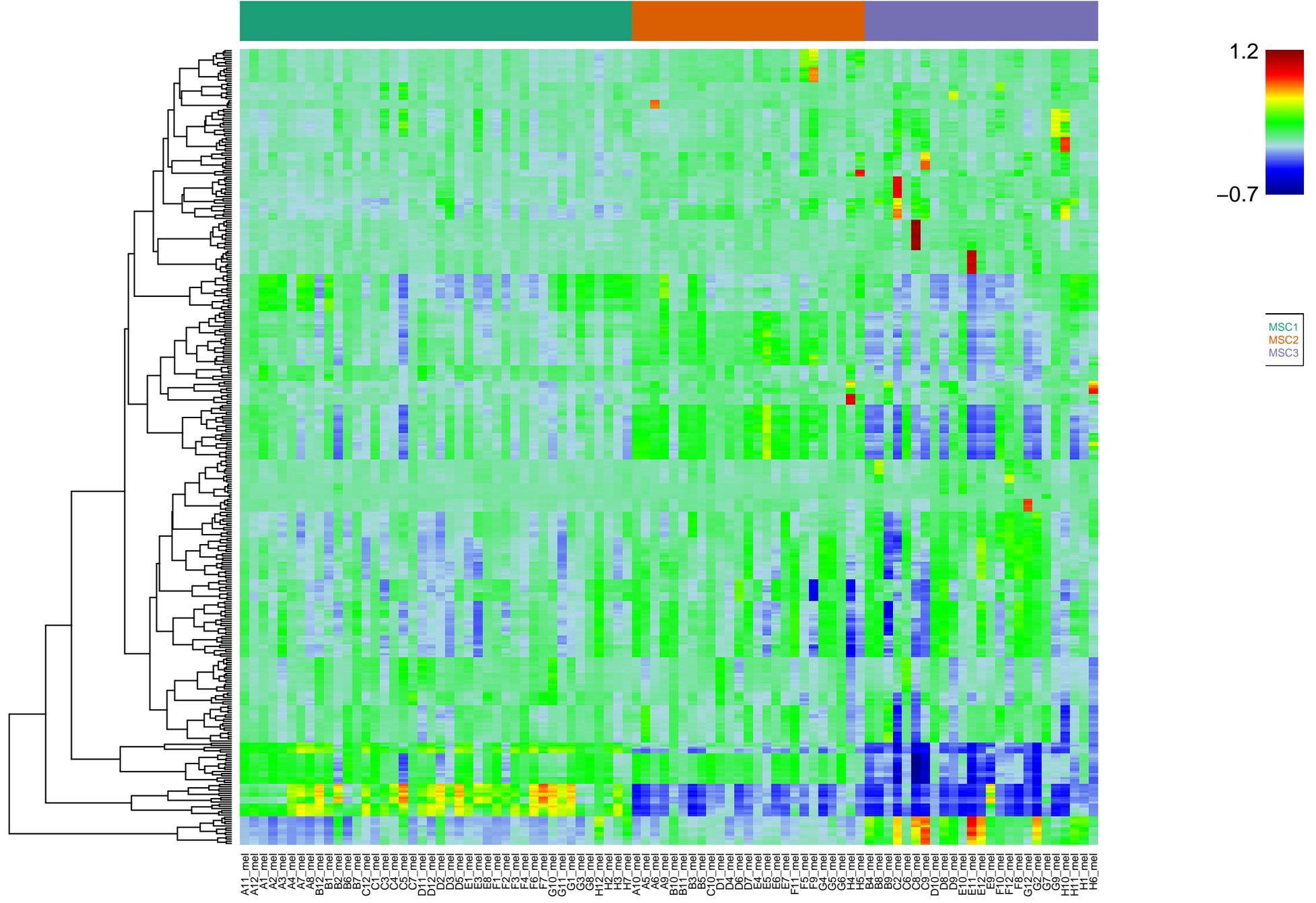


Phylogenetic sample clustering, 365 Spot-Metagenes

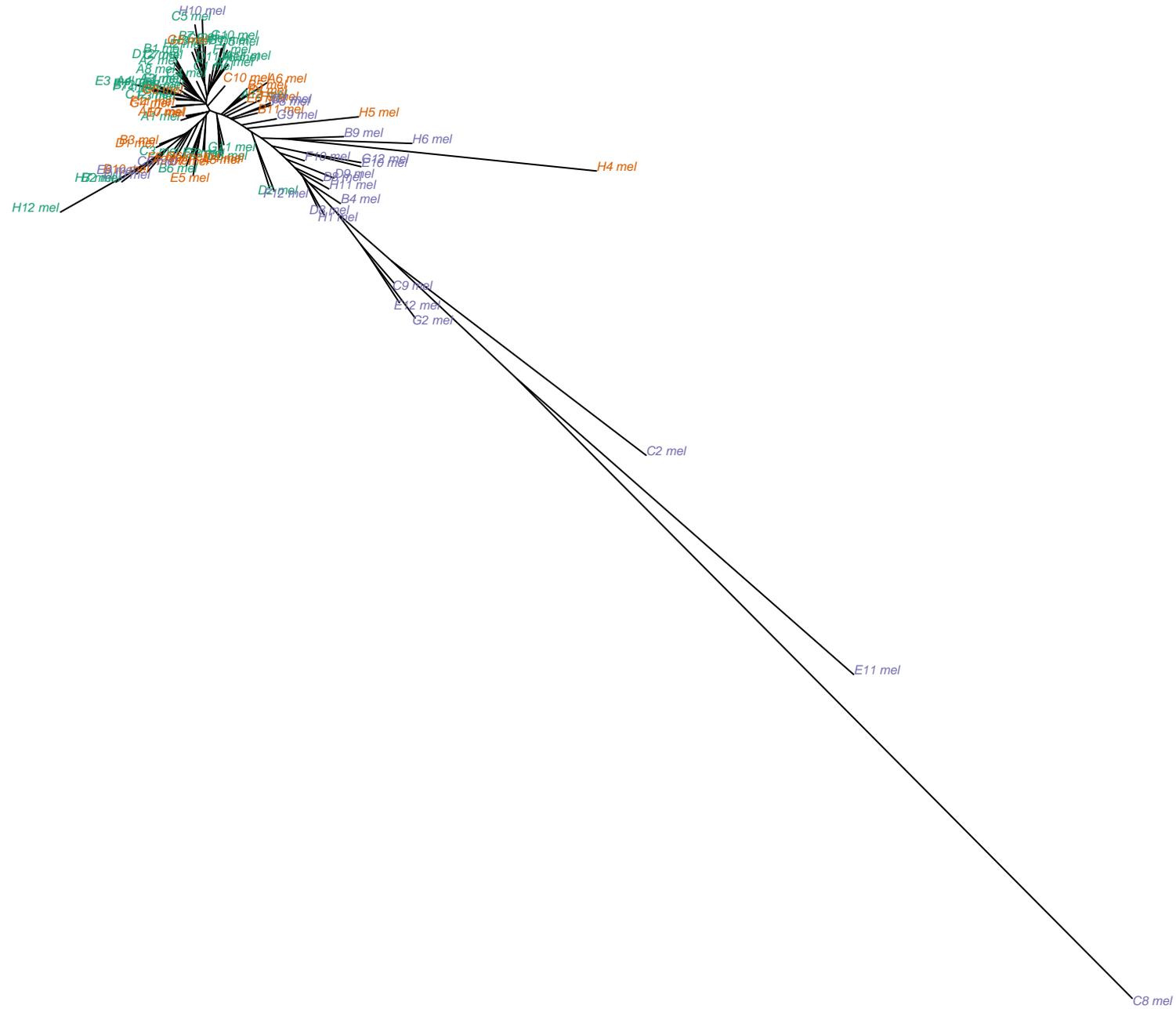


MSC1
MSC2
MSC3

Clustering heatmap, 365 Spot-Metagenes

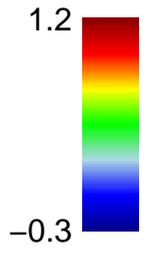
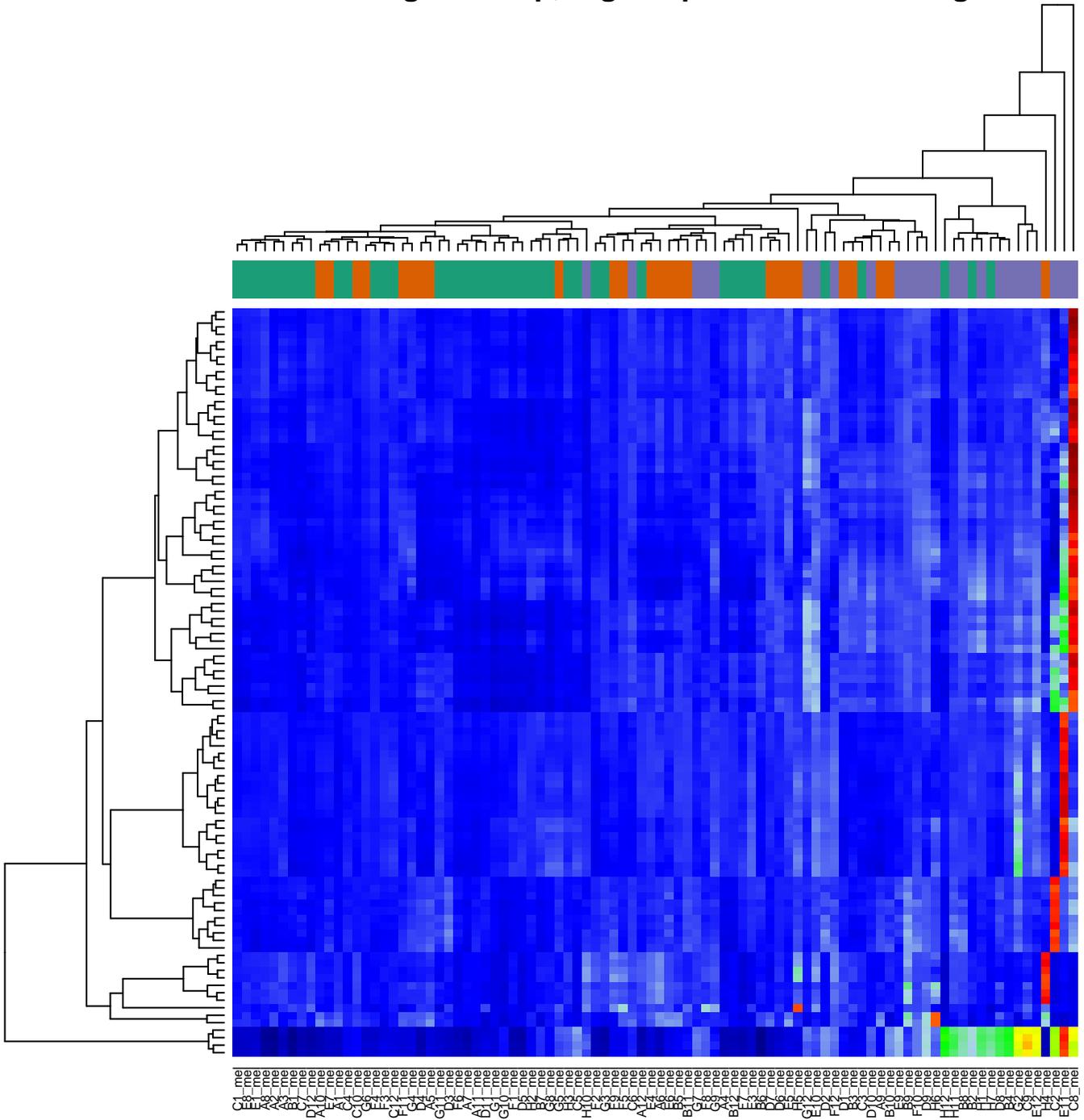


Phylogenetic sample clustering, High Expression: 100 Metagenes



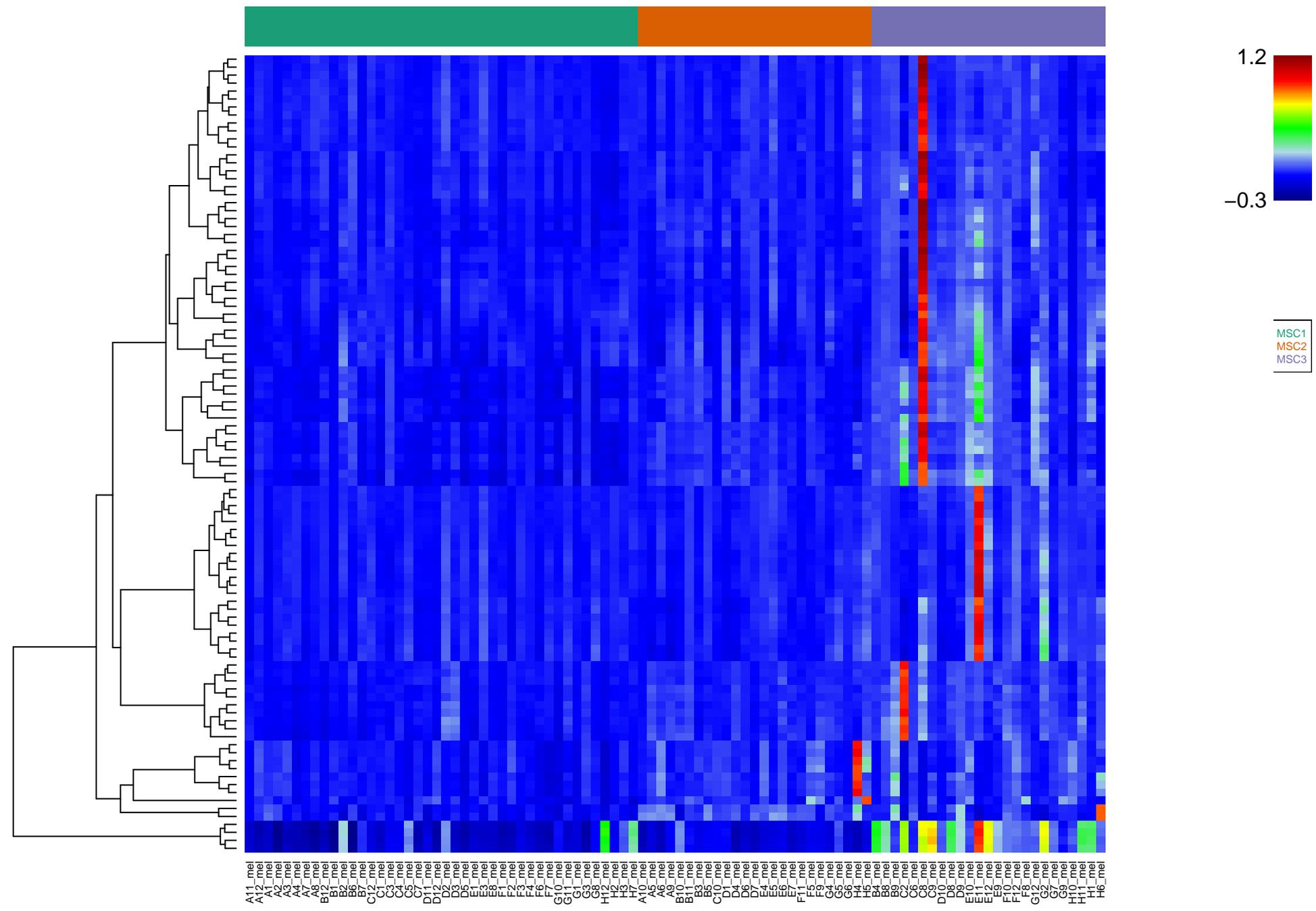
MSC1
MSC2
MSC3

Clustering heatmap, High Expression: 100 Metagenes

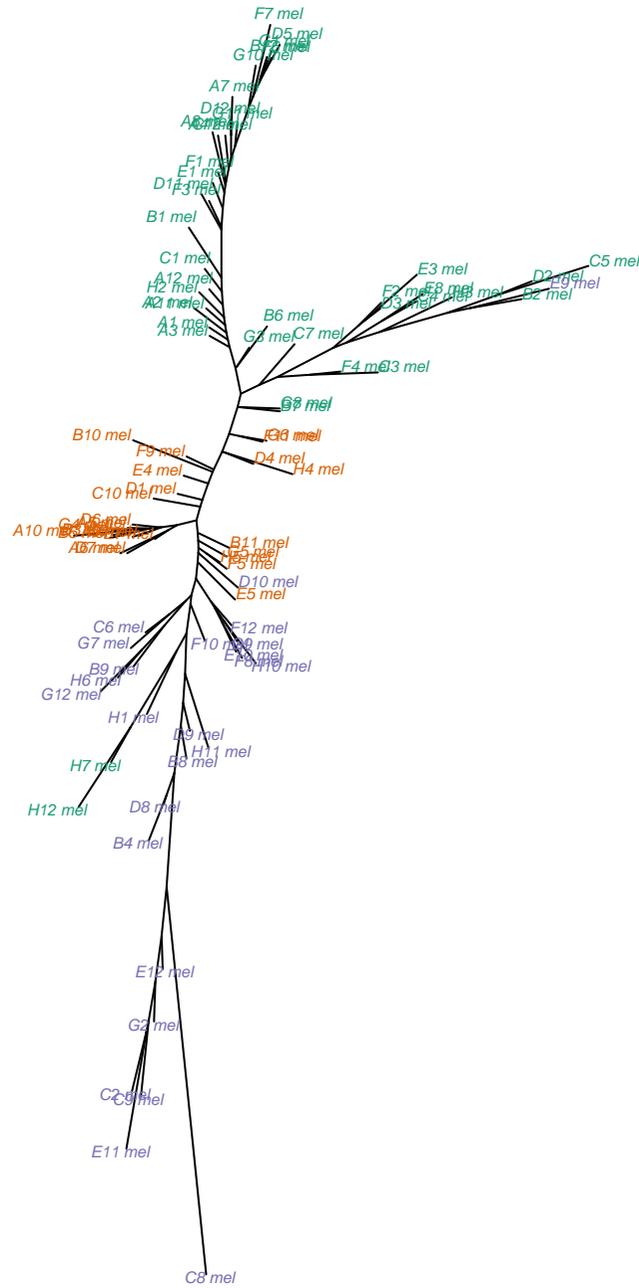


MSC1
MSC2
MSC3

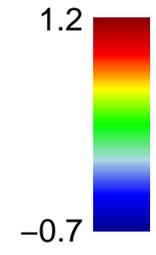
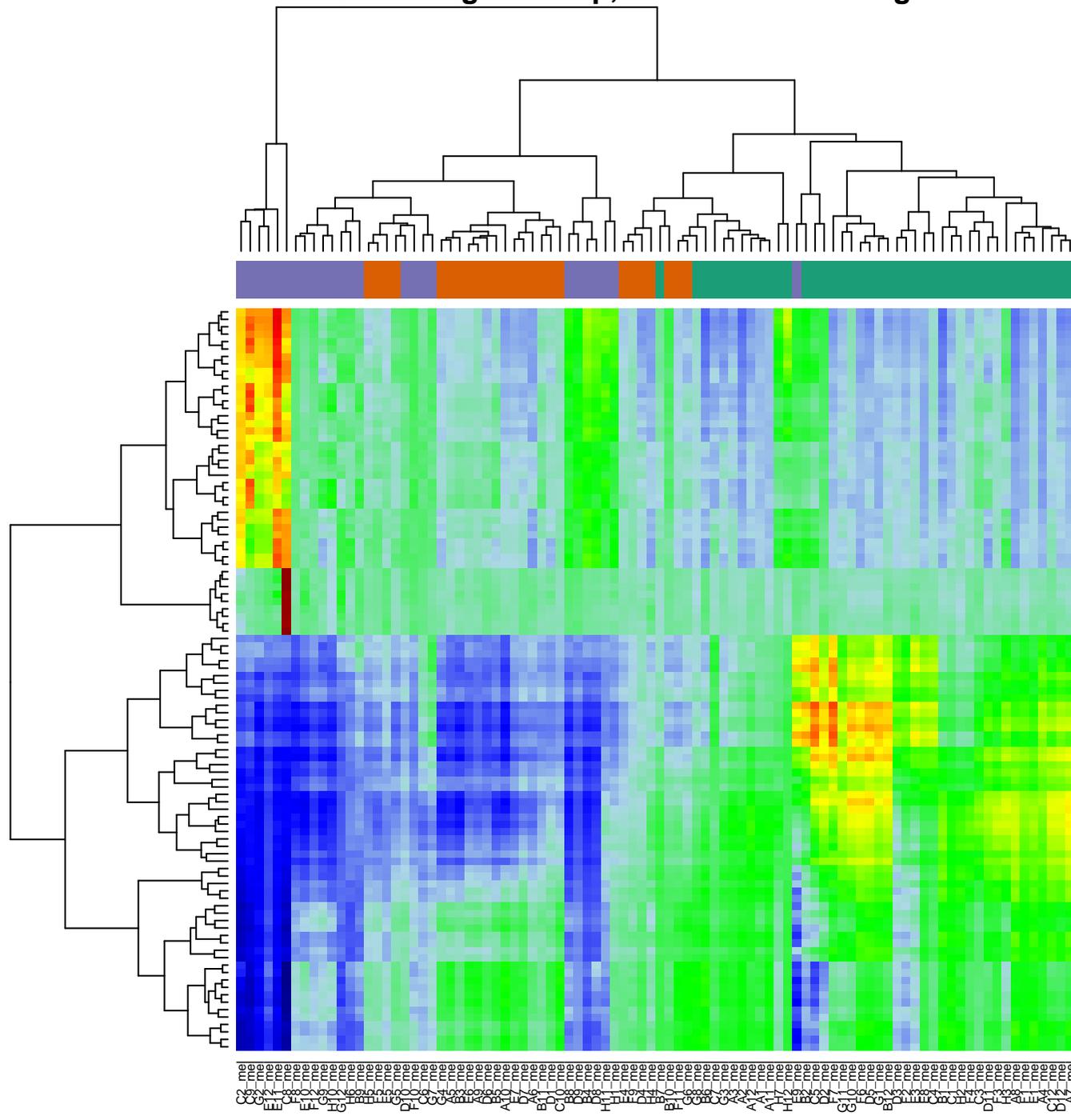
Clustering heatmap, High Expression: 100 Metagenes



Phylogenetic sample clustering, Variance: 100 Metagenes



Clustering heatmap, Variance: 100 Metagenes



MSC1
MSC2
MSC3

Clustering heatmap, Variance: 100 Metagenes

