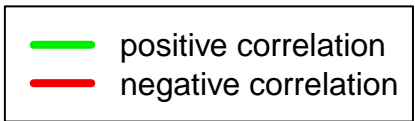
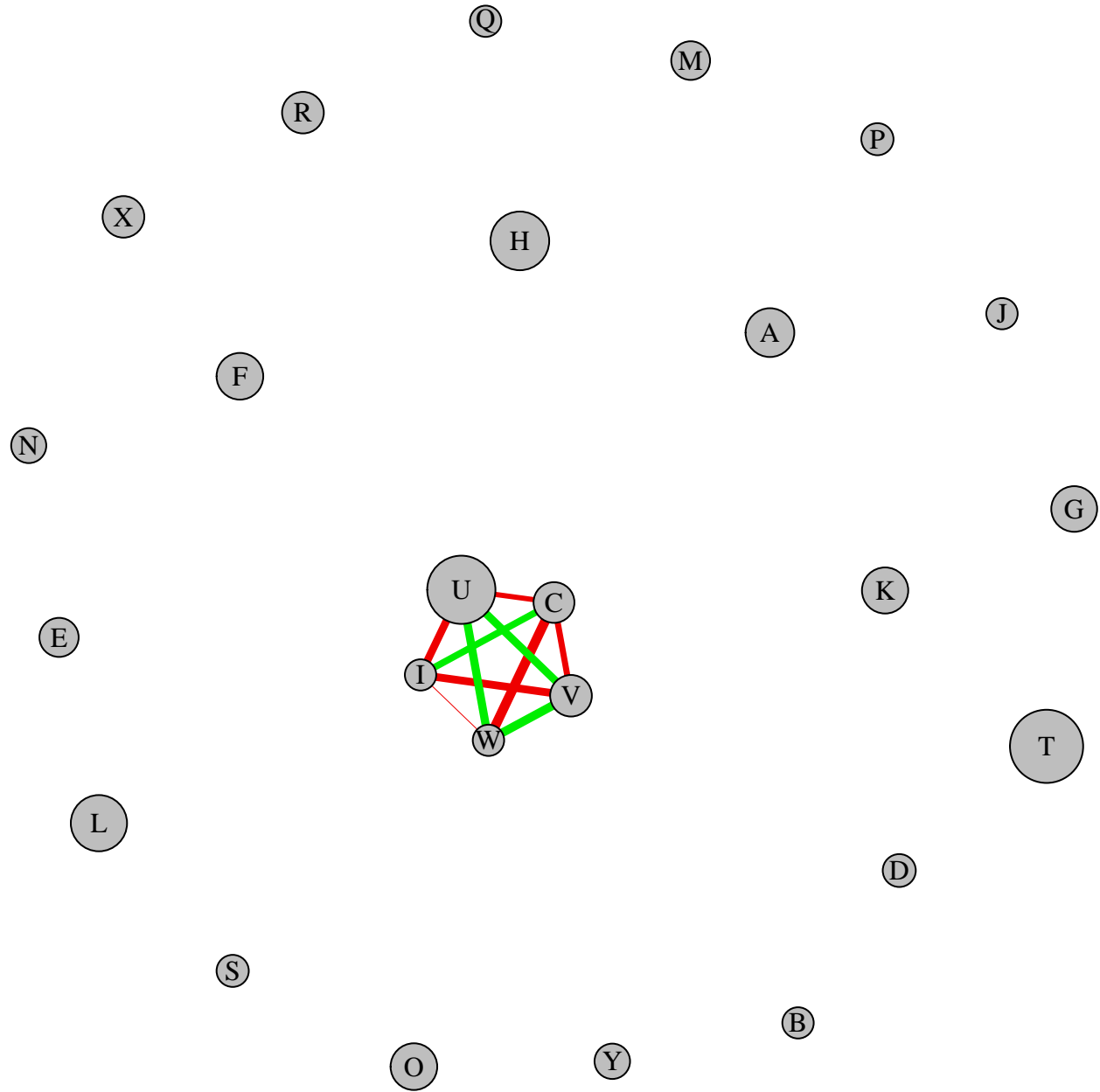
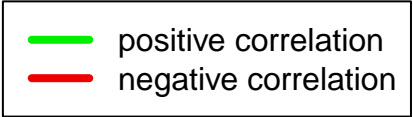
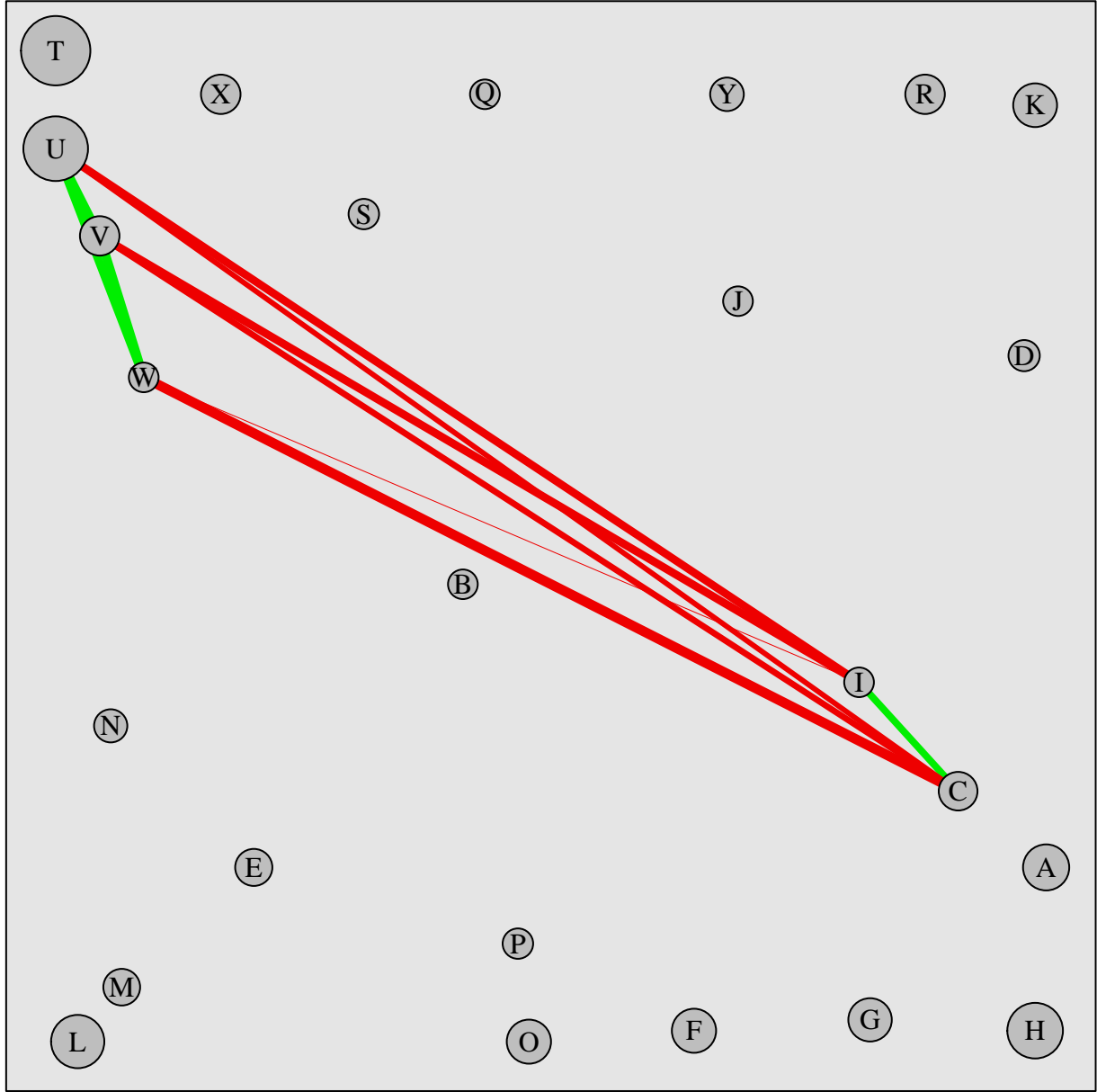
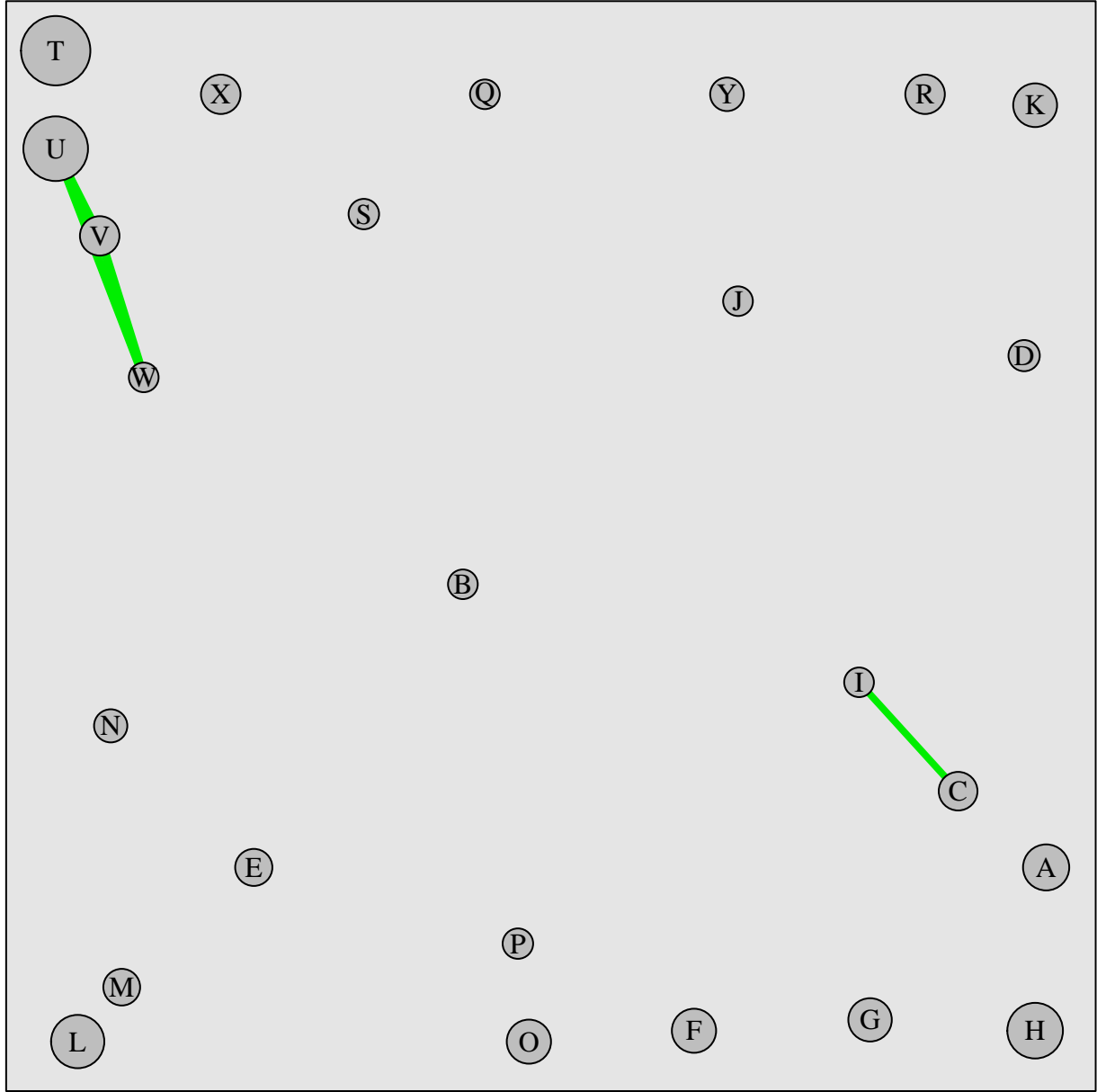


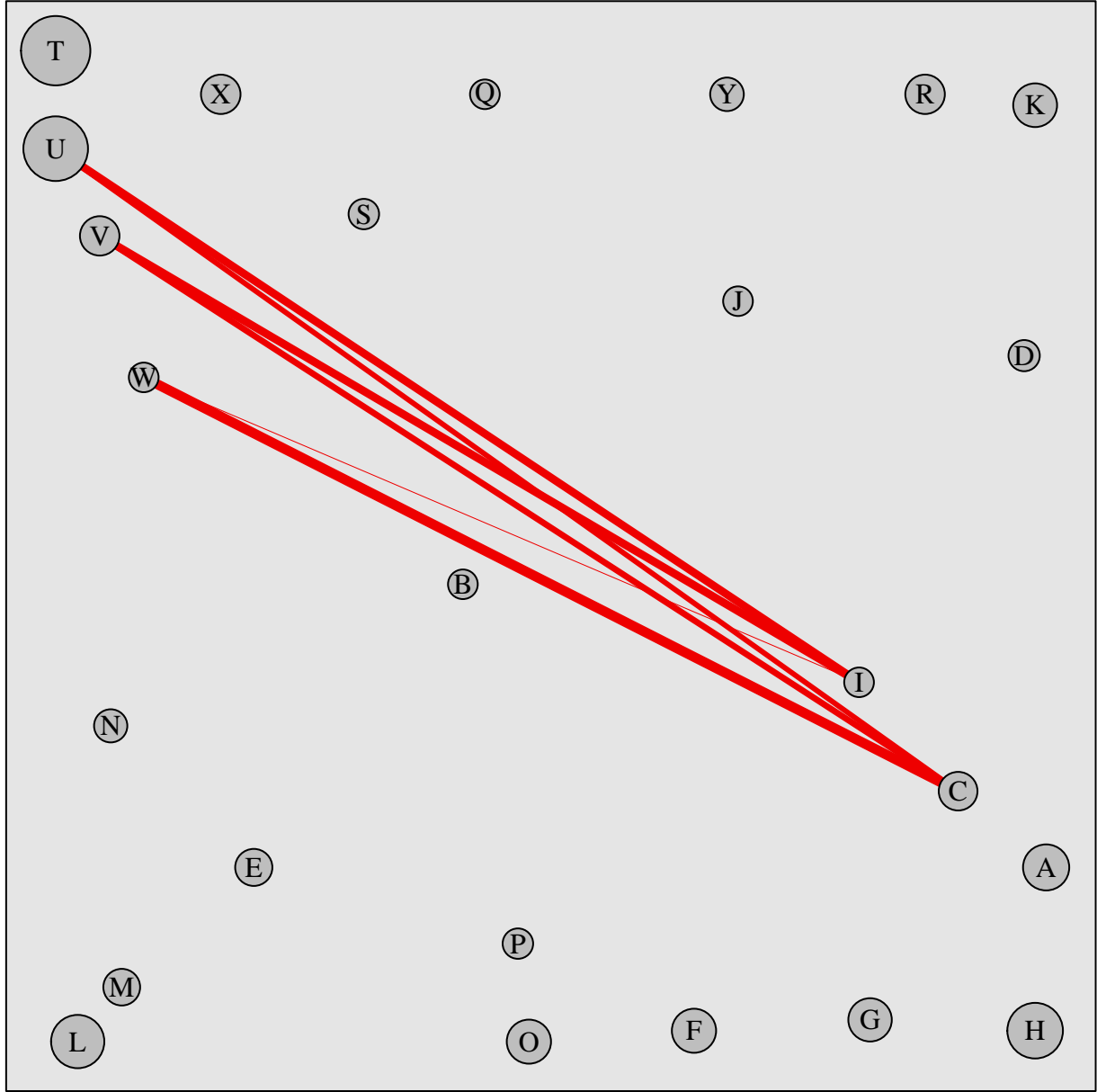
# WTO network






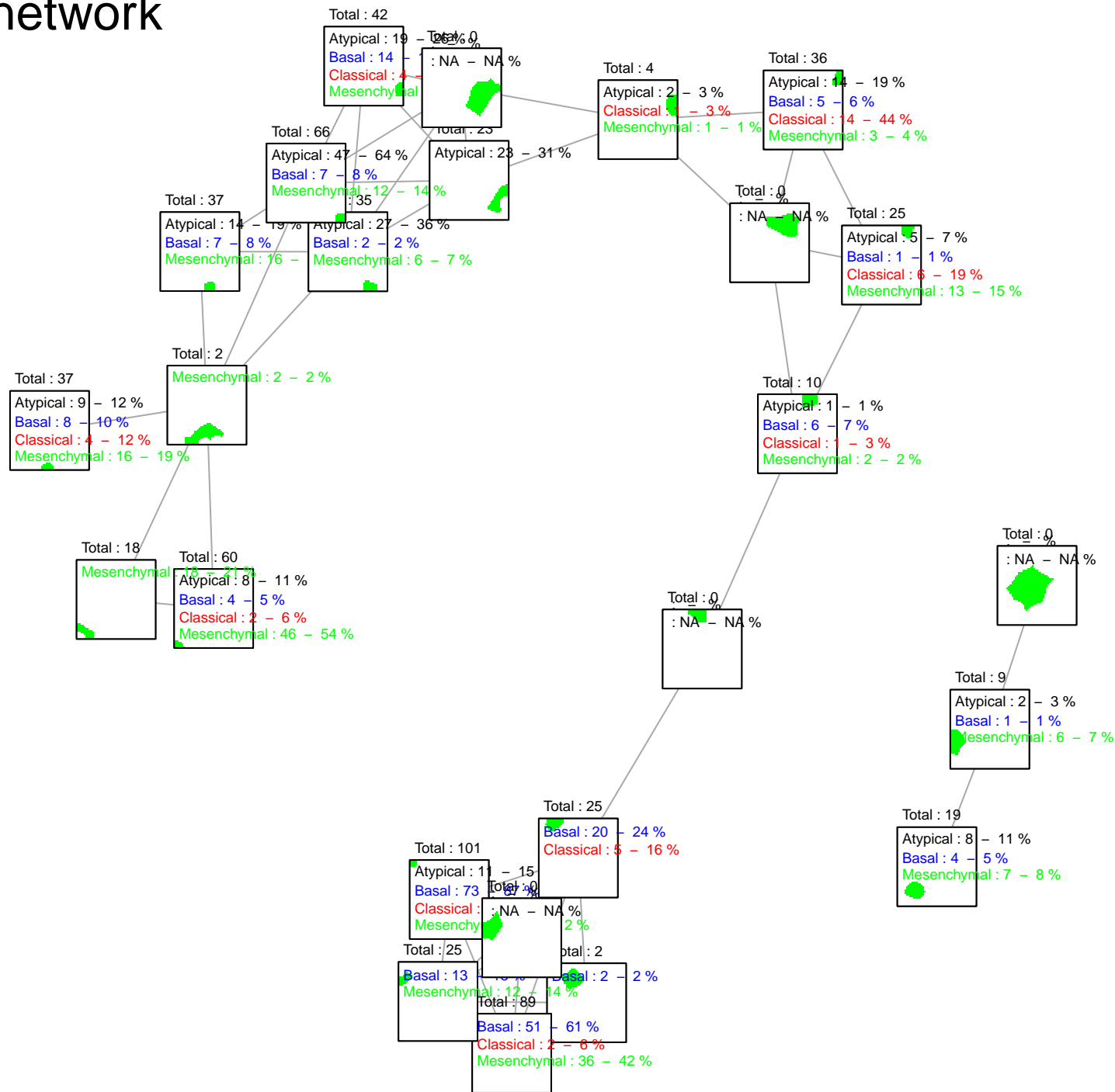


 positive correlation

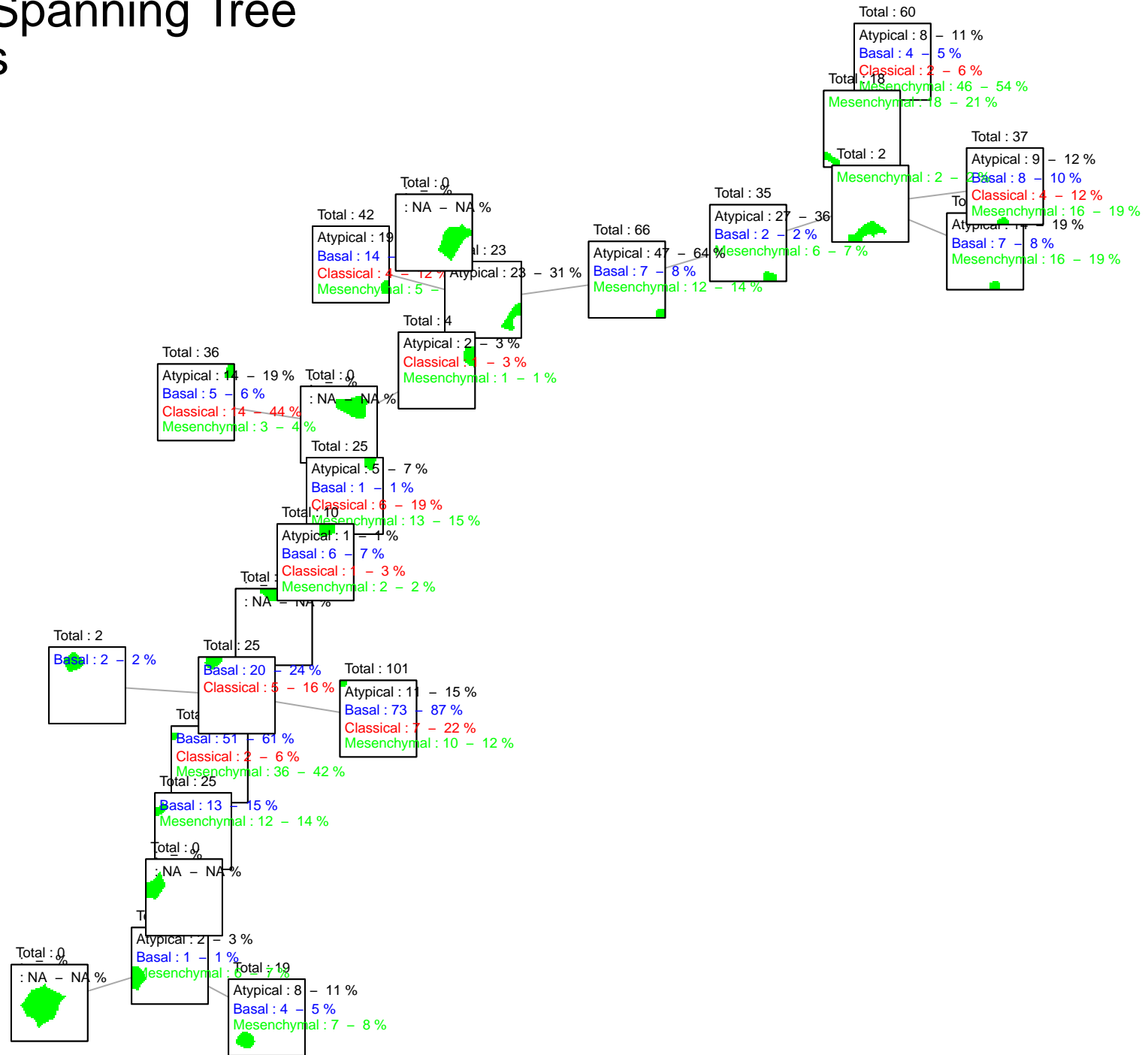


 negative correlation

# Single spot network

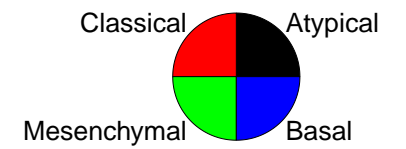
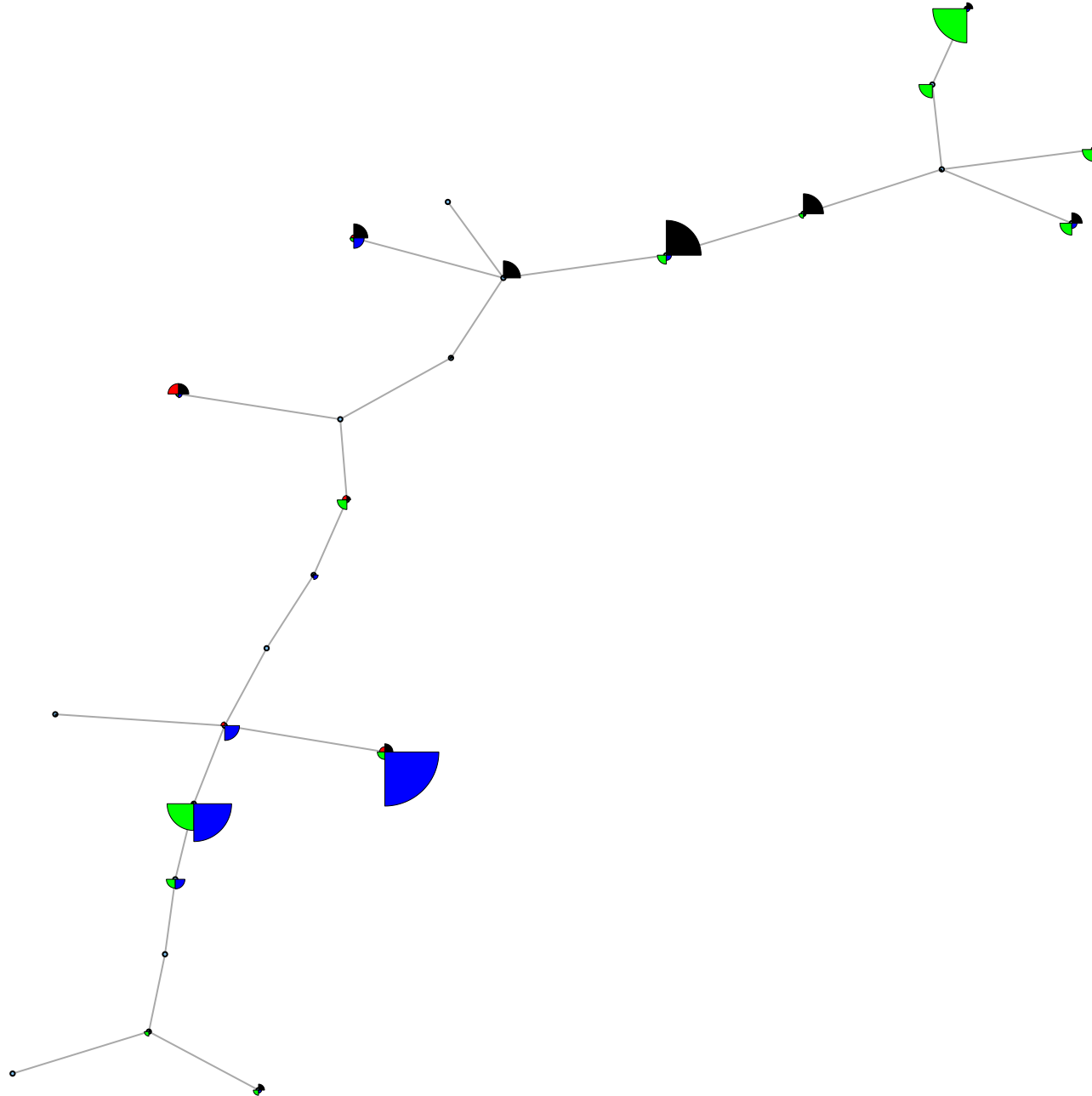


# Correlation Spanning Tree Single Spots



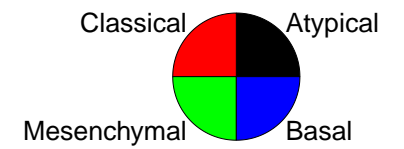
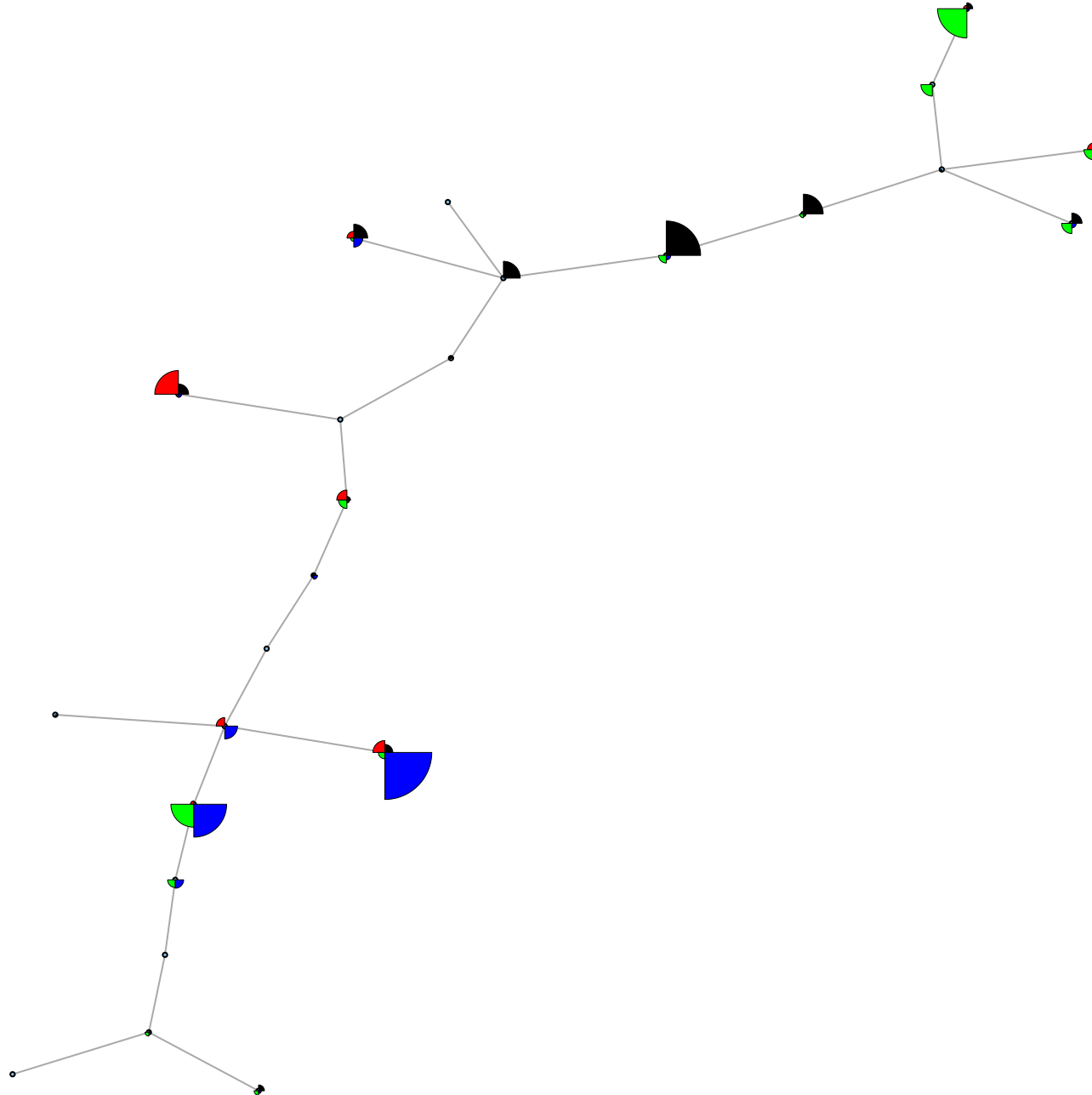
# Correlation Spanning Tree

absolute numbers



# Correlation Spanning Tree

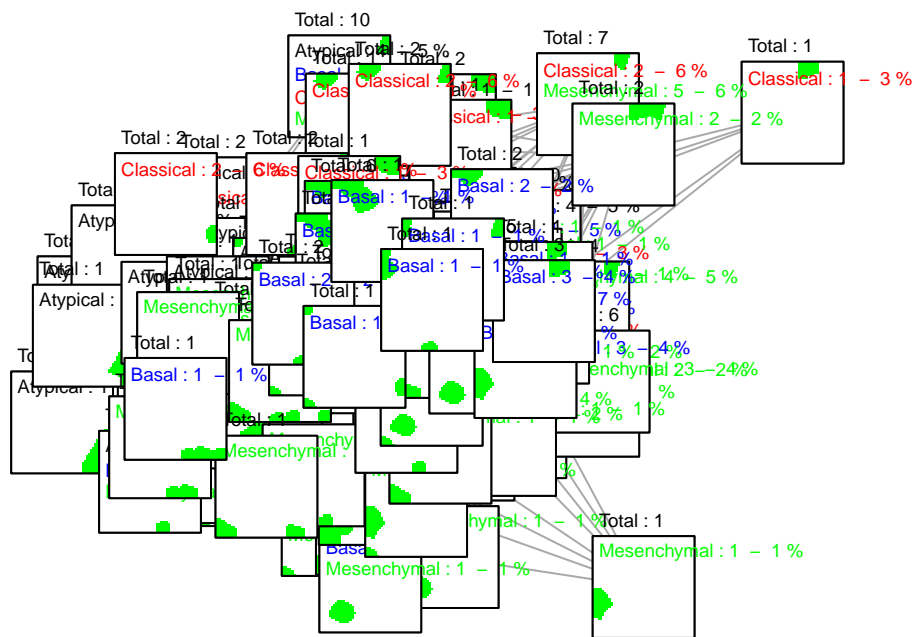
relative to category size



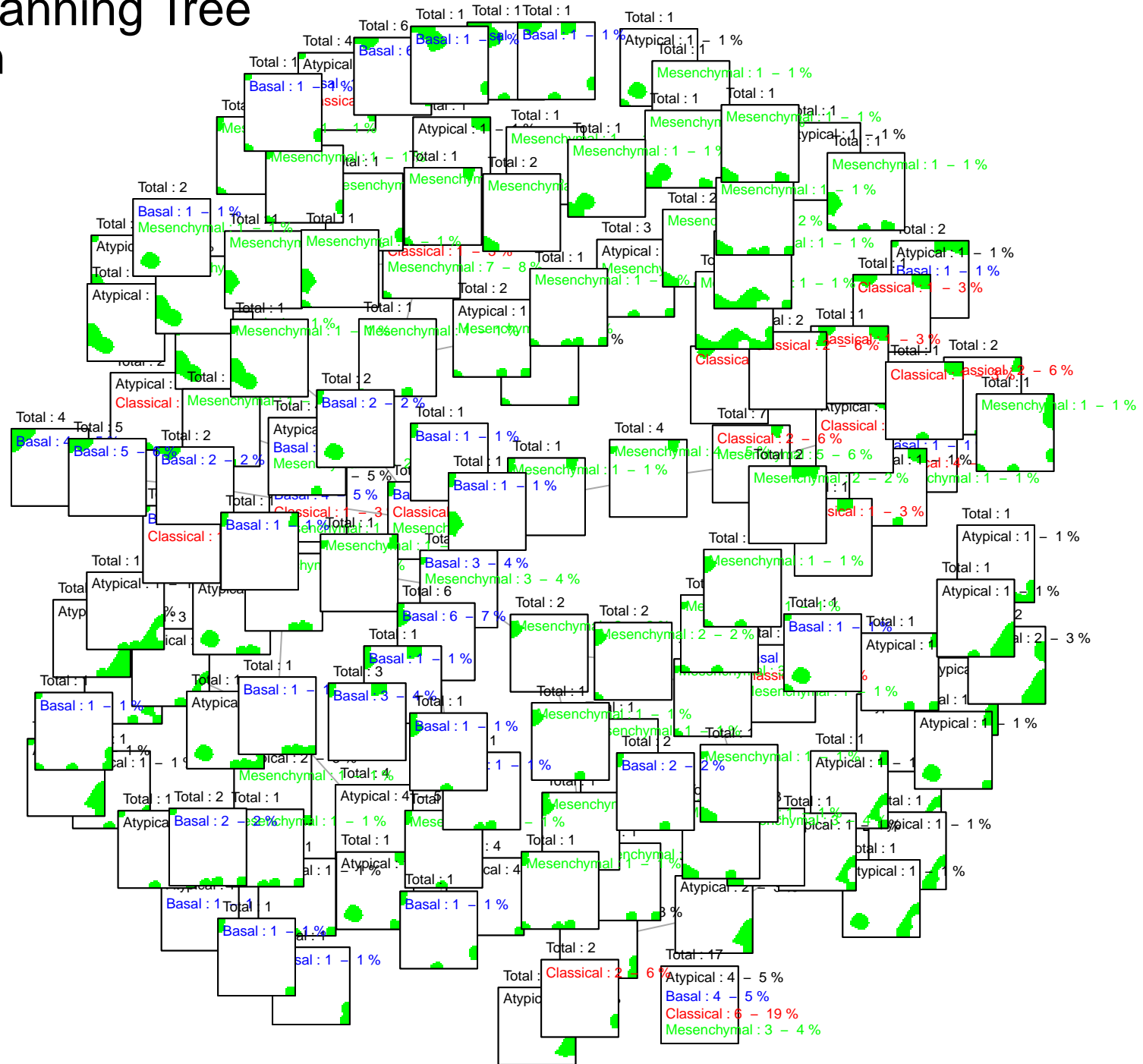


# Spot pattern network

Total : 17  
 Atypical : 4 - 5 %  
 Basal : 4 - 5 %  
 Classical : 6 - 19 %  
 Mesenchymal : 3 - 4 %

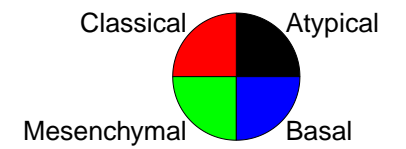
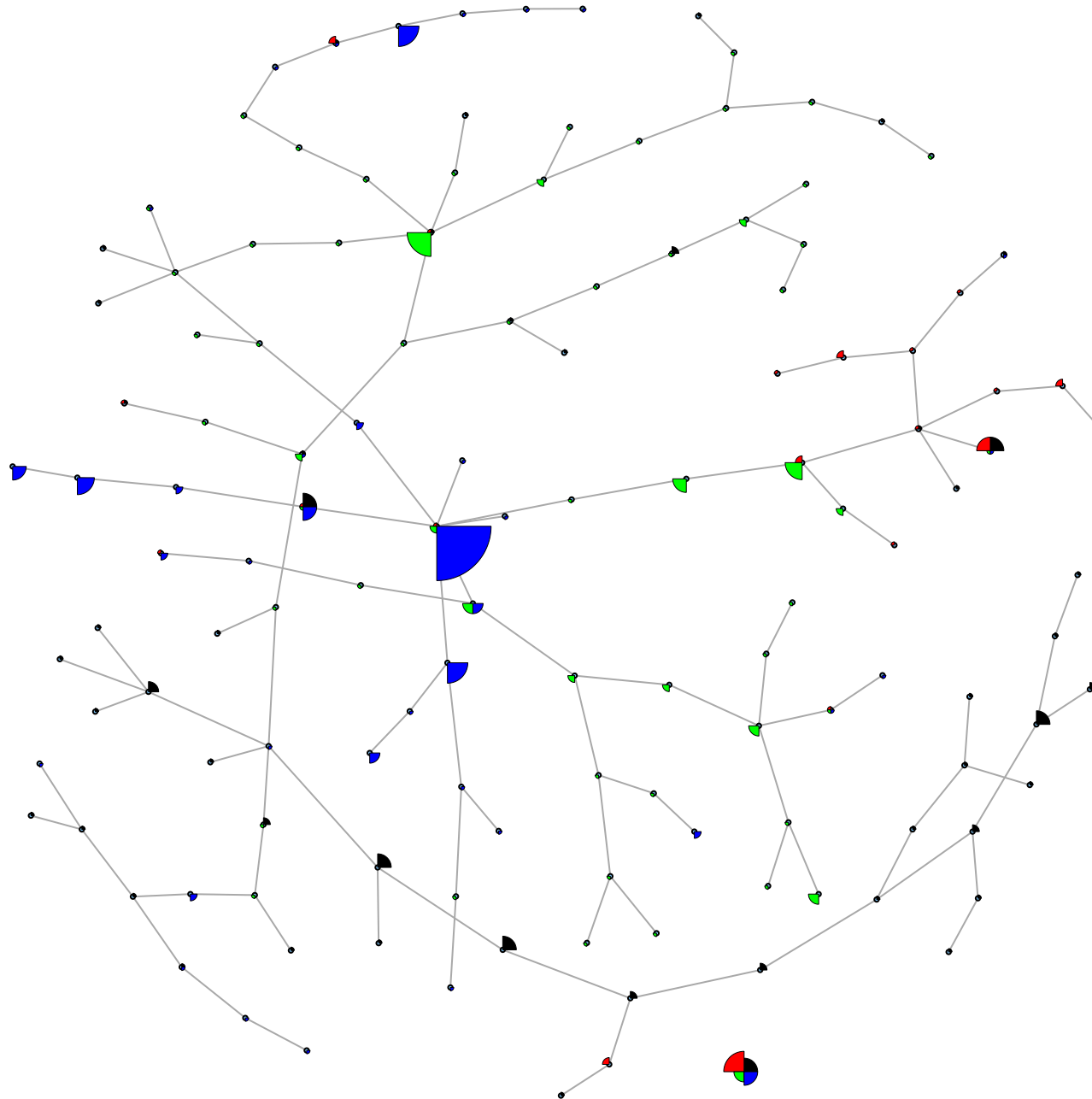


# Distance Spanning Tree Spot Pattern



# Distance Spanning Tree

absolute numbers



# Distance Spanning Tree

relative to category size

