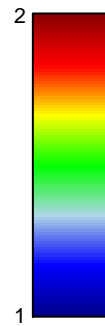
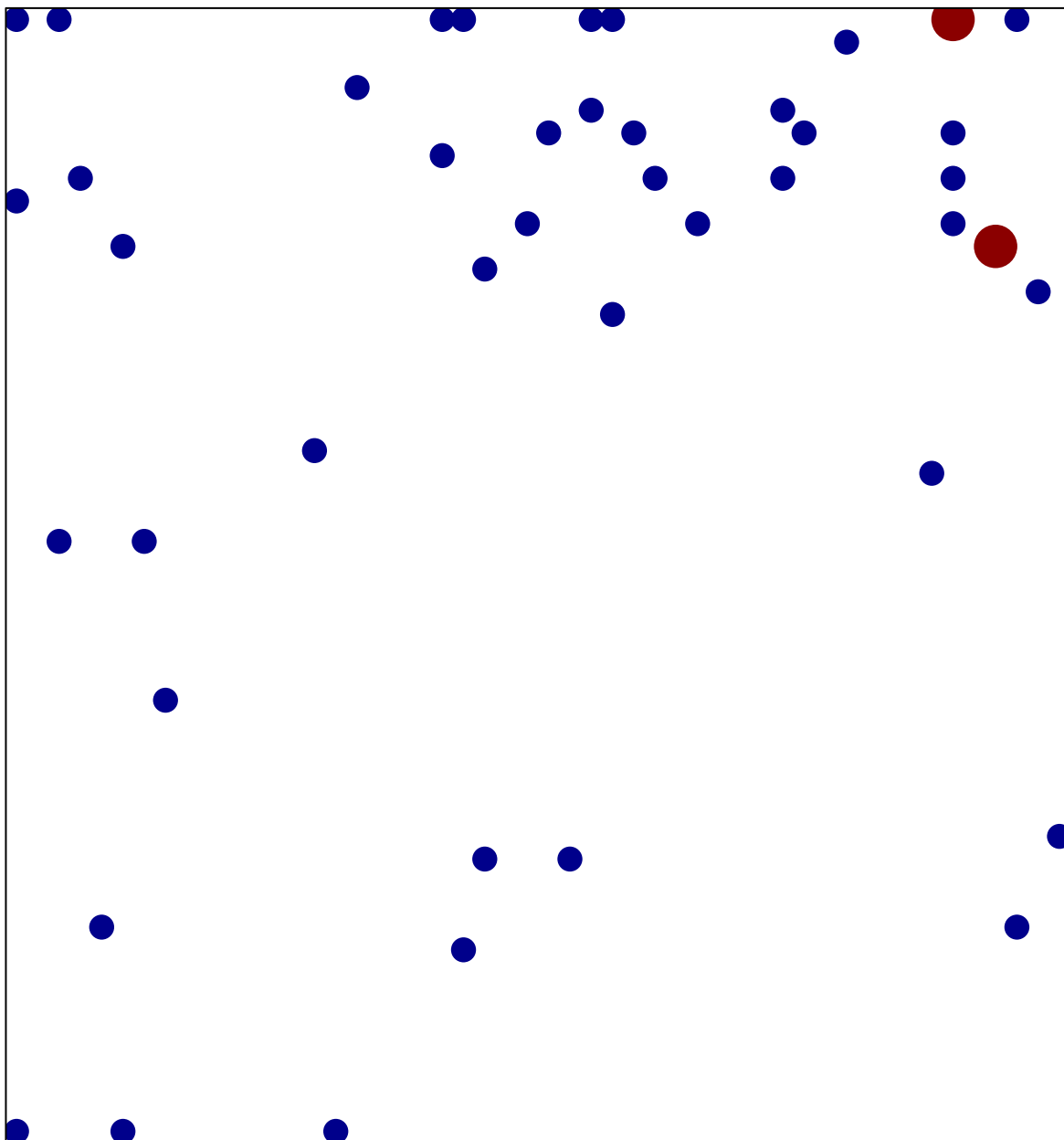
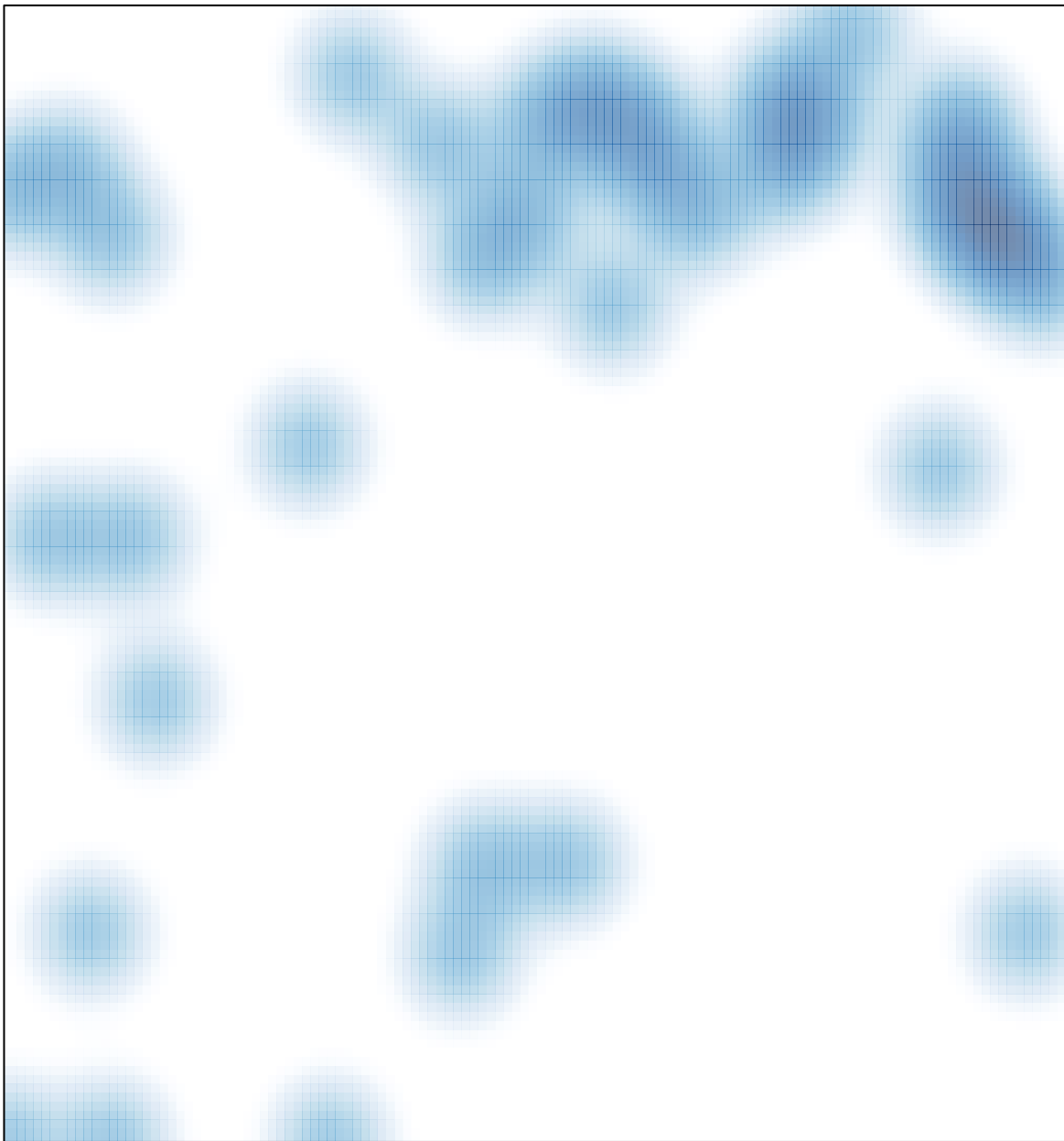


# REACTOME\_P53\_INDEPENDENT\_G1\_S\_DNA\_DAMAGE\_CHECKPOINT



# features = 46  
chi-square p = 0.78

# ME\_P53\_INDEPENDENT\_G1\_S\_DNA\_DAMAGE\_CHE



# features = 46 , max = 2