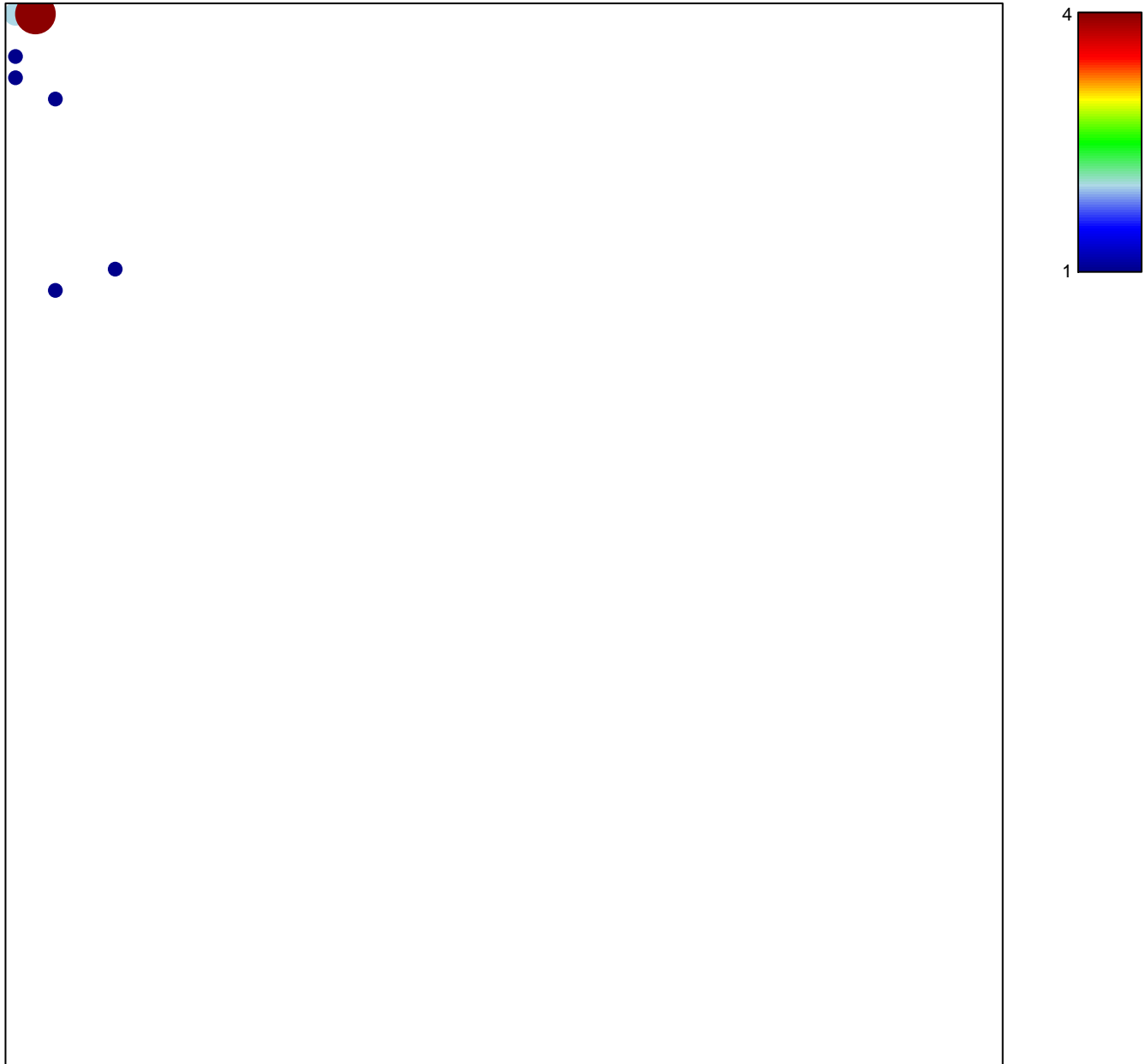
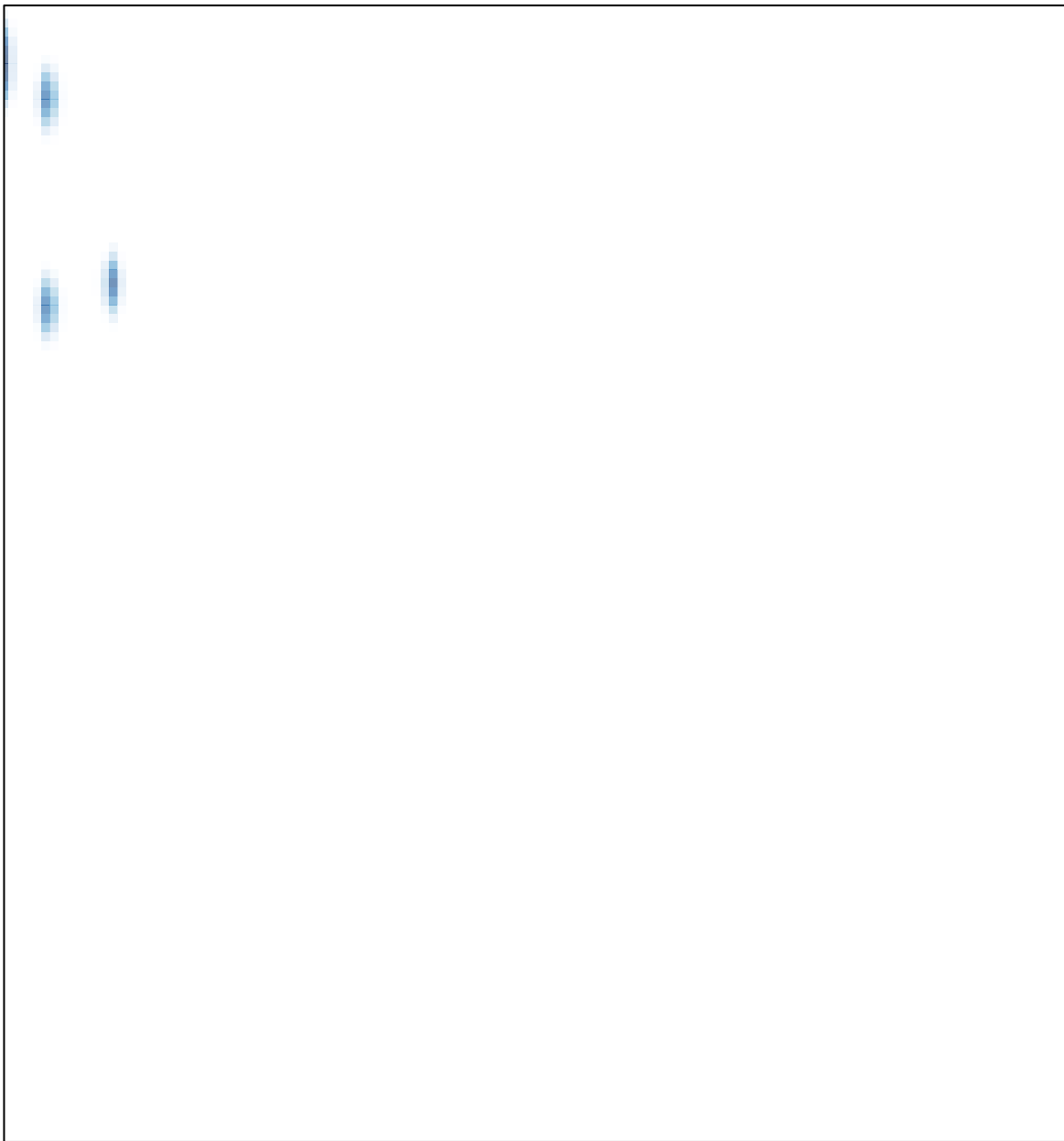


# REACTOME\_REPAIR\_SYNTHESIS\_FOR\_GAP\_FILLING\_BY\_DNA\_POL\_IN\_TC\_NER



# features = 11  
chi-square p = 0.72

# EPAIR\_SYNTHESIS\_FOR\_GAP\_FILLING\_BY\_DNA\_P



# features = 11 , max = 4