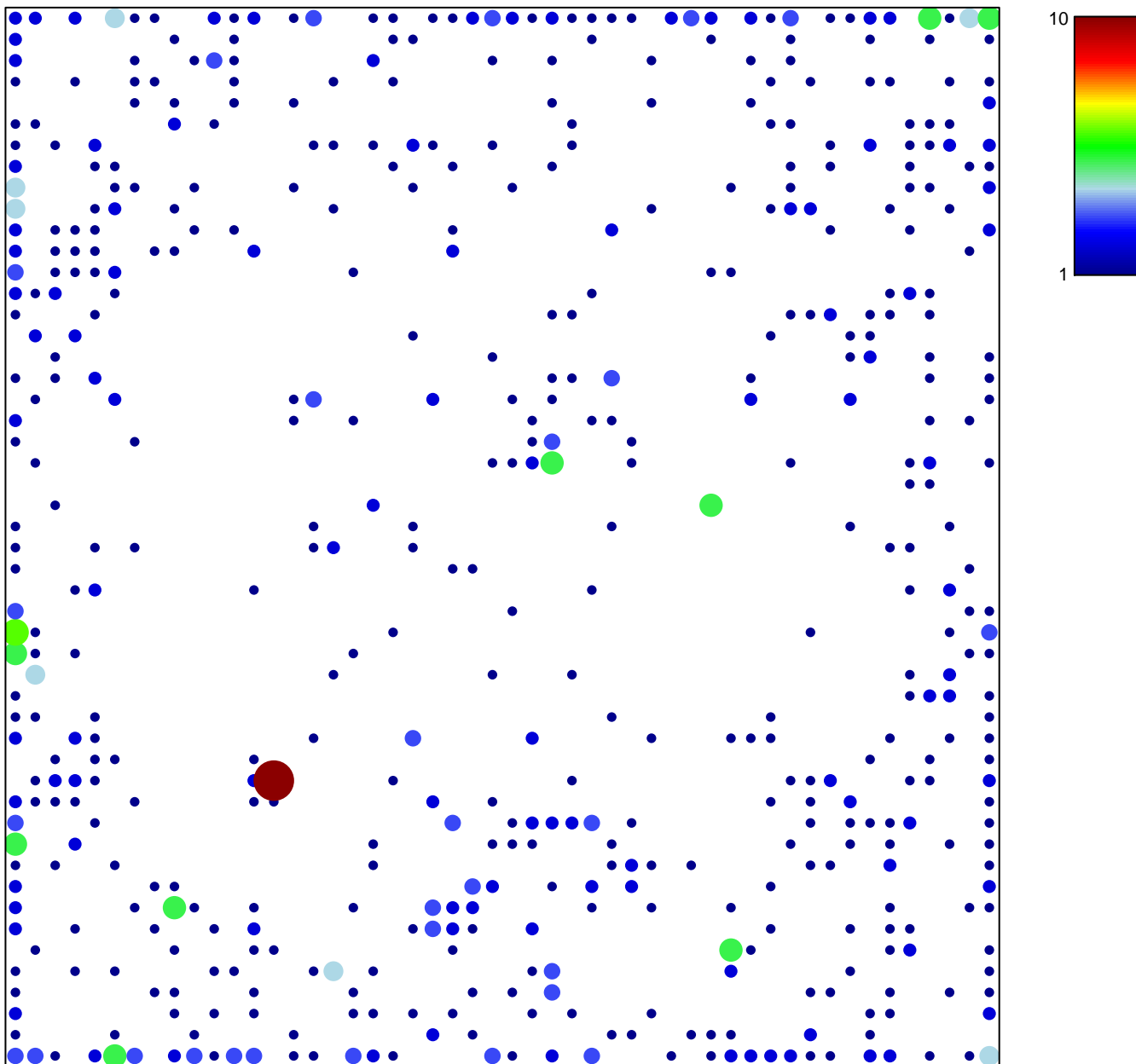
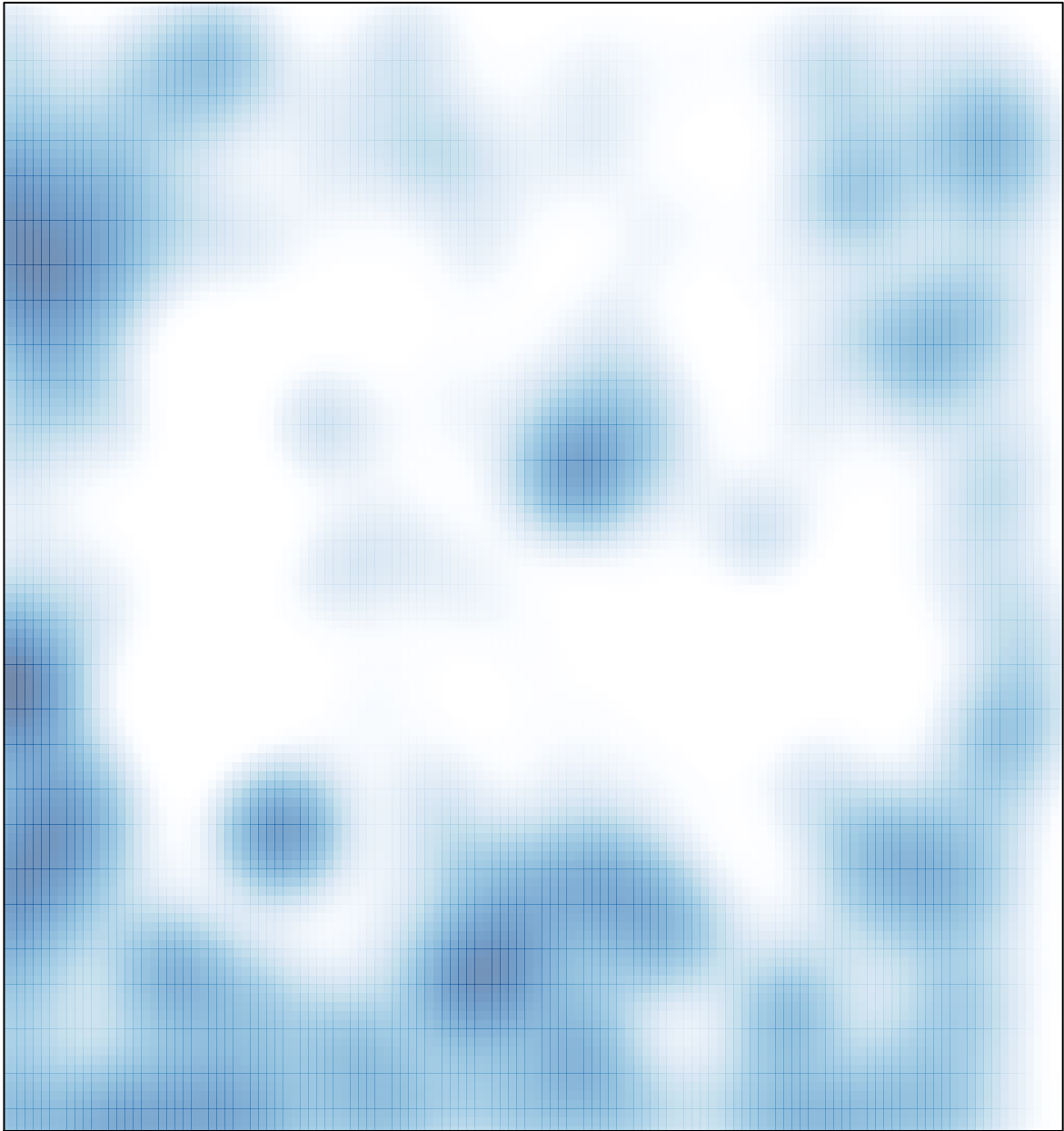


# BYSTRYKH\_HEMATOPOIESIS\_STEM\_CELL\_QTL\_TRANS



# features = 756  
chi-square p = 0.54

# YSTRYKH\_HEMATOPOIESIS\_STEM\_CELL\_QTL\_TRA



# features = 756 , max = 10