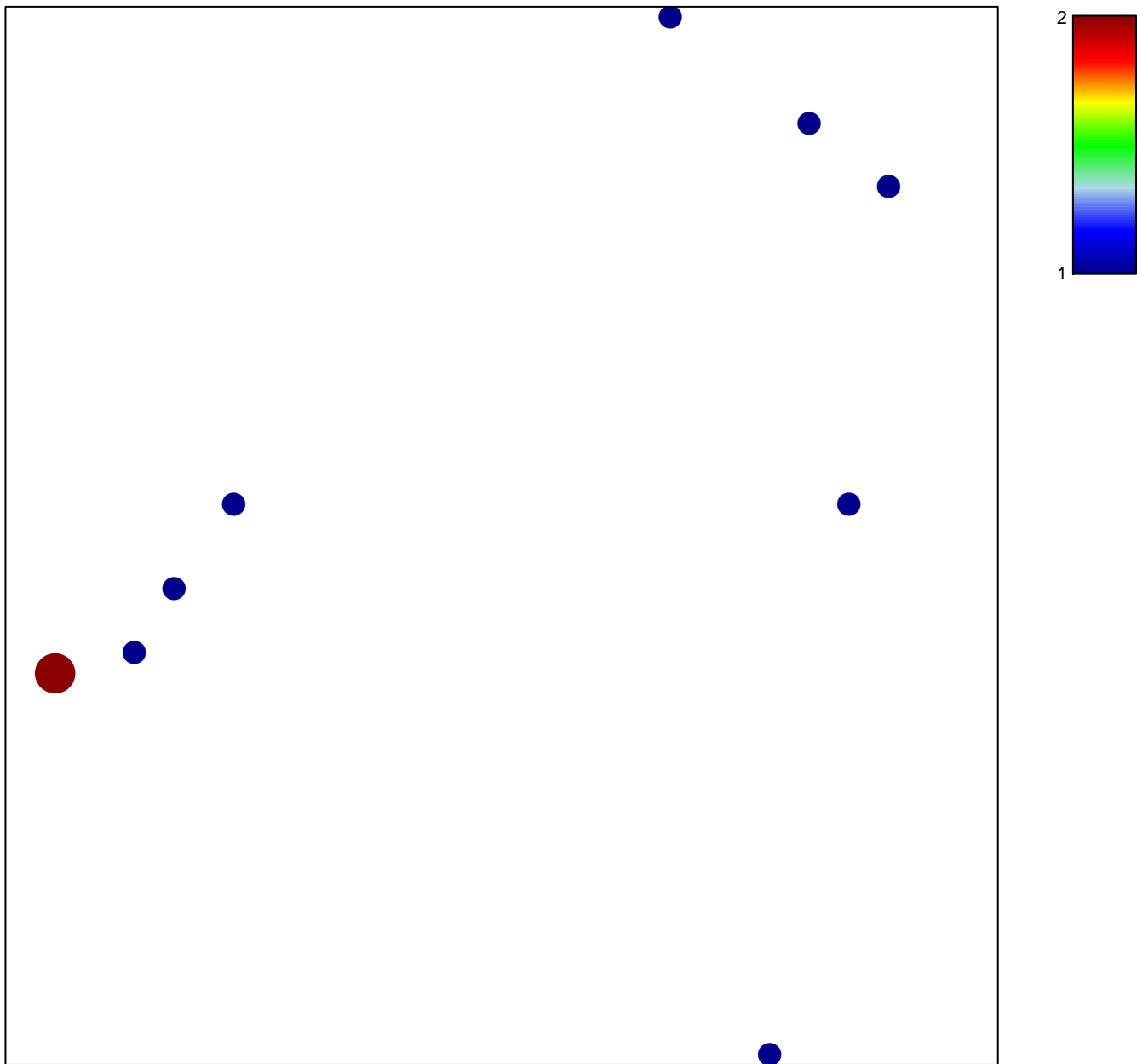
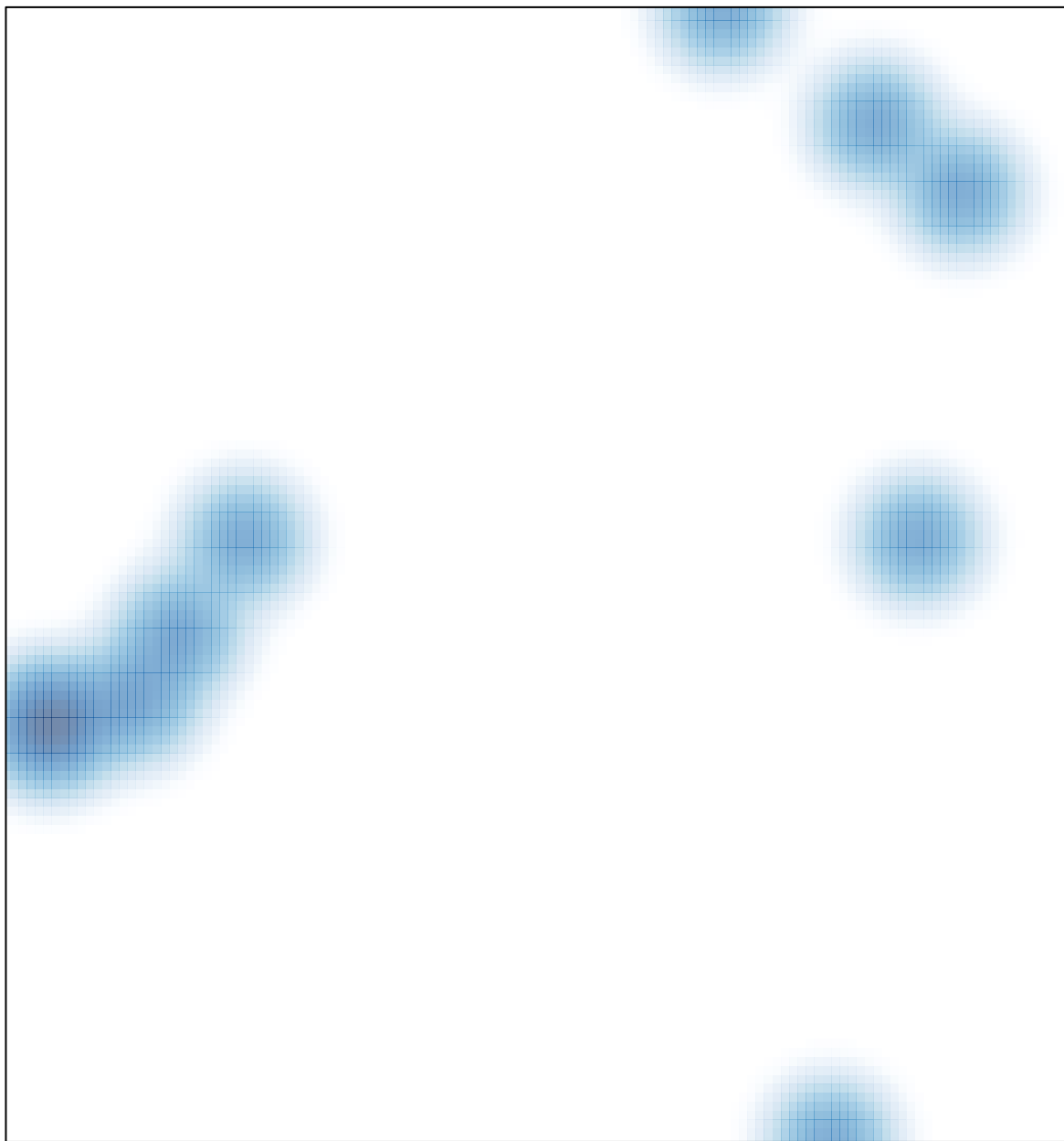


# BASE\_FREE\_SUGAR\_PHOSPHATE\_REMOVAL\_VIA\_THE\_SINGLE\_NUCLEOTIDE\_REPLACE



# features = 10  
chi-square p = 0.83

# \_PHOSPHATE\_REMOVAL\_VIA\_THE\_SINGLE\_NUCL



# features = 10 , max = 2