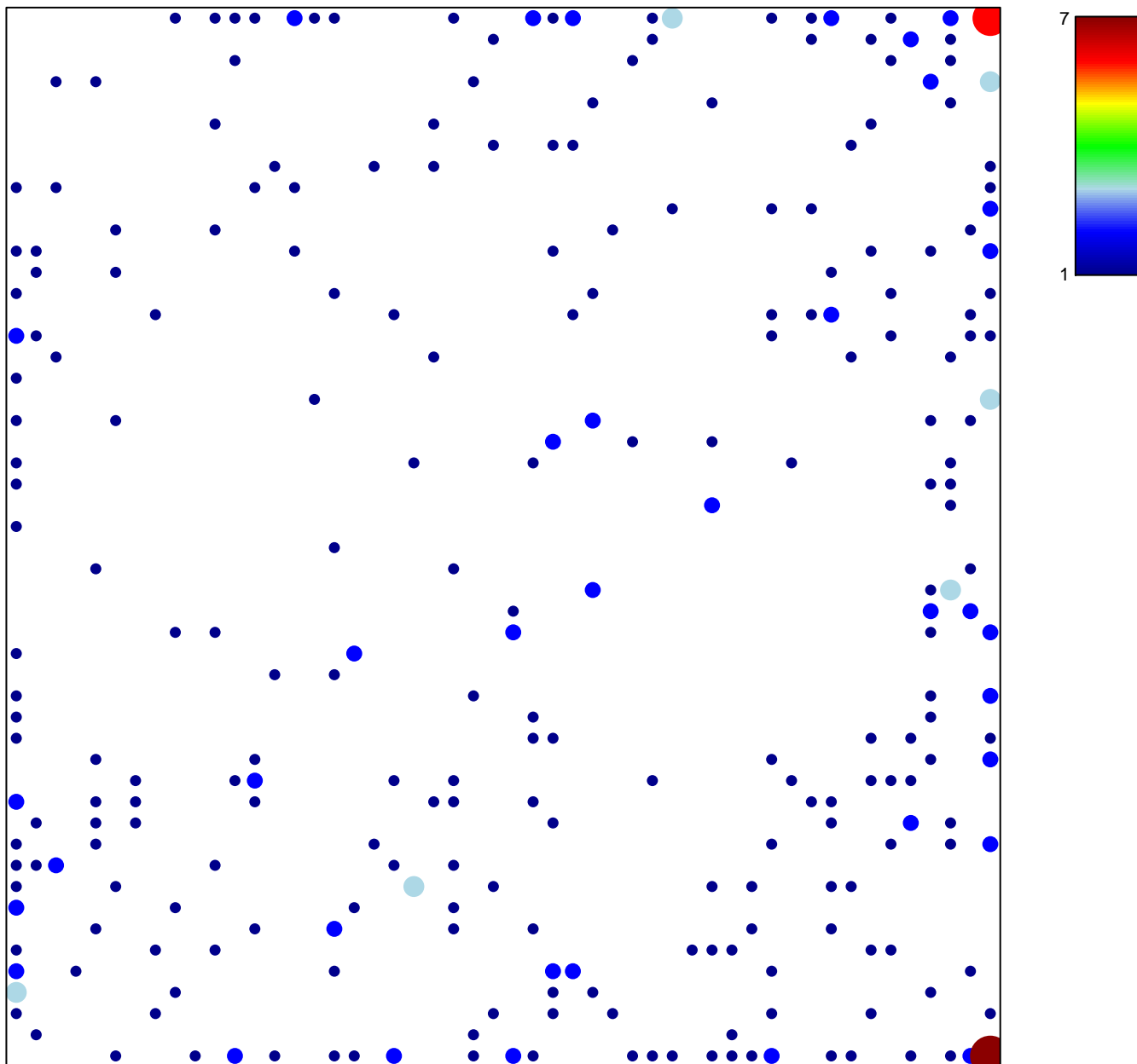
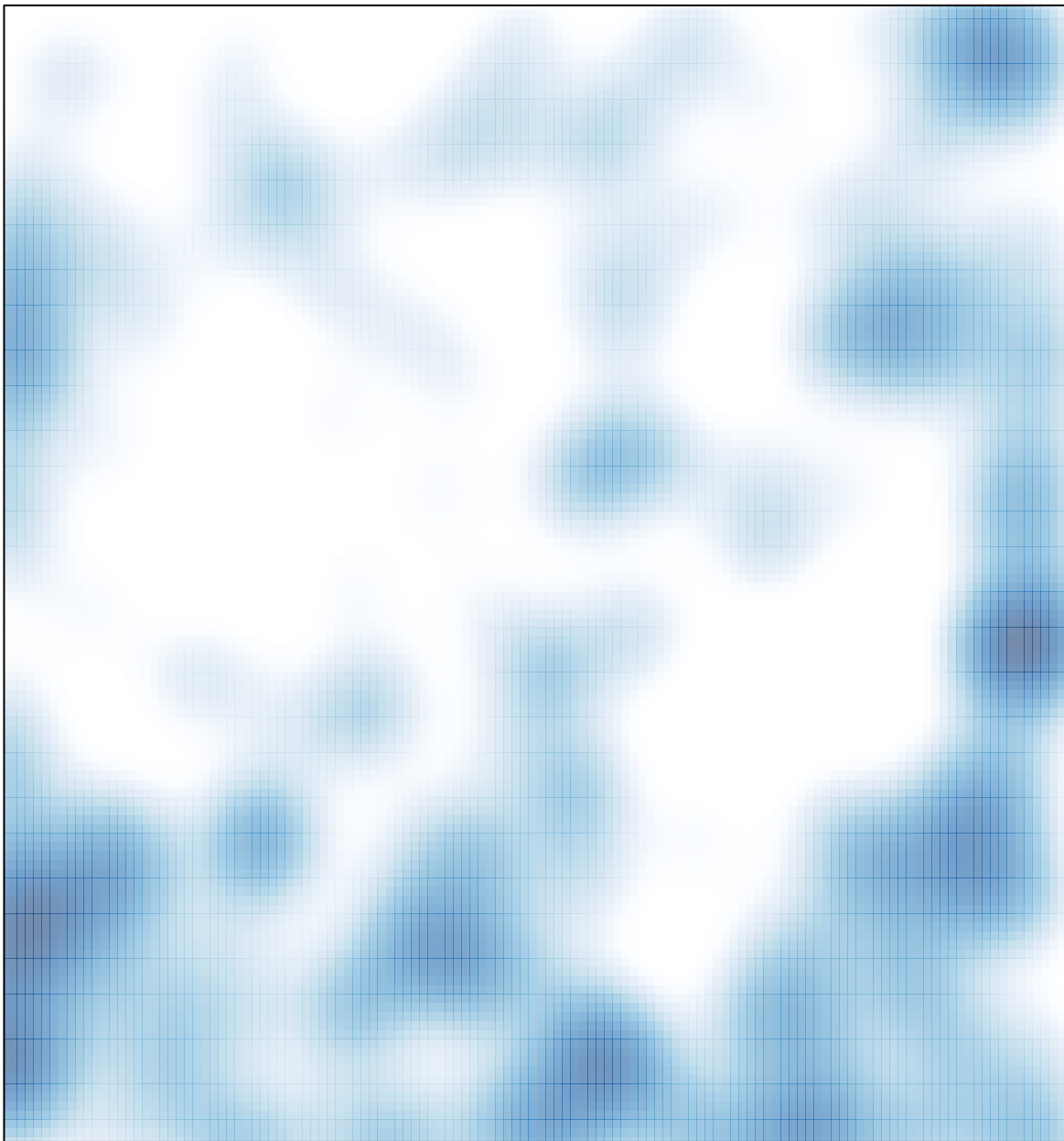


RICKMAN_TUMOR_DIFFERENTIATED_WELL_VS_POORLY_DN



features = 324
chi-square p = 0.64

KMAN_TUMOR_DIFFERENTIATED_WELL_VS_POOR



features = 324 , max = 7