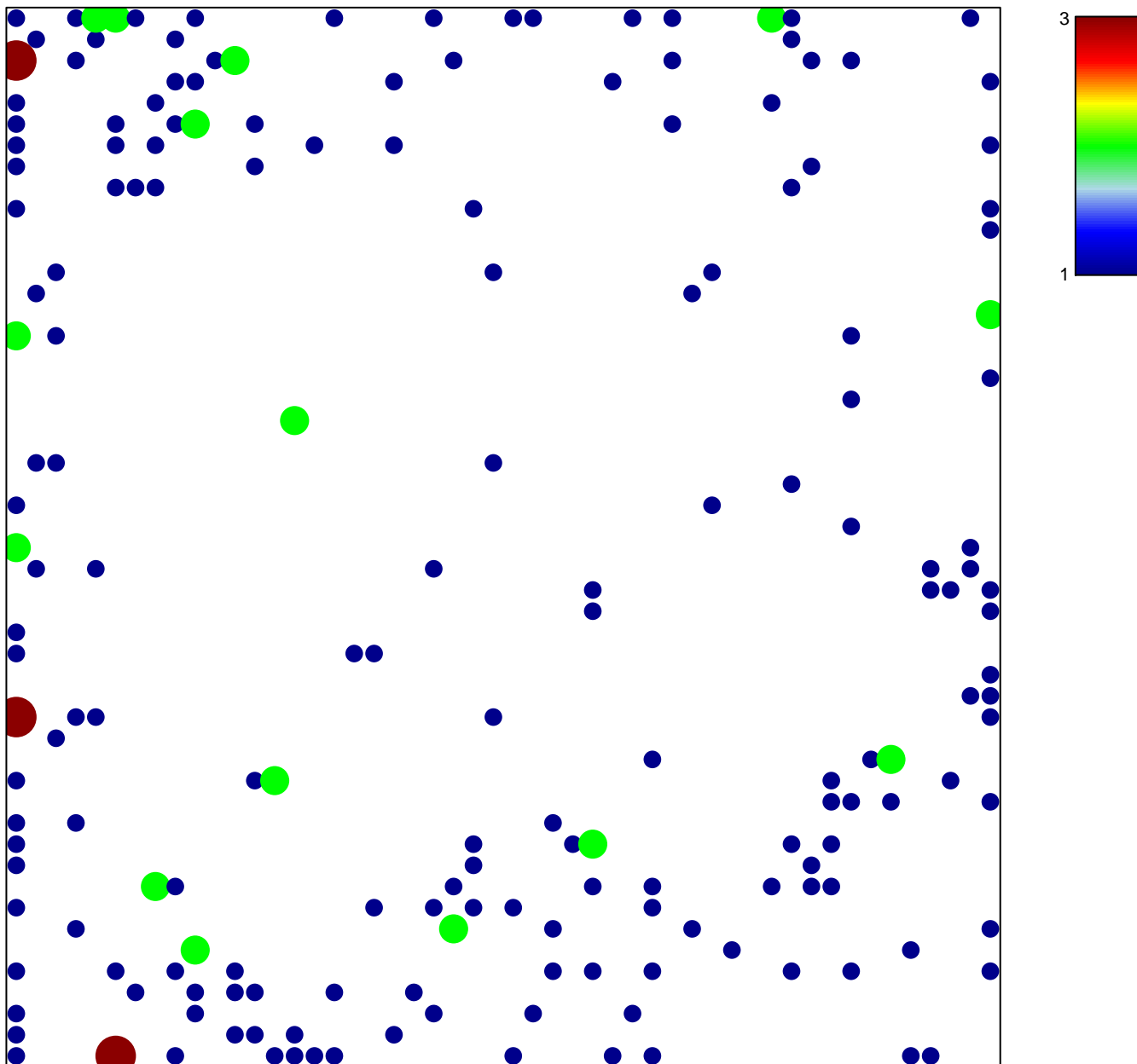
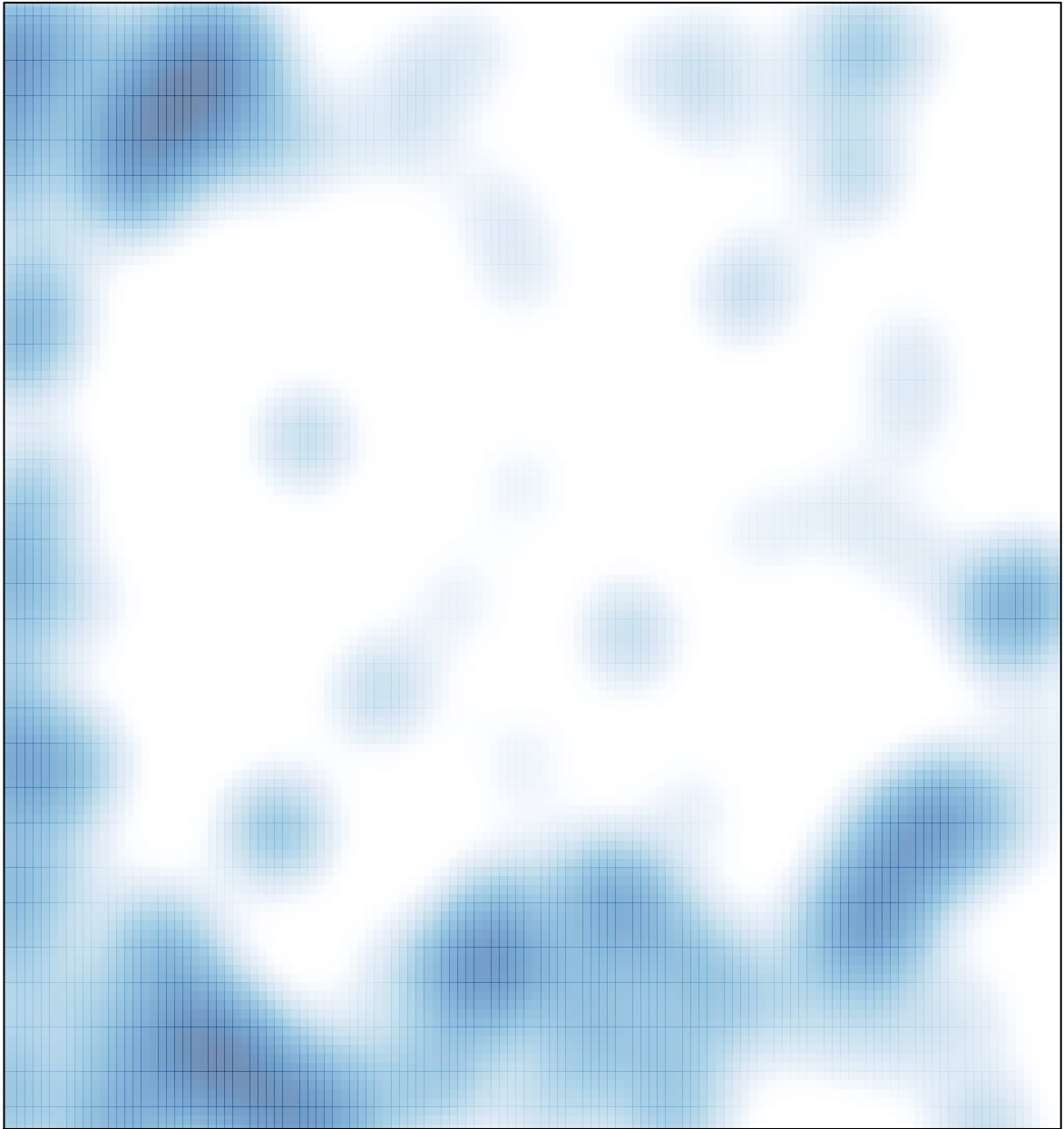


RICKMAN_TUMOR_DIFFERENTIATED_WELL_VS_POORLY_UP



features = 208
chi-square p = 0.71

KMAN_TUMOR_DIFFERENTIATED_WELL_VS_POOR



features = 208 , max = 3