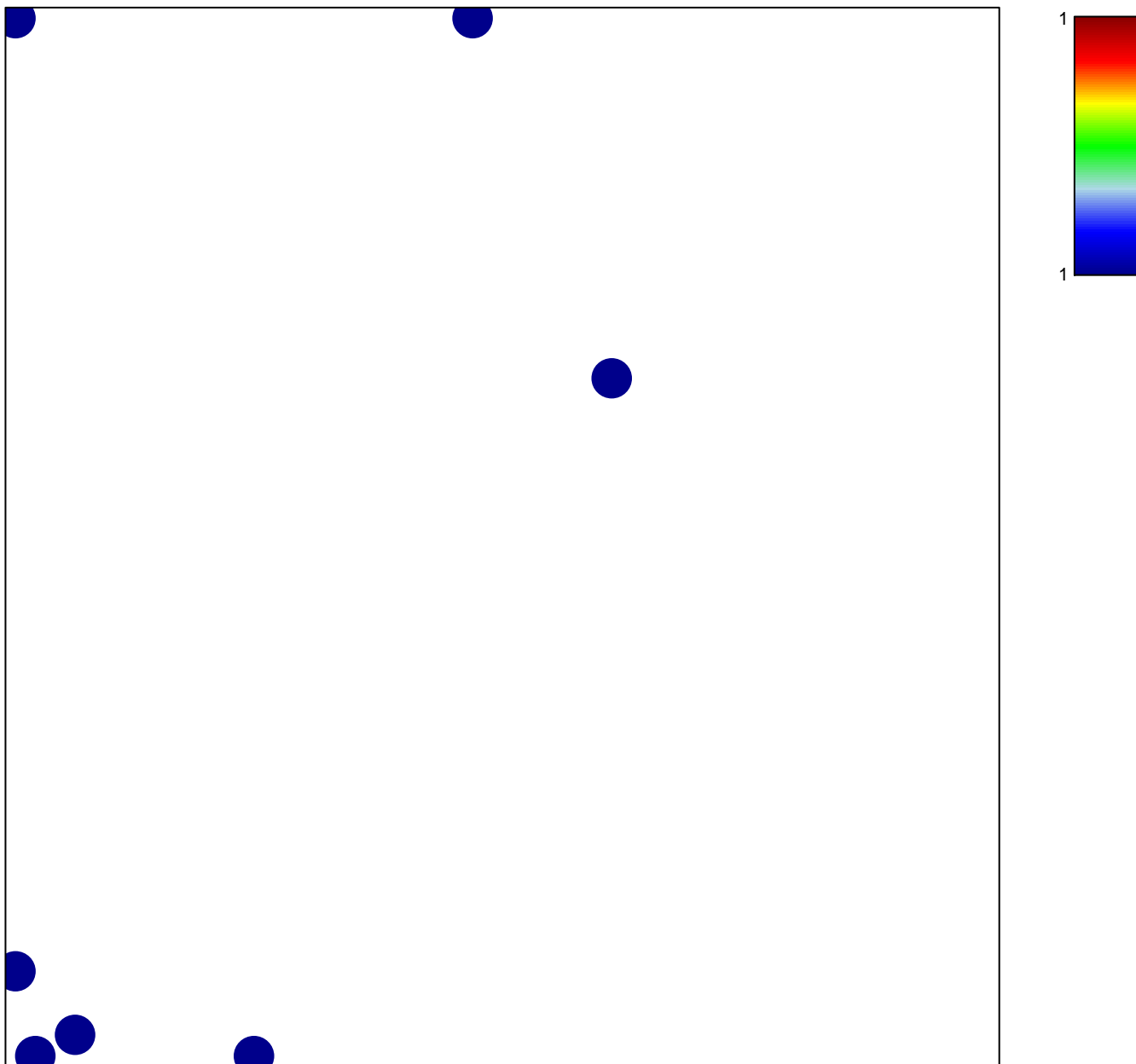
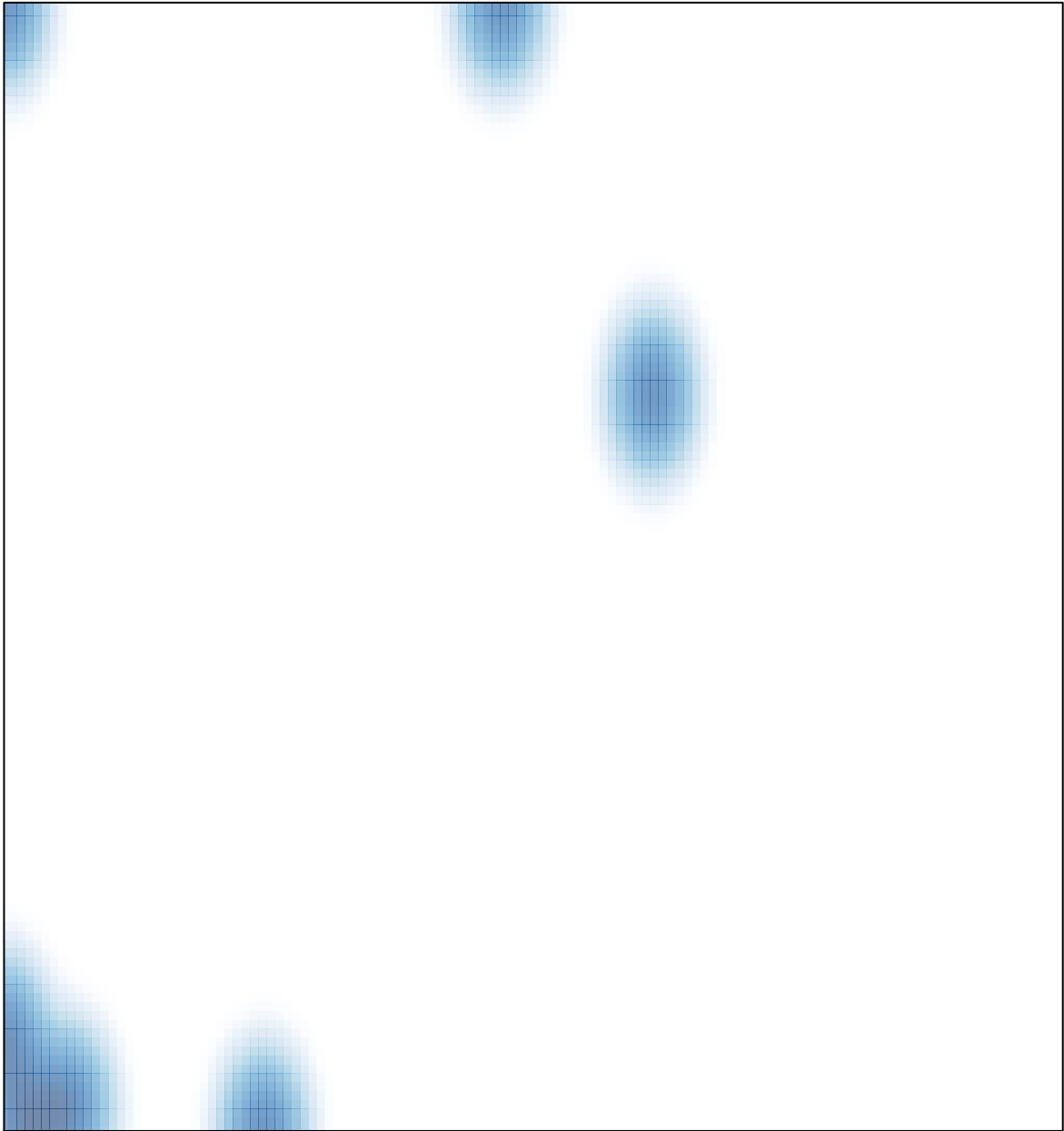


Epithelial-mesenchymal-transition related genes_decreased expression in (GBM) LongTerm



features = 7
chi-square p = 0.82

Normal-transition related genes_decreased expression in



features = 7 , max = 1