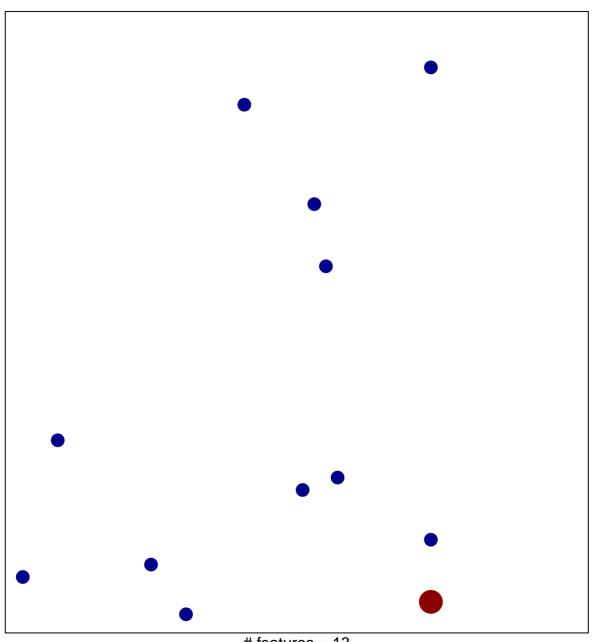
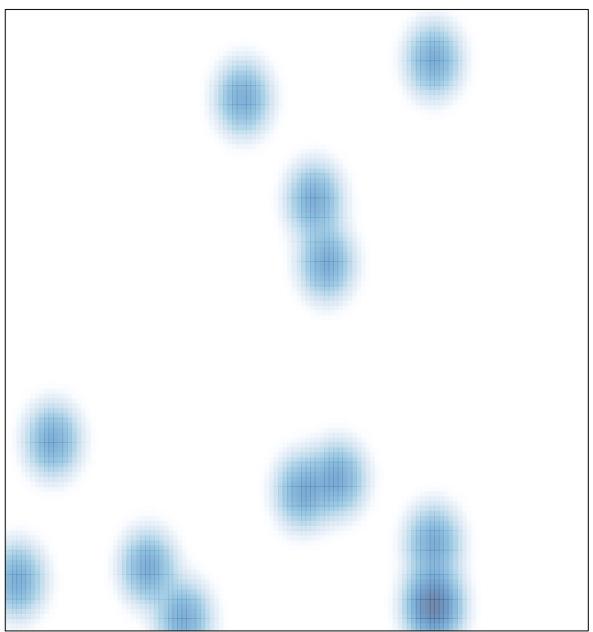
## megakaryocyte development



# features = 13 chi-square p = 0.82

## megakaryocyte development



# features = 13 , max = 2