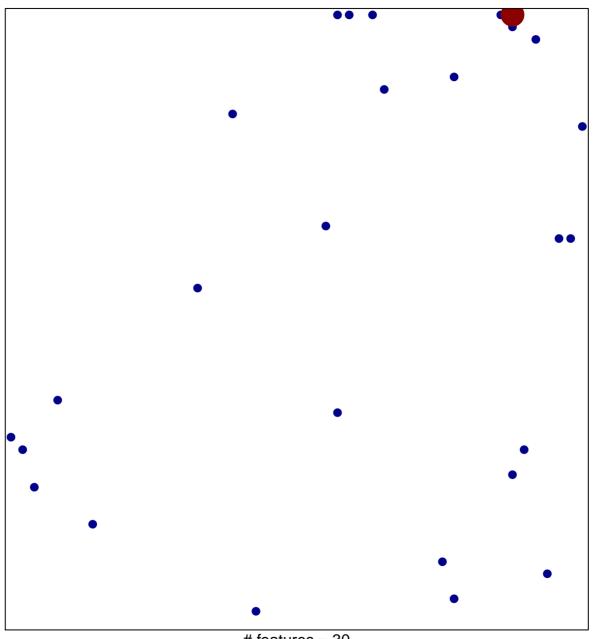
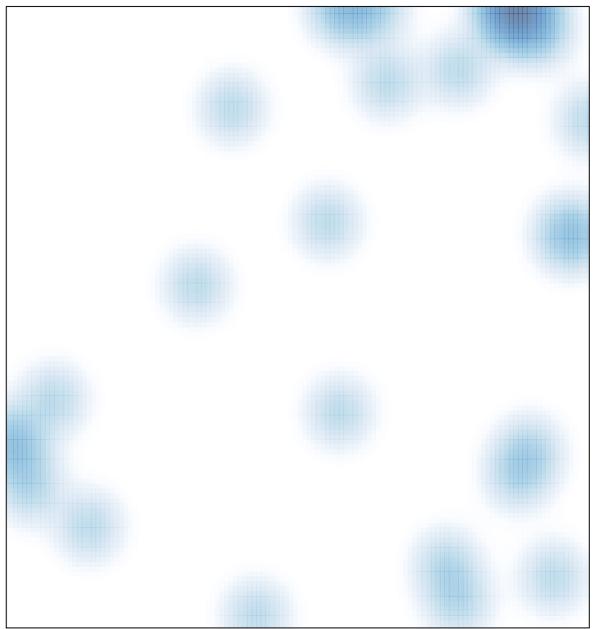
mitotic spindle assembly checkpoint



features = 30 chi-square p = 0.78

mitotic spindle assembly checkpoint



features = 30, max = 4