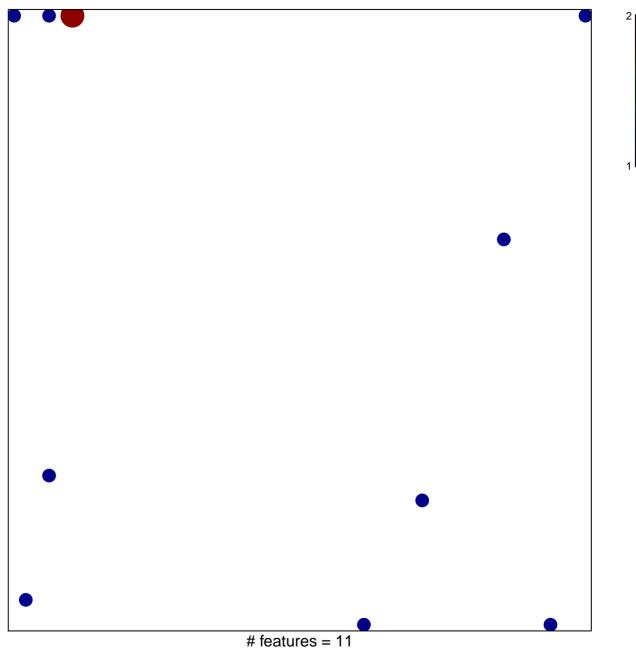
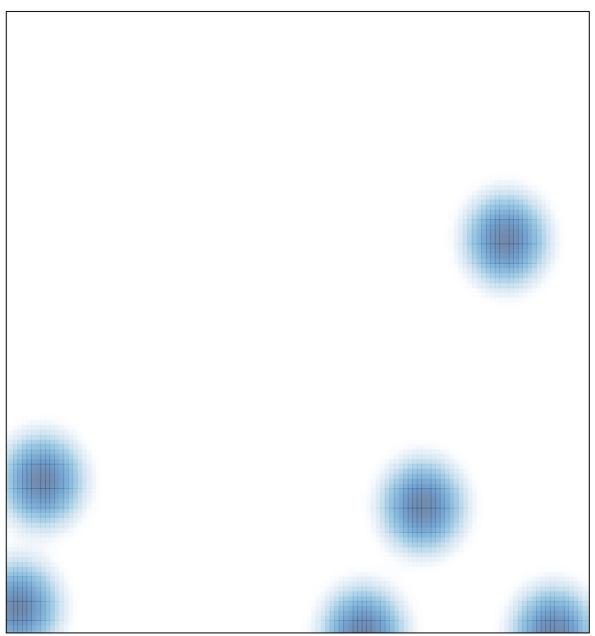
## WU\_APOPTOSIS\_BY\_CDKN1A\_NOT\_VIA\_TP53



# features = 11 chi-square p = 0.78

## WU\_APOPTOSIS\_BY\_CDKN1A\_NOT\_VIA\_TP53



# features = 11, max = 2