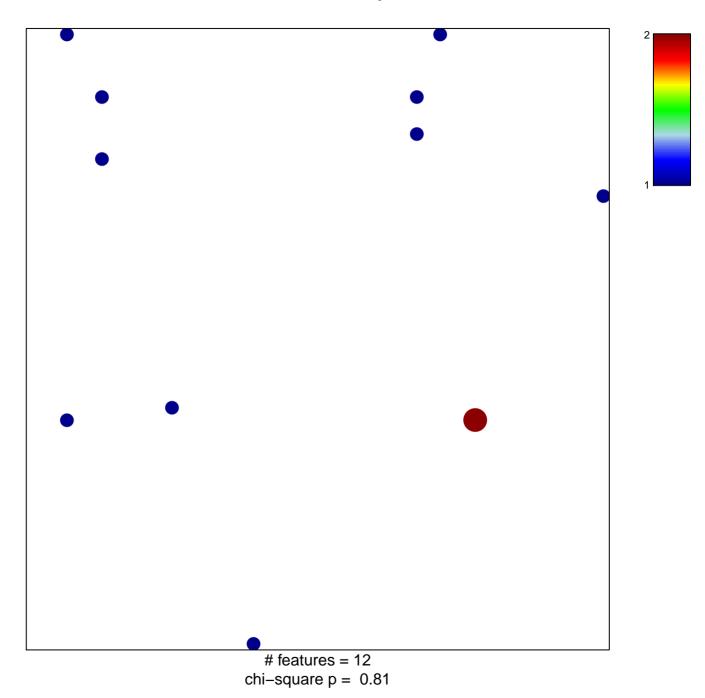
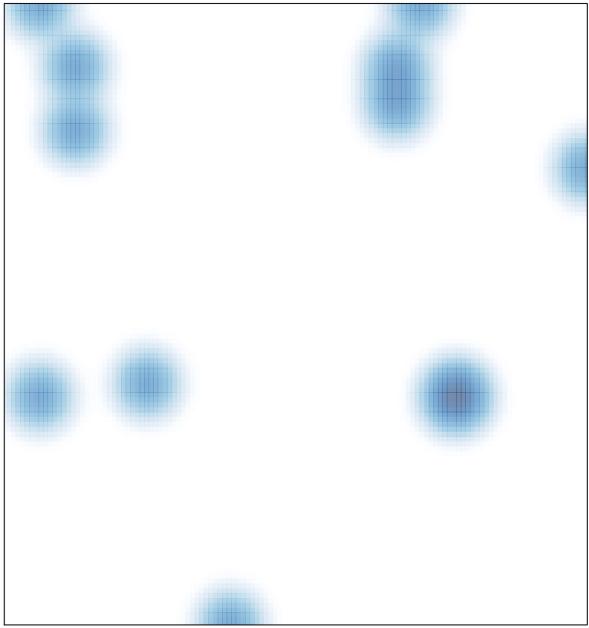
nucleotide metabolic process



## nucleotide metabolic process



# features = 12, max = 2