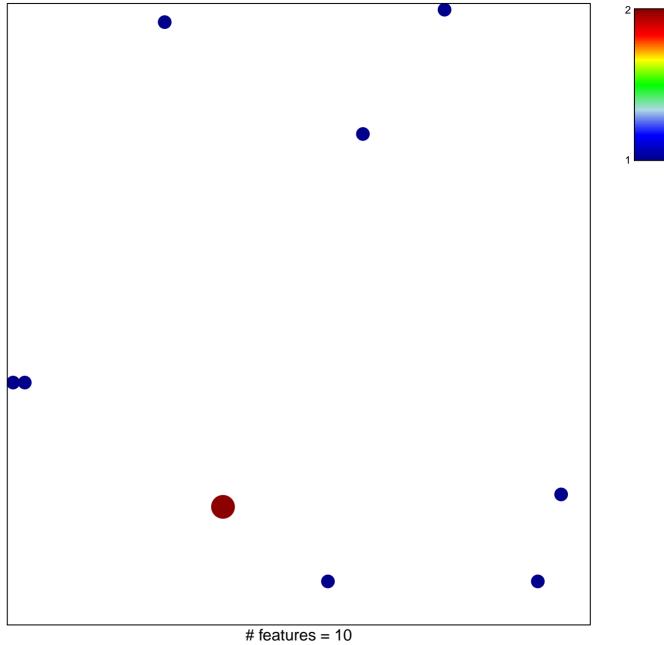
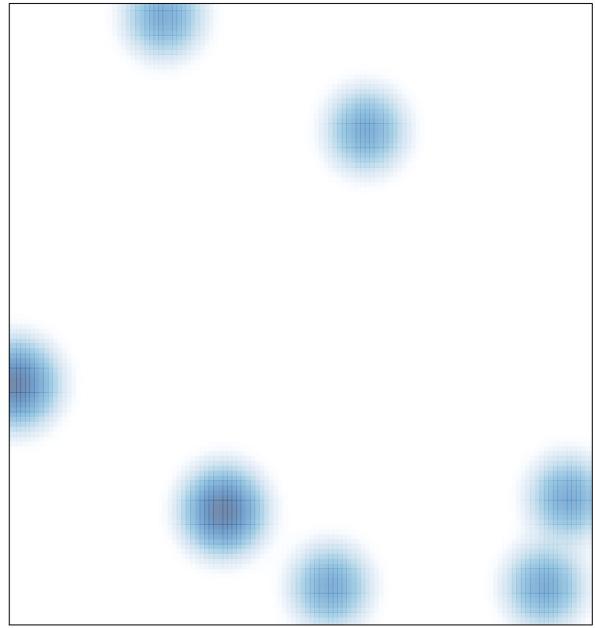
high-density lipoprotein particle remodeling



chi-square p = 0.83

## high-density lipoprotein particle remodeling



# features = 10, max = 2