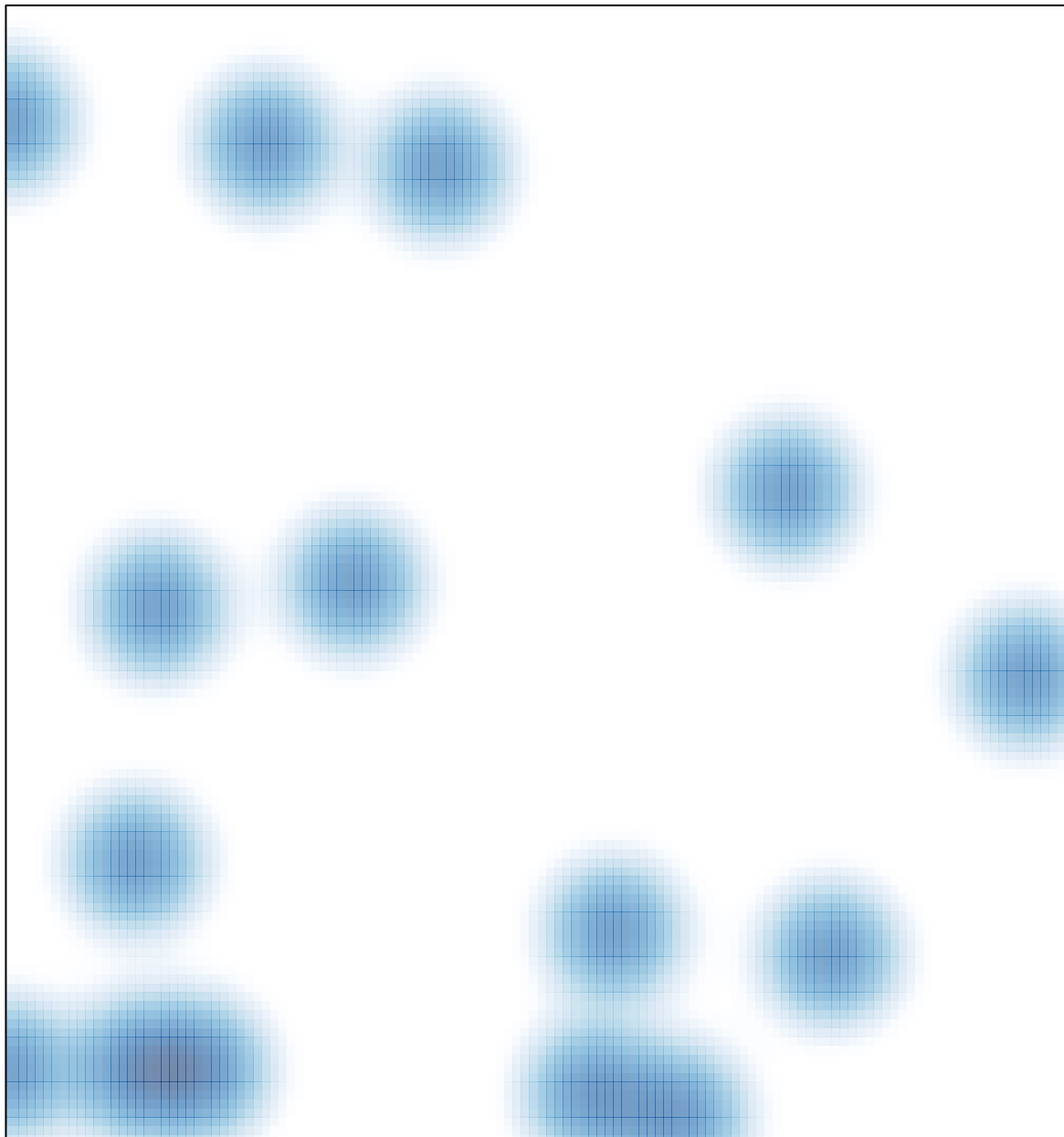


ATP-dependent chromatin remodeling



features = 23 , max = 2