

PAM Metapatients/with_entire_count_matrix/best_k/MyotonicDystrophy_normaliz 50 **-**GSM16564 25 **-**M1656436 0 -Dim2 (26.5%) cluster 2 -25 **-**-50 **-**GSM1*q*56437 50 -50 -100 Dim1 (73.5%)

