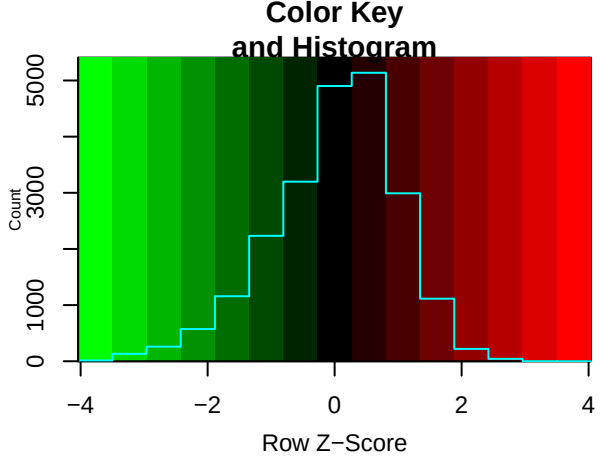
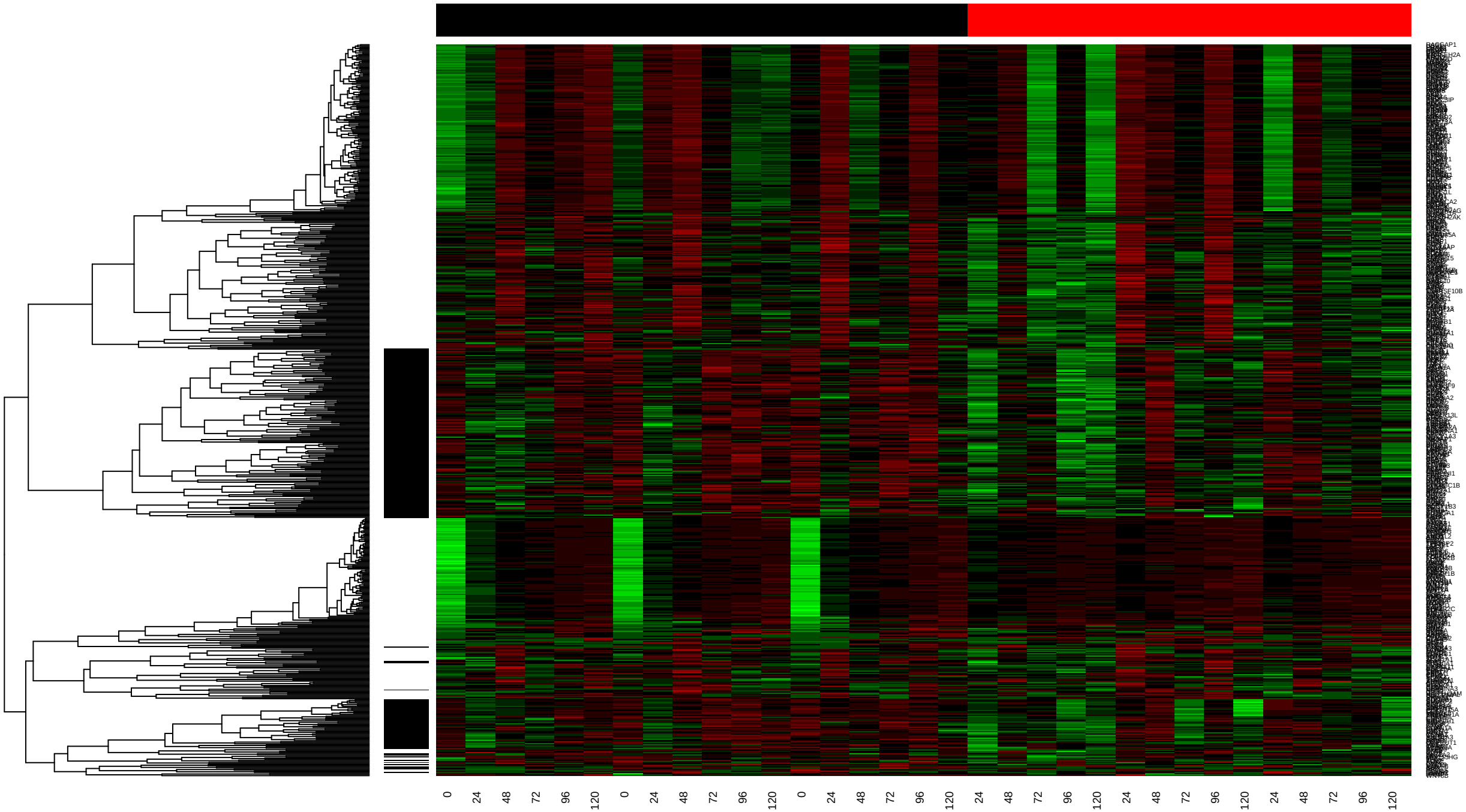
kagohara all simulated data − mitosis genes , half growth overlap with M/S