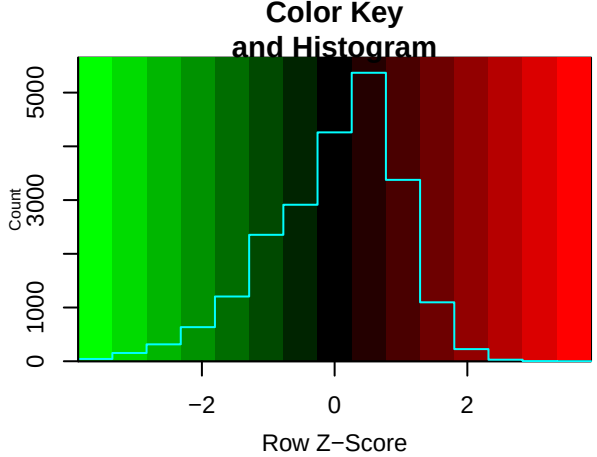
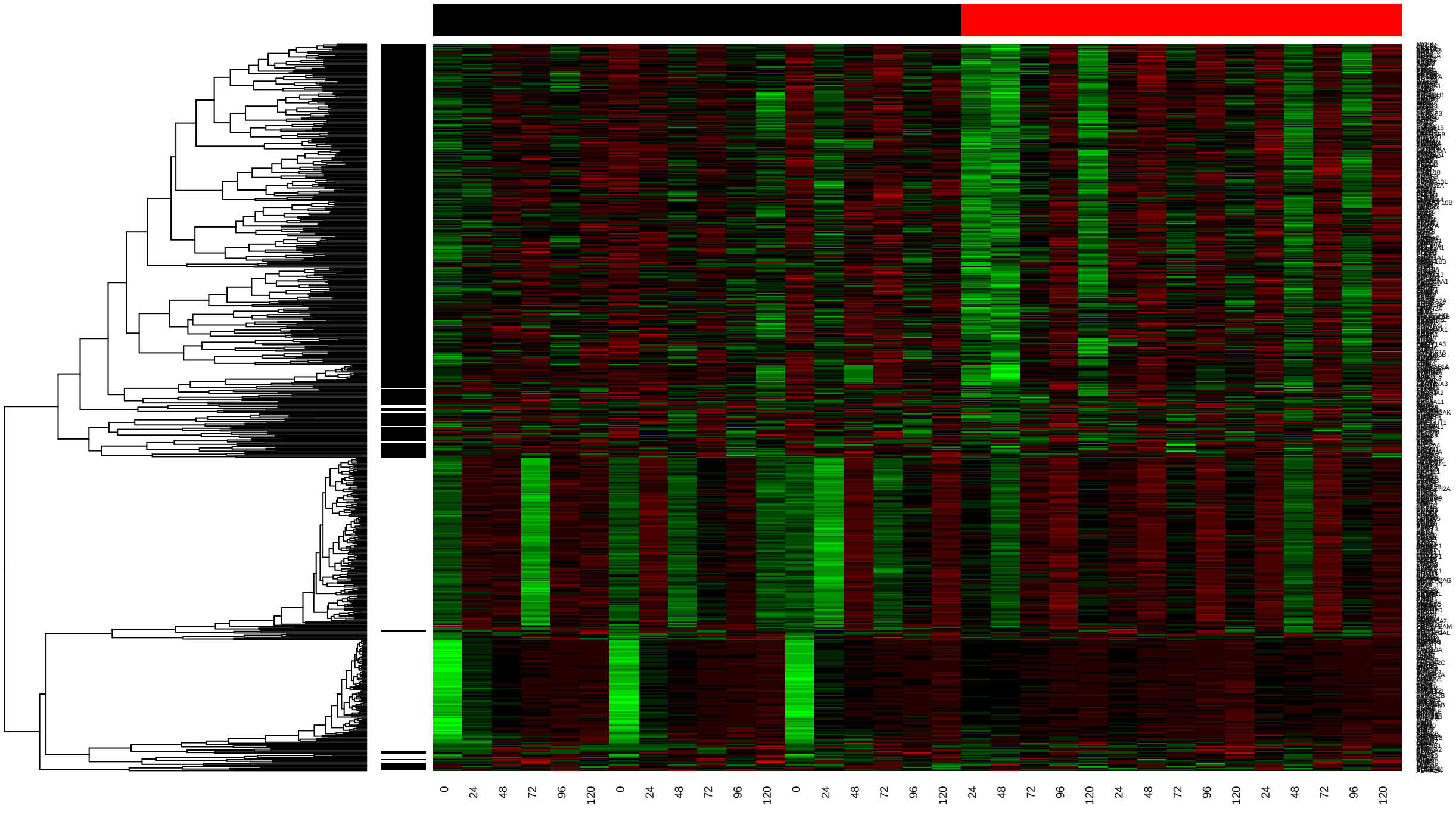
kagohara all simulated data − mitosis genes , full growth overlap with M/S