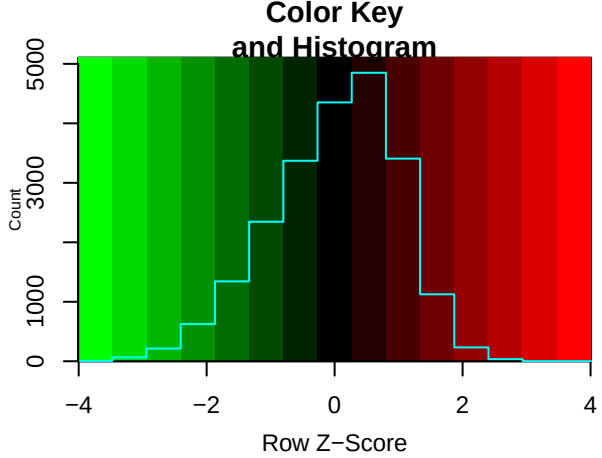
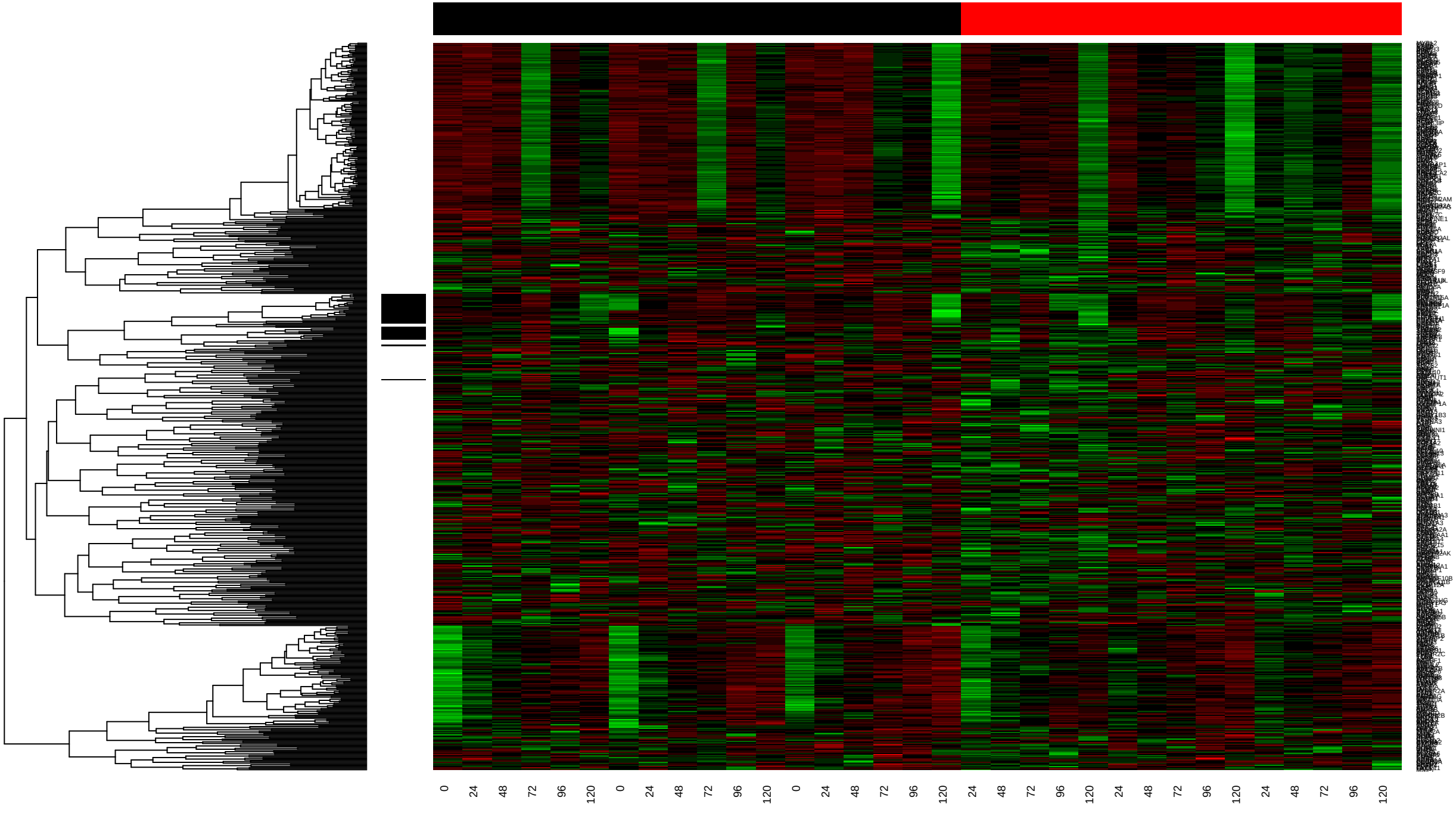
kagohara all simulated data − mitosis genes