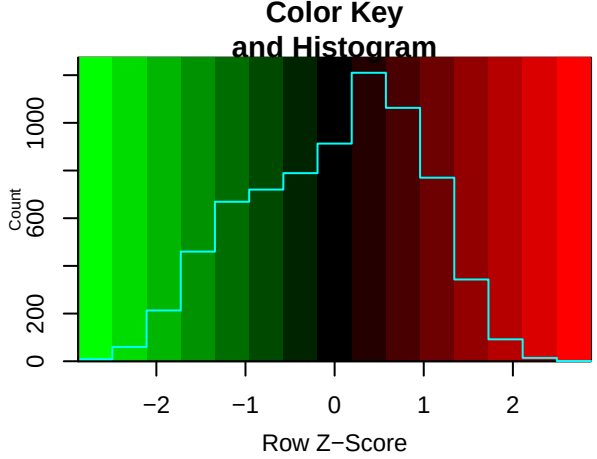
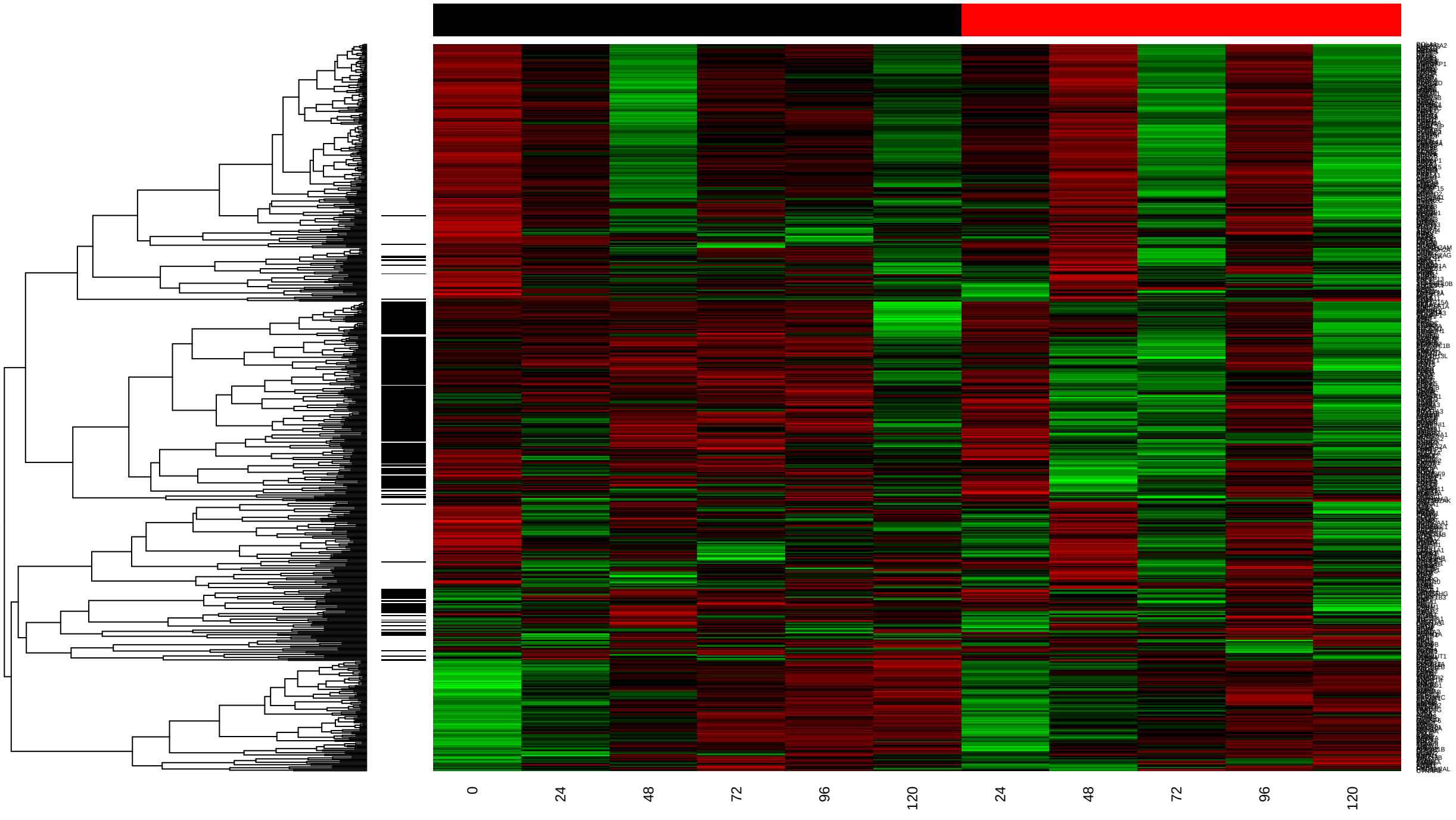
kagohara scc1 simulated data − mitosis genes , half growth overlap with M/S