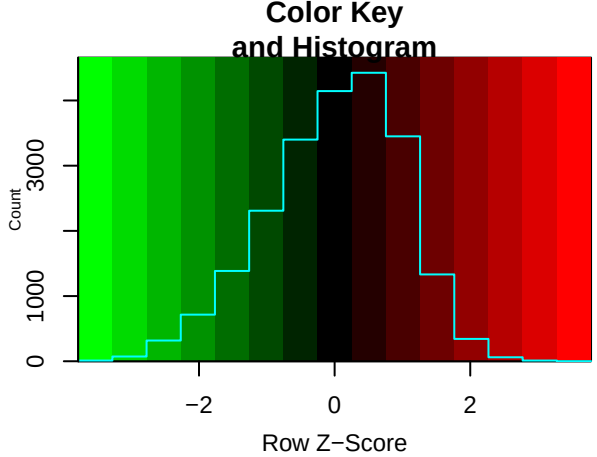
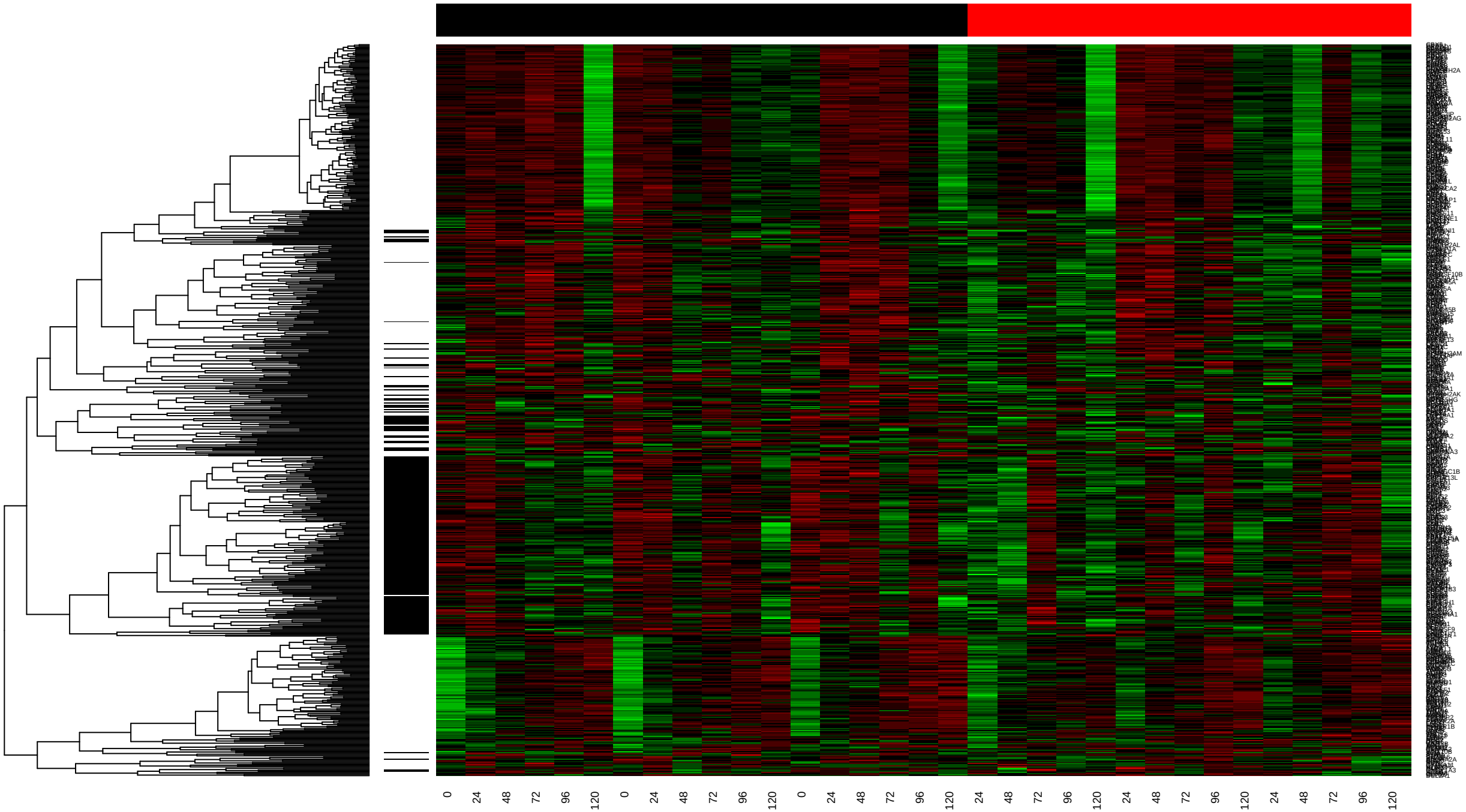
ɟohara all simulated data − mitosis genes , half growth overlap with M/S with cell cycle variance