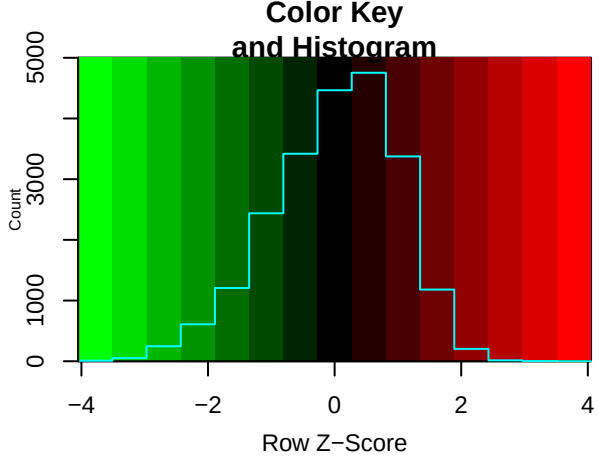
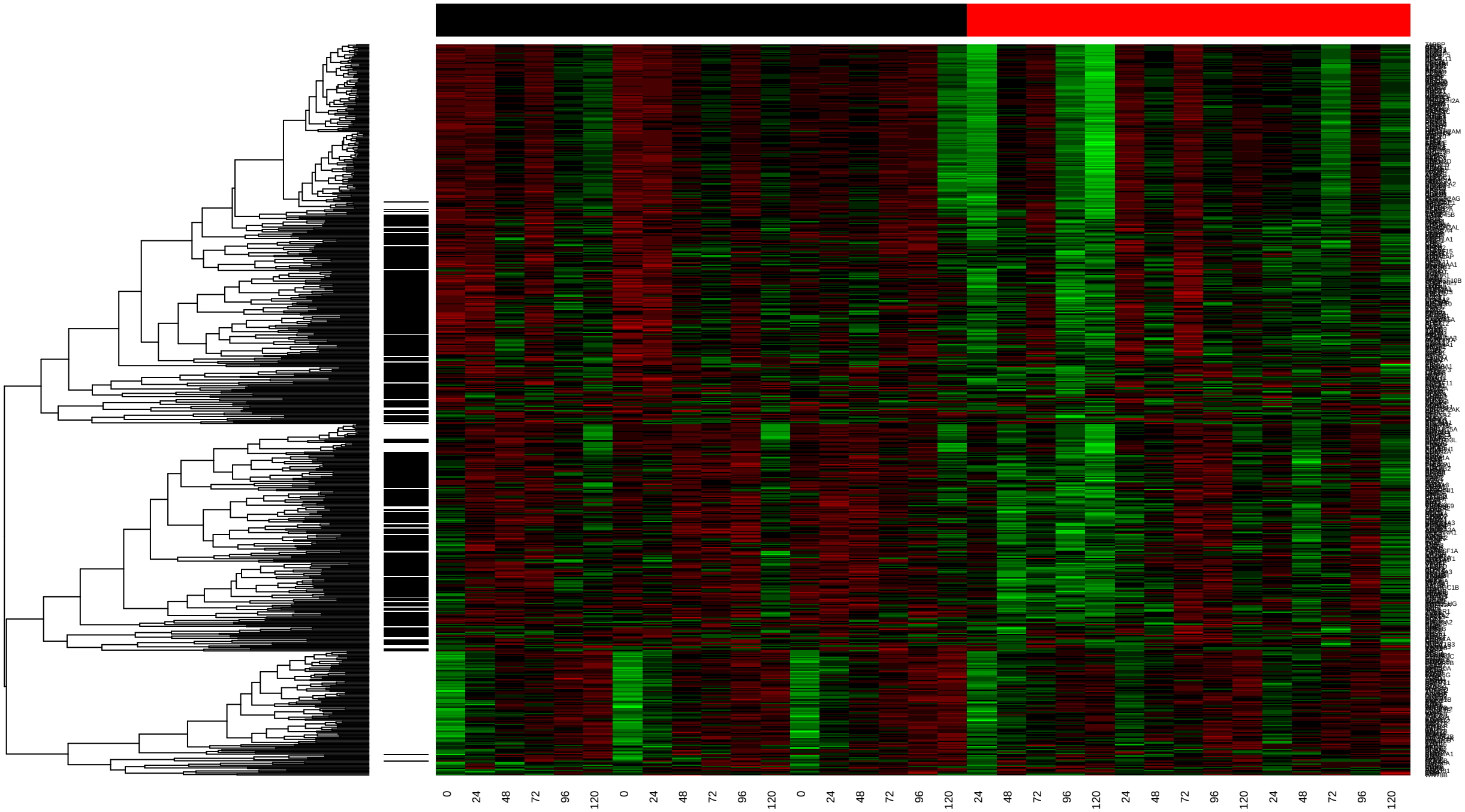
gohara all simulated data − growth genes , half growth overlap with M/S with cell cycle variance