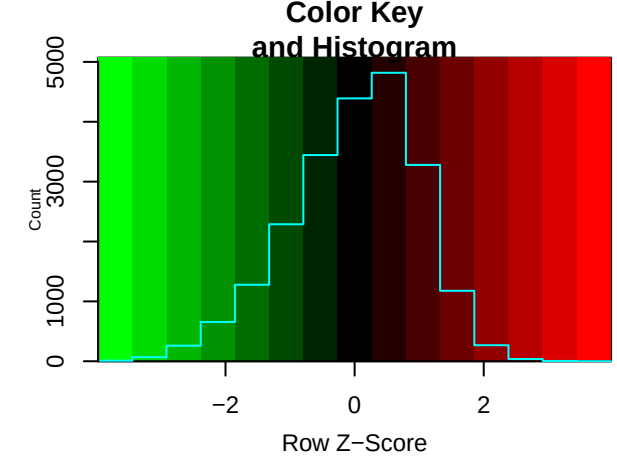
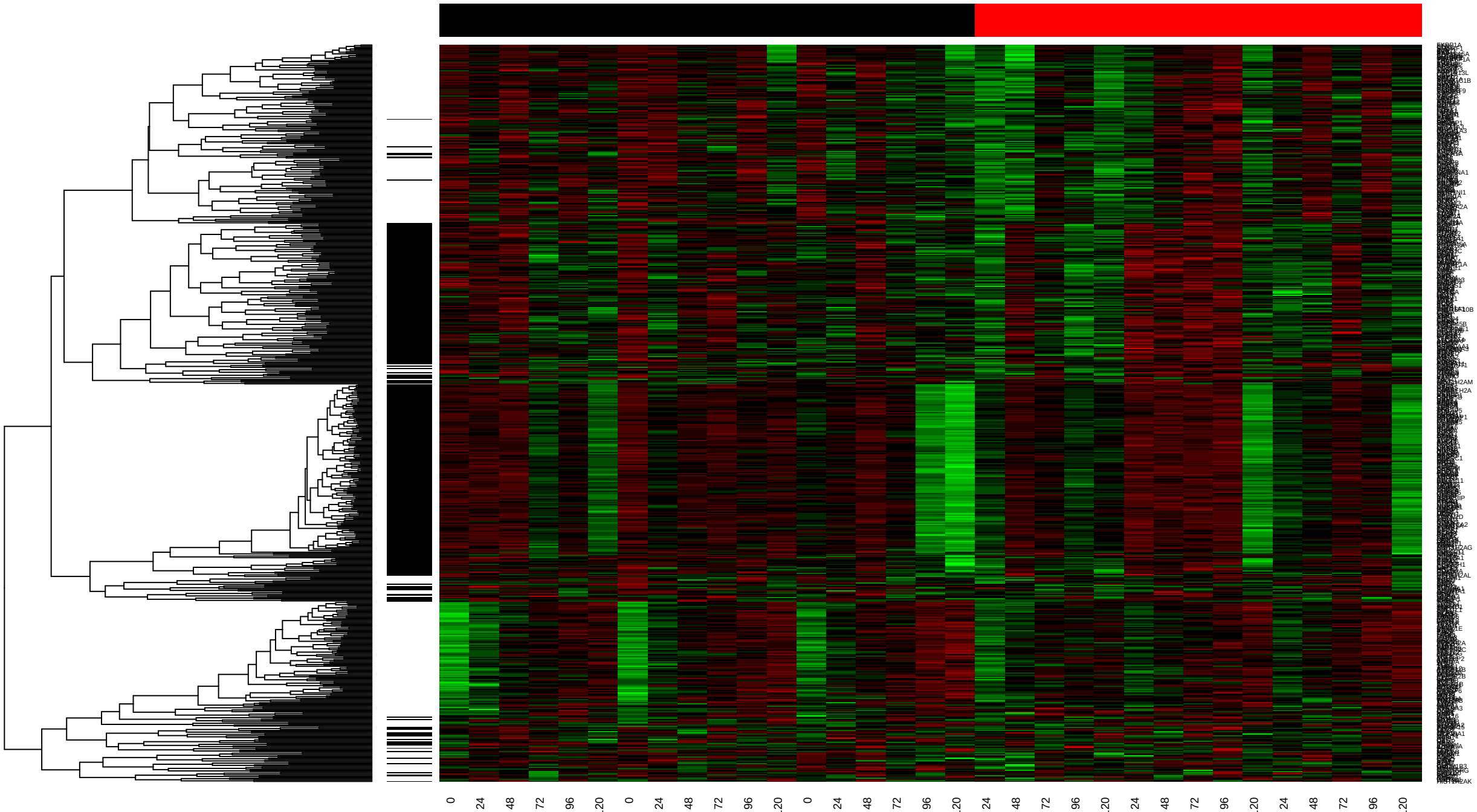
gohara all simulated data − sphase genes , half growth overlap with M/S with cell cycle variance