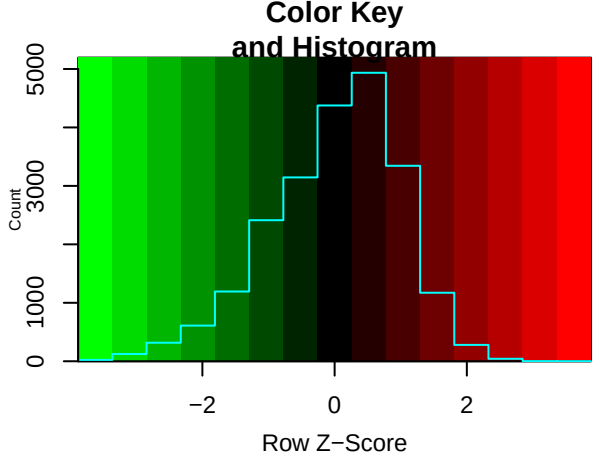
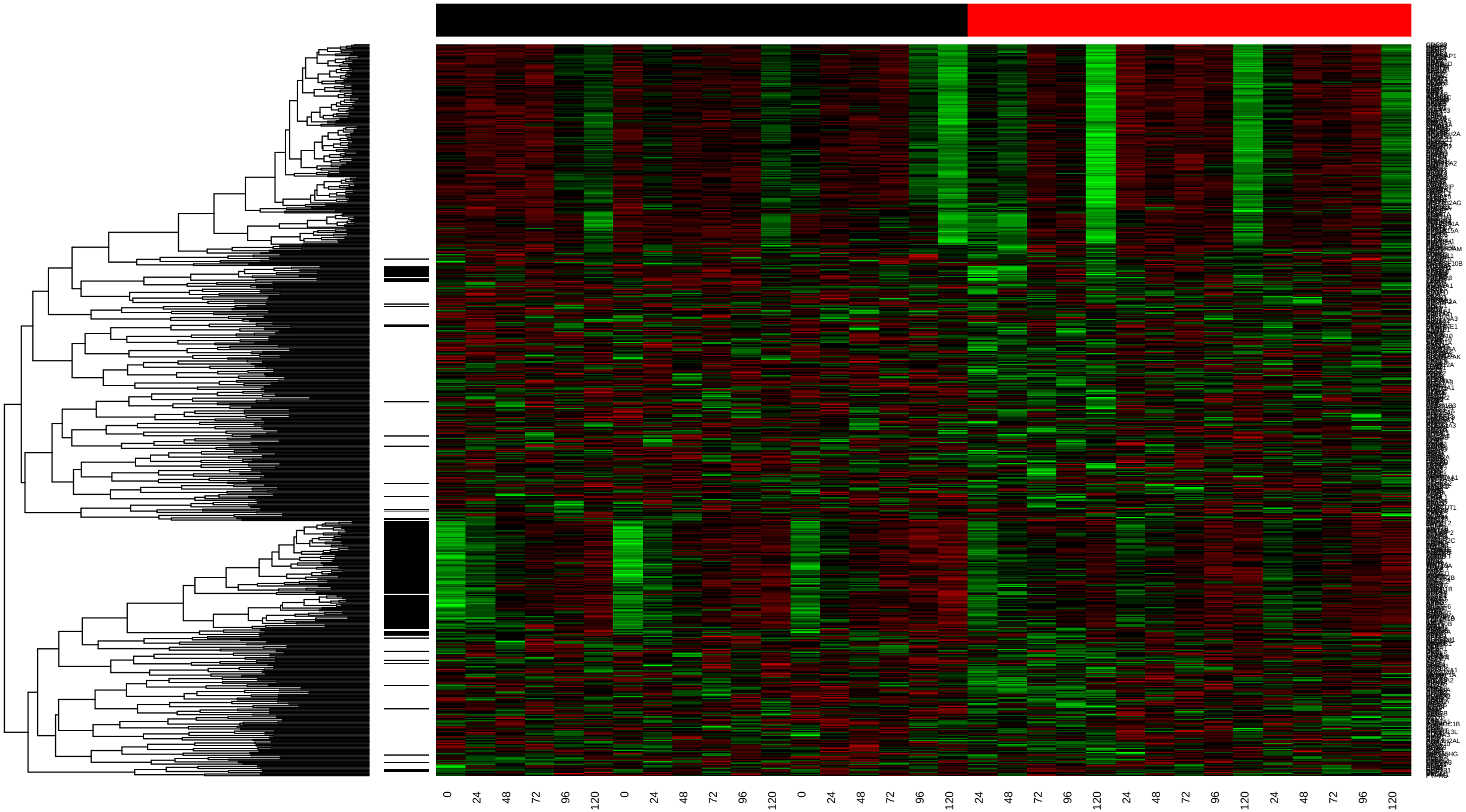
kagohara all simulated data − contact_inhibition genes with cell cycle variance