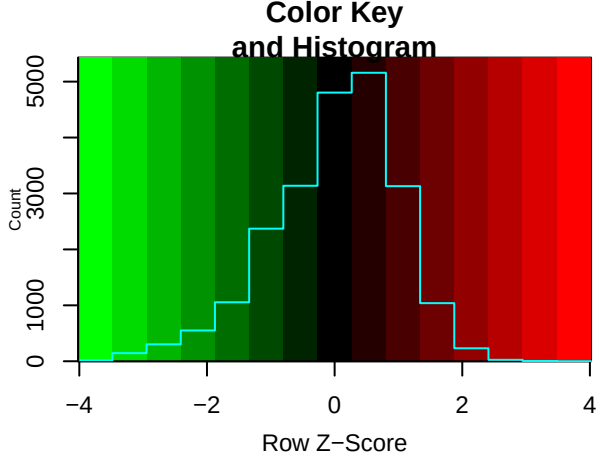
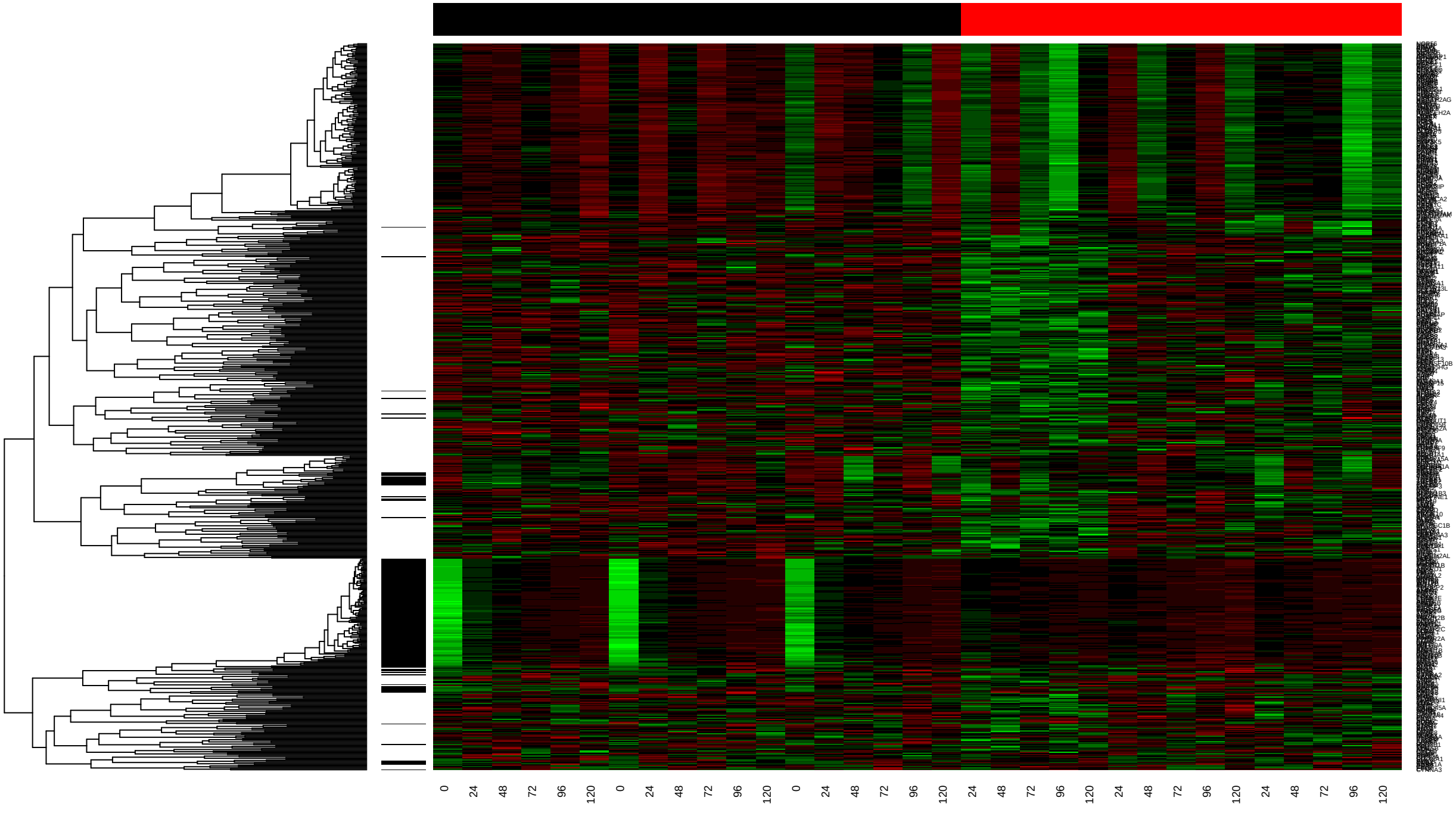
kagohara all simulated data − contact_inhibition genes