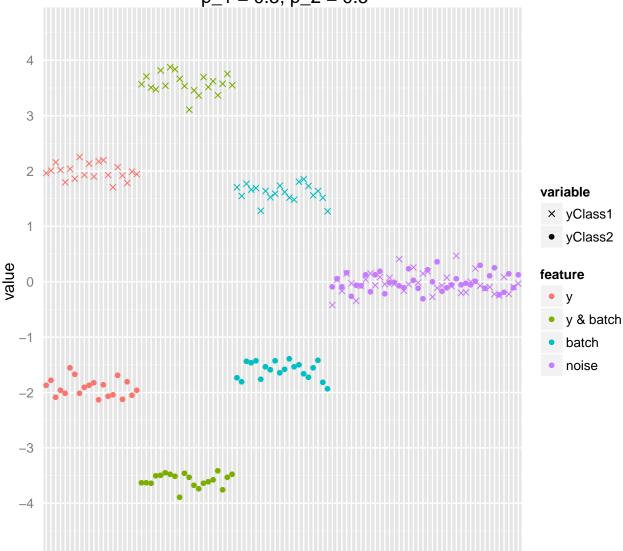
Estimation of primary effects after adjustment pi_1 = 0.9, pi_2 = 0.1 p_1 = 0.5, p_2 = 0.5



Gene