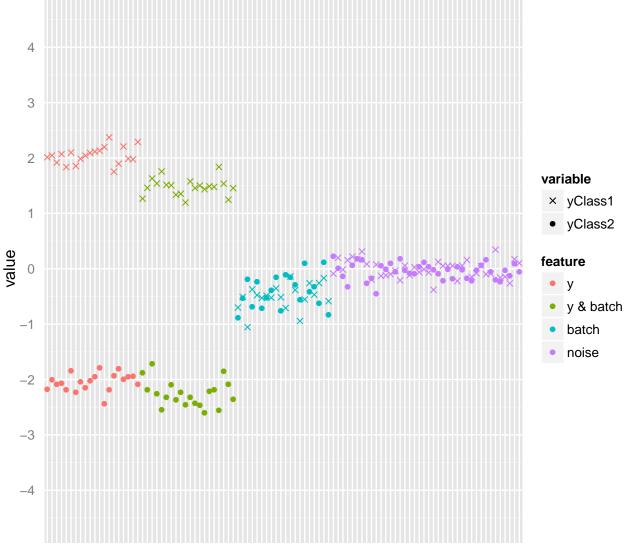
Estimation of primary effects after adjustment pi_1 = 0.4, pi_2 = 0.4 p_1 = 0.4, p_2 = 0.6



Gene