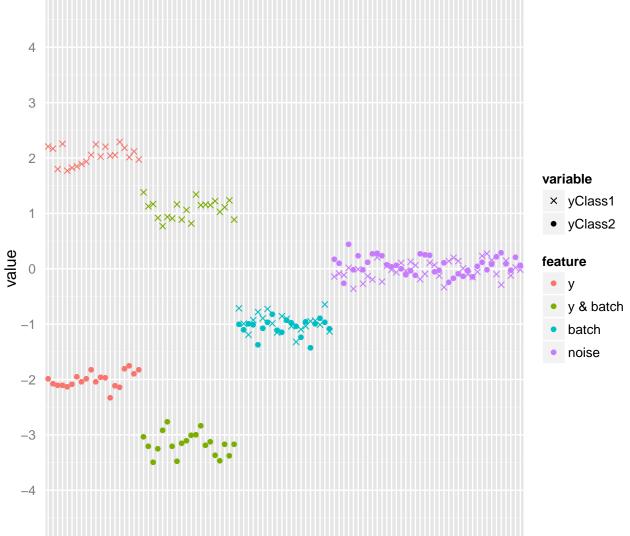
Estimation of primary effects after adjustment pi_1 = 0.4, pi_2 = 0.1 p_1 = 0.25, p_2 = 0.75



Gene