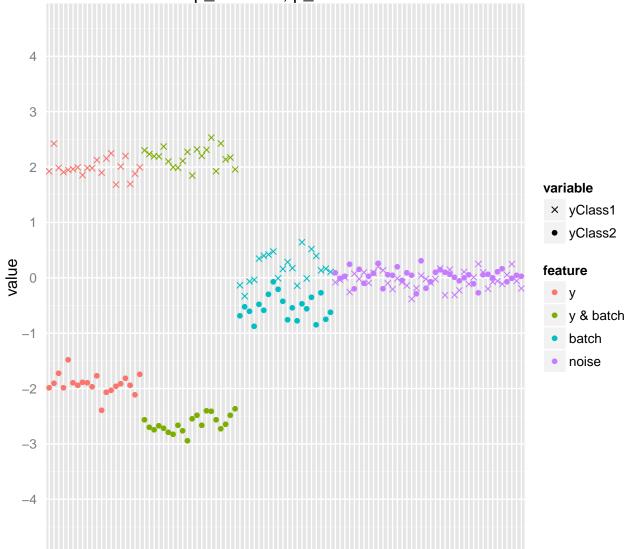
Estimation of primary effects before adjustment pi_1 = 0.55, pi_2 = 0.35 p_1 = 0.45, p_2 = 0.55



Gene