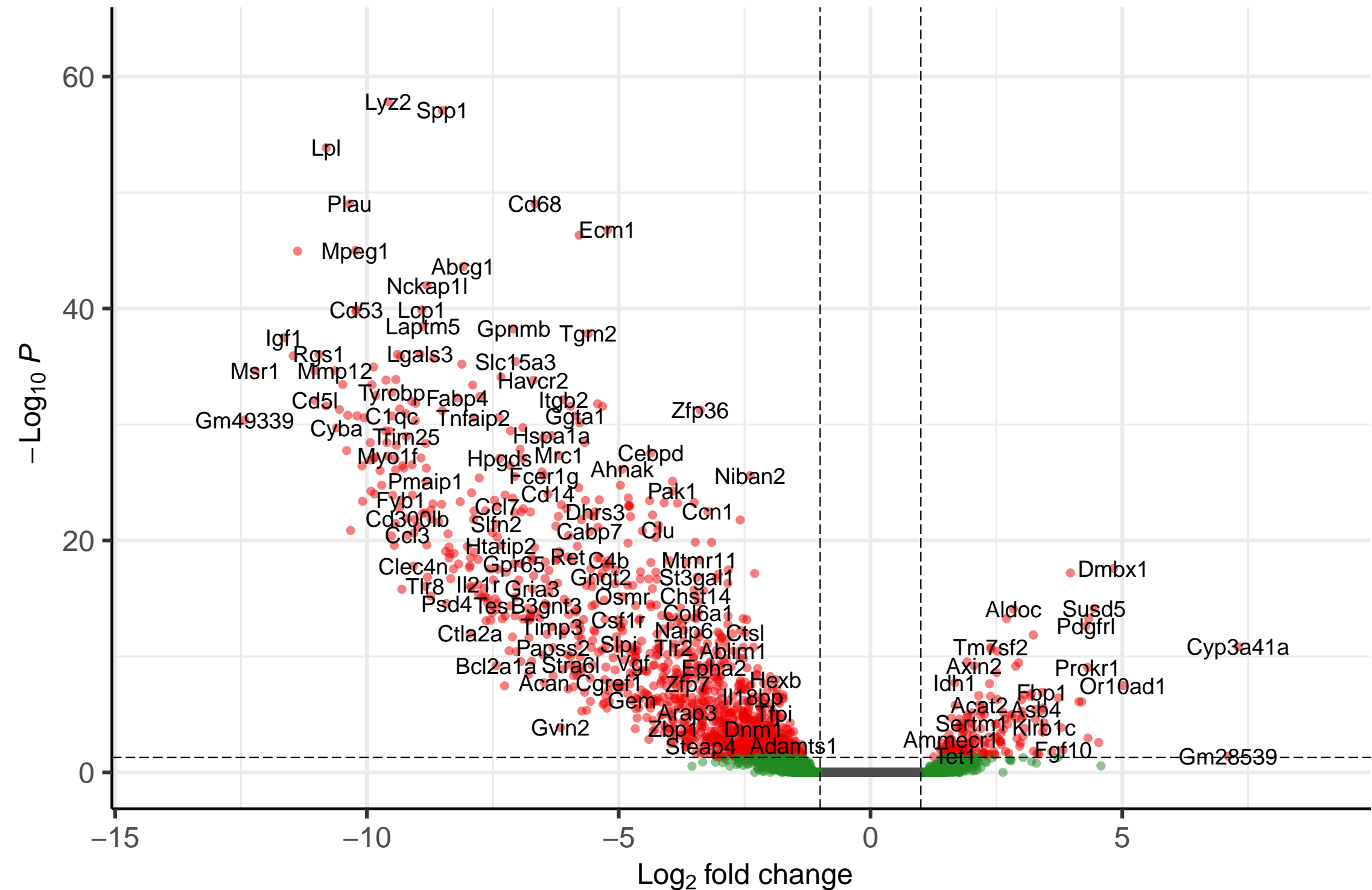


# Full\_Dataset\_protein\_coding\_interaction\_Model\_NS1\_vs\_NS2

Full\_Dataset\_protein\_coding\_interaction

● NS ● Log<sub>2</sub> FC ● p-value and log<sub>2</sub> FC



total = 13689 variables