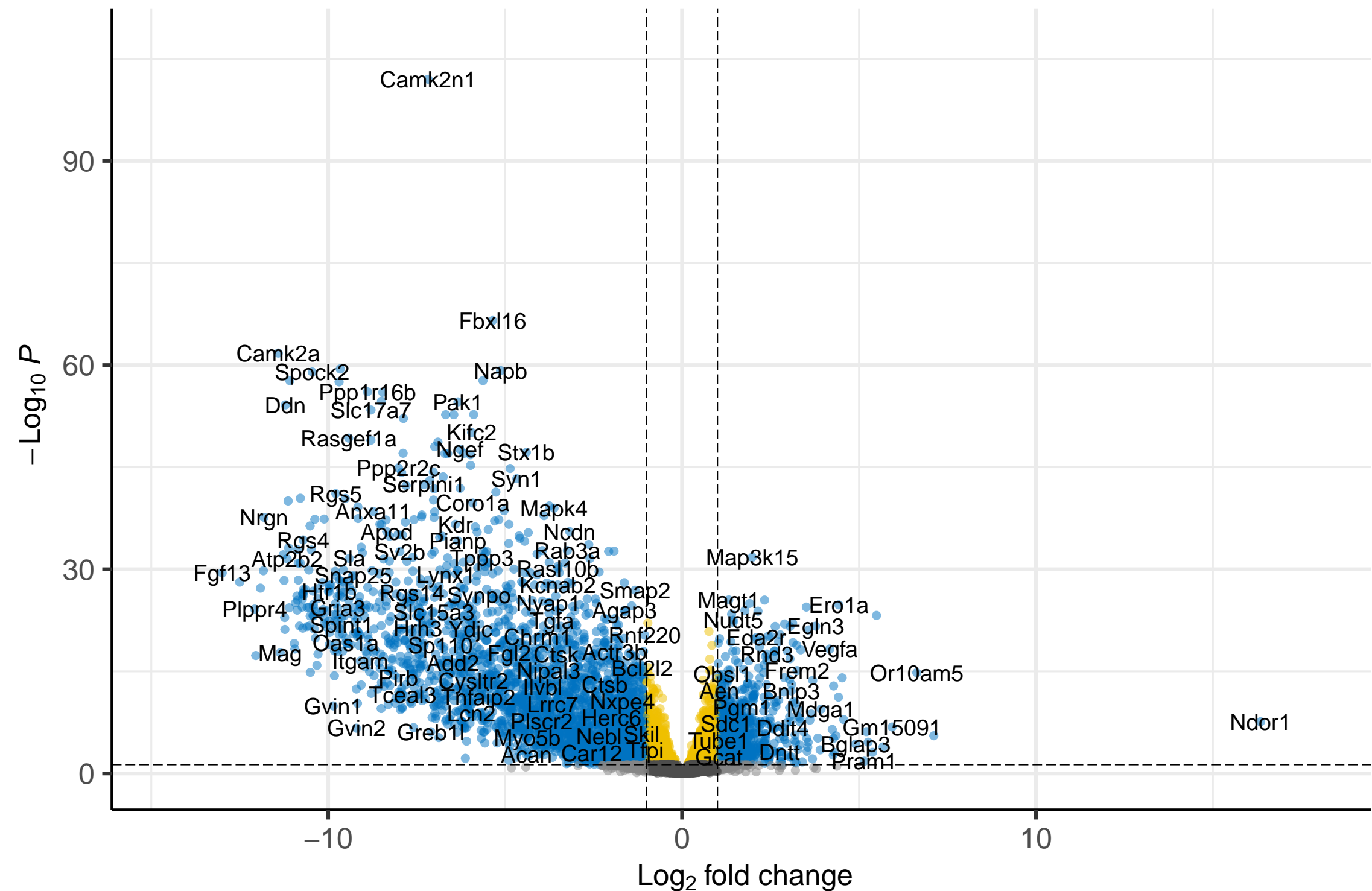


# Full Dataset protein coding interaction Model NS1 vs Tumor

EnhancedVolcano

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



total = 13689 variables