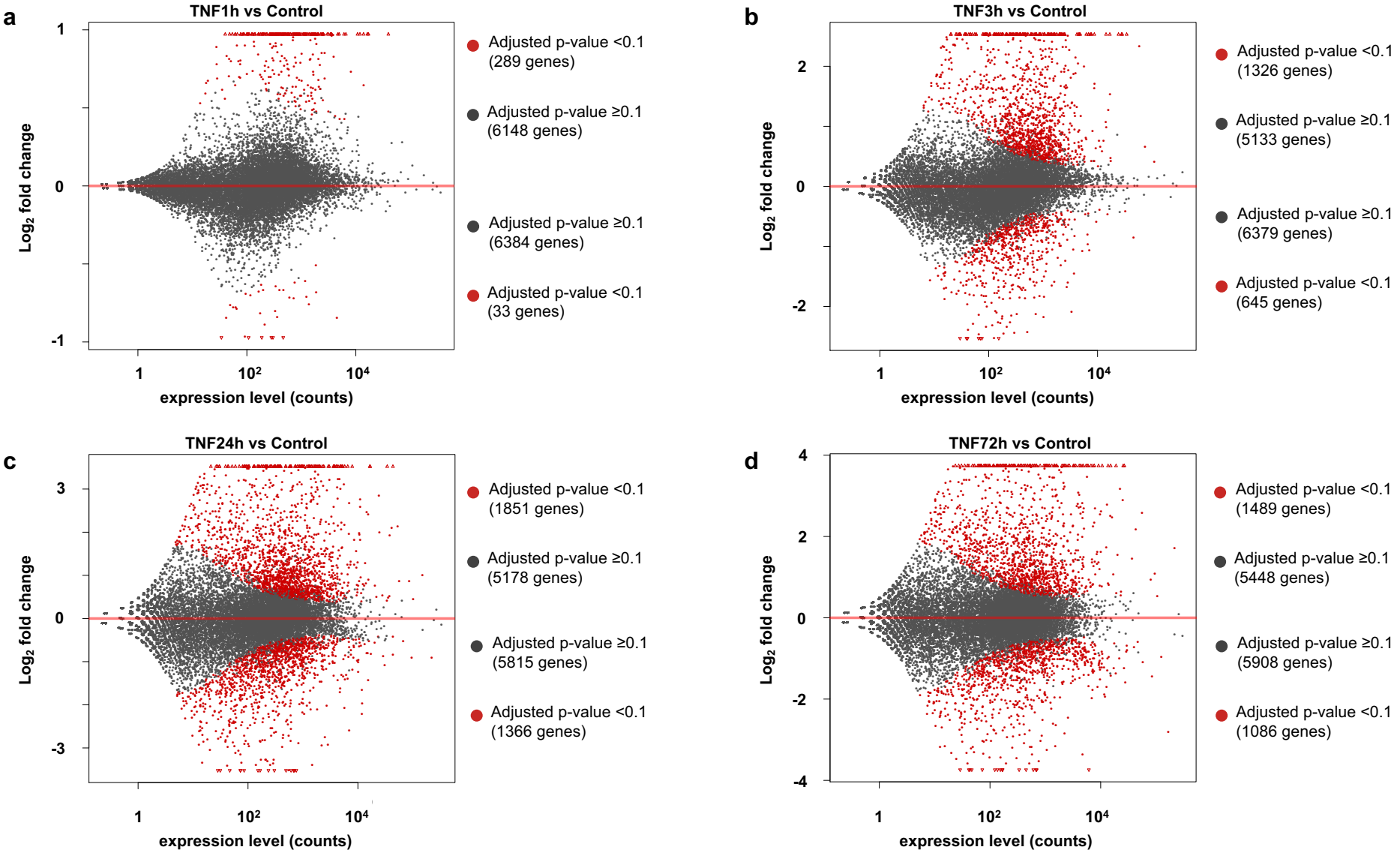
a
TNF1h vs Control
Log2 fold change
expression level (counts)
Adjusted p-value <0.1 (289 genes)
Adjusted p-value ≥0.1 (6148 genes)
Adjusted p-value ≥0.1 (6384 genes)
Adjusted p-value <0.1 (33 genes)
b
TNF3h vs Control
Log2 fold change
expression level (counts)
Adjusted p-value <0.1 (1326 genes)
Adjusted p-value ≥0.1 (5133 genes)
Adjusted p-value ≥0.1 (6379 genes)
Adjusted p-value <0.1 (645 genes)
c
TNF24h vs Control
Log2 fold change
expression level (counts)
Adjusted p-value <0.1 (1851 genes)
Adjusted p-value ≥0.1 (5178 genes)
Adjusted p-value ≥0.1 (5815 genes)
Adjusted p-value <0.1 (1366 genes)
d
TNF72h vs Control
Log2 fold change
expression level (counts)
Adjusted p-value <0.1 (1489 genes)
Adjusted p-value ≥0.1 (5448 genes)
Adjusted p-value ≥0.1 (5908 genes)
Adjusted p-value <0.1 (1086 genes)

S4 Fig