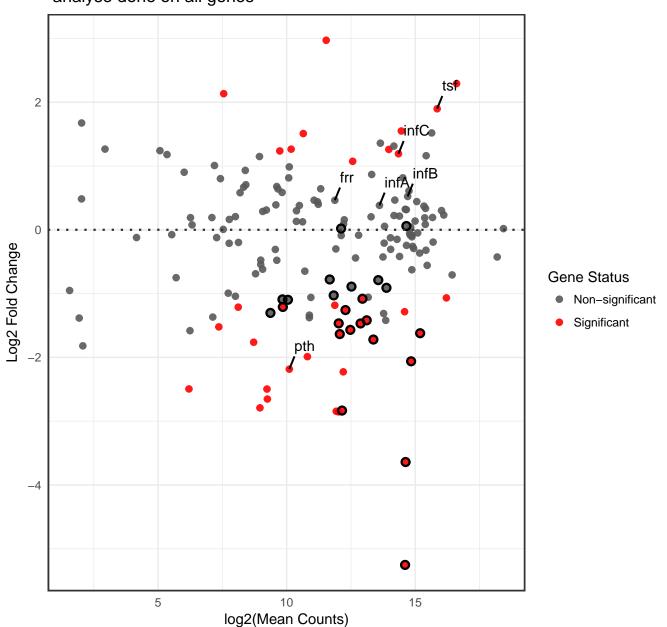


Log2 Fold Change vs Mean Counts analyse done on all genes



Log2 Fold Change vs Mean Counts analyse done on all genes tsf 2.5 infC infA infB Log2 Fold Change 0.0 Gene Status Non-significant Significant pth -2.5 • -5.0 5 10 15 log2(Mean Counts)