



Figure 1 - Supplemental Figure 4: Alternative First Exon usage is the top splicing event when using JuncBASE to identify and quantify alternative splicing events from primary human macrophages stimulated with *Listeria* and *Salmonella* for 24hrs. Significant alternative splicing events identified and categorized by event type using an adjusted p-value cutoff ≤ 0.05 (A) and a cutoff of ≤ 0.25 (B) following *Listeria* infection. Significant alternative splicing events identified and categorized by event type using an adjusted p-value cutoff of ≤ 0.05 (C) and a cutoff of ≤ 0.25 (D) following *Salmonella* infection. Data acquired from GSE73502.