



Figure S2 Number of *Ophiostoma novo-ulmi* RNAseq reads per sample and per gene: (A) Number of reads before (raw reads, dot) and after filtration/cleaning (filtered reads without duplicates, triangle) process present in each of the three conditions. Red dots: means of the three repetitions with standard deviation. No significant differences between conditions for each variable (Fisher's exact test). (B) Distribution of the number of reads (log₁₀ scale) per genes, per conditions (mean of three replicates).