

Supplemental Figure 3. **PFAM domain repertorie in the core part of F.oxysporum genomes.**

The heatmap shows PFAM domains repertorie in the core part of MI39 isolate genome and genomes of other ff. spp. Frequency values associated with each domain were mean-centered by rows. Each row of the heatmap represents log10-transformed frequency values of one PFAM domain across all F. oxysporum formae speciales (yellow, high frequency; brown, low frequency). Ivory color represents missing data points i.e. situations when a domain has not been detected in a pathogen’s genome.