

Table S2. Novel Domain Architectures shared across the Pseudofungi.

| Gene Name | Domain Architecture | Oomycete exemplar (Accession number) | Hypochytrium homologue (Accession number) |
|--|---|---|---|
| 1-phosphatidylinositol 3-phosphate 5-kinase FAB1 | FYVE_like_SF-GAF | <i>Phytophthora nicotianae</i> (KUF83522) | Hypho2016_00014879 |
| Long-chain-fatty-acid--AMP ligase FadD29 | FAAL-PKS_PP-BioF | <i>Phytophthora nicotianae</i> (KUF80336) | Hypho2016_00002377 |
| rhodanese domain-containing dual specificity protein phosphatase | DSPc-FYVE_like_SF- RHOD | <i>Phytophthora nicotianae</i> (KUF77573) | Hypho2016_00006029 |
| putative aminotransferase | AAT_like-dihydrodipicolinate reductase | <i>Phytophthora sojae</i> (XP_009530580) | Hypho2016_00002138 |
| putative callose synthase | Glucan_synthase-Sugar_tr | <i>Phytophthora infestans</i> (XP_002998554) | Hypho2016_00001204 |
| conserved hypothetical | RING-finger-tub | <i>Phytophthora infestans</i> (XP_002898661) | Hypho2016_00011754 |
| AGC/RSK/RSKP90 protein kinase | PH-PH-PH-STKc_AGC | <i>Phytophthora infestans</i> (XP_008915041) | Hypho2016_00001388 |
| phosphatidylinositol 3 and 4-kinase-like protein | PH-FYVE_scVPS27p-PI3Kc_II | <i>Phytophthora sojae</i> (XP_009538551) | Hypho2016_00003770 |
| PAX-interacting protein 1 | TPR_11-Prefoldin_2 | <i>Phytophthora nicotianae</i> (KUF96058) | Hypho2016_00008250 |
| histidine kinase A two component receptor | PAS-PAS-PAS-PAS-PAS-PAS-PAS-PAS-PAS-PAS-PAS-PAS-Histidine kinase-like ATPases | <i>Phytophthora sojae</i> (XP_009526816.1) | Hypho2016_00014859 |
| TKL protein kinase | S_TKc-DEP | <i>Phytophthora parasitica</i> (ETM36095) | Hypho2016_00016538 |
| L-amino adipate-semialdehyde dehydrogenase large subunit | Abhydrolase_6-A_NRPS-SDR_e1 | <i>Phytophthora infestans</i> (XP_002905400) | Hypho2016_00000654 |