Figure S1. A WD40 architecture in *P. patens* is homologous to bacterial sequences. Top panel: screenshot of an BLASTp search against NCBI using the *P. patens* WD40 as query. Note that top hits are all bacterial and fungal sequences. Left panel shows a majority-ruled parsimony tree with maximum-likelihood branch length. Note that the *P. patens* WD40 protein cluster together with bacterial sequences.