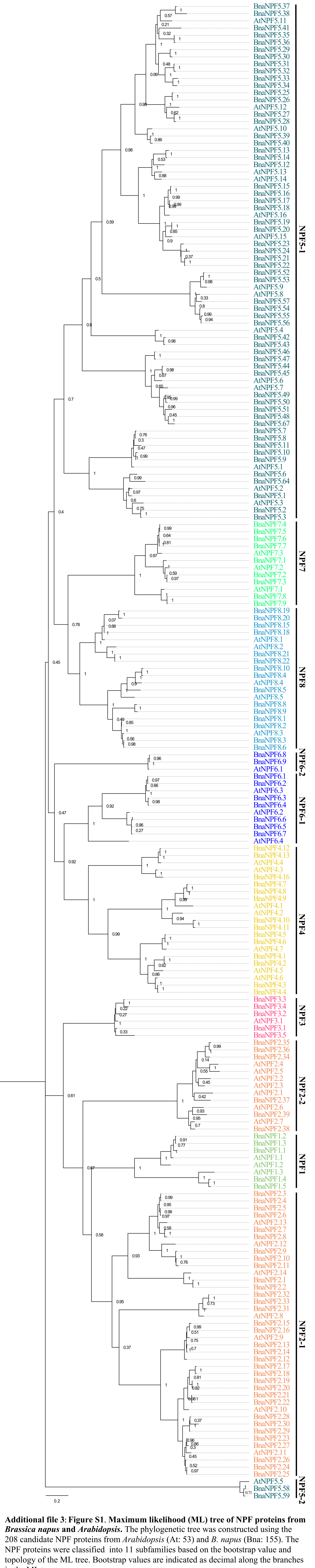




Additional file 3: Figure S1. Maximum likelihood (ML) tree of NPF proteins from *Brassica napus* and *Arabidopsis*. The phylogenetic tree was constructed using the 208 candidate NPF proteins from *Arabidopsis* (At: 53) and *B. napus* (Bna: 155). The NPF proteins were classified into 11 subfamilies based on the bootstrap value and topology of the ML tree. Bootstrap values are indicated as decimal along the branches in the ML tree.