

Supplemental figure 4

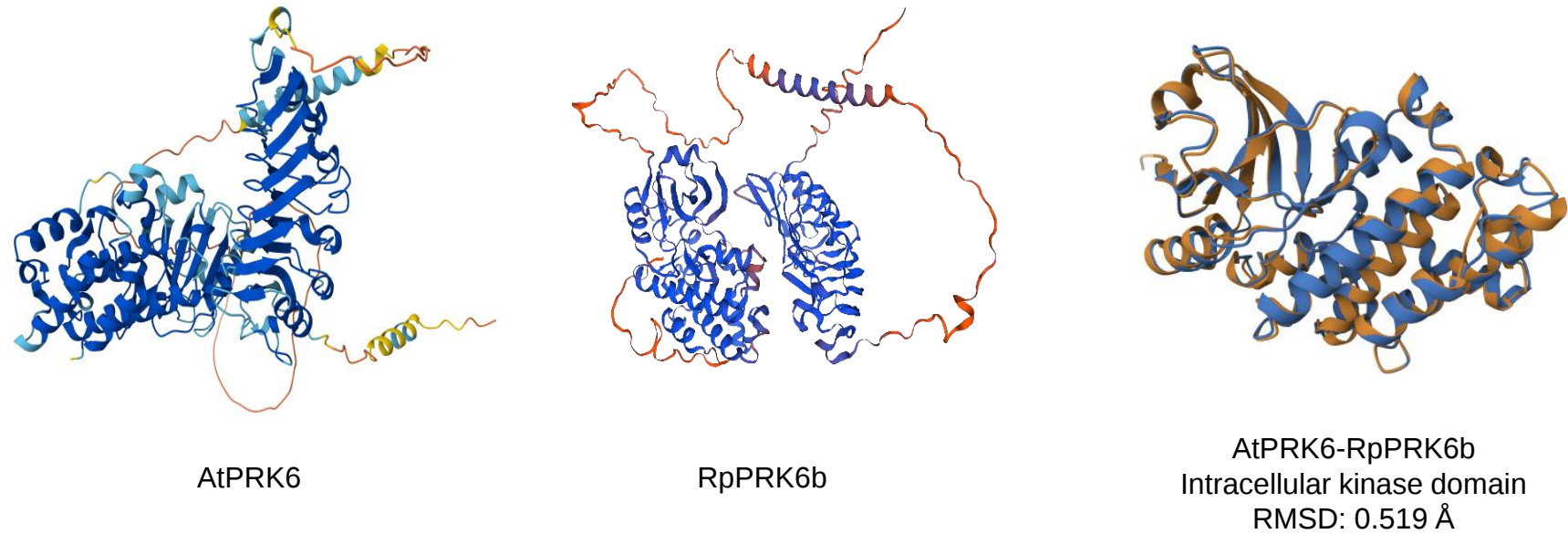


Fig. S4 Predicted structures of PRK6 homologs from *A. thaliana* and *R. pseudoacacia*.

Predicted three-dimensional structures of AtPRK6 (left) and RpPRK6b (middle), and structural superposition of the intracellular kinase domains of AtPRK6 and RpPRK6b (right). Root mean square deviation (RMSD) = 0.519 Å. Structures were predicted with AlphaFold2. Single structures are colored by AlphaFold2 per-residue confidence (left and middle). AtPRK6 is shown in blue and RpPRK6b in brown in the superposition (right).