

Figure S4a: GAP all-in maximum likelihood transcriptome phylogeny: gapA, gapB, mystery gap, and gapC.

To Fig. S4a

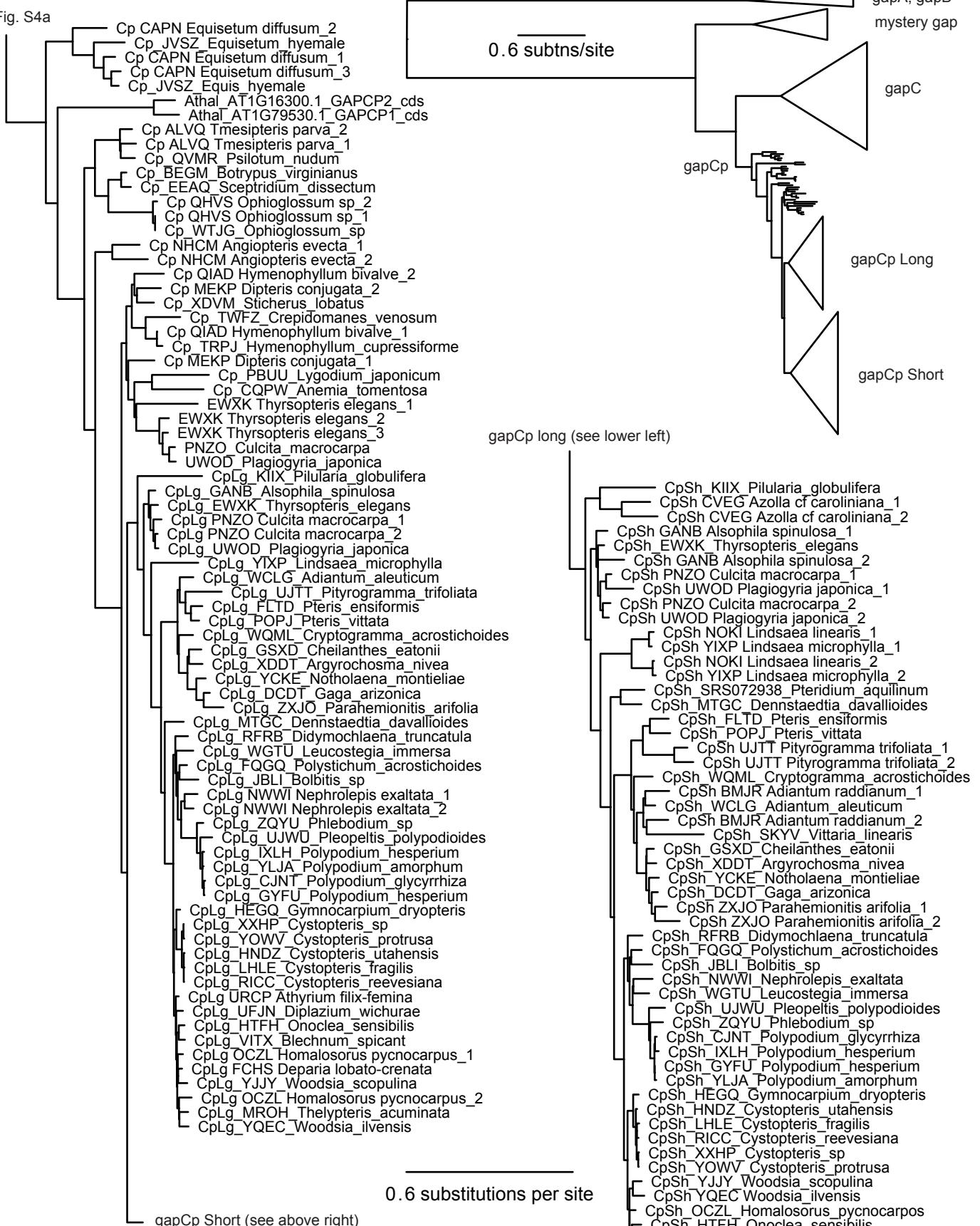


Figure S4b: GAP all-in maximum likelihood transcriptome phylogeny: gapCp (including Cp Short and Cp Long), and a cartoon map of the GAP family phylogeny.