

Figure S2a: CRY all-in maximum likelihood transcriptome phylogeny: preduplication CRY3/4, CRY3, and CRY4.

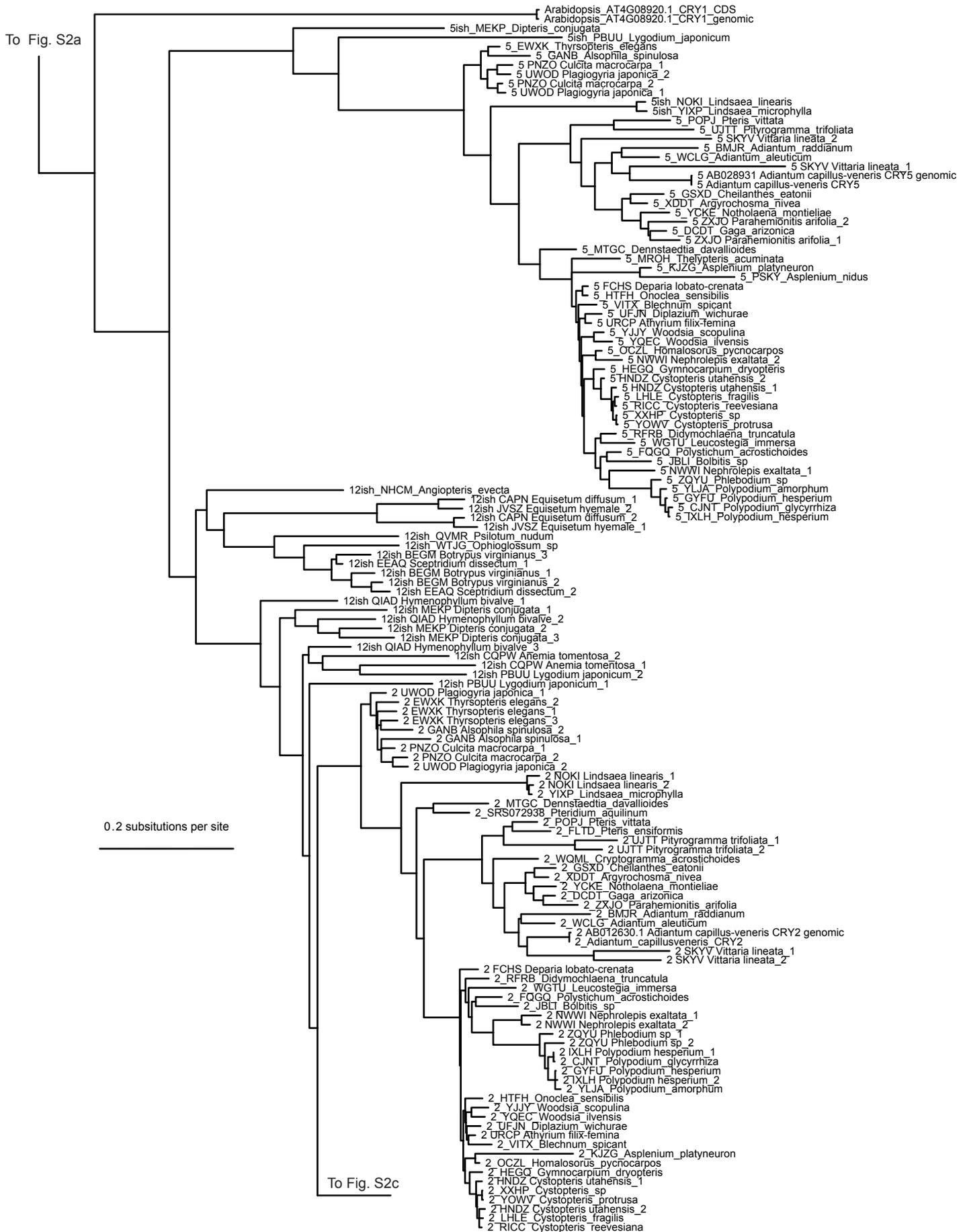


Figure S2b: CRY all-in maximum likelihood transcriptome phylogeny: CRY5, preduplication CRY1/2, and CRY2.

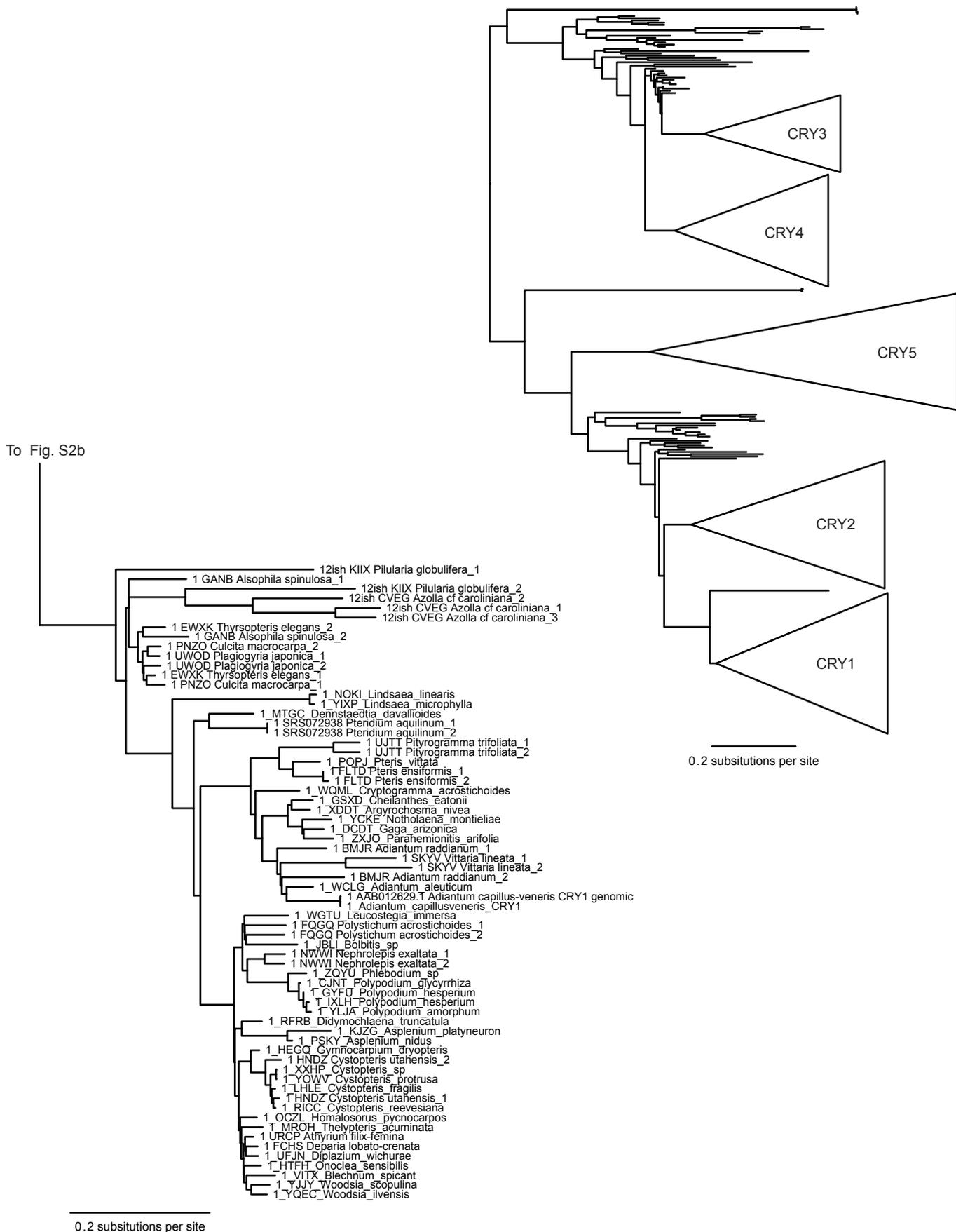


Figure S2c: CRY all-in maximum likelihood transcriptome phylogeny: CRY1, and a cartoon “map” of the entire cryptochrome fern phylogeny.