Supplementary Material to "Molecular evolution and functional divergence of alcohol dehydrogenases in animals, fungi and plants"

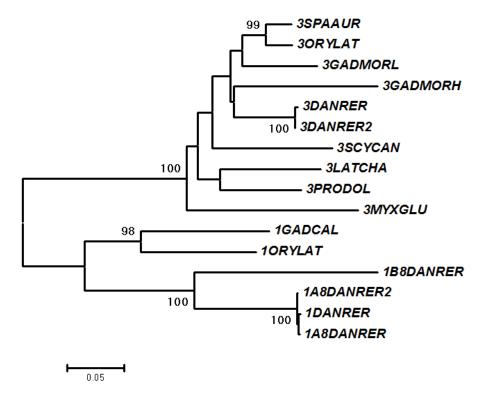


Figure S1 – Evolutionary relationships of fish ADH proteins obtained using the neighbor-joining method, Poisson-corrected amino acid distances, and pairwise deletion of gaps/missing data. Numbers representing bootstrap values higher than 80% are shown. Scale bar indicates levels of sequence divergence.