Figure S2.—Distribution of DNA distance and Ka/Ks (ω) ratio in reciprocal best BLASTP pairs. The gene pairs are from the distantly related genome pair (the two outermost taxa) in each clade as in Figure S.5. DNA distance was measured by DNADIST in the PHYLIP package, while the Ka/Ks ratio was measured by the Yang and Nielsen (2000) method, using yn00 in the PAML package.