**Table S4** **Nuclear *ATPSβ* sequence diversity statistics.** Nuclear-encoded adenosine triphosphate synthetase subunit β (*ATPSβ*) intron sample sizes (*N*, number of haplotypes sequenced), haplotype diversity (*h*), average number of pairwise differences (*k*), nucleotide diversity (*π*), Tajima’s *D*, Fu’s *Fs* statistic, and Ramos-Onsins and Rozas’ *R2*statistic. Standard deviations in parentheses. Boldface indicates statistically significant values. See Table 1 for sample locality abbreviations. NWP: Northwest Pacific; NEP: Northeast Pacific.

|  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- |
| Locality | Region |  *N* | *h* | *π* |  *D* |  *Fs* |  *R2* |
| ERI | NWP | 10 | 0 | 0 | NA |  NA |  NA |
| KHO | NWP |  9 | 0.533 (0.172) | 0.0021 (0.002) |  1.031 |  1.723 |  0.267 |
| STA | NWP |  7 | 0.429 (0.169) | 0.0014 (0.002) | 0.414 |  1.653 | 0.214 |
| PET | NWP |  8 | 0.607 (0.164) | 0.0023 (0.001) | 0.585 |  0.723 | 0.209 |
| KOD | NEP |  8 | 0.250 (0.180) | 0.0004 (0.001) | -1.055 |  -0.182 | 0.331 |
| COR | NEP |  8 | 0.250 (0.180) | 0.0004 (0.001) | -1.055 |  -0.182 | 0.331 |
| JUN | NEP |  8 | 0.250 (0.180) | 0.0004 (0.001) | 0.001 |  0.233 | 0.205 |