**Supplemental Table S3.** Population differentiation estimated by pairwise FST values for microsatellite data before (below diagonal) and after (above diagonal) applying the Excluding Null Allele (ENA) correction method proposed by Chapuis & Estoup (2007).

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
|  | **Cuba** | **Quintana Roo** | **Tabasco** | **Captive-born** |
| **Cuba** | \*\*\*\* | 0.004  (-0.028 - 0.050) | 0.013  (-0.026 - 0.055) | -0.003  (-0.017 - 0.010) |
| **Quintana Roo** | -0.013  (-0.045 - 0.033) | \*\*\*\* | -0.004  (-0.031 - 0.030) | 0.001  (-0.021 - 0.026) |
| **Tabasco** | 0.002  (-0.035 - 0.043) | -0.022  (-0.056 - 0.014) | \*\*\*\* | 0.011  (-0.015 - 0.035) |
| **Captive-born** | -0.008  (-0.029 - 0.013) | -0.006  (-0.029 - 0.027) | 0.002  (-0.020 - 0.022) | \*\*\*\* |

In parenthesis, values for 95% confidence interval after bootstrap resampling over loci.