**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.