Supplementary Table 8. Number of probable first-generation immigrants verified in each population of *S. furcifera* in 2012 and its putative source population at threshold of *a* = 0.05 and *a* = 0.01 (in parentheses).

|  |  |  |  |  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Country | Population | Test statistic1 | Putative source population | | | | | | | | |
| B1 | B2 | B3 | B4 | V | TH | NE | SA1 | LA |
| Bangladesh | B1 | *L*home/ *L*max | - | 3(3) | 1(1) |  |  |  |  |  | 2(2) |
|  | B2 | *L*home/ *L*max |  | - | 2(2) |  |  |  |  | 1(1) | 5(1) |
|  | B3 | *L*home/ *L*max | 2(1) | 1(0) | - | 1(0) | 2(1) | 1(1) | 1(1) | 2(0) | 3(0) |
|  | B4 | *L*home/ *L*max |  |  | 1(0) | - | 1(1) |  |  | 1(0) | 4(2) |
| Vietnam | V | *L*home/ *L*max |  |  |  |  | - |  | 1(1) | 1(0) |  |
| Thailand | TH | *L*home/ *L*max |  |  |  |  |  | - |  | 4(3) |  |
| Nepal | NE | *L*home/ *L*max |  |  | 2(0) | 1(1) |  |  | - | 1(0) | 2(1) |
| Korea | SA1 | *L*home/ *L*max |  |  | 2(1) | 1(1) | 4(1) | 1(0) | 1(1) | - |  |
| Laos | LA | *L*home/ *L*max |  |  | 2(2) | 2(0) |  | 1(0) | 1(0) | 2(1) | - |

The analysis applied the assignment criterion explained by Rannal & Mountain (1997) and the Monte Carlo resampling method reported by Paetkau *et al*. (2004).

1 L= likelihood of immigrant detection (Paetkau *et al*., 2004)