Table 2 Genetic distance (p-distance) based on two mtDNA fragments (cyt.b/ND2. %)

|  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- |
| Taxa/clade | *S. bistrigatus* | *S. subpunctatus* | A (*S. grahami*) | B (*S. collaris*) | C(*S. triangularis*) | D (*S.chinensis*) | *S. collaris* (CAS 240150) |
| *S. bistrigatus* |  | 9.4 | 20.2 | 21.2 | 20.8 | 20.0 | 19.6 |
| *S. subpunctatus* | - |  | 20.2 | 20.9 | 19.9 | 19.7 | 19.9 |
| A (*S. grahami*) | 36.1 | - |  | 12.2 | 12.3 | 12.6 | 16.1 |
| B (S. collaris) | 32.7 | - | 13.3 |  | 8.3 | 13.9 | 16.8 |
| C *(S. triangularis)* | 25.4 | - | 11.9 | 7.9 |  | 14.1 | 15.9 |
| D (*S.chinensis*) | 25.0 | - | 14.9 | 14.0 | 14.7 |  | 15.1 |
| *S. collaris* (CAS 240150) | 24.5 | - | 15.6 | 15.6 | 17.7 | 18.3 |  |