Table S3. Genetic diversity of different populations. The number of samples (n), the number of segregating sites (s), the number of haplotypes (h), the haplotype diversity (Hd), and the nucleotide diversity (π).

|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
| Location | n | s | h | Hd | π |
| Taiwan | 30 | 5 | 7 | 0.618 | 0.00140 |
| Yinggeling, Hainan | 4 | 2 | 3 | 0.833 | 0.00127 |
| Bawangling, Hainan | 9 | 6 | 7 | 0.944 | 0.00316 |
| Guizhou | 12 | 0 | 1 | 0.000 | 0.00000 |
| Zhejiang | 9 | 3 | 4 | 0.806 | 0.00141 |
| Jiangxi | 6 | 0 | 1 | 0.000 | 0.00000 |
| Vietnam | 6 | 7 | 2 | 0.533 | 0.00473 |
| Fujian | 2 | 0 | 1 | 0.000 | 0.00000 |
| Guangdong | 2 | 0 | 1 | 0.000 | 0.00000 |
| Total | 80 | 37 | 24 | 0.899 | 0.01160 |