**Table S3.** Genetic diversity of eight SSR markers within six *Michelia shiluensis* populations.

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
| Locus | Na | Ne | I | PIC | Ho | He |
| MC2 | 5.833 | 4.265 | 1.554 | 0.805 | 0.764 | 0.758 |
| LT106 | 6.500 | 4.226 | 4.605 | 0.836 | 0.596 | 0.756 |
| SGA5 | 6.333 | 5.327 | 1.713 | 0.879 | 0.752 | 0.801 |
| LT116 | 7.667 | 5.585 | 1.767 | 0.909 | 0.635 | 0.797 |
| SGA26 | 7.167 | 4.538 | 1.675 | 0.851 | 0.730 | 0.770 |
| MMA51 | 4.833 | 3.006 | 1.257 | 0.806 | 0.749 | 0.637 |
| MA3-7 | 6.333 | 4.787 | 1.661 | 0.854 | 0.780 | 0.778 |
| WS18 | 3.500 | 2.154 | 0.820 | 0.531 | 0.482 | 0.445 |
| Average | 6.021 | 4.236 | 1.506 | 0.808 | 0.686 | 0.718 |

Notes: Na, number of alleles; Ne, number of effective alleles; I, Shannon index; PIC, polymorphism information content; Ho, observed heterozygosity; He, expected heterozygosity.