|  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- |
| Island | Years | n | H | Number of polymorphic sites (S) | Haplotype diversity (Hd) ± SD | Nucleotide diversity (π) | θ (site/seq) |
| Hawai‘i | 2009-2012 | 68 | 8 | 22 | 0.448 ± 0.072 | 0.0015 | 0.007/4.593 |
|  | 2018-2019 | 68 | 8 | 7 | 0.320 ± 0.073 | 0.0006 | 0.002/1.462 |
| Maui | 2012-2014 | 20 | 4 | 24 | 0.668 ± 0.074 | 0.0130 | 0.010/6.765 |
|  | 2016-2017 | 33 | 3 | 23 | 0.561 ± 0.048 | 0.0070 | 0.008/5.667 |
|  | 2018-2019 | 29 | 5 | 24 | 0.717 ± 0.043 | 0.0164 | 0.009/6.111 |
| O‘ahu | 2013 | 13 | 3 | 4 | 0.295 ± 0.156 | 0.0012 | 0.002/1.289 |
|  | 2014 | 10 | 3 | 18 | 0.511 ± 0.164 | 0.0012 | 0.009/6.363 |
|  | 2015-2016 | 10 | 2 | 21 | 0.356 ± 0.159 | 0.0011 | 0.011/7.423 |
|  | 2017-2018 | 9 | 2 | 21 | 0.222 ± 0.166 | 0.0071 | 0.011/7.726 |
| Kaua‘i | 2008-2019 | 16 | 2 | 3 | 0.125 ± 0.106 | 0.0006 | 0.001/0.904 |