Supplemental Table 2:Haplotype and Nucleotide Diversity for the Fewkes Assemblage, Other Archaeological Turkeys, and Modern Turkeys.

|  |  |  |  |
| --- | --- | --- | --- |
| **Population** | **n=** | **Haplotype diversity** | **Nucleotide diversity** |
| ***h*** | **Standard deviation** | **π** | **Standard deviation** |
| **Archaeological Turkeys** |
| Fewkes Site | 15 | 0.657 | 0.08 | 0.00395 | 0.0015 |
| 5MT3807 Shields Pueblo (1060–1260)a | 22 | 0 |   | 0 |   |
| 5MT765 Sand Canyon Pueblo (1225–1280)a | 9 | 0.222 | 0.166 | 0.00446 | 0.00334 |
| LA12 Arroyo Hondo, component I (1310–1345)a | 29 | 0.135 | 0.085 | 0.0029 | 0.00179 |
| 5MT123 Albert Porter Pueblo (900-1300)a | 14 | 0 |   | 0 |   |
| **Eastern Wild Turkeys (*Meleagris gallopavo* *silvestris*)b** |
| Scotch WMA, Alabama (ESW) | 9 | 0.556 | 0.165 | 0.00279 | 0.00115 |
| Waterhorn WMA, S. Carolina (EWW) | 10 | 0.733 | 0.12 | 0.00609 | 0.00114 |
| Ozark Mts, Missouri (EOM) | 10 | 0.644 | 0.152 | 0.00461 | 0.00142 |
| W. Virginia, various locations (EWV) | 10 | 0.822 | 0.097 | 0.0084 | 0.00148 |
| Black Warrior WMA, Alabama (EBW) | 10 | 0.733 | 0.076 | 0.00513 | 0.00069 |
| Camp Lejeune, N. Carolina (ECL) | 7 | 0.667 | 0.16 | 0.00382 | 0.00147 |

*Note:* Genetic diversity values calculated from sequence data published in: a Kemp et al. [(2017)](https://paperpile.com/c/bcj138/pp8p/?noauthor=1); b Mock et al. [(2002)](https://paperpile.com/c/bcj138/7XA5S/?noauthor=1).