**Table S1. tMRCA estimates comparison among different datasets (WG/SNPs).**

|  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- |
|  | tMRCAa | 95% HPDLb | 95% HPDHc | Year | 95% HPDL | 95% HPDH |
| **Root WGd P1-P2** | 183 | 161 | 207 | 1794 | 1770 | 1816 |
| **P1e** | 65 | 58 | 71 | 1912 | 1906 | 1919 |
| **P2f** | 103 | 89 | 118 | 1874 | 1859 | 1888 |
| **SNPsg P1-P2** | 254 | 216 | 296 | 1723 | 1681 | 1761 |
| **P1** | 68 | 61 | 76 | 1909 | 1901 | 1916 |
| **P2** | 122 | 103 | 143 | 1855 | 1834 | 1874 |

a tMRCA, time of the Most Recent Common Ancestor

b 95%HPDL, Highest Posterior Density Low Interval

c 95%HPDH, Highest Posterior Density High Interval