

Supplemental Figure 3. Median-joining network displaying the relationships between the newly obtained sequences and previously published archaeological (Manin et al. 2018; Speller et al. 2010) and modern (Canales Vergara et al. 2019; Luchini et al. 2001; Mock et al. 2001, 2002; Monteagudo et al. 2013; Padilla-Jacobo et al. 2018; Szalanski et al. 2000) sequences obtained from GenBank. The Mimbres turkeys (in black) are primarily grouped with the aHap1 and aHap2 haplotypes. In this model, missing bases have been kept to estimate the distance between haplotypes.

