

ADDITIONAL DATA FILE 3: Figures S1-S8

Complete Columbian mammoth mitogenome suggests interbreeding with woolly mammoths

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All literature cited refer to those in the main manuscript.

FIGURES

Fig. S1. The Huntington mammoth lower left third molar. Figure from [17] used with permission from the *Journal of Paleontology*.



Fig. S2. Log-transformed amplicon copy number per milligram of substrate vs. fragment length. Open squares are woolly mammoth samples assayed by [24], with regression lines indicated.

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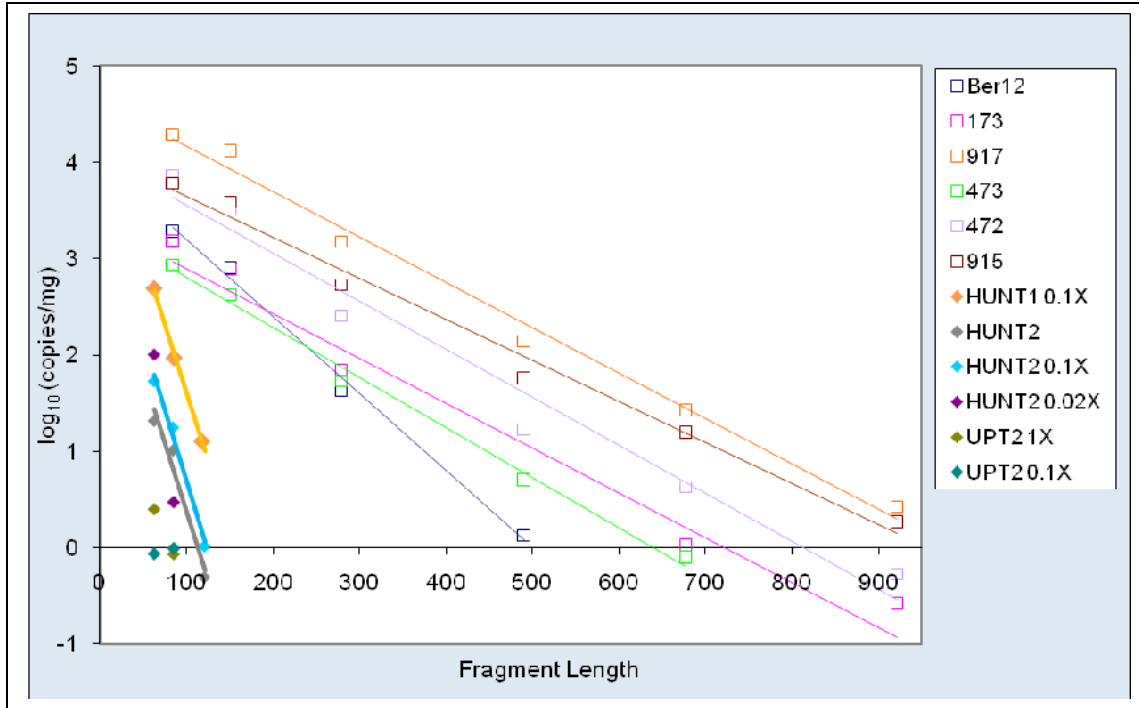


Fig. S3. Maximum clade credibility tree from phylogenetic analysis set 1a, with nodal posterior probabilities indicated. Tip names are preceded by their haplotype as determined by [11] and [12].

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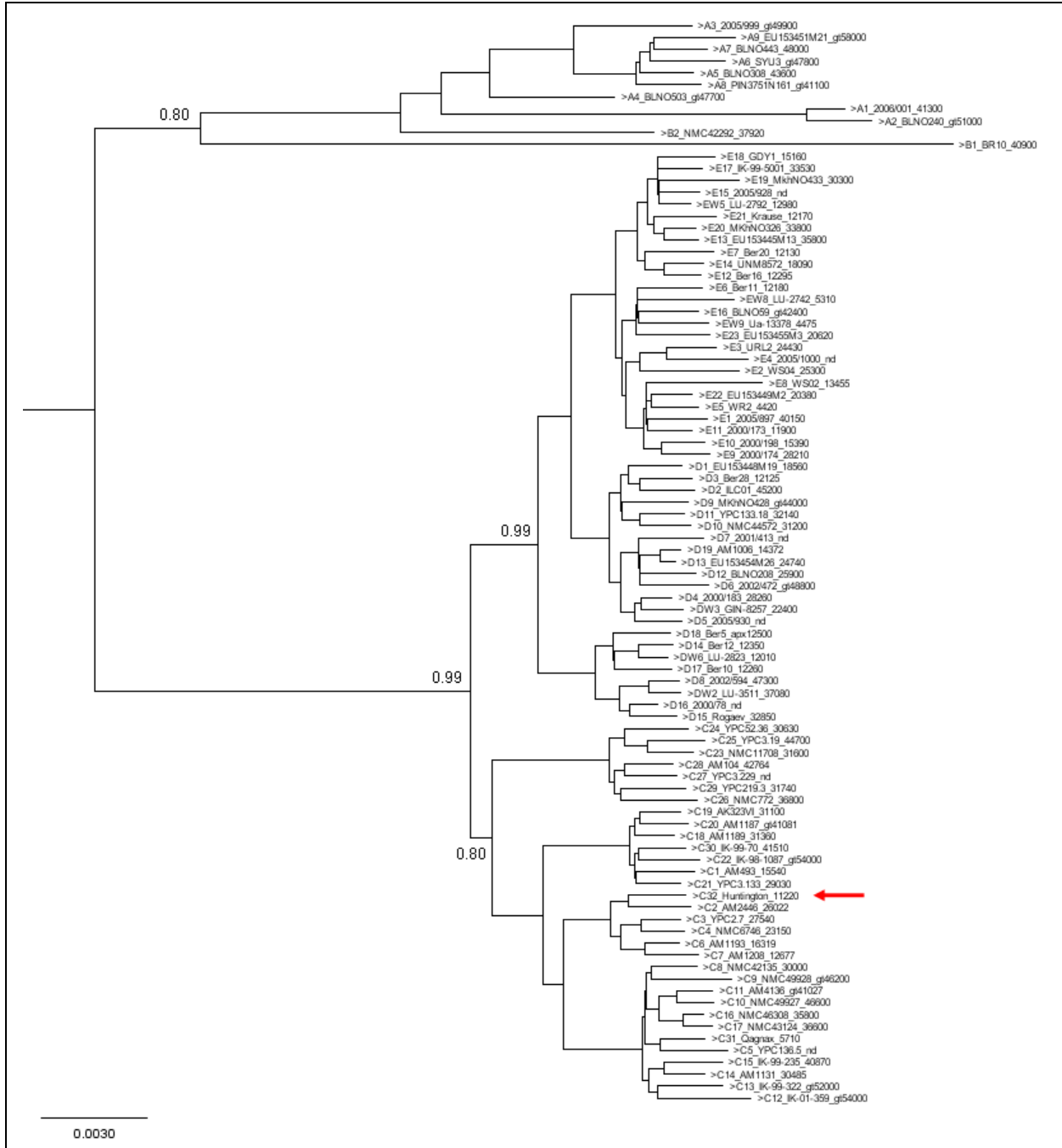


Fig. S5. Maximum clade credibility tree from phylogenetic analysis set 2a, with nodal posterior probabilities indicated. Tip names are preceded by their haplotype as determined by [11] and [12].

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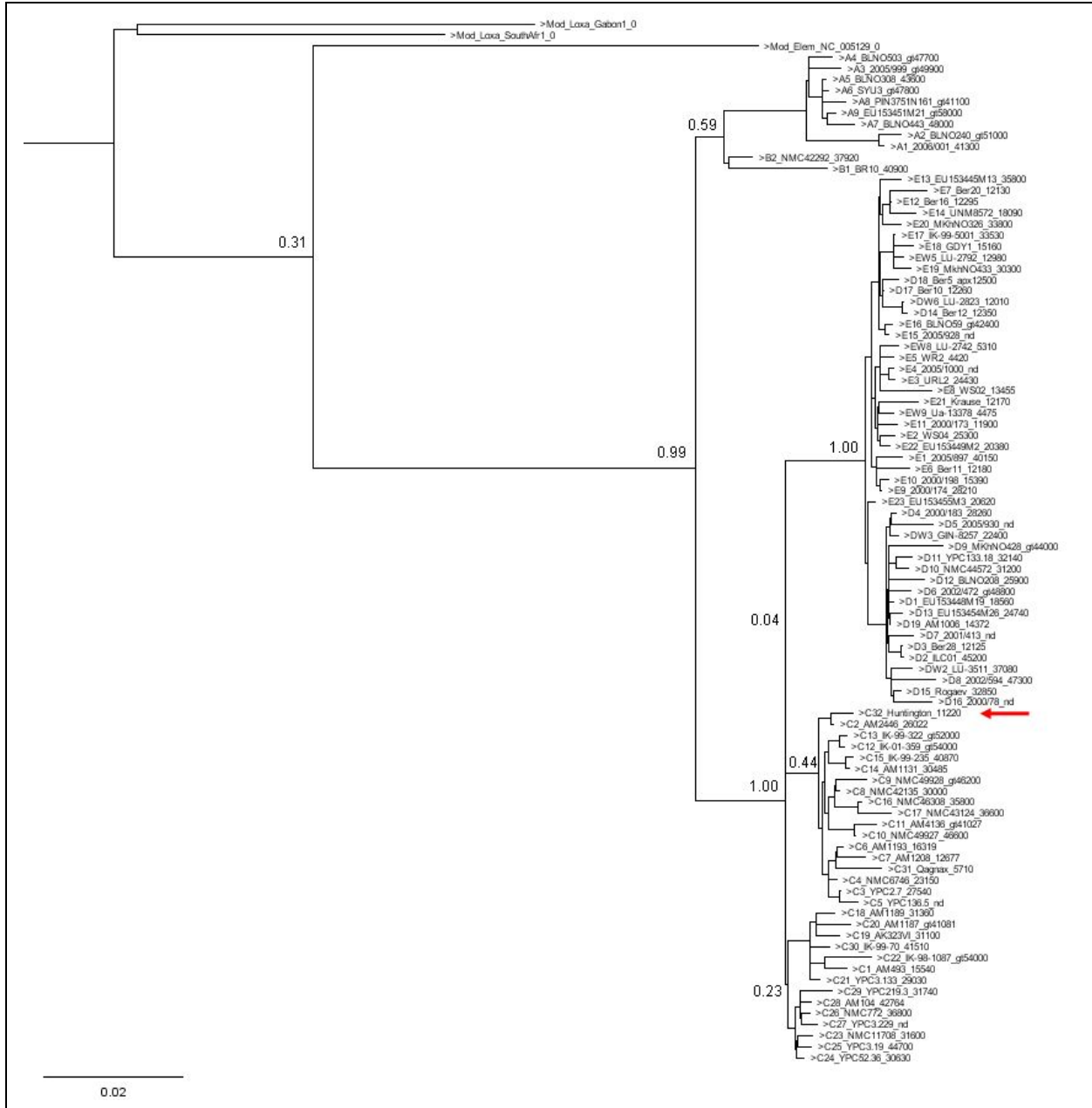


Fig. S6. Maximum clade credibility tree from phylogenetic analysis set 2b, with nodal posterior probabilities indicated. Tip names are preceded by their haplotype as determined by [11] and [12].

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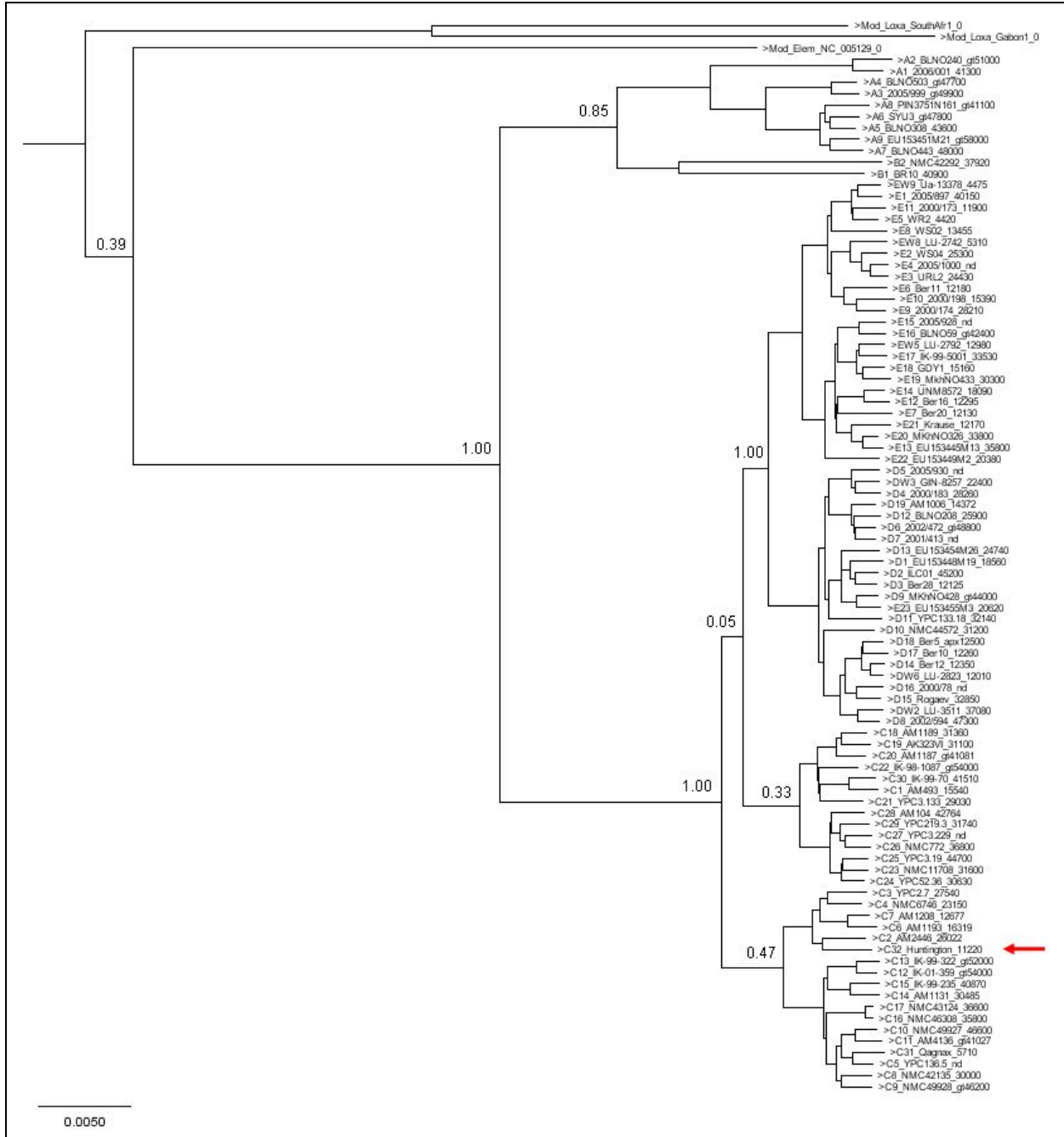


Fig. S7. Maximum clade credibility tree from phylogenetic analysis set 4a, with nodal posterior probabilities indicated. Tip names are preceded by their haplotype as determined by [11] and [12].

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