Supporting Information Figure S5

Revised timeline and distribution of the earliest diverged human maternal lineages in southern Africa

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**Fig. S5. Phylogenetic tree for L0d1c.** Phylogeny was inferred using a total of 279 mtDNA, including all 139 mtDNA in the focus set and 140 additional L0d1c mtDNA from Barbieri et al. 2013. Haplogroups other than L0d1c are “collapsed”. Tips of L0d1c genomes are labeled with the format: Haplogroup (Isolate Name) - Language/Isolate Source [Country], where Isolate Name, Isolate Source, and Country are information included in the corresponding GenBank entries, and Language is the reported spoken language. Tip colours reflect data source: Red=current study, Green=Barbieri et al. 2013, Purple=Schuster et al. 2010, and Aqua=NCBI.