

Figure S2 (A)

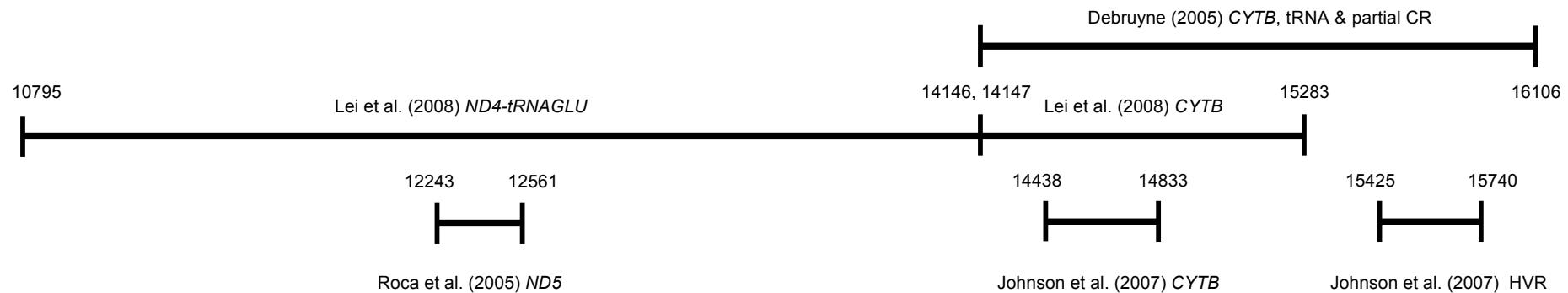


Figure S2 (B)

mtDNA region	DeBruyne (2005) <i>CYTB, tRNA & CR</i> Clade (Haplotype)		Roca et al. (2005) <i>ND5</i> Clade		Johnson et al. (2007) <i>CYTB + HVR</i> Group (Haplotype)		Lei et al. (2008) <i>ND5/6 + CYTB</i> Clade (Haplotype)		Sample ID
S Clade designations									
<i>ND5</i>		AY823346	I				AY781552	Savanna II (LAS1)	14
<i>CYTB</i>				EU116002 (99%)	III (H11)		AY768831	Savanna II (LAS1)	14
<i>CYTB + CR</i>	AY742801	S (Laa Tanzania1)		-	-				
<i>ND5</i>		AY823349	I	EU116004 (99%)	III (H27)		AY781638	Savanna II (LAS1)	365
<i>CYTB</i>				-	-		AY768917	Savanna II (LAS1)	365
<i>CYTB + CR</i>	AY741325	S (Laa Namibia1)							
<i>ND5</i>		AY823348	I	EU116014 (99%)	III (H41)		AY781562	Savanna I (LAS2)	62
<i>CYTB</i>				-	-		AY768841	Savanna I (LAS2)	62
<i>CYTB + CR</i>	AY741070	S (Laa Tanzania2)							
<i>ND5</i>		AY823348	I	EU116015 (99%)	III (H42)		AY781588	Savanna I (LAS2)	157
<i>CYTB</i>				-	-		AY768867	Savanna I (LAS2)	157
<i>CYTB + CR</i>	AY741322 (99%)	S (Laa Zimbabwe5)							
F Clade designations									
<i>ND5</i>		AY823341	II				AY781619	Forest III (LAF6)	218
<i>CYTB</i>				EU116000 (99%)	II (H09)		AY768898	Forest III (LAF6)	218
<i>CYTB + CR</i>	AY742802	F (DRC17)		EU096126 (97%)	I (H20)				
<i>ND5</i>		AY823342	II	EU116000	II (H09)		AY781599	Forest III (LAF5)	182
<i>CYTB</i>				EU096126 (97%)	I (H20)		AY768878	Forest III (LAF5)	182
<i>CYTB + CR</i>	AY741329 (99%)	F (Laa Zimbabwe2)							
<i>ND5</i>		AY823342	II	EU116011 (99%)	I (H37)		AY781602	Forest III (LAF2)	189
<i>CYTB</i>				EU096122 (98%)	II (H33)		AY768881	Forest III (LAF2)	189
<i>CYTB + CR</i>	AY359277(99%)	F (Lac DRC1)							

Note that sequences comprising Groups III and IV of the HVR from Johnson et al. (2007) were from Nyakaana et al. (2002), Eggert et al. (2002), DeBruyne et al. (2003) and DeBruyne (2005) (indicated as a dashes in the table). Note also that the clade designations in the Appendix of Lei et al. (2008) for haplogroups Savanna I and II appear to be reversed from the designations in the main text (the main text designations are used here). For sequence comparisons that did not show 100% match, the percent identity is shown in parentheses.