## Additional file 4: Mutational profiles, haplogroups and GenBank accession codes of the 17 novel complete East Timor mitogenomes

Ochbank accession	Janipic	napiogroup																																				
KJ676776	ET005	"P1e"#	16239T	16266T 1	16357C 73	IG 15	2C 246	ôC 261	33G 315.	1C 508G	750G	1438G 2	706G 3229.1	4659A	4769G 5	964C 6077	T 7028T	8286C	8860G	10118C	10398T 1	1719A 1	13098G 13	722G 14	766T 15258	3G 15326G	15607G											
KJ676787	ET014	P1d	16176T	16266T 1	16357C 16	526A 73	G 212	2C 261	33G 315.	1C 750G	1438G	2706G 3	747T 3834A	4769G	6077T 7	028T 8860	G 9605T	10118C	11626C	11719A	12346T 1	4569A 1	14766T 14	971C 15	326G 15607	7G												
KJ676781	ET017	P1d	16176T	16266T 1	16357C 16	526A 73	G 212	2C 26°	33G 315.	1C 524.1A	524.2C	750G 1	438G 2706G	3747T	3834A 4	769G 6077	T 7028T	8860G	9605T	10118C	11626C 1	1719A 1	12346T 14	569A 14	766T 1497	1C 15326G	15607G											
KJ676777	ET064	P1d	16176T	16266T 1	16357C 73	IG 21	2C 263	3G 31!	15.1C 7500	3 1438G	2706G	4769G 6	077T 7028T	8490C	8519A 8	860G 9922	T 10118C	10128T	11383C	11719A	12346T 1	2372A 1	12711G 12	714C 14	152G 14766	ST 15326G	15479C 1	5607G										
KJ676789	ET072	"P1e"#	16239T	16266T 1	16357C 73	IG 15	2C 246	ôC 261	33G 315.	1C 508G	750G	1438G 2	706G 4659A	4769G	5964C 6	077T 7028	T 8286C	8860G	10118C	10398T	11719A 1	3098G 1	13722G 14	766T 152	258G 15326	6G 15607G												
KJ676782	ET139	P1d	16156A	16169T 1	16176T 16	189Y 16	266T 163	311C 161	6357C 73G	152C	212C	235G 2	63G 315.1C	750G	1406C 1	438G 2706	G 2857C	4533A	4688C	4769G	6018A 6	6077T 7	7028T 82	51A 886	60G 89390	C 9150G	9269T 1	0118C 1	1719A 1234	6T 12561A	14766T	15326G 156	607G 15938T	Г				
KJ676779	ET154	"P1e"#	16239T	16266T ′	16357C 73	IG 15	2C 246	∂C 267	315.°	1C 508G	750G	1438G 2	706G 3229.1	4659A	4769G 5	964C 6077	T 7028T	8286C	8860G	10118C	10398T 1	1719A 1	13098G 13	722G 14	766T 15258	3G 15326G	15607G											
KJ676786	ET156	P1d	16093C	16311C 1	16357C 73	IG 15	OT 263	3G 31!	15.1C 7500	3 1438G	2706G	3474T 4	769G 5444T	6077T	6452A 7	028T 7859.	A 8860G	9051G	10118C	11719A	12346T 1	4383A 1	14766T 15	326G 156	607G													
KJ676784	ET167	P1d	16156A	16169T 1	16176T 16	266T 16	311C 163	357C 730	3G 1520	212C	235G	263G 3	15.1C 524.1A	524.2C	750G 1	438G 2706	G 2857C	4533A	4688C	4769G	6018A 6	6077T 7	7028T 82	51A 88	60G 89390	C 9150G	9269T 1	0118C 1	1719A 1234	6T 12561A	14766T	15038G 153	326G 15607G	3 15938T				
KJ676775	ET232	"P1e"#	16239T	16266T ′	16357C 73	IG 15	2C 246	∂C 267	315.°	1C 508G	750G	1438G 2	706G 4659A	4769G	5964C 6	077T 7028	T 8286C	8287.1C	8287.2C	8287.3C	8860G ′	0118C 1	10398T 11	719A 130	098G 13722	2G 14766T	15258G 1	5326G 1	5607G									
KJ676785	ET284	"P1e"#	16239T	16266T ′	16357C 73	IG 15	2C 246	3C 26°	33G 315.	1C 508G	750G	1438G 2	706G 4659A	4769G	5964C 6	077T 7028	T 8286C	8287.1C	8287.2C	8287.3C	8860G ′	0118C 1	10398T 11	719A 130	098G 13722	2G 14766T	15258G 1	5326G 1	5607G									
KJ676783	ET300	P1d	16093C	16311C '	16357C 73	G 26	3G 315	5.1C 750	50G 1438	3G 1676G	2706G	3474T 4	769G 5444T	6077T	6452A 7	028T 7859.	A 8860G	9051G	10118C	11719A	12346T ′	4383A 1	14766T 15	326G 156	607G													
KJ676790	ET110	M21b	16093C	16223T ′	16311C 16	519C 73	G 152	2C 17/	74T 2630	3 299DE	L 315.1C	489C 7	50G 1438G	2706G	3915A 4	491A 4769	G 4913G	5108C	6286C	6359G	6662G	7028T 7	7861C 87	'01G 886	60G 95400	C 10398G	10400T 1	0873C 1	1482C 1156	0G 11719A	12705T	14766T 147	783C 15043A	15301A 1	5326G			
KJ676774	ET126	D6a	16192T	16223T ′	16274A 16	362C 73	G 263	3G 309	9.1C 315.	1C 489C	709A	750G 1	438G 1719A	2706G	3714G 4	769G 4883	T 5178A	7028T	8473Y	8701G	8860G 9	9540C 1	10398G 10	400T 108	873C 11017	7C 11719A	12654G 1	2705T 1	1551G 1476	6T 14783C	15043A	15301A 153	326G					
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KJ676780	ET257	Q3	16129A	16223T ′	16311C 16	319A 73	G 146	6C 19!	95C 2150	3 263G	309.1C	315.1C 4	89C 519G	750G	1438G 2	706G 4117	C 4335T	4769G	4924A	4928C	5483C 5	5843G 7	7028T 82	00C 870	01G 8790	A 8860G	9540C 9	9656C 9	333C 1039	8G 10400T	10873C	11719A 127	'05T 12930C	3 12940A 1	3500C 14766T	14783C 15043A 1	5172A 15301A 1	15326G
KJ676778	ET095	R9c1b2	16157C	16304C	73G 15	1T 26	3G 315	5.1C 577	73.1C 573.	2C 573.3C	750G	1438G 2	706G 3645C	3970T	4769G 7	028T 7299	G 7861C	8860G	10163A	11002G	11719A ′	2406A 1	12618A 12	858T 139	928C 1476	ST 15326G	15479C											

# postulated novel clade

GenBank accession sample haplogroup