Variable positions in the 467 bp of the cytb gene for the 28 haplotypes of *T. brasiliensis* collected in the study according to sites where they were collected

Нар																		ſ	۱u	cle	oti	de	sit	tes																		Рор	ulat	tion	5		Hap number	Hap frequency	
									1	1		1	1	1	2	2	2 2	2	2	2				3	3	3	3	3	3	3	3	3	3	3	4	4	4	4	4	4									GeneBank code
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Hap_1	A		1	Т	Т	G	i (G	Т	Т		Ą	Т	Т	С	C	21	F,	A	С	А	А	А	Т	Т	0	5	A	G	Т	С	Т	Т	G	A	A	A	- î	P	4			3	1			4	3.77	KT336389
Hap_2	G	i i															0	2																							4	2	7	6	2		21	19.81	KT336390
Нар_3																																				G							1				1	0.94	KT336391
Hap_4	G	i i									(G												С																			4	1			5	4.72	KT336392
Hap_5	G	i.																																							2	12	3	6	5		28	26.42	KT336393
Hap_6	G	i i					1	Ą																																			1				1	0.94	KT336394
Hap_7	G	i .								C	2							. '	Т						C	2.																	1	1			2	1.89	KT336395
Hap_8	G	i.									(G										G		С											G								2	1			3	2.83	KT336396
Hap_9									С		(G												С										А							2	1					3	2.83	KT336397
Hap_10	G	i .			С												0	2																							1						1	0.94	KT336398
Hap_11	G	i .				A					(G																													3						3	2.83	KT336399
Hap_12	G	i.									(G																													1			1	4	Ļ.	6	5.66	KT336400
Hap_13	G	i.																														С									1						1	0.94	KT336401
Hap_14	G	i .		G							(G	С	С										С																	1						1	0.94	KT336402
Hap_15																					G																				1						1	0.94	KT336403
Hap_16	G	i.															0	2																			Т	۰.			1						1	0.94	KT336404
Hap_17	G	i .									(G												С			. (G														1		3	2		6	5.66	KT336405
Hap_18									С		(G												С																				1	2		3	2.83	KT336406
Hap_19	G	i .															0	2													Т													1			1	0.94	KT336407
Hap_20	G	i .																							C																	1		1			2	1.89	KT336408
Hap_21	G	i .									(G				Т	Γ.																									2					2	1.89	KT336409
Hap 22																							G																			5					5	4.72	KT336410
Hap 23	G	i .									(G												С		A	٩			С												2					2	1.89	KT336411
Hap_24																																										1					1	0.94	KT336412
Hap_25																																													1		1	0.94	KT336413
Hap_26																																													1		1	0.94	KT336414
Hap 27																		-	-		-	-						-	-																1		1	0.94	KT336415
Hap_28																		2																	G										1		1	0.94	KT336416
Total					-					-										-	-				-								-	-	-	-	-	-	-	-1	17	27	22	23	19	9	108	100.00	