

Table S4. Measures of haplotype diversity across sampling localities, including the number of samples analyzed (n), the number of polymorphic sites (S), haplotype diversity (h), and nucleotide diversity (π)

Site	Geographic					
	region	n	S	No. haplotypes	h	π
Nosy Mangabe S.R.†	North	9	1	2	0.2222	0.0004
Marotandrano S.R.	North	9	5	2	0.3889	0.0035
Mananara Nord N.P.	North	8	3	2	0.2500	0.0013
Ambatovaky S.R.	North	5	3	2	0.4000	0.0021
Zahamena N.P., S.N.R.	North	11	19	3	0.6364	0.0172
Betampona S.N.R.	North	9	20	2	0.3889	0.0139
Mangerivola S.R.	North	3	0	1	0.0000	0.0000
Mantadia Andasibe N.P.	North	14	9	2	0.4945	0.0080
Torotorofotsy	North	3	9	2	0.6667	0.0107
Maromizaha U.F.	North	4	0	1	0.0000	0.0000
Anosibe an'ala C.F.	North	8	0	1	0.0000	0.0000
Fandriana U.F.	South	11	0	1	0.0000	0.0000
Vatoharanana (Ranomafana N.P.)	South	10	0	1	0.0000	0.0000
Mangevo (Ranomafana N.P.)	South	8	0	1	0.0000	0.0000
Kianjavato U.F.	South	12	0	1	0.0000	0.0000
Vatovavy U.F.	South	10	0	1	0.0000	0.0000
Lakia	South	10	0	1	0.0000	0.0000
Tolongoina U.F.	South	4	0	1	0.0000	0.0000
Manombo S.R.	South	11	0	1	0.0000	0.0000
Overall		159	44	19	0.8557	0.0190

n = no. samples analyzed; S = no. polymorphic sites; h = haplotype diversity; π = nucleotide diversity