

Supporting Table S2: Population genetic parameter estimates at the *APLIA* locus, considered separately for alleles falling in the *APLIA*¹ and *APLIA*² structural classes.

Collection	n ¹	bp ²	π_{tot} ³	θ_{tot} ⁴	TajD ⁵	π_{syn} ⁶	π_{non} ⁷
<i>APLIA</i>¹							
Bancoumana dry	19	1669	0.019	0.032	- 1.734	0.034	0.016
Bancoumana rainy	9	1665	0.048	0.054	- 0.587	0.086	0.039
Toumani-Oulena	6	1669	0.058	0.062	- 0.434	0.103	0.050
Makouchetoum	4	1669	0.054	0.057	- 0.598	0.076	0.051
all pooled	38	1665	0.048	0.048	0.623	0.104	0.106
<i>APLIA</i>²							
Bancoumana dry	0						
Bancoumana rainy	0						
Toumani-Oulena	4	1917	0.017	0.015	0.919	0.150	0.082
Makouchetoum	6	1872	0.023	0.024	- 0.457	0.034	0.018
all pooled	10	1872	0.021	0.023	- 0.542	0.034	0.016

¹number of alleles sequenced, no *APLIA*² alleles were found in the Bancoumana dry population

²locus size, in base pairs, excluding insertions and deletions

³average number of differences per pair of alleles, per nucleotide

⁴Watterson's estimator of the population genetic parameter $4N_e m$

⁵Tajima's D test statistic

⁶average number of difference per pair of alleles, per nucleotide, synonymous sites only

⁷average number of difference per pair of alleles, per nucleotide, nonsynonymous sites only