Supplementary Materials

Title: Fire and grazing determined grasslands of central Madagascar represent ancient

assemblages.

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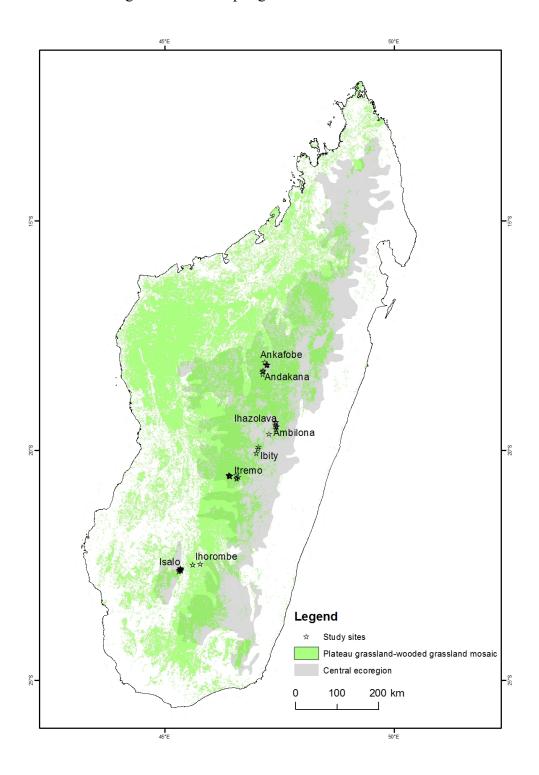
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Supplementary Figures 1 – 4

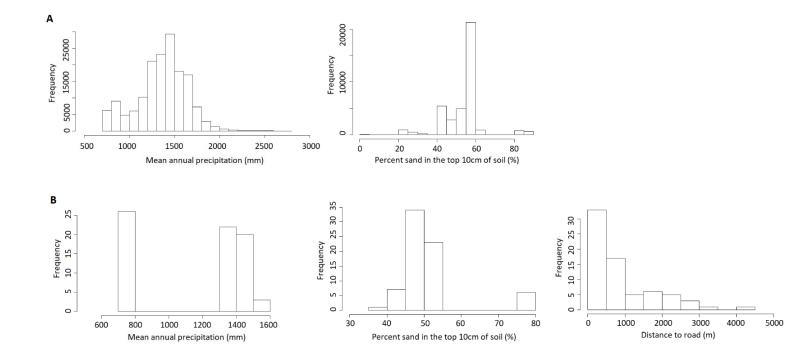
Supplementary Tables 1-3

References

Supplementary Figure 1: *Map of Madagascar depicting limits of grasslands*. The central ecoregion as per Humbert (1955) is shaded grey. Plateau grassland- wooded grassland mosaic distribution as per Moat and Smith (2007) is shaded in green. Locations of study sites are shown as are names of regions where sampling was undertaken.



Supplementary Figure 2: Environmental variables distribution (A) Histograms of mean annual precipitation (Bio_12, Worldclim Global Climate Data version) and percent sand in the top 10 cm soil (Harmonised World Soils Database) across the central ecoregion as mapped by Humbert (1955). (B) Across the 71 study sites, histograms of mean annual precipitation, percent sand in the top 10 cm soil, and distance to road.



Supplementary Table 1: *Table of all grass species encountered.* Table describes: 1) endemicity; 2) number of sites where species were found; 3) maximum number of occurrences per site (out of a maximum of 21); 4) rarity as defined and described in the methods of the main text; and, 5) assemblage group (1 or 2). Assemblage groups are based on residual correlations value between pairs of species as a product of the generalized linear latent variable model described in the main methods. The analysis used only 41 common species and post-hoc assemblage groups were assigned to the rare species.

Genera	Species	Endemic	Number of sites	Maximum	Rare	Assemblage
			of occurrence	number of		group
			(out of 71)	occurrence per		
				site (out of 21)		
Agrostis	elliotii	yes	1	1	yes	2
Alloteropsis	semialata	no	5	5	yes	2
Andropogon	itremoensis	yes	2	5	yes	2
Andropogon	trichozygus	yes	1	23	no	NA
Aristida	rufescens	yes	12	20	no	1
Aristida	similis	yes	1	5	yes	2
Aristida	tenuissima	yes	21	24	no	2
Axonopus	compressus	no	4	7	no	1
Brachiaria	arrecta	no	1	17	no	NA
Brachiaria	subrostrata	yes	3	14	no	1
Brachypodium	madagascariense	yes	1	1	yes	2
Chrysopogon	serrulatus	no	11	20	no	2

Craspedorhachis	africana	no	18	15	no	NA
Ctenium	concinnum	no	5	20	no	2
Cymbopogon	caesius	no	4	5	yes	2
Cynodon	dactylon	no	12	20	no	1
Cyrtococcum	deltoideum	yes	1	1	yes	1
Digitaria	ciliaris	no	4	7	no	1
Digitaria	debilis	no	5	5	yes	NA
Digitaria	longiflora	no	25	21	no	1
Digitaria	pseudodiaginalis	no	6	6	no	2
Digitaria	thouaresiana	no	1	2	yes	NA
Eleusine	indica	no	9	18	no	1
Eragrostis	atrovirens	no	6	19	no	1
Eragrostis	chapelieri	no	1	6	no	NA
Eragrostis	lateritica	yes	17	15	no	1
Eragrostis	racemosa	no	6	13	no	1
Eragrostis	tenella	no	2	5	yes	1
Eragrostis	tenuifolia	no	4	6	no	NA
Eulalia	villosa	no	1	4	yes	NA
Festuca	camusiana	yes	1	1	yes	2
Heteropogon	contortus	no	9	21	no	1
Hyparrhenia	newtonii	no	19	15	no	2

Hyparrhenia	rufa	no	18	21	no	1
Imperata	cylindrica	no	5	5	yes	NA
Loudetia	filifolia	no	11	21	no	2
Loudetia	simplex	no	58	25	no	2
Melinis	minutiflora	no	5	3	yes	2
Melinis	repens	no	4	5	yes	2
Microchloa	kunthii	no	7	8	no	1
Oplismenus	burmanii	no	2	1	yes	2
Panicum	cinctum	yes	12	13	no	2
Panicum	ibitense	yes	4	5	yes	2
Panicum	perrieri	yes	3	2	yes	2
Panicum	subhystrix	yes	5	5	yes	2
Panicum	umbellatum	yes	23	21	no	1
Paspalum	scrobiculatum	no	16	18	no	1
Pennisetum	pseudotriticoides	yes	5	13	no	2
Pogonarthria	squarrosa	no	2	2	yes	2
Schizachyrium	brevifolium	no	3	16	no	NA
Schizachyrium	exile	no	7	12	no	1
Schizachyrium	sanguineum	no	49	22	no	2
Setaria	pumila	no	14	20	no	1
Setaria	sphacelata	no	1	1	yes	1

Sporobolus	centrifugus	no	21	18	no	1
Sporobolus	paniculatus	no	3	7	no	1
Sporobolus	piliferus	no	1	1	yes	2
Sporobolus	pyramidalis	no	5	20	no	1
Stenotaphrum	oostachyum	yes	6	15	no	1
Stenotaphrum	unilaterale	yes	1	4	yes	1
Styppeiochloa	hitchcockii	yes	1	1	yes	2
Trachypogon	spicatus	no	45	25	no	2
Tricanthecium	brazzavillense	no	2	5	yes	2
Tricholaena	monache	no	3	5	yes	2
Tristachya	humbertii	yes	4	17	no	1
Tristachya	isalensis	yes	5	9	no	2
Urelytrum	agropyroides	no	6	19	no	2

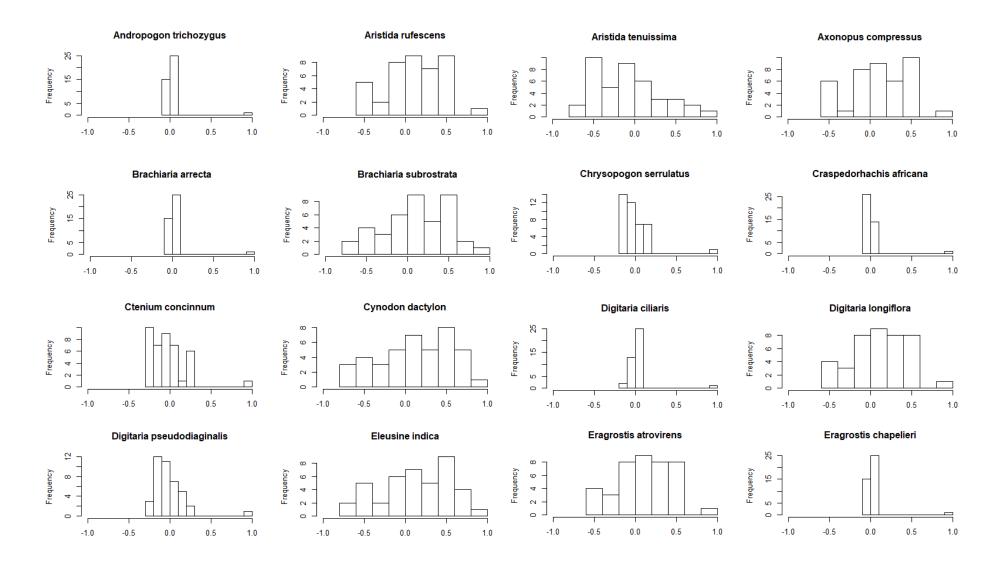
Supplementary Table 2: Description of five measured traits alongside collection method, related function and literature references. The five traits are: 1) leaf table height (H_{LT}, cm); 2) leaf thickness (LT, cm); 3) leaf size: leaf width to leaf length ratio (LW/LL); 4) growth form (mat forming, rambling, caespitose); and, 5) bulk density (BD, g/cm³)]).

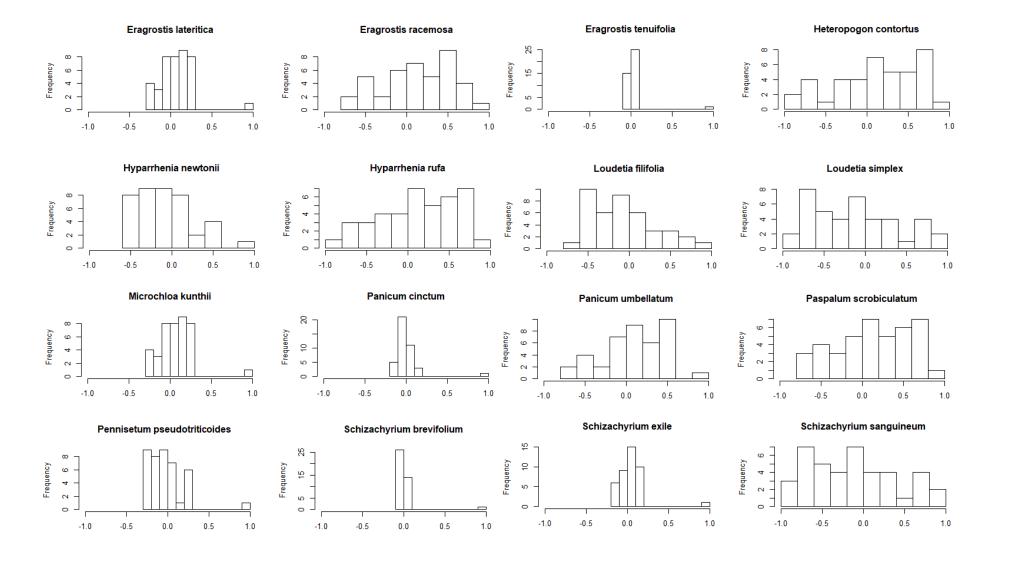
Traits	Collection method	Related function	References
Leaf table height (H _{LT} ,	The height visually estimated to	Plant height is a key functional trait with consequences for	Westoby, 1998;
cm)	correspond to the c. 80 th quantile of leaf	light competition in frequently burnt environment. Tall	Dìaz et al., 2016;
	biomass was measured on three	grasses are effective competitors for light, often associated	D'Antonio &
	individuals per species.	with high total biomass and are more flammable which	Vitousek, 1992;
		reinforce a fire feedback to increase flammability. Tall	Rossiter et al., 2003;
		grasses are "fire resistors and grazer avoiders". Short grasses	Archibald et al.,
		have low proportion of stem material and are relatively	2019; Hempson et
		higher-quality forage.	al., 2015
Leaf thickness (LT, cm)	Leaf thickness was measured on three	Leaf thickness is related to its toughness and digestibility.	Theron and Booysen,
	fully expanded leaves on each of three	Toughness is among the most important mechanical	1966; Coley, 1983;
	individuals per species.	attributes influencing grazing. Thick, tough leaves are less	Wilson et al., 1983;
		digestible to herbivores. They are hypothesized to have high	
		carbon content to make grasses more flammable. Thinner soft	
		leaves are more palatable and attract grazers.	

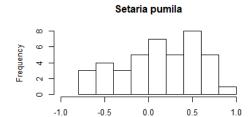
Leaf size: leaf width to	Leaf width and length were measured on	Large versus small leaves are grazing and fire attraction traits	Stobbs, 1973;
leaf length ratio	the same three leaves per individual per	respectively. Large leaves are more palatable and preferred	Archibald et al.,
(LW/LL)	species for leaf thickness measurement.	by grazers by reducing foraging time. Small leaves arranged	2019; Schwilk, 2015
		in an aerated canopy ignite easily and burn intensely, i.e.	
		more flammable.	
Growth form (mat	Growth form was recorded for each	Mat-forming habit with culms growing laterally is a grazing	Hempson et al., 2015
forming, rambling,	species.	adaptation trait. With this growth form, most of the	; Linder et al., 2018 ;
caespitose)		meristematic tissues are kept below grazing depth, allowing	Dìaz et al., 2007.
		grasses to resist intense grazing. In contrast, caespitose	Hempson et al., 2019
		grasses with erect culms can protect their meristematic tissue	; Archibald et al.,
		from fire damage with intravaginal buds protected within	2019
		basal leaf sheaths or underground, and tillers tightly	
		clustered. Caespitose growth form can be associated with	
		"generalist tolerator" and "avoider" life histories as well.	
		Rambling species are characterized by culms with an	
		architecture in between prostrate and upright, which are	
		better light competitors than mat-forming species but less	
		than caespitose species.	
Bulk density (BD,	Bulk density is the ratio between plant	Species with high bulk density attract grazers with a high	Hempson et al.,
g/cm ³)	biomass and volume. It is calculated by	density of palatable leaves clustered in the canopy which	2019; Coughenour,
	dividing the total aboveground biomass	promote grazing. Intermediate bulk density promotes fire	1985.
	by an estimate of the grass canopy	spread with enough fuel to burn and sufficient air flow for	
	volume. Volume was calculated using	combustion.	
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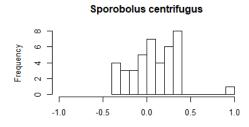
measures of the tuft basal diameter (D_B), leaf table height (H_{LT}) and leaf table diameter (D_{LT} , diameter at H_{LT}). For caespitose grasses, volume (V) was calculated using the formula for a truncated cone: $V = \pi / 3 * H_{LT} * ((D_B /$ $(2)^2 + (D_{LT}/2)^2 + D_B * D_{LT}$). For matforming grasses, a square of the individual(s) was marked out using a spade, and the volume was calculated as a cube: $V = D_B * D_{LT} * H_{LT}$. Aboveground biomass was determined on three individuals per species by clipping, drying (at 60°C for 72 h) and weighing (using a scale with two decimal place scale) the parts of the individual for which the volume estimate was made.

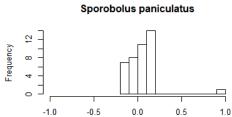
Supplementary Figure 3: *Histograms of residual correlations value, estimated from a generalized latent variable model for each species.* Model incorporates mean annual precipitation, presence/absence of fire, distance to road and a single latent variable. Values range from -1 to +1 and species with residual correlations ranging from -0.1 to +0.1 represent a lack of any association and were not classified into assemblages.

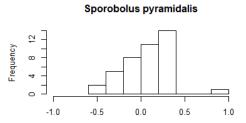




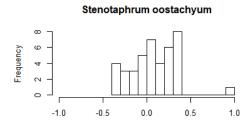


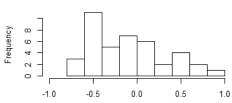




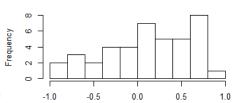


Tristachya isalensis

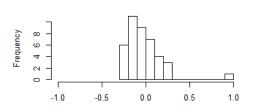


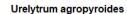


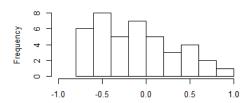
Trachypogon spicatus



Tristachya humbertii



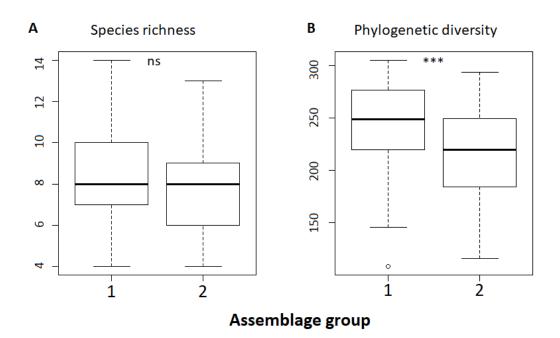




Supplementary Table 3: Table of Akaike Information Criterion (AIC) values derived from generalized latent variable models. Values correspond to the different environmental covariates' association used in the models of grass species frequency data in addition to a single unobserved predictor (latent variable). AIC values were sorted from the lowest to the highest and the model with mean annual precipitation (MAP), distance to road, presence/ absence of fire was kept for interpretation.

Environmental covariates used for the model	AIC values
MAP + distance to road + presence/ absence fire	4904.07
MAP + distance to road + presence/ absence fire + percent sand	4906.25
MAP + presence/ absence fire	4923.8
MAP + distance to road	5011.67
MAP + distance to road + percent sand	5016.9
MAP	5040.02
MAP + percent sand	5043.26
distance to road + presence/ absence fire + percent sand	5168.96
presence/ absence fire + percent sand	5179.44
distance to road + presence/ absence fire	5193.85
presence/ absence fire	5199.39
distance to road	5348.65
distance to road + percent sand	5356.67
percent sand	5363.98
null model	5393.02

Supplementary Figure 4. Grass species richness and phylogenetic diversity across assemblage group. Assemblage groups (1 and 2) are based on residual correlations values between pairs of species as a product of the generalized linear latent variable model described in the main methods. No significant differences were found between species richness but phylogenetic diversity differed significantly between the two groups (GLM, P < 0.001).



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