

1 **Elephant rewilding affects landscape openness and fauna habitat across a 92-year period**

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13 ***Appendix S2***

14 **Journal:** Ecological Applications

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17 **Appendix S2. Method used to select detection functions for Distance sampling of tree density.**

18 **Section S1:** Description of the method used to select detection functions for Distance sampling.

19 To identify the most parsimonious method to tree estimate density, three models with different
20 detection functions were compared using Akaike Information Criterion (AIC) values: half-normal
21 key function and cosine adjustment, a uniform key function and cosine adjustment and a hazard-rate
22 key function and simple polynomial adjustment. To account for differences in detectability between
23 reserves with different elephant reintroduction times (and subsequent differences in vegetation
24 density / cover), model selection was conducted separately for the reserve without elephants and the
25 reserves with a younger elephant reintroduction time (i.e. “thick” vegetation; 1995, 2003) and the
26 reserves with an older elephant reintroduction time (i.e. sparse vegetation; 1972, 1927). The function
27 with the lowest AIC score was used to estimate density. All models falling within two AIC points
28 of the “best” model were deemed equivalent. Where this occurred, we attempted to use the same
29 detection function between transects (Table SI.3).

30 **Table S1:** Akaike Information Criterion (AIC) values for Distance sampling models used to
31 estimate the density of trees bearing small- (a), medium- (b) and large-sized (c) tree hollows (HBT)
32 and the density of all trees (d), *Senegalia nigrescens* (e) and *Sclerocarya birrea* (f) trees. Different
33 detection functions were fit for sites occurring at three reserves with “thick” vegetation (maximum
34 transect width 20 m) and two reserves with “sparse” vegetation (maximum transect 40 m). *
35 indicates the detection function used to estimate density.

Detection function	Transect width	
	20 m	40 m
<i>a) Small HBT</i>		
Uniform cosine	592.31	210.36

Half-normal cosine	591.45*	210.79*
Hazard-rate simple polynomial	592.73	210.64
<i>b) Medium HBT</i>		
Uniform cosine	363.31	294.26
Half-normal cosine	362.46	293.90
Hazard-rate simple polynomial	358.99*	291.68*
<i>c) Large HBT</i>		
Uniform cosine	63.10*	130.14*
Half-normal cosine	63.47	130.77
Hazard-rate simple polynomial	65.18	131.18
<i>d) All trees</i>		
Uniform cosine	3601.41*	935.17*
Half-normal cosine	3606.32	934.88
Hazard-rate simple polynomial	3621.14	934.92
<i>e) Senegalia nigrescens</i>		
Uniform cosine	2202.66	326.19
Half-normal cosine	2200.63*	326.67*
Hazard-rate simple polynomial	2209.97	327.21
<i>f) Sclerocarya birrea</i>		
Uniform cosine	1114.27	373.12
Half-normal cosine	1114.04*	373.96*
Hazard-rate simple polynomial	1117.84	373.82
