

Table S1 – AIC model selection (Val Maira and Capanne di Marcarolo)

SD + NDVI + Altitude + WildRuminants_TO + WildBoar_TO + Mesocarnivores_TO	204
SD + NDVI + Altitude + s(WildRuminants_TO) + s(WildBoar_TO) + s(sMesocarnivores1)	198
SD + NDVI + Altitude + s(WildRuminants_TO, Altitude, k=4) + s(WildBoar_TO, Altitude, k=4) + sMesocarnivores1*Altitude	202
SD + NDVI + Altitude + s(WildRuminants_TO, Altitude, by=as.factor(Site), k=4) + s(WildBoar_TO, Altitude, by=as.factor(Site), k=4) + sMesocarnivores1*as.factor(Site)	187

Table S2 – Model output

Family: Tweedie(p=1.512)
Link function: log

Formula:
Numero1 ~ SD1 + NDVI1 + Altitudine + s(sRoe1, Altitudine, by = as.factor(Site), k = 4) + s(sBoar1, Altitudine, by = as.factor(Site), k = 4) + sMesocarnivores1

Parametric coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	0.0000000	0.0000000	NAN	NAN
SD1	-0.0660683	0.3100705	-0.213	0.8322
NDVI1	0.1788460	0.2148964	0.832	0.4096
Altitudine	-0.0008294	0.0004866	-1.705	0.0951 .
sMesocarnivores1	-0.0245347	0.1954541	-0.126	0.9007

Signif. codes: 0 ‘***’ 0.001 ‘**’ 0.01 ‘*’ 0.05 ‘.’ 0.1 ‘ ’ 1

Approximate significance of smooth terms:

	edf	Ref.df	F	p-value
s(sRoe1,Altitudine):as.factor(Site)1	2.000	2.000	4.486	0.0167 *
s(sRoe1,Altitudine):as.factor(Site)2	2.923	2.992	4.438	0.0104 *
s(sBoar1,Altitudine):as.factor(Site)1	1.793	1.955	2.281	0.0852 .
s(sBoar1,Altitudine):as.factor(Site)2	1.000	1.000	6.364	0.0152 *

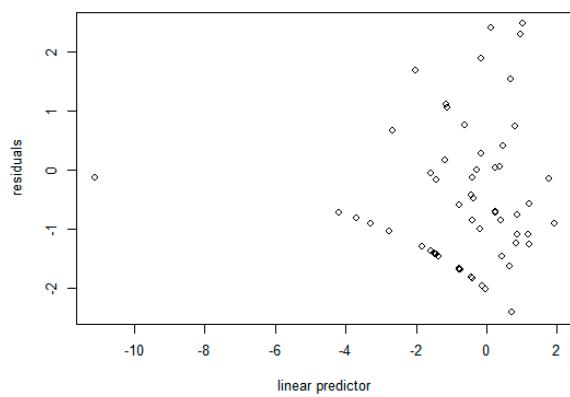
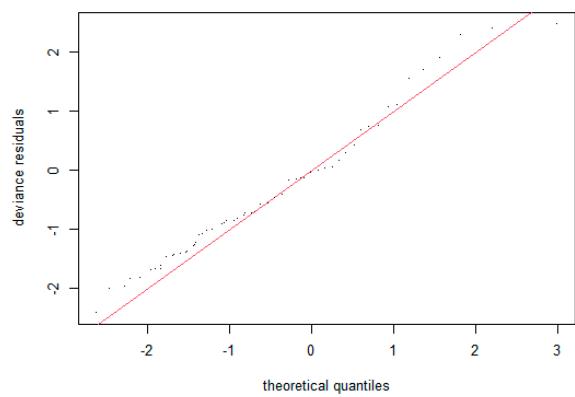
Signif. codes: 0 ‘***’ 0.001 ‘**’ 0.01 ‘*’ 0.05 ‘.’ 0.1 ‘ ’ 1

Rank: 14/15

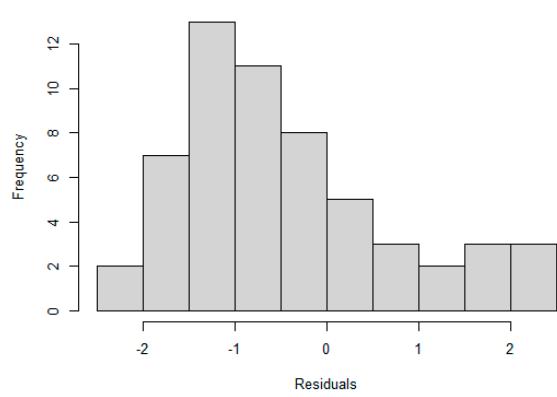
R-sq.(adj) = 0.101 Deviance explained = 43.4%
-REML = 91.609 Scale est. = 1.6992 n = 57

Figure S1 – Gam model check

Resids vs. linear pred.



Histogram of residuals



Response vs. Fitted Values

