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Supporting Information

Table S1 Locations of the six study sites. Latitude, longitude and altitude are given for the centre of each 40 x 40 m study plot. Latitude and longitude are given in decimal degrees to 5 decimal places, and therefore are accurate to within approximately 1 m. Aspect and slope were calculated from Intermap NEXTMap 5 data, accessed through Strava.

Site	Treatment	Latitude (°)	Longitude (°)	Altitude (m asl)	Aspect	Slope (%)
F1	Burned	37.17713	-7.86082	261	E	20
F2	Burned	37.19641	-7.86006	372	WNW	30
F3	Burned	37.19848	-7.85699	449	W	22.5
NF1	Unburned	37.16919	-7.86516	340	NE	14
NF2	Unburned	37.17334	-7.86506	317	W	12
NF3	Unburned	37.18149	-7.86588	245	NW	10

Table S2 Summary of the plant families identified on floristic surveys, with the number of species and the total number of 1 x 1 m quadrats in which members of each family was recorded. *Apiaceae* includes one flower type not identified beyond family level, and therefore was represented by *at least* four species

Family	No. species	No. quadrats
Adoxaceae	1	4
Amaryllidaceae	2	17
Apiaceae	4*	29
Asparagaceae	2	3
Asteraceae	14	379
Boraginaceae	2	39
Brassicaceae	1	3
Campanulaceae	2	25
Caprifoliaceae	2	16
Caryophyllaceae	2	8
Cistaceae	4	167
Ericaceae	3	26
Euphorbiaceae	1	4
Fabaceae	10	285
Gentianaceae	1	36
Geraniaceae	1	2
Iridaceae	2	11
Lamiaceae	3	200
Linaceae	1	1
Orchidaceae	1	1
Papaveraceae	1	2
Plantaginaceae	1	30
Primulaceae	1	24
Ranunculaceae	2	30
Resedaceae	2	23
Rosaceae	2	12
Solanaceae	1	39
Thymelaeaceae	1	8

Table S3 R packages used during analysis. Packages were loaded into at least one script during the analytical process but may not have formed part of the final analysis.

Package	Citation
AICcmodavg	Mazerolle, M.J. (2016) AICcmodavg: Model selection and multimodel inference based on (Q)AIC(c). R package version 2.1-0. https://cran.r-project.org/package=AICcmodavg .
arm	Gelman, A. & Su, Y.-S. (2015) arm: Data Analysis Using Regression and Multilevel/Hierarchical Models. R package version 1.8-6. https://CRAN.R-project.org/package=arm .
bipartite	Dormann, C.F., Gruber B. & Fruend, J. (2008) Introducing the bipartite Package: Analysing Ecological Networks. <i>R News</i> , 8 , 8–11.
car	Fox, J. & Weisberg, S. (2011) <i>An {R} Companion to Applied Regression</i> . Second Edition. Sage, Thousand Oaks, CA, USA.
data.table	Dowle, M., Srinivasan, A., Short, T., Lianoglou, S., Saporta, R. & Antonyan, E. (2015) data.table: Extension of Data.frame. R package version 1.9.6. https://CRAN.R-project.org/package=data.table .
effects	Fox, J. (2003) Effect Displays in R for Generalised Linear Models. <i>Journal of Statistical Software</i> , 8 , 1–27.
ggmap	Kahle, D. & Wickham, H. (2013) ggmap: Spatial Visualization with ggplot2. <i>The R Journal</i> , 5 , 144–161.
ggplot2	Wickham, H. (2009) <i>ggplot2: Elegant Graphics for Data Analysis</i> . Springer-Verlag, New York, USA.
glmmADMB	Fournier, D.A., Skaug, H.J., Ancheta, J., Ianelli, J., Magnusson, A., Maunder, M., Nielsen, A. & Sibert, J. (2012) AD Model Builder: using automatic differentiation for statistical inference of highly parameterized complex nonlinear models. <i>Optimization Methods and Software</i> , 27 , 233–249.
gridExtra	Auguie, B. (2016) gridExtra: Miscellaneous Functions for "Grid" Graphics. R package version 2.2.1. https://CRAN.R-project.org/package=gridExtra .
lme4	Bates, D., Maechler, M., Bolker, B. & Walker, S. (2015). Fitting Linear Mixed-Effects Models Using lme4. <i>Journal of Statistical Software</i> , 67 , 1–48.
MASS	Venables, W. N. & Ripley, B. D. (2002) <i>Modern Applied Statistics with S</i> . Fourth Edition. Springer, New York, USA.
plyr	Wickham, H. (2011) The Split-Apply-Combine Strategy for Data Analysis. <i>Journal of Statistical Software</i> , 40 , 1–29.
RColorBrewer	Neuwirth, E. (2014) RColorBrewer: ColorBrewer Palettes. R package version 1.1-2. https://CRAN.R-project.org/package=RColorBrewer .
reshape2	Wickham, H. (2007) Reshaping Data with the reshape Package.

Journal of Statistical Software, **21**, 1–20.

RVAideMemoire	Hervé, M. (2016) RVAideMemoire: Diverse Basic Statistical and Graphical Functions. R package version 0.9-56. https://CRAN.R-project.org/package=RVAideMemoire .
scales	Wickham, H. (2016) scales: Scale Functions for Visualization. R package version 0.4.1. https://CRAN.R-project.org/package=scales .
svglite	Wickham, H., Henry, L., Luciani, T.J., Decorde, M. & Lise, V. (2016) svglite: An 'SVG' Graphics Device. R package version 1.2.0. https://CRAN.R-project.org/package=svglite .
tidyr	Wickham, H. (2016) tidyr: Easily Tidy Data with `spread()` and `gather()` Functions. R package version 0.5.0. https://CRAN.R-project.org/package=tidyr .
vegan	Oksanen, J., Blanchet, F.G., Kindt, R., Legendre, P., Minchin, P.R., O'Hara, R.B., Simpson, G.L., Solymos, P., Stevens, M.H.H. & Wagner, H. (2016) vegan: Community Ecology Package. R package version 2.3-5. https://CRAN.R-project.org/package=vegan .

Table S4 Summary of captured insects according to Order, Family, Genus, number of identified species, number of morphotypes and the total number of individuals.

Order	Family	Genus	Identified species	No. morphotypes	No. individuals
Coleoptera				3	7
	Cerambycidae			4	8
			<i>Nustera distigma</i>		1
	Cetoniidae			3	10
		Protaetia		1	2
			<i>Protaetia opaca</i>		2
			<i>Oxythyrea funesta</i>		89
			<i>Tropinota squalida</i>		14
	Chrysomelidae			3	22
			<i>Chrysolina americana</i>		1
			<i>Dicladispa testacea</i>		1
			<i>Lachnaia hirta</i>		1
	Cleridae			1	1
			<i>Trichodes octopunctatus</i>		1
	Coccinellidae			1	4
			<i>Coccinella septempunctata</i>		5
	Curculionidae			1	3
	Dasytidae			2	17
	Elateridae			4	19
	Malachiidae			1	2
	Meloidae			4	9
	Mordellidae			1	3
	Oedemeridae			4	8
	Tenebrionidae			2	3
			<i>Heliothaurus ruficollis</i>		82
Diptera				1	2
	Bombyliidae			3	7
	Empididae			2	2
	Syrphidae			3	5
	Tachinidae			3	5
	Tipulidae			1	1
Hemiptera				2	2
	Acanthosomatidae			1	1
	Pentatomidae			1	1
			<i>Graphosoma lineatum</i>		2
Heteroptera	Miridae			1	1
Hymenoptera	Andrenidae			2	2
		Andrena		3	9
		Panurgus		1	1
	Apidae			1	1
		Eucera		1	1
		Xylocopa		1	1
			<i>Apis mellifera</i>		79
			<i>Bombus terrestris</i>		2
			<i>Bombus hortorum</i>		2
	Crabronidae			1	1
	Halictidae			3	5
		Lasioglossum		3	4
	Colletidae	Hylaeus		1	1
	Ichneumonidae			1	1
	Megachilidae			1	1
		Rhodanthidium		1	1
	Sphecidae			1	1
	Vespidae			2	2
			<i>Polistes dominula</i>		4
			<i>Vespa crabro</i>		3
			<i>Vespula vulgaris</i>		5

Order	Family	Genus	Identified species	No. morphotypes	No. individuals
Lepidoptera				1	1
	Erebidae		<i>Utetheisa pulchella</i>		1
	Gelechiidae		<i>Dichomeris lamprostoma</i>		1
	Geometridae		<i>Aspitates ochrearia</i>		1
			<i>Scotopteryx peribolata</i>		1
	Hesperiidae	Thymelicus		1	1
	Lasiocampidae		<i>Trichiura ilicis</i>		1
	Lycaenidae		<i>Aricia cramera</i>		2
			<i>Lampides boeticus</i>		1
			<i>Leptotes pirithous</i>		8
			<i>Satyrium ilicis</i>		2
			<i>Plebejus argus</i>		1
			<i>Polyommatus icarus</i>		2
	Noctuidae		<i>Autographa gamma</i>		1
			<i>Synthimia fixa</i>		1
	Nymphalidae		<i>Coenonympha pamphilus</i>		7
			<i>Lasiommata megera</i>		1
			<i>Maniola jurtina</i>		20
			<i>Melanargia ines</i>		5
			<i>Pararge aegeria</i>		1
			<i>Pyronia cecilia</i>		2
			<i>Pyronia tithonus</i>		2
			<i>Vanessa cardui</i>		2
	Papilionidae		<i>Iphiclides feisthamelii</i>		1
			<i>Papilio machaon</i>		1
	Pieridae		<i>Colias croceus</i>		2
			<i>Euchloe crameri</i>		6
			<i>Pieris brassicae</i>		1
			<i>Pieris rapae</i>		4
			<i>Pontia daplidice</i>		1
	Pterophoridae		<i>Amblyptilia acanthadactyla</i>		1
	Pyralidae		<i>Endotricha flammealis</i>		1
	Tortricidae			1	1
Neuroptera				1	1
	Chrysopidae			1	1
Unidentified				10	15
Totals	44	9	47	91	572

Table S5 – Number of pollen grains found on insects, according to species and family, grouped by season and by treatment.

Pollen species	Family	Total Count	Spring Count	Summer count	Autumn count	Winter count	Burned count	Unburned count
<i>Allium paniculatum</i>	Amaryllidaceae	50	3	0	0	47	49	1
<i>Anagallis arvensis</i>	Primulaceae	536	434	11	12	79	407	129
<i>Anarrhinum bellidifolium</i>	Plantaginaceae	4746	4535	118	1	92	67	4679
<i>Andryala integrifolia</i>	Asteraceae	5756	4289	1	445	1021	5590	166
<i>Anthyllis vulneraria</i>	Fabaceae	611	529	39	12	31	224	387
<i>Apiaceae</i>	Apiaceae	4407	4296	29	14	68	3252	1155
<i>Arbutus unedo</i>	Ericaceae	81	0	1	79	1	79	2
<i>Calluna vulgaris</i>	Ericaceae	197	159	5	0	33	167	30
<i>Campanula lusitanica</i>	Campanulaceae	308	243	16	5	44	165	143
<i>Carlina racemosa</i>	Asteraceae	2	2	0	0	0	2	0
<i>Centaureum erythraea</i>	Gentianaceae	54	32	2	10	10	14	40
<i>Centranthus calcitrapae</i>	Valerianaceae	139	84	0	2	53	139	0
<i>Chamaemelum mixtum</i>	Asteraceae	1856	1676	6	1	173	49	1807
<i>Cistus crispus</i>	Cistaceae	4147	3982	1	36	128	279	3868
<i>Cistus ladanifer</i>	Cistaceae	1788	1082	3	25	678	1159	629
<i>Cistus salviifolius</i>	Cistaceae	7304	2319	4	28	4953	1792	5505
<i>Cistus sp.</i>	Cistaceae	4309	939	373	0	2997	3911	398
<i>Coleostephus myconis</i>	Asteraceae	15432	13592	5	1754	81	15034	398
<i>Cynara cardunculus</i>	Asteraceae	530	312	186	0	32	38	492
<i>Daphne gnidium</i>	Thymelaeaceae	200	142	12	17	29	53	147
<i>Daucus carota</i>	Apiaceae	377	70	233	9	65	312	65
<i>Echium plantagineum</i>	Boraginaceae	262	239	15	0	8	209	53
<i>Erica arborea</i>	Ericaceae	46	16	0	1	29	31	15
<i>Eryngium campestre</i>	Apiaceae	61	32	10	15	4	61	0
<i>Euphorbia exigua</i>	Euphorbiaceae	14	14	0	0	0	0	14
<i>Euphorbia sp.</i>	Euphorbiaceae	2	2	0	0	0	2	0
<i>Galactites tomentosus</i>	Asteraceae	3547	2717	1	6	823	3493	54
<i>Genista triacanthos</i>	Fabaceae	2015	467	0	768	780	2015	0
<i>Gladiolus italicus</i>	Iridaceae	2	0	0	0	2	2	0
<i>Helychrysum stoechas</i>	Asteraceae	4178	2295	360	109	1414	1790	2388
<i>Jasione montana</i>	Campanulaceae	2382	722	50	34	1576	2118	264
<i>Lavandula stoechas</i>	Lamiaceae	11574	10051	26	5	1492	2873	8701
<i>Leontodon taraxacoides</i>	Asteraceae	5512	3661	216	0	1635	2131	3381
<i>Leucojum autumnale</i>	Amaryllidaceae	134	22	5	102	5	121	13
<i>Lithodora prostrata</i>	Boraginaceae	4172	169	0	27	3976	4169	3
<i>Logfia gallica</i>	Asteraceae	32	31	0	0	1	13	19
<i>Lotus parviflorus</i>	Fabaceae	16	0	0	0	16	16	0
<i>Lotus subbiflorus</i>	Fabaceae	1073	1073	0	0	0	0	1073
<i>Ornithogalum broteroi</i>	Asparagaceae	265	253	5	4	3	245	20
<i>Pulicaria odora</i>	Asteraceae	9292	8744	263	232	53	5695	3597

Pollen species	Family	Total Count	Spring Count	Summer count	Autumn count	Winter count	Burned count	Unburned count
<i>Pulicaria paludosa</i>	Asteraceae	228	150	3	27	48	133	95
<i>Ranunculus bullatus</i>	Ranunculaceae	4	1	3	0	0	0	4
<i>Ranunculus gramineus</i>	Ranunculaceae	1125	633	124	0	368	660	465
<i>Ranunculus sp.</i>	Ranunculaceae	9	9	0	0	0	9	0
<i>Reseda media</i>	Resedaceae	726	615	20	62	29	647	79
<i>Sanguisorba minor</i>	Rosaceae	6	6	0	0	0	0	6
<i>Sanguisorba verrucosa</i>	Rosaceae	376	361	0	8	7	133	243
<i>Scilla autumnalis*</i>	Asparagaceae	15	0	0	0	15	15	0
<i>Scilla monophyllos*</i>	Asparagaceae	42	6	7	2	27	34	8
<i>Silene gallica</i>	Caryophyllaceae	80	60	1	7	12	42	38
<i>Solanum nigrum</i>	Solanaceae	252	127	12	90	23	107	145
<i>Spergularia purpurea</i>	Caryophyllaceae	52	15	10	7	20	42	10
<i>Stachys arvensis</i>	Lamiaceae	85	84	0	0	1	84	1
<i>Thymus mastichina</i>	Lamiaceae	194	174	0	3	17	92	102
<i>Trifolium arvense</i>	Fabaceae	5660	5374	8	213	65	5551	109
<i>Tuberaria guttata</i>	Cistaceae	26866	25960	301	3	602	34	26832
<i>Ulex argenteus</i>	Fabaceae	8846	1837	3041	0	3968	7425	1421
<i>Ulex eriocladus</i>	Fabaceae	9225	144	2	5351	3728	6534	2691
<i>Ulex sp.</i>	Fabaceae	17	0	0	0	17	17	0
<i>Urginea maritima</i>	Asparagaceae	176	163	3	6	4	170	6
<i>Viburnum tinus</i>	Adoxaceae	29	7	0	0	22	20	9
Total pollen count		151422						

* - Species identified outside the quadrats during floristic surveys.

Table S6 R software outputs showing summary of analyses of the effects of burning and season over consecutive sampling periods on the abundance (A) and species richness (B) of insects. Intercept value represents unburned sites in autumn, sampling period 0, and is the natural logarithm of the estimate, so $e^{(\text{intercept})}$ gives the true estimated value. For other levels of each variable, estimated value = $e^{(\text{intercept})} \times e^{\text{ES}}$, where ES = effect size for that level from the statistical model, so e^{ES} is the multiplicative effect of the parameter in question.

A. Insect Abundance

```
Generalized linear mixed model fit by maximum likelihood (Laplace
Approximation) [glmerMod]
Family: poisson ( log )
Formula: Count ~ Treatment * Order + (1 | Site)
Data: dframe3
```

AIC	BIC	logLik	deviance	df.resid
575.6	623.9	-268.8	537.6	75

Scaled residuals:

Min	1Q	Median	3Q	Max
-3.2066	-0.9665	-0.2279	0.5000	4.8677

Random effects:

Groups Name	Variance	Std.Dev.
Site (Intercept)	6.545e-19	8.09e-10

Number of obs: 94, groups: Site, 6

Fixed effects:

	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	1.6529	0.1459	11.332	< 2e-16 ***
TreatmentFire	0.4420	0.1915	2.309	0.020968 *
Order2	-0.9598	0.4335	-2.214	0.026836 *
Order3	-0.7366	0.3482	-2.115	0.034408 *
Order4	-0.2666	0.2509	-1.063	0.287894
Order5	0.7894	0.1891	4.174	2.99e-05 ***
Order6	-1.2962	0.3482	-3.722	0.000197 ***
Order7	-0.9598	0.4335	-2.214	0.026836 *
Order8	0.2640	0.2251	1.173	0.240957
Order9	0.8594	0.2198	3.910	9.22e-05 ***
TreatmentFire:Order2	-0.4420	0.5730	-0.771	0.440458
TreatmentFire:Order3	-1.3583	0.6856	-1.981	0.047556 *
TreatmentFire:Order4	0.6731	0.3078	2.187	0.028735 *
TreatmentFire:Order5	-0.5653	0.2599	-2.175	0.029637 *
TreatmentFire:Order6	-0.3467	0.4770	-0.727	0.467351
TreatmentFire:Order7	-0.1543	0.5156	-0.299	0.764672
TreatmentFire:Order8	-0.1077	0.2892	-0.372	0.709677
TreatmentFire:Order9	-0.2028	0.2915	-0.696	0.486605

Single term deletions

Model:

```
Count ~ Treatment * Order + (1 | Site)
      Df   AIC   LRT Pr(Chi)
<none>      575.63
Treatment:Order  8 583.27 23.645 0.002628 **
```

B. Insects Species Richness

Generalized linear mixed model fit by maximum likelihood (Laplace Approximation) [glmerMod]

Family: poisson (log)

Formula: SpeciesRichness ~ Treatment * Order + (1 | Site)

Data: dframe3

AIC	BIC	logLik	deviance	df.resid
383.5	431.9	-172.8	345.5	75

Scaled residuals:

Min	1Q	Median	3Q	Max
-1.6209	-0.7071	-0.1075	0.6299	2.1930

Random effects:

Groups Name	Variance	Std.Dev.
Site (Intercept)	0	0

Number of obs: 94, groups: Site, 6

Fixed effects:

	Estimate	Std. Error	z value	Pr(> z)	
(Intercept)	1.35812	0.16903	8.035	9.38e-16	***
TreatmentFire	0.08880	0.24080	0.369	0.71231	
Order2	-0.66498	0.44186	-1.505	0.13233	
Order3	-0.66498	0.39188	-1.697	0.08972	.
Order4	-0.15415	0.28031	-0.550	0.58236	
Order5	0.46103	0.23579	1.955	0.05055	.
Order6	-1.10681	0.37374	-2.961	0.00306	**
Order7	-0.66498	0.44186	-1.505	0.13234	
Order8	0.12348	0.27208	0.454	0.64994	
Order9	0.25131	0.30861	0.814	0.41544	
TreatmentFire:Order2	-0.08880	0.59131	-0.150	0.88063	
TreatmentFire:Order3	-0.78194	0.71855	-1.088	0.27650	
TreatmentFire:Order4	-0.06899	0.40842	-0.169	0.86585	
TreatmentFire:Order5	-0.23398	0.34097	-0.686	0.49258	
TreatmentFire:Order6	-0.08880	0.52934	-0.168	0.86678	
TreatmentFire:Order7	0.13435	0.53974	0.249	0.80343	
TreatmentFire:Order8	-0.41772	0.39506	-1.057	0.29034	
TreatmentFire:Order9	0.33865	0.41003	0.826	0.40886	

Single term deletions

Model:

SpeciesRichness ~ Treatment * Order + (1 | Site)

	Df	AIC	LRT	Pr(Chi)
<none>		383.54		
Treatment:Order	8	372.20	4.663	0.7929

Model:

SpeciesRichness ~ Treatment + Order + (1 | Site)

	Df	AIC	LRT	Pr(Chi)
<none>		372.20		
Treatment	1	370.21	0.010	0.9221
Order	8	427.64	71.439	2.539e-12 ***

Table S7 R software outputs showing summary of analyses of the effects of burning and season over consecutive sampling periods on pollen transport (A. pollen load and B. pollen type). Intercept value represents unburned sites in the first spring, and is the base 10 logarithm of the estimate, so $10^{(\text{intercept})}$ gives the true estimated value. For other levels of each variable, estimated value = $10^{(\text{intercept})} \times 10^{\text{ES}}$, where ES = effect size for that level from the statistical model, so 10^{ES} is the multiplicative effect of the parameter in question.

A. Pollen Load

```
Linear mixed model fit by REML ['lmerMod']
Formula: log10(PollenLoad + 1) ~ Treatment + Order + Treatment * Order +
(1 | Site)
Data: dframe1
```

REML criterion at convergence: 1218.6

Scaled residuals:

Min	1Q	Median	3Q	Max
-2.9451	-0.6275	-0.0238	0.5986	3.4846

Random effects:

Groups	Name	Variance	Std.Dev
Site	(Intercept)	0.02495	0.157
	Residual	0.47211	0.687

Number of obs: 572, groups: Site, 6

Fixed effects:

	Estimate	Std. Error	t value
(Intercept)	1.55145	0.13554	11.446
TreatmentFire	0.59399	0.18436	3.222
Order2	-0.32662	0.29960	-1.090
Order3	-0.42639	0.23960	-1.780
Order4	0.03083	0.17249	0.179
Order5	0.38767	0.12998	2.983
Order6	0.16892	0.23944	0.705
Order7	-0.19814	0.29847	-0.664
Order8	0.15243	0.15654	0.974
Order9	0.61369	0.15102	4.064
TreatmentFire:Order2	-0.62100	0.39595	-1.568
TreatmentFire:Order3	-0.63886	0.47125	-1.356
TreatmentFire:Order4	-0.48986	0.21160	-2.315
TreatmentFire:Order5	-0.76111	0.17953	-4.239
TreatmentFire:Order6	-0.80604	0.32800	-2.457
TreatmentFire:Order7	-0.05769	0.35494	-0.163
TreatmentFire:Order8	-0.31886	0.20047	-1.591
TreatmentFire:Order9	-0.82712	0.20123	-4.110

Single term deletions

Model:

```
log10(PollenLoad + 1) ~ Treatment + Order + Treatment * Order +
(1 | Site)
```

	Df	AIC	LRT	Pr(Chi)
<none>		1226		
Treatment:Order	8	1238	28.009	0.0004725 ***

B. Pollen Type

Linear mixed model fit by REML ['lmerMod']

Formula:

$\log_{10}(\text{PollenTypes} + 1) \sim \text{Treatment} + \text{Order} + \text{Treatment} * \text{Order} +$
(1 | Site)

Data: dframe1

REML criterion at convergence: -175.1

Scaled residuals:

Min	1Q	Median	3Q	Max
-3.5314	-0.6221	0.0362	0.7192	2.4624

Random effects:

Groups	Name	Variance	Std.Dev.
Site	(Intercept)	0.001888	0.04345
	Residual	0.038166	0.19536

Number of obs: 572, groups: Site, 6

Fixed effects:

	Estimate	Std. Error	t value
(Intercept)	0.63246	0.03798	16.654
TreatmentFire	0.20680	0.05159	4.008
Order2	-0.02242	0.08518	-0.263
Order3	-0.12098	0.06812	-1.776
Order4	-0.02393	0.04904	-0.488
Order5	-0.01950	0.03696	-0.528
Order6	0.04339	0.06808	0.637
Order7	0.07695	0.08486	0.907
Order8	-0.02602	0.04450	-0.585
Order9	0.24951	0.04294	5.811
TreatmentFire:Order2	-0.27207	0.11257	-2.417
TreatmentFire:Order3	-0.25821	0.13399	-1.927
TreatmentFire:Order4	-0.16031	0.06016	-2.665
TreatmentFire:Order5	-0.13779	0.05104	-2.700
TreatmentFire:Order6	-0.31653	0.09326	-3.394
TreatmentFire:Order7	-0.22185	0.10092	-2.198
TreatmentFire:Order8	-0.21946	0.05699	-3.851
TreatmentFire:Order9	-0.27552	0.05721	-4.816

Table S8 – Mean temperatures taken between 12 – 16 pm on each day of fieldwork.

Year	Dates	Mean Temperatures (°C)
2013	16 Apr	27.8
	14 May	27.5
	11 Jun	27.8
	09 Jul	31.5
	24 Sep	32.3
	05 Nov	23.6
	09 Dec	19.3
2014	28 Jan	15.4
	19 Feb	20.3
	25 Mar	24.5
	29 Apr	33.8
	03 Jun	34.6
	15 Jul	37.4
	12 Aug	34.2
	23 Sep	24.0
	28 Oct	25.7
	02 Dec	22.5
2015	21 Jan	14.8
	25 Feb	19.3
	25 Mar	20.2
	05 May	27.5

Additional graphs

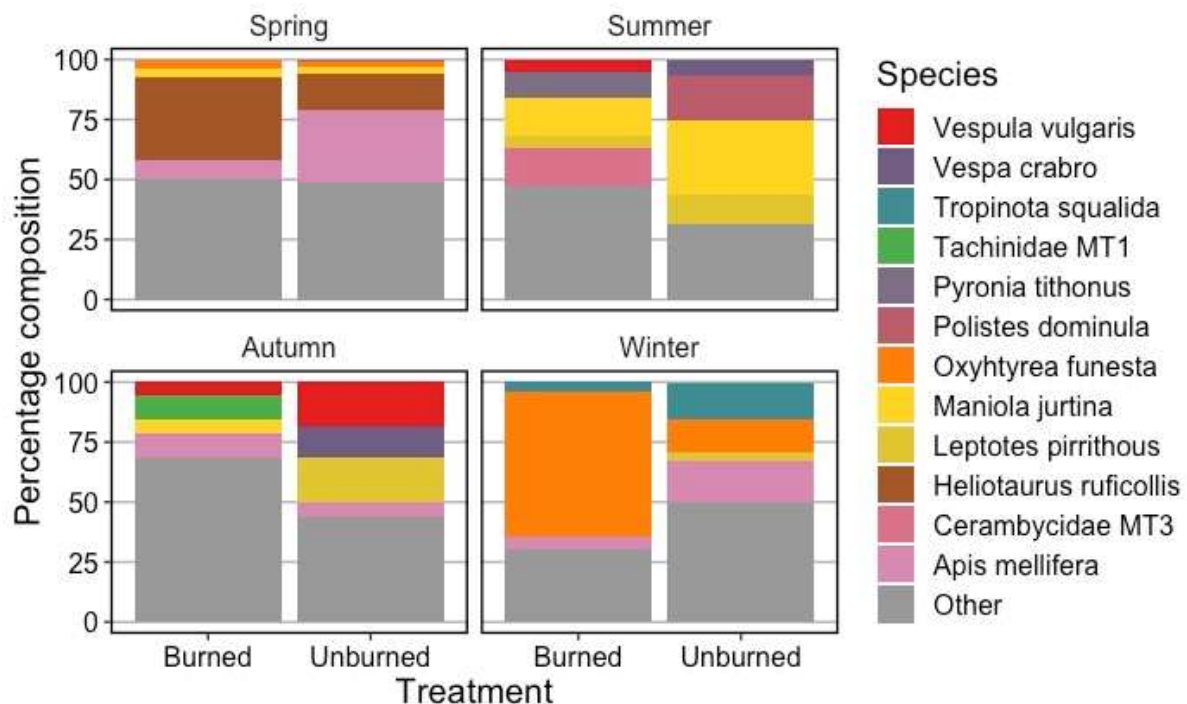


Figure a. Assemblage composition by species of insects caught in burned and unburned areas of Serra-do-Caldeirão, Portugal, and across seasons. Species never comprising >10% of individuals in any combination of season and treatment are grouped as “Others”, and all other families are shown independently.

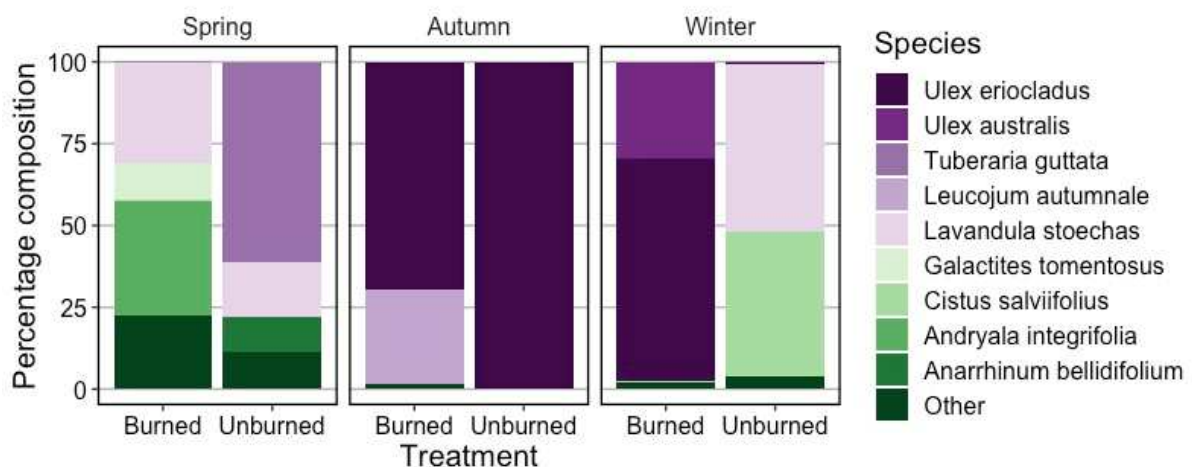


Figure b. Assemblage composition by families of pollen carried by *Apis mellifera* caught in burned and unburned plots of Serra-do-Caldeirão, Portugal, and across seasons. Families never comprising >7% of individuals in any combination of season and treatment are grouped as “Other”, and all other families are shown independently.

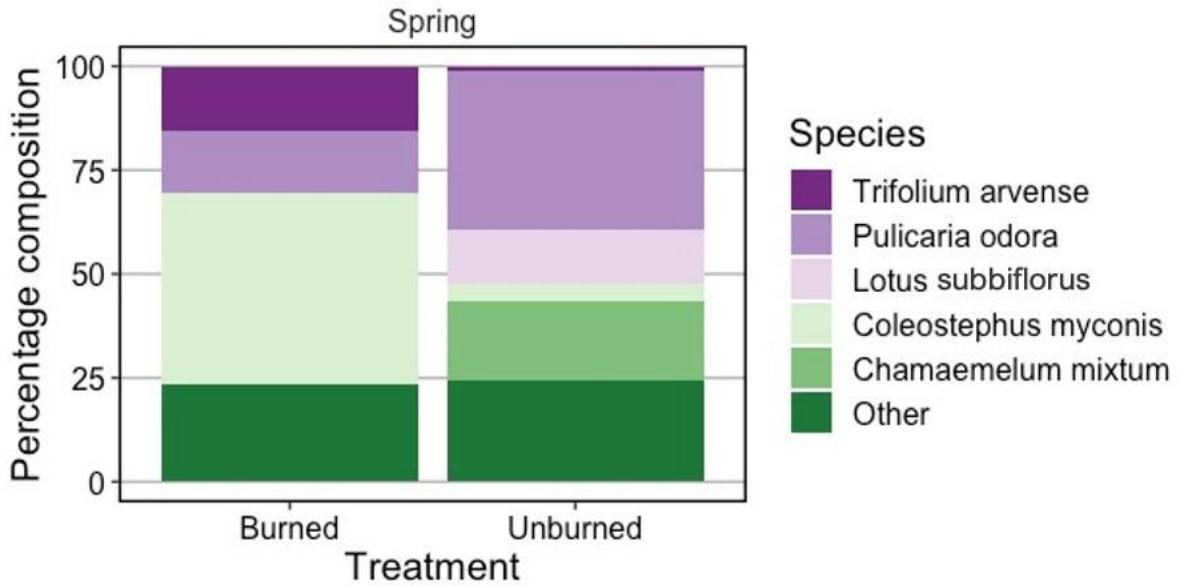


Figure c. Assemblage composition by families of pollen carried by *H. rufficollis* caught in burned and unburned plots of Serra-do-Caldeirão, Portugal, and across seasons. Families never comprising >7% of individuals in any combination of season and treatment are grouped as “Other”, and all other families are shown independently.

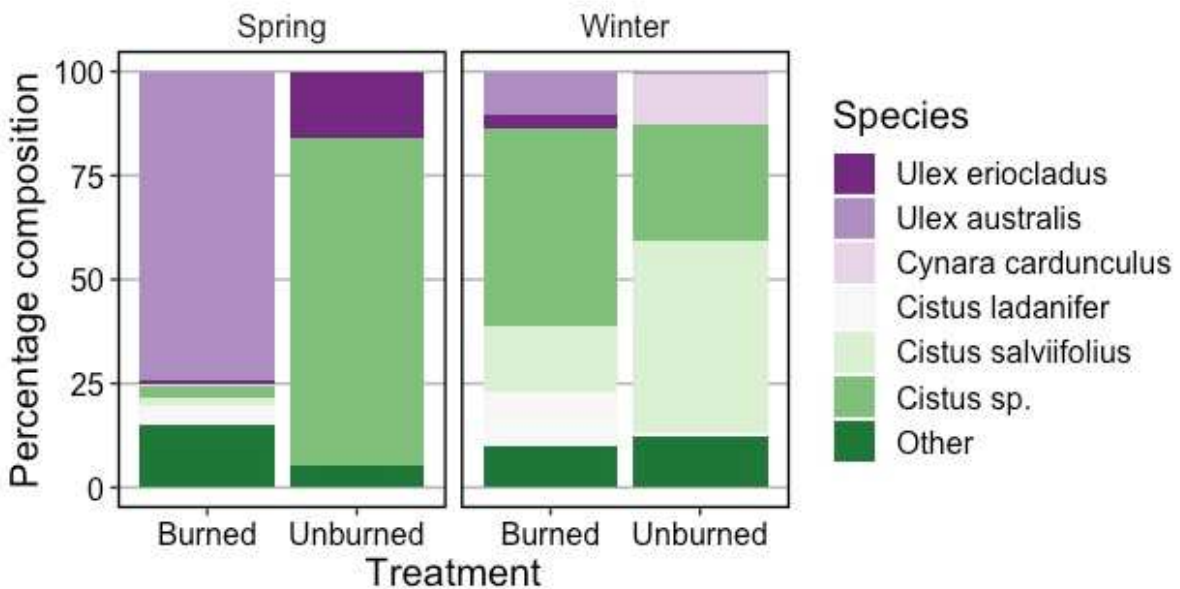


Figure d. Assemblage composition by families of pollen carried by *O. funesta* caught in burned and unburned plots of Serra-do-Caldeirão, Portugal, and across seasons. Families never comprising >7% of individuals in any combination of season and treatment are grouped as “Other”, and all other families are shown independently.