

Supplementary Materials

Local and Landscape Effects on Carrion-Associated Rove Beetle (Coleoptera: Staphylinidae) Communities in German Forests

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The supplementary material is structured in sections Materials and methods and Results.

Materials and methods

Table S1: Environmental variables included in the analyses undertaken on the BExIS platform (Biodiversity Exploratories Information System, <https://www.bexitis.uni-jena.de>).

Variable	Variable type	Categories or description of variable	unit	Data source information
region	categorial	Schwäbische Alb (ALB), Hainich-Dün (HAI), Schorfheide-Chorin (SCH)	-	ID: 20826, version: 1.14.10, owner: Nieschulze, Schulze, Fischer, Ayasse, Weisser, Ostrowski, König-Ries
SMI index	continuous	silvicultural management intensity (SMI) index (from 0 to 1) 0 = undisturbed 1 = disturbed	-	ID: 17746, version: 1.2.2; owner: Schall & Ammer
mineral soil pH	continuous	pH-measurement in the mineral forest soil (0 - 10 cm)	-	ID: 19067, version: 4.1.2; owner: Schöning, Klötzing, Schäfer, Gan, Schrumpf, Trumbore
soil temperature	continuous	forest soil temperature at 10 centimeters below surface	°C	ID: 19007, version: 1.0.5; owner: Wöllauer, Hänsel, Nauss, Forteva
silt content	continuous	amount of medium silt (particle size 0.0063 - 0.02 mm) in forest soil samples	g/kg soil	ID: 14686, version: 1.9.6; owner: Schöning, Solly, Klötzing, Trumbore, Schrumpf
understory proportion	continuous	forest understory proportion up to 2 m above the ground	-	ID: 17066, version: 1.1.3; owner: Schall & Ammer
Shannon plants	continuous	Shannon diversity index for all vascular plants in forests (based on vegetation relevés on all 400 m ² forest sites by visual estimation)	-	ID: 16806, version: 1.2.2; owner: Grassein & Fischer
litter cover	continuous	forest litter cover in summer	%	ID: 6240, version: 1.6.16; owner: Boch, Socher, Mueller, Prati, Fischer
deadwood cover	continuous	forest deadwood cover	%	ID: 6240, version: 1.6.16; owner: Boch, Socher, Mueller, Prati, Fischer

Results

Table S2: List of species of all collected rove beetles on decomposing piglet cadavers at 65 forest study sites in three distinct study regions in descending order by total abundance. Additional information for each species is given for subfamily, size class, ecological niche, and food preference [1–9].

Species name (author, year)	Subfamily	Size class	Ecological niche	Food	Total abundance	Abundance in region 'ALB'	Abundance in region 'HAI'	Abundance in region 'SCH'	Abbreviation for analyzes
<i>Tachinus pallipes</i> (Gravenhorst, 1806)	Tachyporinae	III	de, sap	z	404	182	160	62	T_pal
<i>Bisnius fimetarius</i> (Gravenhorst, 1802)	Staphylininae	III	de, cp, fu	z	338	132	161	45	B_fim
<i>Philonthus addendus</i> (Sharp, 1867)	Staphylininae	IV	de	z	310	12	16	282	P_add
<i>Anotylus mutator</i> (Lohse, 1963)	Oxytelinae	II	de, cp	sa (?)	206	79	127	0	A_mut
<i>Philonthus succicola</i> (Thomson, 1860)	Staphylininae	V	de, cp	z	185	7	4	174	P_suc
<i>Tachinus laticollis</i> (Gravenhorst, 1802)	Tachyporinae	II	de, sap	z	166	133	26	7	T_lat
<i>Philonthus tenuicornis</i> (Mulsant & Rey, 1853)	Staphylininae	IV	de, cp	z	124	6	5	113	P_ten
<i>Ontholestes tessellatus</i> (Geoffroy, 1785)	Staphylininae	V	de, cp, ca	z	123	73	21	29	O_tes
<i>Philonthus splendens</i> (Fabricius, 1793)	Staphylininae	V	de, cp, sap	z	109	2	0	107	P_spl
<i>Philonthus decorus</i> (Gravenhorst, 1802)	Staphylininae	V	de, hyg, mu	z	91	25	25	41	P_dec
<i>Philonthus marginatus</i> (Müller, 1764)	Staphylininae	IV	de, cp	z	88	13	0	75	P_mar
<i>Megarthrus depressus</i> (Paykull, 1789)	Proteininae	I	de, cp, hu, fu	sa, m (?)	64	60	2	2	M_dep

<i>Ontholestes murinus</i> (Linné, 1758)	Staphylininae	V	de, ca	z	61	1	0	60	O_mur
<i>Omalium septentrionis</i> (Thomson, 1857)	Omaliinae	II	de, hu, fu	z (?)	41	34	7	0	O_sep
<i>Ocypus olens</i> (Müller, 1764)	Staphylininae	V	hyg, hu	z	30	0	0	30	O_ole
<i>Atheta monticola</i> (Thomson, 1852)	Aleocharinae	I	de, sap	z (?)	27	14	12	1	A_mon
<i>Philonthus nitidus</i> (Fabricius, 1787)	Staphylininae	V	de, cp	z	20	0	0	20	P_nit
<i>Quedius cinctus</i> (Paykull, 1790)	Staphylininae	IV	de, cp	z	15	13	2	0	Q_cin
<i>Atheta dadopora</i> (Thomson, 1867)	Aleocharinae	I	myc, st, de	z (?)	15	11	1	3	A_dad
<i>Aleochara curtula</i> (Goeze, 1777)	Aleocharinae	II-IV	de, cp	pa, z	14	4	1	9	A_cur
<i>Atheta crassicornis</i> (Fabricius, 1793)	Aleocharinae	II	de, sap	z (?)	14	6	7	1	A_cra
<i>Lordithon lunulatus</i> (Linné, 1760)	Tachyporinae	III	myc, fu	z	14	0	13	1	L_lun
<i>Staphylinus erythropterus</i> (Linné, 1758)	Staphylininae	V	hyg, hu	z	14	0	0	14	S_ery
<i>Atheta castanoptera</i> (Mannerheim, 1830)	Aleocharinae	II	de, myc, fu	z (?)	13	9	4	0	A_cas
<i>Megarthrus stercorarius</i> (Mulsant & Rey, 1878)	Proteininae	I	de, cp, fu	sa, m (?)	11	11	0	0	M_ster
<i>Philonthus varians</i> (Paykull, 1789)	Staphylininae	III	de, cp	z	11	1	1	9	P_var
<i>Atheta fungi</i> (Gravenhorst, 1806)	Aleocharinae	I	de, hu	z	10	10	0	0	A_fun.1

<i>Ocyphus tenebricosus</i> (Gravenhorst, 1846)	Staphylininae	V	hyg, hu	z	10	10	0	0	O_ten
<i>Atheta hansseni</i> (Strand, 1943)	Aleocharinae	I	de, fu	z (?)	10	7	3	0	A_han
<i>Atheta britanniae</i> (Bernhauer & Scheerpeltz, 1926)	Aleocharinae	I	de, myc	z (?)	9	9	0	0	A_bri
<i>Proteinus brachypterus</i> (Fabricius, 1792)	Proteininae	I	de, myc, hu	sa, m (?)	9	9	0	0	P_bra
<i>Atheta sodalis</i> (Erichson, 1837)	Aleocharinae	I	de, myc, fu	z (?)	8	7	1	0	A_sod
<i>Megarthrus nitidulus</i> (Kraatz, 1857)	Proteininae	I	de, cp, fu	sa, m (?)	8	8	0	0	M_nit
<i>Omalium rivulare</i> (Paykull, 1789)	Omaliinae	II	de, sap, fu	sa	8	0	8	0	O_riv
<i>Quedius lateralis</i> (Gravenhorst, 1806)	Staphylininae	V	de, sap, su, fu	z	8	6	1	1	Q_lat
<i>Atheta cinnamoptera</i> (Thomson, 1856)	Aleocharinae	I	de, cp	z (?)	7	4	3	0	A_cin
<i>Quedius lucidulus</i> (Erichson, 1839)	Staphylininae	III	de, hu	z	7	7	0	0	Q_luc
<i>Tachinus proximus</i> (Kraatz, 1855)	Tachyporinae	III/IV	de, cp	z	7	7	0	0	T_pro
<i>Atheta laevana</i> (Mulsant & Rey, 1852)	Aleocharinae	I	de, cp, si	z (?)	6	4	2	0	A_lae
<i>Anotylus sculpturatus</i> (Gravenhorst, 1806)	Oxytelinae	II	de, cp	sa	5	3	2	0	A_scu
<i>Atheta picipes</i> (Thomson, 1856)	Aleocharinae	I	de, fu, co	z (?)	5	4	0	1	A_pic
<i>Pella cognata</i> (Märkel, 1842)	Aleocharinae	III	myr, hu	sa (?)	5	5	0	0	P_cog

<i>Atheta intermedia</i> (Thomson, 1852)	Aleocharinae	II	de, cp	z (?)	4	0	4	0	A_int
<i>Rugilus rufipes</i> (Germar, 1836)	Paederinae	III	de, hyg	z	4	1	1	2	R_ruf
<i>Atheta boreella</i> (Brundin, 1948)	Aleocharinae	I	de, hyg	z (?)	4	2	0	2	A_bor
<i>Omalium rugatum</i> (Mulsant & Rey, 1880)	Omaliinae	II	de	z (?)	3	3	0	0	O_rug
<i>Philonthus politus</i> (Linné, 1758)	Staphylininae	IV/V	de, ca	sa (?)	3	0	0	3	P_pol
<i>Platystethus arenarius</i> (Geoffroy, 1785)	Oxytelinae	II/III	de, cp	z	3	3	0	0	P_are
<i>Quedius mesomelinus</i> <i>skoraszewskyi</i> (Korge, 1960)	Staphylininae	IV	de, ni	z	3	0	3	0	Q_mes
<i>Aleochara stichai</i> (Likovsky, 1965)	Aleocharinae	II	de	pa, z	2	2	0	0	A_sti
<i>Atheta ravilla</i> (Erichson, 1839)	Aleocharinae	I	de, fu, ni	z (?)	2	2	0	0	A_rav
<i>Ocalea picata</i> (Stephens, 1832)	Aleocharinae	II/III	de, hyg, mu	z (?)	2	2	0	0	O_pic
<i>Othius subuliformis</i> (Stephens, 1833)	Staphylininae	III	de, hu	sa, z (?)	2	2	0	0	O_sub
<i>Placusa tachyporoides</i> (Walzl, 1838)	Aleocharinae	I	co	m	2	2	0	0	P_tac
<i>Xantholinus laevigatus</i> (Jacobsen, 1849)	Staphylininae	IV	de, hyg	z	2	2	0	0	X_lae
<i>Bisnius pseudoparcus</i> (Brunne, 1976)	Staphylininae	I	de	z	2	0	0	2	B_pse

<i>Acrotona parvula</i> (Mannerheim, 1830)	Aleocharinae	I	de, cp	z (?)	1	0	0	1	A_par
<i>Aleochara funebris</i> (Wollaston, 1864)	Aleocharinae	II	de	pa, z	1	1	0	0	A_fun
<i>Aleochara intricata</i> (Mannerheim, 1830)	Aleocharinae	II/III	de, sap	pa, z	1	0	0	1	A_intr
<i>Anotylus inustus</i> (Gravenhorst, 1806)	Oxytelinae	II	de, cp, xer	sa, z (?)	1	1	0	0	A_inu
<i>Anthobium atrocephalum</i> (Gyllenhal, 1827)	Omaliinae	II	de, hu, fu	sa, z (?)	1	1	0	0	A_atr
<i>Atheta gagatina</i> (Baudi di Selve, 1848)	Aleocharinae	I	de, myc, fu	z (?)	1	0	0	1	A_gag
<i>Atheta paracrassicornis</i> (Brundin, 1954)	Aleocharinae	I	de, myc	z (?)	1	1	0	0	A_par.1
<i>Autalia longicornis</i> (Scheerpeltz, 1947)	Aleocharinae	I	de, myc	z (?)	1	1	0	0	A_lon
<i>Geostiba circellaris</i> (Gravenhorst, 1806)	Aleocharinae	I	hyg, hu	z (?)	1	1	0	0	G_cir
<i>Lathrobium brunnipes</i> (Fabricius, 1793)	Paederinae	IV	de, hyg	z	1	0	0	1	L_bru
<i>Megarthrus denticollis</i> (Beck, 1817)	Proteininae	I	de, cp, hu	sa, m (?)	1	1	0	0	M_den
<i>Megarthrus prosseni</i> (Schatzmayr, 1904)	Proteininae	I	de, cp, fu	sa, m (?)	1	0	0	1	M_pro
<i>Ocyphus brunnipes</i> (Fabricius, 1781)	Staphylininae	V	de, hu, hyg	z	1	0	0	1	O_bru
<i>Oxypoda brevicornis</i> (Stephens, 1832)	Aleocharinae	I/II	z (?)	z	1	1	0	0	O_bre
<i>Oxytelus laqueatus</i> (Marsham, 1802)	Oxytelinae	II/III	de, cp	sa	1	0	1	0	O_laq

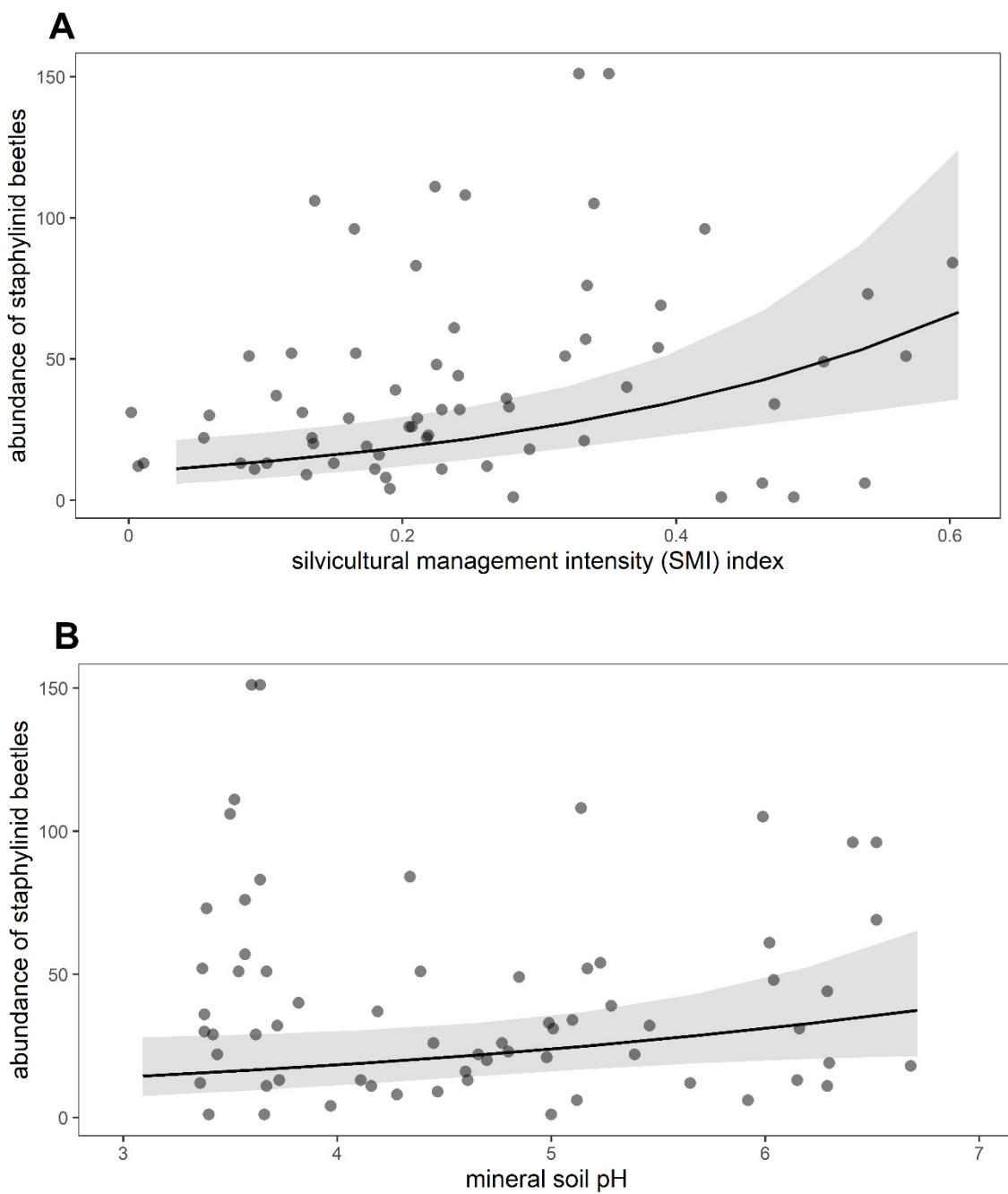
<i>Pella lugens</i> (Gravenhorst, 1802)	Aleocharinae	II	myr	sa	1	1	0	0	P_lug
<i>Philonthus laminatus</i> (Creutzer, 1799)	Staphylininae	IV	de, sap	z	1	0	0	1	P_lam
<i>Quedius fuliginosus</i> (Gravenhorst, 1802)	Staphylininae	V	de, hu, hyg	z	1	0	0	1	Q_ful
<i>Rugilus mixtus</i> (Lohse, 1956)	Paederinae	III	de	z	1	1	0	0	R_mix
<i>Tachinus rufipes</i> (Linné, 1758)	Tachyporinae	III	de, sap	z	1	1	0	0	T_ruf
<i>Bolitochara tecta</i> (Assing, 2014)	Aleocharinae	III	myc, si, pp, bo, xde	z	1	0	1	0	B_tec
<i>Gyrohypnus punctulatus</i> (Paykull, 1789)	Staphylininae	III/IV	de	z	1	0	1	0	G_pun
<i>Pella humeralis</i> (Gravenhorst, 1806)	Aleocharinae	III	myr, hu	sa, z (?)	1	1	0	0	P_hum
<i>Platarea brunnea</i> (Fabricius, 1798)	Aleocharinae	II	de, hu, xer	z	1	0	1	0	P_bru

ALB = Schwäbische Alb, HAI = Hainich-Dün, SCH = Schorfheide-Chorin

Key for 'size class': I= 1 - 3 mm, II= 3 - 4.5 mm, III= 4.5 - 7 mm, IV= 7 - 11 mm, V= > 11 mm

Key for 'ecological niche' (and explanation [lives on or in]): bo= boleticolous (pored fungi), ca= cadavericolous (carrión), cp= coprophilic (feces), co= corticolous (bark), de= detriticolous (detritus), fu= fungicolous (fungi), hu= humicolous (humus), hyg= hygrophilic (humidity), mu= muscicolous (moss), myc= mycetophilic (mushrooms), myr= myrmecophilic (ants), ni= nidicolous (bird-nest), pp= polyporicolous (sponges), sap= saprophilic (decomposing material), si= silvicolous (forests), st= stercolicolous (dung), su= succicolous (plant sap), xer= xerophilic (drought), xde= xylodetriticolous (woody detritus)

Key for 'food' (and explanation [feeds on]): z= zoophagous (animal material), s= saprophagous (decomposing material), p= parasitoid (lives as parasitoid), (?)= unpublished information (G.D.) based on literature [1-9] of near-related species.



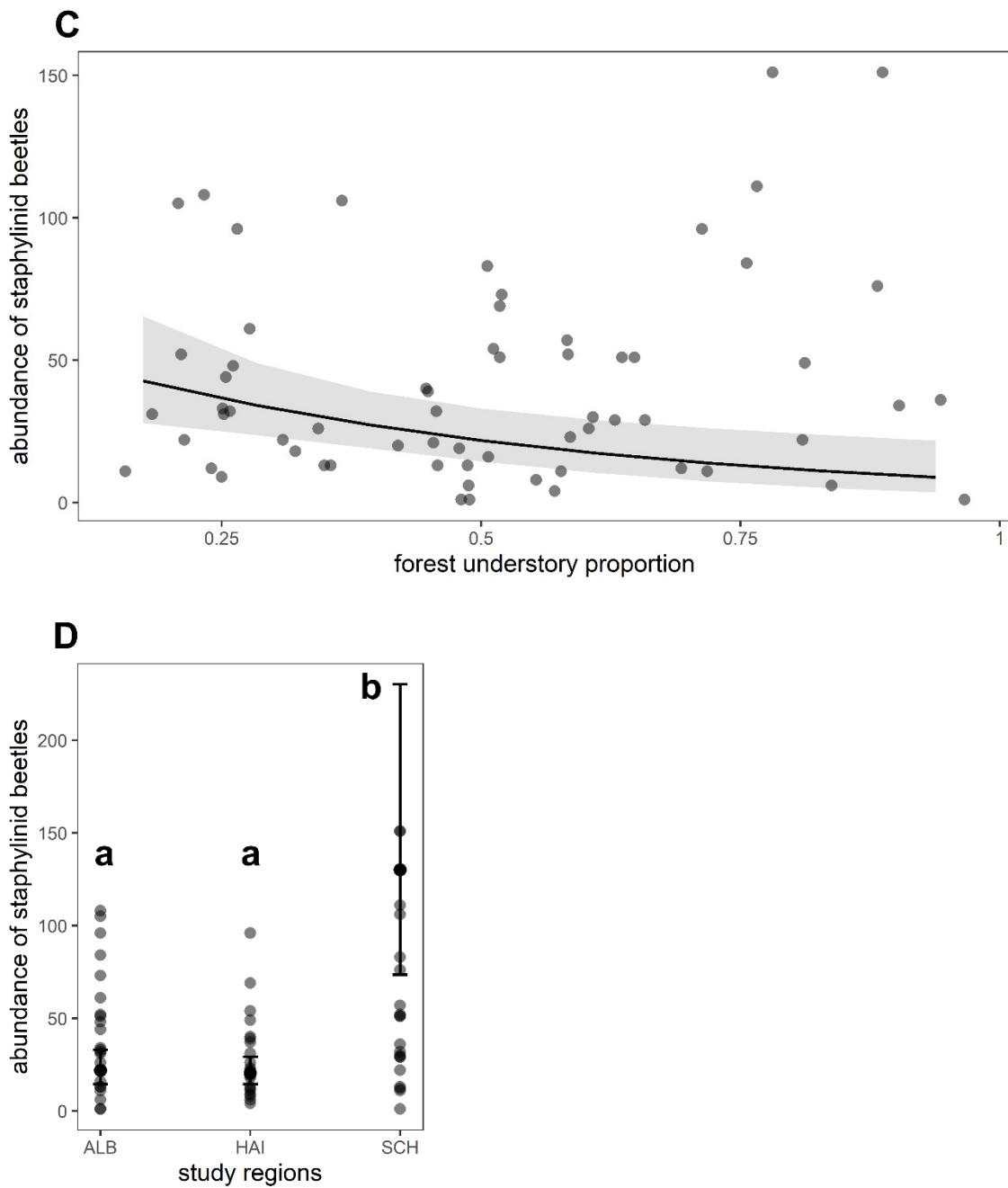
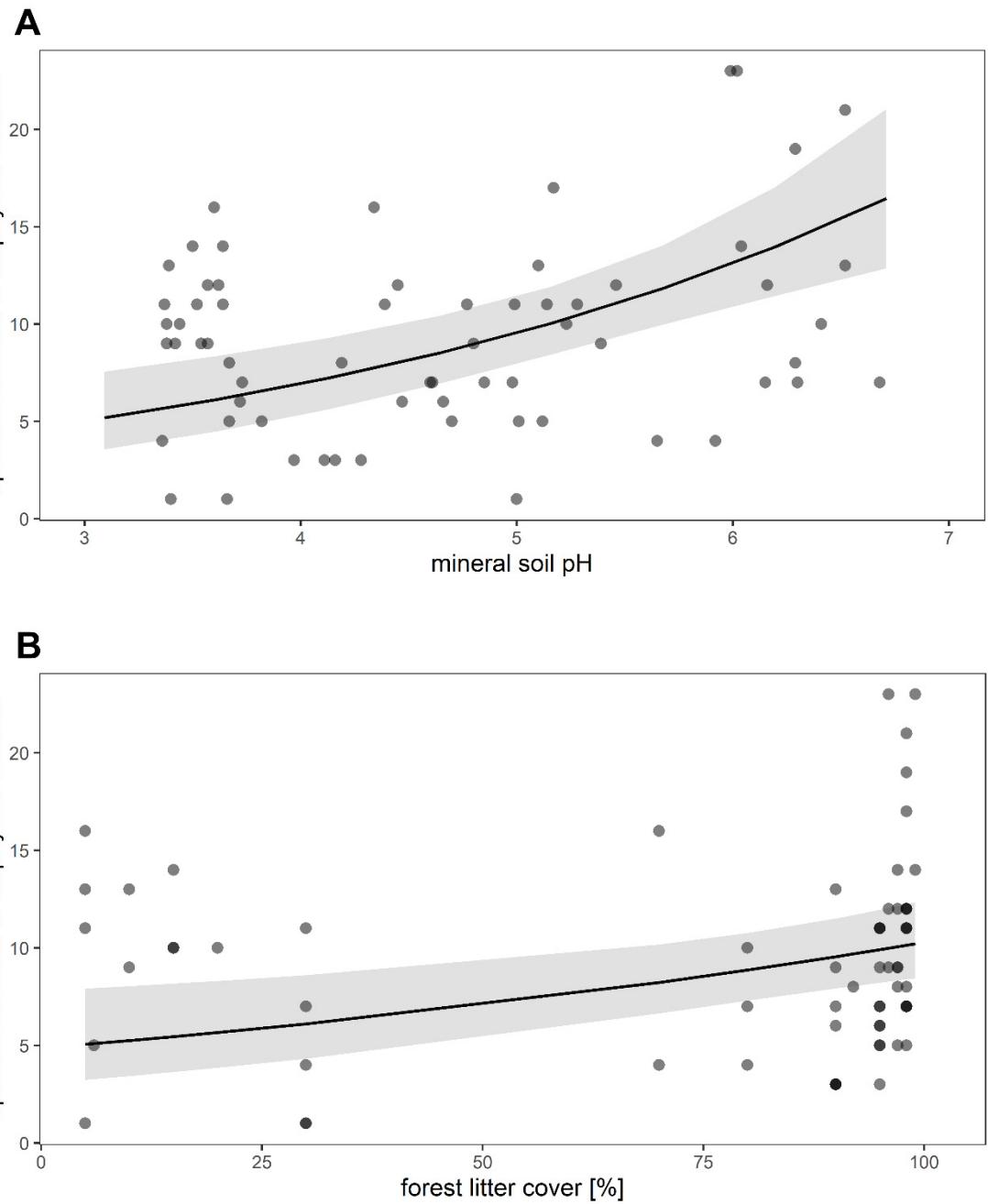


Figure S1 A – D: Marginal effects of multiple regression between total abundance of rove beetles on piglet cadavers at 65 forest sites and A) forest management intensity, B) mineral soil pH, C) forest understory proportion, and D) region. For A - C) regression lines and 95% confidence intervals for negative binomial GLMs are shown. For D) box plots show the median and interquartile range for negative binomial GLMs. Different letters indicate significant differences among groups with estimated marginal means post-hoc tests ($p < 0.05$). ALB = Schwäbische Alb, HAI = Hainich-Dün, SCH = Schorfheide-Chorin.



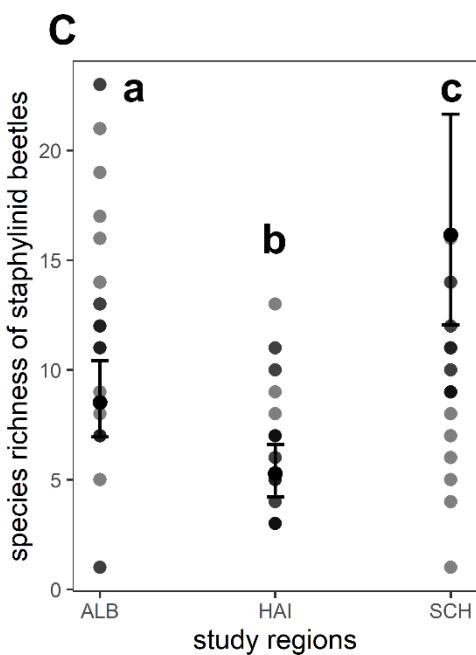
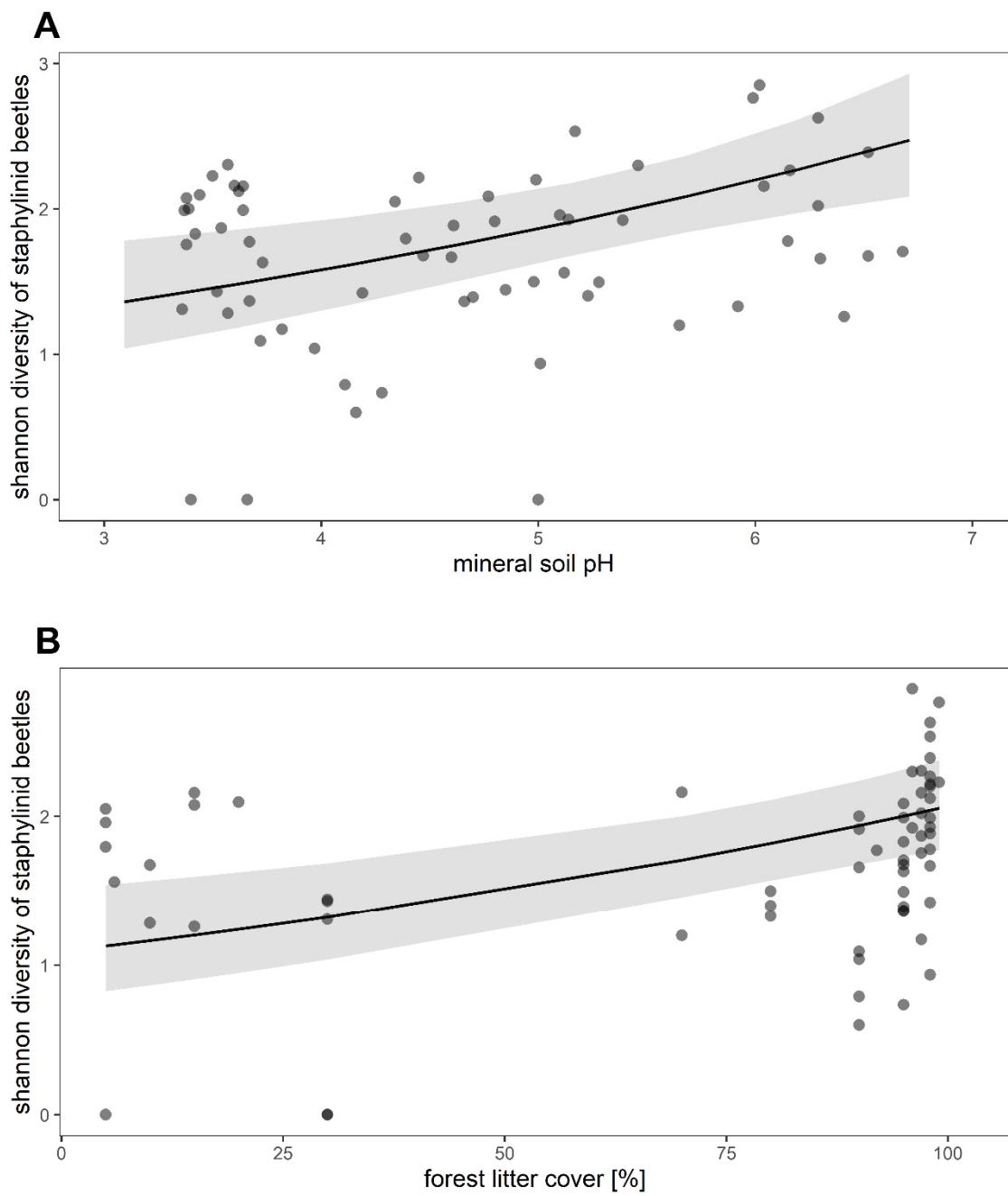


Figure S2 A – C: Marginal effects of multiple regression between species richness of rove beetles on piglet cadavers at 65 forest sites and A) mineral soil pH, B) forest litter cover, and C) region. For A - B) regression lines and 95% confidence intervals for negative binomial GLMs are shown. For C) box plots show the median and interquartile range for negative binomial GLMs. Different letters indicate significant differences among groups with estimated marginal means post-hoc tests ($p < 0.05$). ALB = Schwäbische Alb, HAI = Hainich-Dün, SCH = Schorfheide-Chorin.



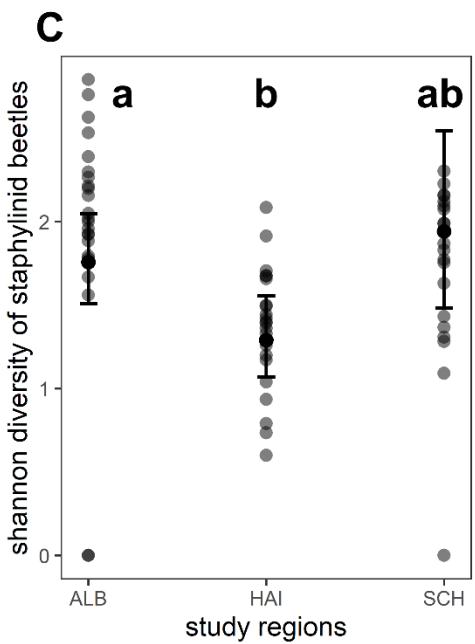


Figure S3 A – C: Marginal effects of multiple regression between Shannon diversity of rove beetles on piglet cadavers at 65 forest sites and A) mineral soil pH, B) forest litter cover, and C) region. For A - B) regression lines and 95% confidence intervals for Gaussian GLMs are shown. For C) box plots show the median and interquartile range for Gaussian GLMs. Different letters indicate significant differences among groups with estimated marginal means post-hoc tests ($p < 0.05$). ALB = Schwäbische Alb, HAI = Hainich-Dün, SCH = Schorfheide-Chorin.

Table S3: Summary of statistical results of the redundancy analysis (RDA). Significant ($p < 0.05$) values are given in bold.

Partitioning of variance			
	Total variance	Proportion	
Total	0.645	1.000	
Constrained	0.245	0.38	
Unconstrained	0.4	0.621	
Canonical axes			
	Axis 1	Axis 2	Axis 3
Eigenvalue	0.161	0.06	0.017
Percentage explained	25.01	9.33	2.66
Cumulative percentage variance			
for species	25.01	34.33	36.99
for species-environment relation	65.9	90.5	97.5
ANOVA (forward tests for axes)			
Variance	0.161	0.06	0.017
F value	24.178	9.017	2.571
p value	0.001	0.001	0.006
Correlations			
region HAI	0.536	0.571	-0.613
region SCH	-0.984	0.16	0.061
SMI	0.003	-0.74	-0.584
Coefficients			
region HAI	0.011	0.174	-0.248
region SCH	-0.291	0.027	-0.184
SMI	-0.019	-0.083	-0.097

p = significance level, SMI = silvicultural management intensity index,
ALB = Schwäbische Alb, HAI = Hainich-Dün, SCH = Schorfheide-Chorin.

Table S4: Results of pairwise ANOSIM analyses showing the effects of region and SMI on the community composition of rove beetles on piglet cadavers at 65 forest study sites. Significant ($p < 0.05$) p -values are given in bold.

Test	<i>R statistic</i>	<i>p</i>	Test	<i>R statistic</i>	<i>p</i>
Global for 'region'	0.529	0.001	Global for 'SMI'	0.177	0.001
ALB vs. HAI	0.176	0.002	MEDIUM vs. HIGH	0.164	0.04
ALB vs. SCH	0.687	0.001	MEDIUM vs. LOW	0.142	0.001
HAI vs. SCH	0.752	0.001	HIGH vs. LOW	0.3	0.003

ALB = Schwäbische Alb, HAI = Hainich-Dün, SCH = Schorfheide-Chorin, SMI = silvicultural management intensity index.

Table S5: Results of SIMPER analysis showing those species that contributed most to the similarity of the rove beetle community composition within regions: average abundance (%) of characteristic species in each region (ALB, HAI, SCH), their contribution to the within-group similarity (%), and the cumulative total (%) of contributions (90% cut-off).

Region ALB			
Average similarity: 28.59 %			
Species	Abundance	Contribution	Cumulative
<i>T. pallipes</i>	7.91	22.8	22.8
<i>B. fimetarius</i>	5.74	18.02	40.82
<i>T. laticollis</i>	5.78	16.47	57.29
<i>O. tessellatus</i>	3.17	10.66	67.96
<i>A. mutator</i>	3.43	9.43	77.38
<i>O. septentrionis</i>	1.48	6.81	84.19
<i>M. depressus</i>	2.61	4.22	88.41
<i>P. decorus</i>	1.09	2.89	91.31

Region HAI			
Average similarity: 29.72 %			
Species	Abundance	Contribution	Cumulative
<i>B. fimetarius</i>	7.32	39.39	39.39
<i>A. mutator</i>	5.77	36.6	75.98
<i>T. pallipes</i>	7.27	7.91	83.89
<i>P. decorus</i>	1.14	6.47	90.37

Region SCH			
Average similarity: 33.06 %			
Species	Abundance	Contribution	Cumulative
<i>P. addendus</i>	14.1	33.53	33.53
<i>P. succicola</i>	8.7	26.62	60.16
<i>P. splendens</i>	5.35	11.08	71.24
<i>P. tenuicornis</i>	5.65	7.94	79.18
<i>O. olens</i>	1.5	3.65	82.83
<i>O. tessellatus</i>	1.45	3.39	86.22
<i>P. marginatus</i>	3.75	2.73	88.95
<i>O. murinus</i>	3	2.69	91.63

ALB = Schwäbische Alb, HAI = Hainich-Dün, SCH = Schorfheide-Chorin.

Table S6: Results of SIMPER analysis showing those species that contributed most to the dissimilarity of the rove beetle community composition between regions: average abundance (%) of characteristic species for each region (ALB, HAI, SCH), their contribution to the between-group dissimilarity (%), and the cumulative total (%) of contributions (90% cut-off).

Groups ALB & HAI

Average dissimilarity = 76.39 %

Species	Group ALB	Group HAI	Contribution	Cumulative
	Abundance	Abundance		
<i>T. pallipes</i>	7.91	7.27	17.78	17.78
<i>A. mutator</i>	3.43	5.77	14.33	32.11
<i>B. fimetarius</i>	5.74	7.32	14.29	46.4
<i>T. laticollis</i>	5.78	1.18	8.87	55.28
<i>O. tessellatus</i>	3.17	0.95	6.03	61.31
<i>M. depressus</i>	2.61	0.09	4.31	65.61
<i>P. decorus</i>	1.09	1.14	4.15	69.76
<i>O. septentrionis</i>	1.48	0.32	3.52	73.29
<i>P. addendus</i>	0.52	0.73	2.28	75.57
<i>A. monticola</i>	0.61	0.55	1.9	77.47
<i>A. crassicornis</i>	0.26	0.32	1.17	78.64
<i>A. castanoptera</i>	0.39	0.18	1.04	79.67
<i>O. tenebricosus</i>	0.43	0	1.01	80.69
<i>P. tenuicornis</i>	0.26	0.23	1	81.68
<i>O. rivulare</i>	0	0.36	0.93	82.62
<i>A. dadopora</i>	0.48	0.05	0.91	83.53
<i>A. fungi</i>	0.43	0	0.9	84.42
<i>P. succicola</i>	0.3	0.18	0.85	85.27
<i>Q. cinctus</i>	0.57	0.09	0.84	86.11
<i>L. lunulatus</i>	0	0.59	0.81	86.92
<i>P. marginatus</i>	0.57	0	0.81	87.73
<i>A. sodalis</i>	0.3	0.05	0.74	88.46
<i>A. britanniae</i>	0.39	0	0.7	89.16
<i>A. hansseni</i>	0.3	0.14	0.67	89.83
<i>P. brachypterus</i>	0.39	0	0.66	90.49

Groups ALB & SCH

Average dissimilarity = 90.42 %

Species	Group ALB	Group SCH	Contribution	Cumulative
	Abundance	Abundance		
<i>P. addendus</i>	0.52	14.1	15.61	15.61
<i>P. succicola</i>	0.3	8.7	10.67	26.28
<i>T. pallipes</i>	7.91	3.1	9.2	35.49
<i>B. fimetarius</i>	5.74	2.25	6.85	42.33
<i>T. laticollis</i>	5.78	0.35	6.16	48.49
<i>P. splendens</i>	0.09	5.35	6.12	54.61

<i>P. tenuicornis</i>	0.26	5.65	5.58	60.19
<i>A. mutator</i>	3.43	0	4.21	64.4
<i>O. tessellatus</i>	3.17	1.45	3.46	67.86
<i>P. marginatus</i>	0.57	3.75	3.41	71.27
<i>P. decorus</i>	1.09	2.05	3.23	74.5
<i>M. depressus</i>	2.61	0.1	2.92	77.42
<i>O. murinus</i>	0.04	3	2.83	80.25
<i>O. olens</i>	0	1.5	2.39	82.63
<i>O. septentrionis</i>	1.48	0	2.34	84.98
<i>P. nitidus</i>	0	1	1.23	86.21
<i>S. erythropterus</i>	0	0.7	1.15	87.36
<i>A. monticola</i>	0.61	0.05	0.91	88.27
<i>A. dadopora</i>	0.48	0.15	0.69	88.96
<i>O. tenebricosus</i>	0.43	0	0.66	89.63
<i>A. fungi</i>	0.43	0	0.6	90.23

Groups HAI & SCH

Average dissimilarity = 92.11 %

Species	Group HAI	Group SCH	Contribution	Cumulative
	Abundance	Abundance		
<i>P. addendus</i>	0.73	14.1	16.8	16.8
<i>P. succicola</i>	0.18	8.7	11.59	28.39
<i>A. mutator</i>	5.77	0	10.67	39.06
<i>B. fimetarius</i>	7.32	2.25	10.66	49.72
<i>T. pallipes</i>	7.27	3.1	9.02	58.74
<i>P. splendens</i>	0	5.35	6.65	65.39
<i>P. tenuicornis</i>	0.23	5.65	5.93	71.32
<i>P. decorus</i>	1.14	2.05	3.98	75.3
<i>P. marginatus</i>	0	3.75	3.39	78.7
<i>O. murinus</i>	0	3	3.03	81.73
<i>O. tessellatus</i>	0.95	1.45	2.78	84.51
<i>O. olens</i>	0	1.5	2.62	87.13
<i>T. laticollis</i>	1.18	0.35	1.7	88.83
<i>P. nitidus</i>	0	1	1.34	90.17

ALB = Schwäbische Alb, HAI = Hainich-Dün, SCH = Schorfheide-Chorin.

Table S7: Results of SIMPER analysis showing those species that contributed most to the similarity of the rove beetle community composition within silvicultural management intensity (SMI) levels: average abundance (%) of characteristic species at each SMI level (LOW, MEDIUM, HIGH), their contribution to the within-group similarity (%), and the cumulative total (%) of contributions (90% cut-off).

SMI level 'LOW'

Average similarity: 24.59 %

Species	Abundance	Contribution	Cumulative
<i>A. mutator</i>	6.08	40.69	40.69
<i>B. fimetarius</i>	3.92	23.7	64.39
<i>P. decorus</i>	1.5	8.39	72.78
<i>T. pallipes</i>	1.73	4.47	77.25
<i>P. addendus</i>	2.15	4.46	81.71
<i>A. monticola</i>	0.46	3.11	84.82
<i>O. tessellatus</i>	0.92	2.55	87.37
<i>P. succicola</i>	2	2.52	89.89
<i>O. septentrionis</i>	0.69	2.5	92.39

SMI level 'MEDIUM'

Average similarity: 20.90 %

Species	Abundance	Contribution	Cumulative
<i>T. pallipes</i>	6.9	17.93	17.93
<i>B. fimetarius</i>	5.72	16.62	34.55
<i>P. addendus</i>	8.48	15.81	50.36
<i>P. succicola</i>	4.48	10.49	60.85
<i>T. laticollis</i>	2.72	7.51	68.36
<i>O. tessellatus</i>	2.66	5.52	73.88
<i>A. mutator</i>	1.48	4.28	78.16
<i>P. tenuicornis</i>	2.76	3.85	82.01
<i>P. decorus</i>	1.66	3.83	85.84
<i>P. marginatus</i>	2.69	2.23	88.07
<i>M. depressus</i>	1.45	2.13	90.2

SMI level 'HIGH'

Average similarity: 25.20 %

Species	Abundance	Contribution	Cumulative
<i>T. pallipes</i>	15.9	37.81	37.81
<i>B. fimetarius</i>	7	23.34	61.15
<i>O. tessellatus</i>	2.2	19.7	80.85
<i>T. laticollis</i>	6.2	11.75	92.6

Table S8: Results of SIMPER analysis showing those species that contributed most to the dissimilarity of the rove beetle community composition between silvicultural management intensity (SMI) levels: average abundance (%) of characteristic species for each SMI level (LOW, MEDIUM, HIGH), their contribution to the between-group dissimilarity (%), and the cumulative total (%) of contributions (90% cut-off).

Groups MEDIUM & HIGH

Average dissimilarity = 81.82 %

Species	Group MEDIUM	Group HIGH	Contribution	Cumulative
	Abundance	Abundance		
<i>T. pallipes</i>	6.9	15.9	19.72	19.72
<i>B. fimetarius</i>	5.72	7	11.62	31.34
<i>P. addendus</i>	8.48	0.8	11.08	42.42
<i>T. laticollis</i>	2.72	6.2	8.07	50.5
<i>P. succicola</i>	4.48	0.3	6.55	57.05
<i>O. tessellatus</i>	2.66	2.2	4.85	61.89
<i>P. tenuicornis</i>	2.76	0.1	2.98	64.87
<i>P. decorus</i>	1.66	0.4	2.82	67.69
<i>M. depressus</i>	1.45	0.7	2.81	70.5
<i>A. mutator</i>	1.48	0.5	2.8	73.29
<i>P. marginatus</i>	2.69	0	2.68	75.98
<i>P. splendens</i>	1.86	0	1.95	77.92
<i>O. murinus</i>	1.79	0	1.86	79.78
<i>O. septentrionis</i>	0.72	0.2	1.59	81.38
<i>O. olens</i>	0.76	0	1.39	82.77
<i>A. monticola</i>	0.31	0.6	1.31	84.08
<i>S. erythropterus</i>	0.48	0	1.02	85.1
<i>A. crassicornis</i>	0.28	0.2	0.99	86.08
<i>O. tenebricosus</i>	0.14	0.5	0.98	87.06
<i>L. lunulatus</i>	0.07	0.9	0.91	87.97
<i>Q. cinctus</i>	0.28	0.6	0.84	88.81
<i>A. intermedia</i>	0.07	0.2	0.78	89.59
<i>A. dadopora</i>	0.38	0.2	0.75	90.34

Groups MEDIUM & LOW

Average dissimilarity = 81.71 %

Species	Group MEDIUM	Group LOW	Contribution	Cumulative
	Abundance	Abundance		
<i>P. addendus</i>	8.48	2.15	12.24	12.24
<i>A. mutator</i>	1.48	6.08	10.79	23.03
<i>B. fimetarius</i>	5.72	3.92	10.31	33.34
<i>T. pallipes</i>	6.9	1.73	9.76	43.09
<i>P. succicola</i>	4.48	2	7.75	50.85
<i>T. laticollis</i>	2.72	0.96	4.48	55.33
<i>P. tenuicornis</i>	2.76	1.65	4.48	59.81

<i>P. splendens</i>	1.86	2.12	4.22	64.03
<i>P. decorus</i>	1.66	1.5	4.11	68.14
<i>O. tessellatus</i>	2.66	0.92	3.86	72
<i>P. marginatus</i>	2.69	0.38	2.99	74.99
<i>M. depressus</i>	1.45	0.58	2.83	77.82
<i>O. murinus</i>	1.79	0.35	2.22	80.04
<i>O. septentrionis</i>	0.72	0.69	2.19	82.23
<i>O. olens</i>	0.76	0.31	1.76	83.99
<i>A. monticola</i>	0.31	0.46	1.32	85.32
<i>S. erythropterus</i>	0.48	0	0.97	86.29
<i>P. nitidus</i>	0.41	0.31	0.88	87.17
<i>A. dadopora</i>	0.38	0.08	0.7	87.87
<i>A. castanoptera</i>	0.28	0.12	0.65	88.52
<i>A. fungi</i>	0.28	0.08	0.64	89.16
<i>A. crassicornis</i>	0.28	0.15	0.61	89.77
<i>O. rivulare</i>	0.03	0.27	0.59	90.35

Groups HIGH & LOW

Average dissimilarity = 86.41 %

Species	Group HIGH	Group LOW	Contribution	Cumulative
	Abundance	Abundance		
<i>T. pallipes</i>	15.9	1.73	20.77	20.77
<i>A. mutator</i>	0.5	6.08	14.91	35.68
<i>B. fimetarius</i>	7	3.92	12.23	47.91
<i>T. laticollis</i>	6.2	0.96	8.39	56.3
<i>O. tessellatus</i>	2.2	0.92	4.82	61.12
<i>P. addendus</i>	0.8	2.15	4.45	65.57
<i>P. decorus</i>	0.4	1.5	4.03	69.61
<i>P. succicola</i>	0.3	2	3.5	73.1
<i>P. splendens</i>	0	2.12	3.42	76.52
<i>P. tenuicornis</i>	0.1	1.65	2.53	79.04
<i>O. septentrionis</i>	0.2	0.69	1.86	80.91
<i>A. monticola</i>	0.6	0.46	1.58	82.49
<i>M. depressus</i>	0.7	0.58	1.38	83.87
<i>A. crassicornis</i>	0.2	0.15	1.13	85
<i>O. tenebricosus</i>	0.5	0.04	1.12	86.12
<i>L. lunulatus</i>	0.9	0.12	1.05	87.17
<i>A. intermedia</i>	0.2	0	0.93	88.1
<i>O. rivulare</i>	0	0.27	0.76	88.86
<i>Q. cinctus</i>	0.6	0.04	0.75	89.61
<i>O. olens</i>	0	0.31	0.65	90.26

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