



Digital supplementary material to

DUNN, J.D., HODAPP, D., MENZEL, F. & KOHLMEIER, P. 2024: Larval chemical cues induce rapid changes in foraging preferences of ant workers (Hymenoptera: Formicidae). – Myrmecological News 34: 71-79.

The content of this digital supplementary material was subject to the same scientific editorial processing as the article it accompanies. However, the authors are responsible for copyediting and layout.

Data S1: Per capita honey and fly foraging activity in subcolonies with and without brood. (separate .txt file)

Data S2: Number of observations foragers and non-foragers were found on brood. These data were originally produced in Kohlmeier & al. (2018). (separate .txt file)

Data S3: Per capita honey and fly foraging activity in colonies exposed to larval CHCs or hexane. (separate .txt file)

Data S4: Raw data of GC-MS analysis of larva and worker CHCs. (separate .txt file)

Script S1: R script for the statistical analysis. (separate .R file)

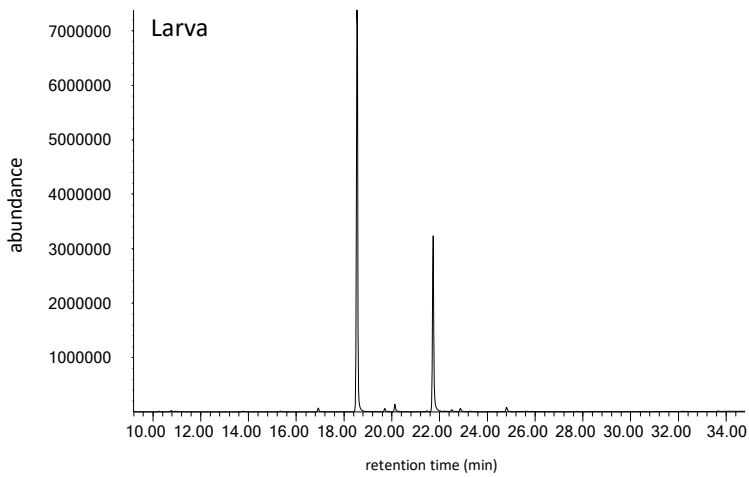
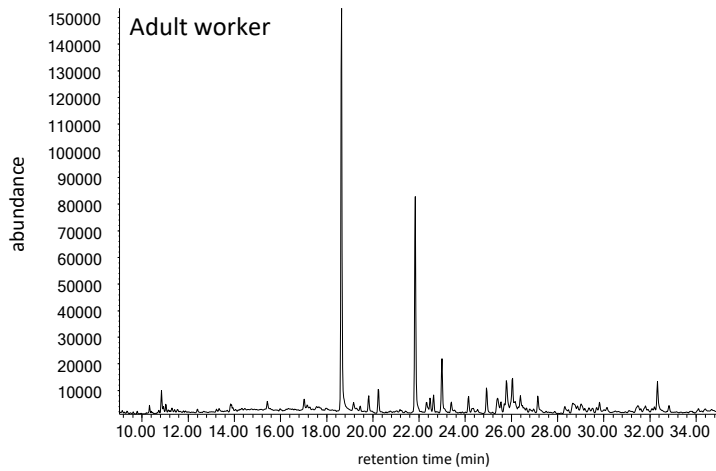


Fig. S1 Representative GC-MS data from CHC extracts. The figure shows extracts of a single adult worker (above) and a single larva (below).

Table S1 List of the compounds found in *Temnothorax longispinosus* adults and la

	Retention time	Retention Index	Adults	Larvae	diagnostic ions
<i>n</i> - C25	15.4	2500	0.79 ± 0.1	0.05 ± 0.02	---
<i>n</i> - C26	17	2600	0.85 ± 0.08	1.21 ± 0.22	---
<i>n</i> - C27	18.6	2700	29.41 ± 1.27	63.02 ± 1.16	---
11-;13-MeC27	19.14	2733	0.84 ± 0.1	0.18 ± 0.03	168 196 224 252
7-MeC27	19.26	2740	0.07 ± 0.02	0 ± 0	112 309
5-MeC27	19.42	2750	0.21 ± 0.03	0.11 ± 0.02	85 337
11,15-DiMeC27	19.6	2762	0.19 ± 0.05	0.03 ± 0.01	168 196 238 267
3-MeC27	19.8	2775	1.75 ± 0.33	1.22 ± 0.17	57 365
<i>n</i> - C28	20.22	2800	2.71 ± 0.15	1.58 ± 0.16	---
unknown hydrocarbon	20.34	2809	0.04 ± 0.02	0.17 ± 0.04	---
12-;13-MeC28	20.64	2827	0.02 ± 0.01	0.04 ± 0.01	182 196 239 252
unknown hydrocarbon	20.74	2833	0.26 ± 0.03	0.5 ± 0.1	---
unknown hydrocarbon	20.88	2841	0.07 ± 0.01	0.04 ± 0.01	---
unknown hydrocarbon	21.02	2850	0.09 ± 0.01	0.06 ± 0.01	---
4-MeC28	21.14	2857	0.34 ± 0.04	0.12 ± 0.03	71 365
3-MeC28	21.22	2862	0.1 ± 0.03	0.12 ± 0.03	57 379
C29-ene	21.38	2872	0.12 ± 0.01	0.05 ± 0.01	---
<i>n</i> - C29	21.8	2900	15.54 ± 0.86	22.98 ± 0.77	---
11-;13-;15-MeC29	22.3	2929	2.39 ± 0.28	0.3 ± 0.04	168 196 224 253 281
7-MeC29	22.46	2940	0.74 ± 0.08	0.16 ± 0.03	112 337
5-MeC29	22.6	2949	0.85 ± 0.09	0.7 ± 0.08	85 365
11,15-;13,17-DiMeC29	22.74	2958	0.97 ± 0.29	0.08 ± 0.02	168 196 224 238 267 294
3-MeC29	22.96	2972	5.14 ± 0.31	1.55 ± 0.14	57 393
5,x-DiMeC29 ¹	23.08	2980	0.07 ± 0.03	0.02 ± 0.01	85 379
<i>n</i> - C30	23.34	3000	1.07 ± 0.09	0.29 ± 0.08	---
3,7-DiMeC29*	23.5	3007	0.09 ± 0.04	0 ± 0	57 407 127 337 168 182 196 210
11-;12-;13-;14-MeC30	23.84	3029	0.41 ± 0.04	0.04 ± 0.01	252 267 281 294
unknown hydrocarbon	23.98	3038	0.09 ± 0.05	0.03 ± 0.01	---
3-MeC30	24.12	3047	0.53 ± 0.1	0.11 ± 0.02	57 407
x,y-DiMeC30 ¹	24.3	3058	0.83 ± 0.05	0.21 ± 0.03	---
unknown hydrocarbon	24.5	3071	0.12 ± 0.02	0.04 ± 0.01	---
<i>n</i> - C31	24.9	3100	1.78 ± 0.13	1.1 ± 0.09	---
9-;11-;13-;15-MeC31	25.36	3128	3.67 ± 0.32	0.36 ± 0.05	140 168 196 224 239 281 308 337
7-MeC31	25.52	3139	0.43 ± 0.07	0.18 ± 0.02	112 365
unknown hydrocarbon	25.66	3148	0 ± 0	0.15 ± 0.04	---
7,11-DiMeC31	25.82	3159	5.33 ± 0.62	0.61 ± 0.11	112 182 309 379

3-MeC31	26	3171	1.57 ± 0.18	0.22 ± 0.03	47 421
11,15,19-TriMeC31	26.12	3179	0.34 ± 0.11	0.13 ± 0.04	168 196 238 267 308 337
7,15,21-TriMeC31	26.3	3192	0.68 ± 0.19	0.19 ± 0.06	112 168 239 267 337 392
3,11-DiMeC31 ²	26.46	3202	0.54 ± 0.09	0.1 ± 0.02	57 182 308 453
3,25-DiMeC31 ³	26.62	3213	0.03 ± 0.01	0 ± 0	47 112 379 435
3,x,y-TriMeC31 ¹	26.84	3227	0.58 ± 0.07	0 ± 0	57 449
unknown hydrocarbon	26.98	3237	0.16 ± 0.08	0.08 ± 0.02	---
3-MeC32	27.14	3247	0.92 ± 0.1	0.15 ± 0.03	57 435
unknown hydrocarbon	27.34	3260	0.15 ± 0.05	0.04 ± 0.01	---
unknown hydrocarbon	27.84	3293	0.13 ± 0.02	0.11 ± 0.02	---
					168 196 224
11-;13-;15-;17-MeC33	28.28	3324	1.83 ± 0.21	0.15 ± 0.02	252
					281 309 337
unknown hydrocarbon	28.44	3335	0.05 ± 0.02	0.07 ± 0.01	---
x,y-DiMeC33 ⁴	28.68	3352	4.38 ± 0.41	0.32 ± 0.04	---
5,x-DiMeC33 ¹	28.98	3373	0.88 ± 0.1	0.11 ± 0.03	85 435
unknown hydrocarbon	29.16	3385	0.05 ± 0.02	0.01 ± 0	---
3,9-;3,11-DiMeC33	29.34	3398	0.39 ± 0.04	0 ± 0	57 154 182 337 365 463
unknown hydrocarbon	29.7	3425	0.48 ± 0.06	0.03 ± 0.01	---
unknown hydrocarbon	29.88	3438	0.16 ± 0.07	0.02 ± 0	---
unknown hydrocarbon	30.06	3452	0.43 ± 0.05	0.03 ± 0.01	---
unknown hydrocarbon	30.5	3485	0.08 ± 0.02	0.03 ± 0.01	---
unknown hydrocarbon	30.66	3497	0 ± 0	0.03 ± 0.01	---
11-;13-;15-MeC35	31.06	3526	0.39 ± 0.06	0.07 ± 0.01	168 196 224 309 337 365
11,x-;13,x-;15,x-DiMeC35	31.44	3553	2.1 ± 0.22	0.12 ± 0.02	168 196 351 379
unknown hydrocarbon	31.58	3563	0 ± 0	0.1 ± 0.02	---
unknown hydrocarbon	31.74	3575	0.61 ± 0.09	0.05 ± 0.01	---
unknown hydrocarbon	31.88	3585	0 ± 0	0.01 ± 0	---
unknown hydrocarbon	32.08	3600	0.27 ± 0.04	0.02 ± 0	---
unknown hydrocarbon	32.16	3605	0.01 ± 0.01	0.01 ± 0	---
unknown hydrocarbon	32.3	3615	3.27 ± 0.18	0 ± 0	---
unknown hydrocarbon	32.48	3629	0.1 ± 0.04	0.01 ± 0.01	---
unknown hydrocarbon	32.78	3650	0.24 ± 0.03	0.02 ± 0	---
unknown hydrocarbon	33.06	3671	0.04 ± 0.01	0.04 ± 0.01	---
unknown hydrocarbon	33.34	3691	0 ± 0	0.01 ± 0	---
unknown hydrocarbon	33.7	3718	0.17 ± 0.02	0.16 ± 0.01	---
11,x-;13,x-;15,x-DiMeC37 ¹	34.06	3747	0.74 ± 0.07	0.09 ± 0.01	168 196 224 351 379 407
unknown hydrocarbon	34.34	3769	0.22 ± 0.04	0.03 ± 0.01	---
unknown hydrocarbon	34.68	3796	0.06 ± 0.01	0.03 ± 0.01	---

¹position of methyl groups indicated with "x" or "y" is unknown

²possibly other 3,X-DiMe compounds

³identification tentative

⁴x = 11,13 and 15; y = 17, 19 and 21; combination of first and second methyl group unknown

Table S2: Overview of the tests run to confirm model fit and to check for potential deviations from uniformity

Model	Figure	KS test	Dispersion test	Outlier test	Within-group deviation from uniformity
glm(diff_food~foodsource)	2A	p = 0.63572	p = 0.96	p = 0.26214	ns
glm(sqrt(trips.chc)~treatment.chc*food.chc)	3B+C	p = 0.13176	p = 0.952	p = 1	ns
lmer(nAlk~Stage + (1 Population))	4B	p = 0.52007	p = 0.904	p = 0.41958	ns
lmer(CHCs[,3]~Stage) : n-C27	4B	p = 0.7799	p = 0.96	p = 1	ns
lmer(CHCs[,18]~Stage) : n-C29	4B	p = 0.9666	0.976	p = 1	ns
lmer(sqrt(CHCs[,9])~Stage) : n-C28	None	p = 0.36998	p = 0.888	p = 0.41958	ns
lmer(sqrt(CHCs[,32])~Stage) : n-C31	None	p = 0.89748	p = 0.912	p = 1	ns