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Supplemental Information

Ascaroside Signaling Is Widely

Conserved among Nematodes

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Supplemental Inventory

1. Supplemental Figures and Tables

Figure S1, related to Figure 1

Figure S2, related to Figure 2

Figure S3, related to Figure 3

Table S1

Table S2

2. Supplemental Experimental Procedures

3. Supplemental References

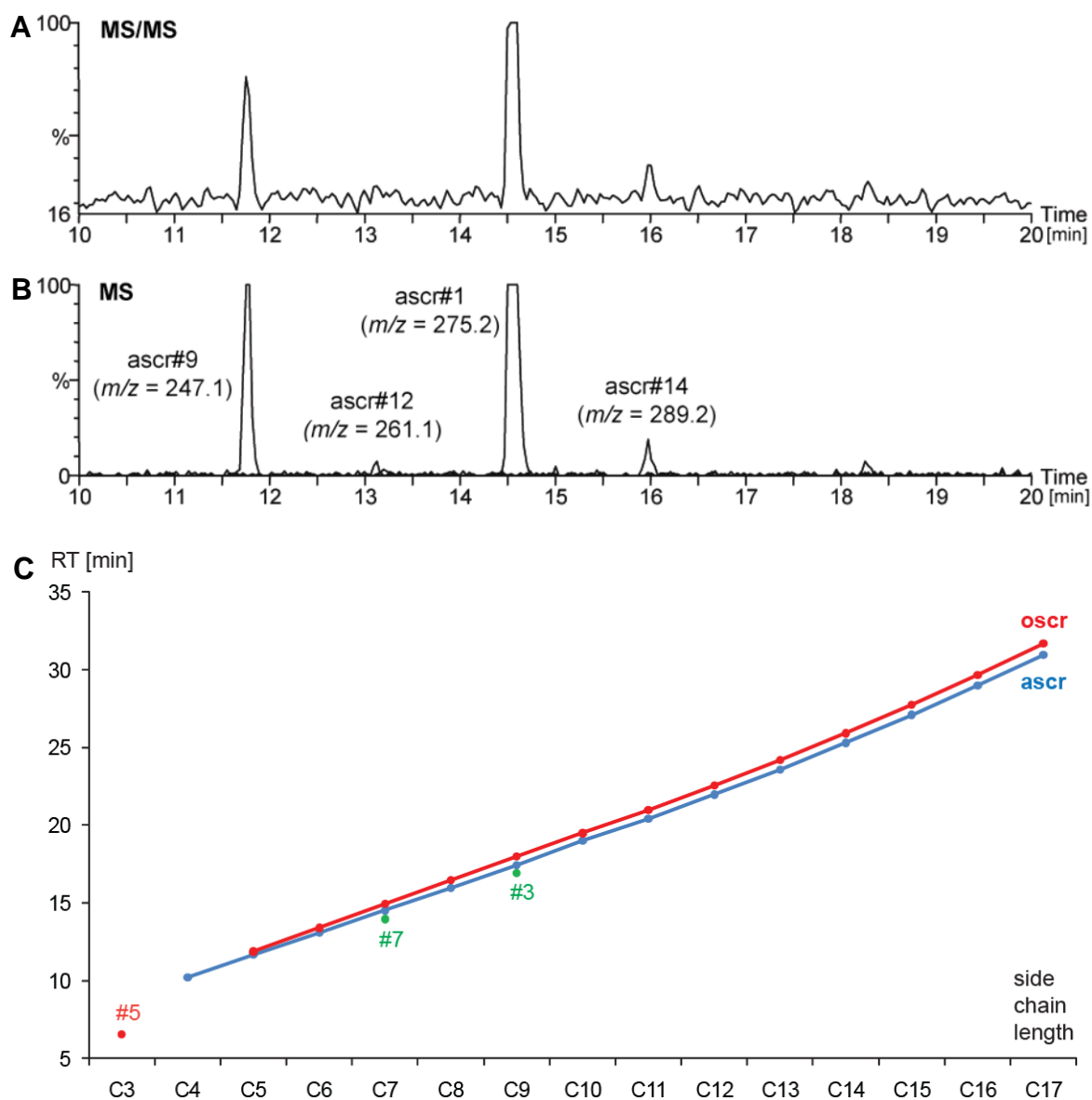


Figure S1. Identification of Ascarosides, Related to Figure 1

(A) Total ion current (TIC) chromatogram from LC-MS/MS screen for precursor ions of m/z 73.0 reveals ascaroside signals in worm water extract of adult *S. glaseri*.

(B) Corresponding ion traces for ascr#9, ascr#12, ascr#1, and ascr#14 from LC-MS measurement.

(C) HPLC retention times of ascarosides (ascr's) and omega-ascarosides (oscr's), side chains range from 3–17 carbons, derived from analysis of wild-type and peroxisomal beta-oxidation mutants of *C. elegans* and synthetic standards. See Figure 1.

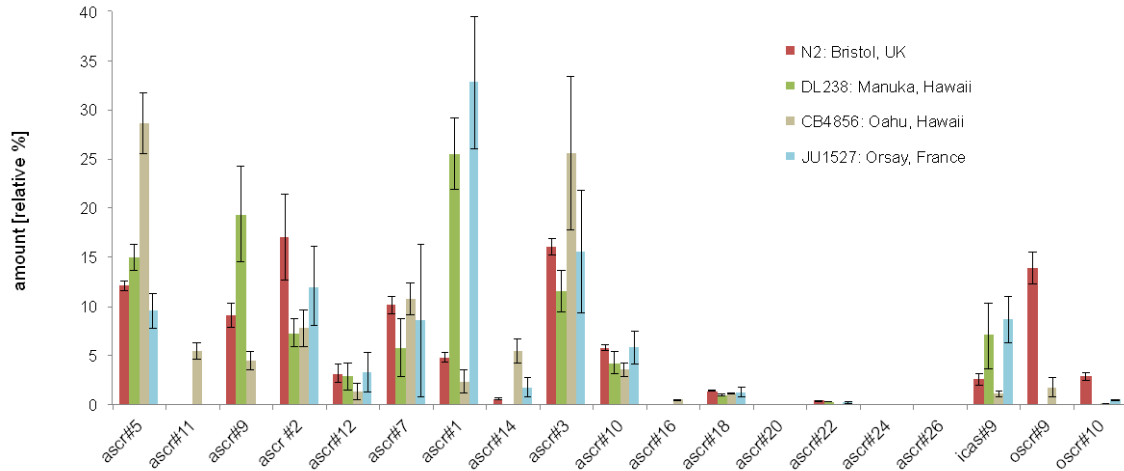


Figure S2. HPLC-MS Analysis of Worm Media Samples Obtained from Several Strains of *C. elegans*, Related to Figure 2

Error bars are S.D. See Figure 2.

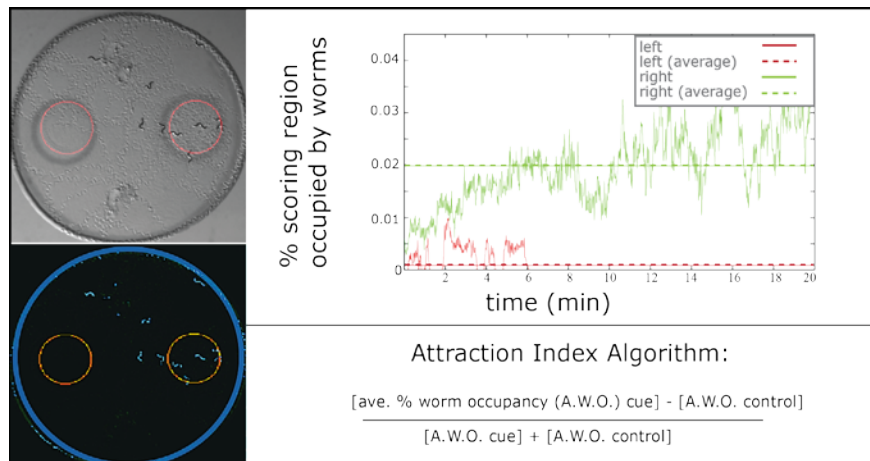


Figure S3. Automated Software Scoring Worm Retention in Areas Conditioned by Ascarosides, Related to Figure 3

Our software detects the presence of worms in both regions and provides an output of the percentage of each scoring region occupied by worms, which is calculated once per second during the entire trial. The output is a plot of worm occupancy ratio vs. time for both scoring regions (right panel). We adopted the scoring index described by Bargmann *et al.* (1993) [47]. See Figure 3.

Supplemental Experimental Procedures

2.1. Automated Software (for Retention Assay): A video camera attached to the microscope produces a digital video stream, which is then analyzed. The ratio of time the average worm spends in each region of interest is calculated for every trial. For ease of implementation, we assumed that all worms in a single experiment are roughly the same size. We thus counted worm pixels instead of whole worms, allowing us to take into account fractions of a worm in the region of interest. It also eliminated the need for a shape-based worm identification algorithm, and allowed each frame to be analyzed independently. We applied a band-pass filter to each frame to eliminate the effect of uneven lighting and also accentuate the worms against the background. The worm was then identified after thresholding the filtered image. Throughout each experiment, we know the locations and sizes of the regions of interest. Through the filtering described above, we know which pixels are occupied by worms and which ones are not. We were then able to calculate the ratio of worm-pixels to all pixels inside the region of interest to produce the worm-occupancy ratio. This calculation is done for every frame, giving us a plot output of worm-occupancy ratio vs. time for each region.

2.2. Culture of Nematodes and Collection of worm Exudates: *C. elegans*, *Rhabditis sp.*, *O. tipulae*, *C. sp.7*, *P. pacificus*, *P. redivivus*, *Koernia sp.*, and *P. strongyloides* were maintained on standard 6 cm agar plates with *E. coli* OP50. To grow large amounts, we grew them in liquid culture using S complete medium with *E. coli* HB101 at 20°C at 250 rpm in an incubator shaker. The worms were exposed to several wash and filtration steps using S Basal to remove bacteria. The worms were collected between the washes by centrifugation at 3000 rpm for 5 minutes. We then incubated the worms at 1 worm/ μ L in S Basal for 6 hours. The worms were then filtered out and the remaining supernatant was passed through a 0.22 μ m filter before being stored at -20°C.

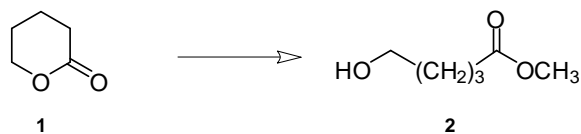
S. carpocapsae, *S. glaseri*, *S. riobrave*, and *H. bacteriophora* were maintained as described by Hallem, E.A. et al. (2011) [50]. Briefly, several last-instar *Galleria mellonella* larvae were placed on a 6 cm Petri dish with a 55 mm Whatman 1 filter paper. Approximately 500-1000 IJs were placed on this filter paper. After 7 days, the insect cadavers were placed on an inverted 6 cm Petri plate carrying a Whatman 1 filter paper and immersed in a small volume of ddH₂O in a 10 cm Petri dish. Emerging IJs were collected and washed several times as described above, before being incubated in ddH₂O at 1 worm/ μ L for 6 hours. Adults were dissected out of insect cadavers after 3-4 days post-infection and washed as described above and incubated at 1 worm/10 μ L (given their large size) or grown on OP50 for use on the holding assay.

N. brasiliensis, *A. suum*, *P. penetrans*, and *Romanomermis spp.* were provided by collaborators (Table S1). *N. brasiliensis* was incubated in 0.85% saline at 1 worm/ 10 μ L (given their large size) at 30°C. *A. suum* were washed in sterile saline before being incubated in DMEM for 3, 6, and 13 hours at 37°C. The supernatants were tested individually, then subsequently pooled for combined analysis. *P. penetrans* and *Romanomermis spp.* were incubated in ddH₂O at 20°C at 250 rpm overnight.

2.3. Worm Sample Preparation: Worm water samples were lyophilized, extracted with 2 x 1-10 ml methanol, and filtered over cotton wool. Extracts were concentrated in vacuum and resulting residues were resuspended in 150 μ L methanol and filtered.

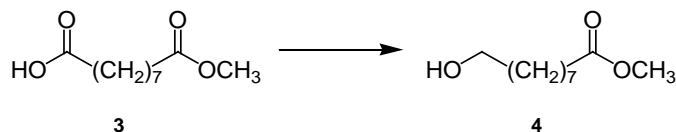
2.4. Syntheses of Ascarosides:

Methyl 5-hydroxypentanoate



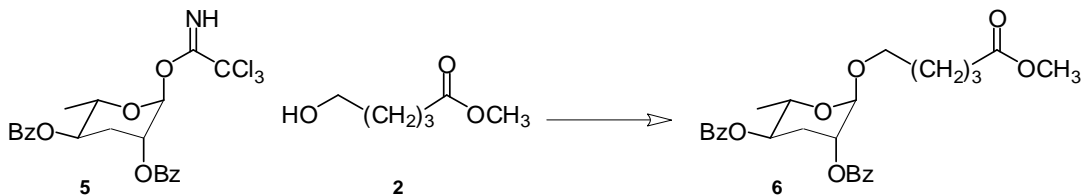
Freshly distilled δ -valerolactone (**1**) (856 mg) in methanol (17 ml) was treated with conc. H_2SO_4 (100 μl) and refluxed for 12 h [51]. The solution was cooled to $-20\text{ }^\circ\text{C}$, treated with NaHCO_3 (80 mg), and stirred for 10 min, after which the mixture was filtered and the solvent removed in vacuum. The residue was taken up in dichloromethane (10 ml), dried over Na_2SO_4 , filtered, and concentrated in vacuum to give methyl 5-hydroxypentanoate (**2**) (1071 mg, 94%) as a colorless oil, which was used directly without any further purification. ^1H NMR (400 MHz, acetone- d_6): δ 1.55 (2H, m), 1.68 (2H, m), 2.35 (2H, t, $J = 7.4$ Hz), 3.57 (2H, m), 3.64 (3H, s); ^{13}C NMR (100 MHz, acetone- d_6): δ 22.2, 32.9, 34.1, 51.5, 61.9, 174.2.

Methyl 9-hydroxynonanoate



A solution of nonanedioic acid monomethyl ester (**3**) (923 mg, 4.6 mmol) in dry tetrahydrofuran (3 ml) at $-20\text{ }^\circ\text{C}$ was treated with 1 M BH_3 in tetrahydrofuran (4.6 ml, 4.6 mmol) over 10 min [52]. After stirring at RT for 4 h, the reaction was quenched with 0.77 M aqueous K_2CO_3 solution (10 ml) at $0\text{ }^\circ\text{C}$. The product was extracted with diethyl ether (3 x 20 ml), washed with saturated aqueous NaCl solution, dried over Na_2SO_4 , and concentrated in vacuum to afford methyl 9-hydroxynonanoate (**4**) (850 mg, 99% yield) as a colorless oil which was used directly without any further purification. ^1H NMR (400 MHz, chloroform- d_1): δ 1.27-1.37 (8H, m), 1.50-1.66 (4H, m), 2.29 (2H, t, $J = 7.5$ Hz), 3.62 (2H, t, $J = 6.5$ Hz), 3.66 (3H, s).

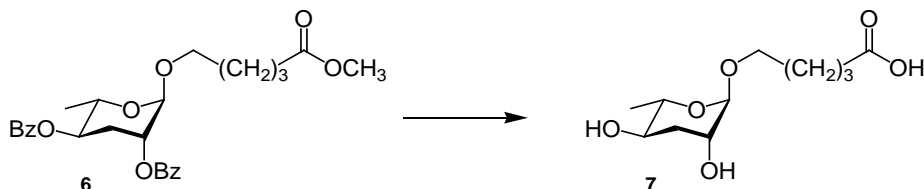
Methyl 5-(3'*R*,5'*R*-dibenzoyloxy-6'*S*-methyl-(2*H*)-tetrahydropyran-2-yloxy)-pentanoate



A solution of 2,4-di-*O*-benzoyl-ascarylose-1-(2,2,2-trichloroacetimidate) (**5**) (132 mg, 263 μmol , prepared according to Butcher, R.A. et al. (2007) [12]) and **2** (125 mg, 950 μmol) in dry dichloromethane (3 ml) at $0\text{ }^\circ\text{C}$ was treated with trimethylsilyloxy triflate (5 μl). After 3 h the solution was washed with saturated aqueous NaHCO_3 solution (0.5 ml), dried over Na_2SO_4 , and concentrated in vacuum. Flash column chromatography on silica gel using a 20 – 40% (v/v) ethyl acetate gradient in *n*-hexane afforded **6** (66.8 mg, 142 μmol , 47%) as a colorless oil. ^1H NMR (400 MHz, acetone- d_6): δ 1.28 (3H, d, $J = 6.2$ Hz), 1.67-1.80

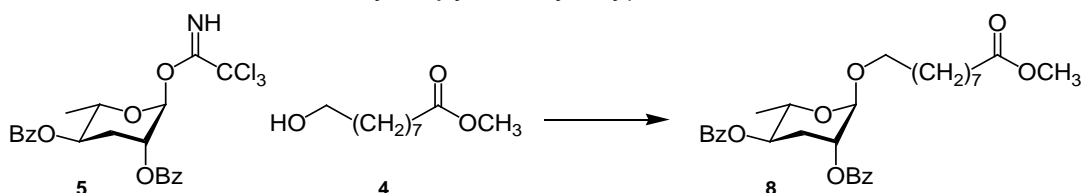
(4H, m), 2.23 (1H, m), 2.40 (2H, t, $J = 6.9$ Hz), 2.48 (1H, m), 3.58 (1H, m), 3.64 (3H, s), 3.83 (1H, m), 4.13 (1H, dq, $J = 9.8$ Hz, $J = 6.0$ Hz), 4.87 (1H, s.br), 5.15 (1H, ddd, $J = 11.0$ Hz, $J = 10.4$ Hz, $J = 4.5$ Hz), 5.18 (1H, s.br), 7.50-7.60 (4H, m), 7.62-7.72 (2H, m), 8.05 (2H, d, $J = 7.5$ Hz), 8.11 (2H, d, $J = 7.5$ Hz); ^{13}C NMR (100 MHz, acetone- d_6): δ 18.3, 22.5, 29.6, 30.4, 34.0, 51.5, 67.5, 67.9, 71.4, 71.5, 97.0, 129.4, 129.5, 130.3, 130.4, 131.0, 131.0, 134.1, 134.2, 165.9, 166.0, 174.0.

5-(3'*R*,5'*R*-Dihydroxy-6'*S*-methyl-(2*H*)-
tetrahydropyran-2-yloxy)-pentanoic acid (oscr#9)



A solution of **6** (66.8 mg, 142 μmol) in dry tetrahydrofuran (0.5 ml) was added to a mixture of LiOH (28 mg, 1.4 mmol) and water (0.6 ml) in 1,4-dioxane (4 ml). After stirring at 66 $^{\circ}\text{C}$ for 2 h the solution was acidified with glacial acetic acid and concentrated in vacuum. Flash column chromatography on silica gel using a 5 – 30% (v/v) methanol gradient in dichloromethane containing 1% glacial acetic acid afforded oscr#9 (**7**) (26 mg, 105 μmol , 74%) as a colorless oil. ^1H NMR (400 MHz, methanol- d_4): δ 1.22 (3H, d, $J = 6.0$ Hz), 1.58-1.72 (4H, m), 1.77 (1H, ddd, $J = 13.1$ Hz, $J = 11.1$ Hz, $J = 3.2$ Hz), 1.95 (1H, ddt, $J = 13.1$ Hz, $J = 3.7$ Hz, $J = 0.9$ Hz), 2.33 (2H, t, $J = 7.2$ Hz), 3.43 (1H, dt, $J = 9.6$ Hz, $J = 6.0$ Hz), 3.47 – 3.59 (2H, m), 3.71 (1H, dt, $J = 9.8$ Hz, $J = 6.2$ Hz), 3.77 (1H, m), 4.50 (1H, s); ^{13}C NMR (100 MHz, methanol- d_4): δ 18.1, 23.0, 30.1, 34.7, 36.0, 67.9, 68.3, 69.4, 70.9, 100.4, 177.5; ESI-MS (negative mode) $m/z = 247.1$ [M-H].

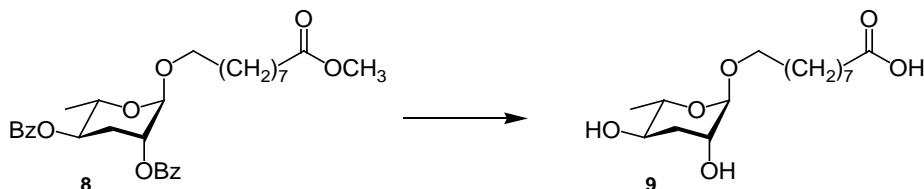
Methyl 9-(3'*R*,5'*R*-dibenzoyloxy-6'*S*-methyl-(2*H*)-
tetrahydropyran-2-yloxy)-nonanoate



A solution of 2,4-di-*O*-benzoyl-ascarylose-1-(2,2,2-trichloroacetimide) (**5**) (132 mg, 263 μmol , prepared according to Butcher, R.A. et al. (2007) [12]) and **4** (112.8 mg, 600 μmol) in dry dichloromethane (3 ml) at 0 $^{\circ}\text{C}$ was treated with trimethylsilyloxy triflate (5 μl). After 3 h the solution was washed with saturated aqueous NaHCO_3 solution (0.5 ml), dried over Na_2SO_4 and concentrated in vacuum. Flash column chromatography on silica gel using 20–40% (v/v) ethyl acetate gradient in *n*-hexane afforded **8** (99.3 mg, 190 μmol , 72% yield) as a colorless oil. ^1H NMR (400 MHz, acetone- d_6): δ 1.28 (3H, d, 6.2 Hz), 1.30 – 1.40 (6H, m), 1.40 – 1.49 (2H, m), 1.56 – 1.72 (2H, m), 2.22 (1H, ddd, $J = 13.6$ Hz, $J = 11.5$ Hz, $J = 3.2$ Hz), 2.30 (2H, t, $J = 7.5$ Hz), 2.46 (1H, m), 3.55 (1H, dt, $J = 9.8$ Hz, $J = 6.5$ Hz), 3.60 (3H, s), 3.81 (1H, dt, $J = 9.6$ Hz, $J = 6.6$ Hz), 4.13 (1H, dq, $J = 9.7$ Hz, $J = 6.2$ Hz), 4.86 (1H, s.br), 5.15 (1H, ddd, $J = 11.4$ Hz, $J = 9.8$ Hz, $J = 4.6$ Hz), 5.18 (1H, m),

7.50 – 7.60 (4H, m), 7.63 – 7.71 (2H, m), 8.04 (2H, m), 8.11 (2H, m); ^{13}C NMR (100 MHz, acetone- d_6): δ 18.3, 25.6, 26.8, 29.7, 29.9, 30.0, 30.2, 30.4, 34.4, 51.4, 67.4, 68.2, 71.4, 71.5, 97.0, 129.4, 129.5, 130.2, 130.3, 130.9, 131.0, 134.1, 134.2, 165.9 (2C), 174.3.

9-(3'*R*,5'*R*-Dihydroxy-6'*S*-methyl-(2*H*)-
tetrahydropyran-2-yloxy)-nonanoic acid (oscr#10)



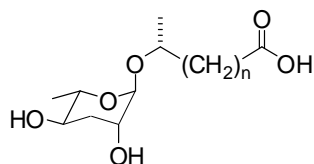
A solution of **8** (99.3 mg, 190 μmol) in tetrahydrofuran (500 μl) was added to a mixture of LiOH (38 mg, 1.9 mmol) and water (800 μl) in 5 ml 1,4-dioxane (5 ml). After stirring at 66 $^{\circ}\text{C}$ for 3 h the solution was acidified with acetic acid and concentrated in vacuum. Flash column chromatography on silica gel using a 5–30% (v/v) methanol gradient in dichloromethane containing 1% glacial acetic acid afforded **9** (49 mg, 161 μmol , 85%) as a colorless oil. ^1H NMR (400 MHz, methanol- d_4): δ 1.22 (3H, d, $J = 6.1$ Hz), 1.32-1.43 (8H, m), 1.56-1.63 (4H, m), 1.77 (1H, ddd, $J = 13.1$ Hz, $J = 11.1$ Hz, $J = 3.2$ Hz), 1.96 (1H, ddt, $J = 13.1$ Hz, $J = 3.7$ Hz, $J = 0.9$ Hz), 2.28 (2H, t, $J = 7.4$ Hz), 3.41 (1H, dt, $J = 9.6$ Hz, $J = 6.2$ Hz), 3.49 – 3.59 (2H, m), 3.68 (1H, dt, $J = 9.8$ Hz, $J = 5.5$ Hz), 3.76 (1H, m), 4.49 (1H, s); ^{13}C NMR (100 MHz, methanol- d_4): δ 17.3, 25.2, 26.4, 28.0, 29.3, 29.5, 29.6, 30.5, 34.1, 61.1, 67.4, 68.5, 69.9, 99.4, 176.8; ESI-MS (negative ion mode) $m/z = 303.2$ [M-H].

Table S1. Nematode Strains

Species	Strain	Source or References
<i>Nippostrongylus brasiliensis</i>	no designation	Edward G. Platzer
<i>Heterorhabditis bacteriophora</i>	M31e	Sternberg collection
<i>Rhabditis sp.</i>	AF5	Caenorhabditis Genetics Center
<i>Oscheius tipulae</i>	PS1305	Sternberg collection
<i>Oscheius dolichuroides</i>	PS1727	Sternberg collection
<i>Oscheius carolinensis</i>	no designation	Sternberg collection
<i>Caenorhabditis elegans</i>	N2	Sternberg collection
<i>Caenorhabditis sp.7</i>	JU1325	Marie-Anne Felix
<i>Ascaris suum</i>	no designation	Robin Gasser
<i>Pristionchus pacificus</i>	RS2333	Ralf Sommer
<i>Koernia sp.</i>	RS1982	Ralf Sommer
<i>Pratylenchus penetrans</i>	3/C	Antoon T. Ploeg
<i>Panagrellus redivivus</i>	PS2298	Sternberg collection
<i>Pelodera strongyloides</i>	DF5022	Caenorhabditis Genetics Center
<i>Steinernema carpocapsae</i>	ALL	Sternberg collection
<i>Steinernema scapterisci</i>	no designation	Sternberg collection
<i>Steinernema riobrave</i>	no designation	Sternberg collection
<i>Steinernema glaseri</i>	VII	S. Patricia Stock
<i>Romanomermis iyengari</i>	no designation	Edward G. Platzer
<i>Romanomermis culicivorax</i>	3B4	Edward G. Platzer
<i>Caenorhabditis elegans</i>	DL238	J. Scott Cameron
<i>Caenorhabditis elegans</i>	CB4856	Caenorhabditis Genetics Center
<i>Caenorhabditis elegans</i>	JU1527	Marie-Anne Felix

All nematode strains used in this study are listed here; species without a strain identification are labeled “no designation”.

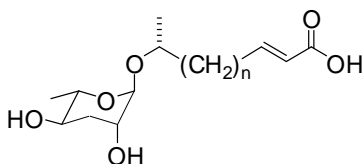
Table S2A. Structures and HPLC-ESI-MS Data of (ω -1)-Oxygenated Saturated Ascarosides



Side chain (n)	SMID ¹	Molecular formula	Molecular mass [g/mol]	<i>m/z</i> [M-H] ⁻
1	ascr#11	C ₁₀ H ₁₈ O ₆	234.1103	233.10
2	ascr#9	C ₁₁ H ₂₀ O ₆	248.1260	247.12
3	ascr#12	C ₁₂ H ₂₂ O ₆	262.1416	261.13
4	ascr#1	C ₁₃ H ₂₄ O ₆	276.1573	275.15
5	ascr#14	C ₁₄ H ₂₆ O ₆	290.1729	289.17
6	ascr#10	C ₁₅ H ₂₈ O ₆	304.1886	303.18
7	ascr#16	C ₁₆ H ₃₀ O ₆	318.2042	317.20
8	ascr#18	C ₁₇ H ₃₂ O ₆	332.2199	331.21
9	ascr#20	C ₁₈ H ₃₄ O ₆	346.2355	345.23
10	ascr#22	C ₁₉ H ₃₆ O ₆	360.2512	359.24
11	ascr#24	C ₂₀ H ₃₈ O ₆	374.2668	373.26
12	ascr#26	C ₂₁ H ₄₀ O ₆	388.2825	387.28

¹SMID: Small Molecule Identifier for small molecules identified from *C. elegans* and other nematodes. The Small Molecule Metabolite Identifier database (www.smid-db.org) is an electronic resource maintained by Frank Schroeder and Lukas Mueller at the Boyce Thompson Institute in collaboration with WormBase. The purpose of this database is to introduce searchable, gene-style identifiers, "SMIDs", for all small molecules newly identified from *C. elegans* and other nematodes.

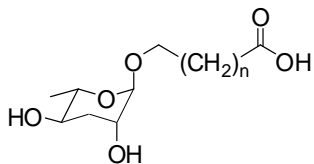
Table S2B. Structures and HPLC-ESI-MS Data of (ω -1)-Oxygenated Unsaturated Ascarosides



Side chain (n)	SMID ¹	Molecular formula	Molecular mass [g/mol]	<i>m/z</i> [M-H] ⁻
2	ascr#7	C ₁₃ H ₂₂ O ₆	274.1416	273.13
4	ascr#3	C ₁₅ H ₂₆ O ₆	302.1729	301.17

¹SMID: Small Molecule Identifier for small molecules identified from *C. elegans* and other nematodes. The Small Molecule Metabolite Identifier database (www.smid-db.org) is an electronic resource maintained by Frank Schroeder and Lukas Mueller at the Boyce Thompson Institute in collaboration with WormBase. The purpose of this database is to introduce searchable, gene-style identifiers, "SMIDs", for all small molecules newly identified from *C. elegans* and other nematodes.

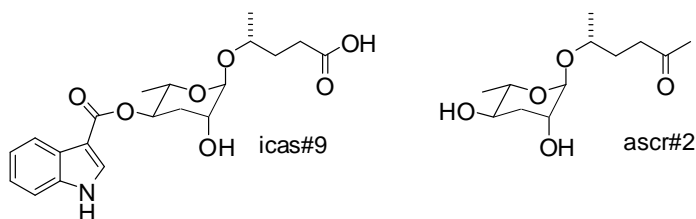
Table S2C. Structures and HPLC-ESI-MS Data of (ω)-Oxygenated Saturated Ascariosides



Side chain (n)	SMID ¹	Molecular formula	Molecular mass [g/mol]	<i>m/z</i> [M-H] ⁻
1	ascr#5	C ₉ H ₁₆ O ₆	220.0947	219.09
3	oscr#9	C ₁₁ H ₂₀ O ₆	248.1260	247.12
7	oscr#10	C ₁₅ H ₂₈ O ₆	304.1886	303.18
9	oscr#18	C ₁₇ H ₃₂ O ₆	332.2199	331.21

¹SMID: Small Molecule Identifier for small molecules identified from *C. elegans* and other nematodes. The Small Molecule Metabolite Identifier database (www.smid-db.org) is an electronic resource maintained by Frank Schroeder and Lukas Mueller at the Boyce Thompson Institute in collaboration with WormBase. The purpose of this database is to introduce searchable, gene-style identifiers, "SMIDs", for all small molecules newly identified from *C. elegans* and other nematodes.

Table S2D. Structures and HPLC-ESI-MS Data of Indole Ascaroside icas#9 and ascr#2



SMID ¹	Molecular formula	Molecular mass [g/mol]	<i>m/z</i>
icas#9	C ₂₀ H ₂₅ NO ₇	391.1631	390.16 [M-H] ⁻
ascr#2	C ₁₂ H ₂₂ O ₅	246.1467	269.14 [M+Na] ⁺

¹SMID: Small Molecule Identifier for small molecules identified from *C. elegans* and other nematodes. The Small Molecule Metabolite Identifier database (www.smid-db.org) is an electronic resource maintained by Frank Schroeder and Lukas Mueller at the Boyce Thompson Institute in collaboration with WormBase. The purpose of this database is to introduce searchable, gene-style identifiers, "SMIDs", for all small molecules newly identified from *C. elegans* and other nematodes.

Supplemental References

50. Hallem, E.A., Dillman, A.R., Hong, A.V., Zhang, Y., Yano, J.M., DeMarco, S.F., and Sternberg, P.W. (2011) A sensory code for host seeking in parasitic nematodes. *Curr Biol*, 21(5), 377-383.
51. Huckstep M., Taylor R.J.K., and Caton M.P.L. (1982) A convenient method of preparing the leukotriene precursor methyl 5-oxopentanoate. *Synthesis*, 10, 881-882.
52. Kai K., Takeuchi, J., Kataoka, T., Yokoyama, M., and Watanabe, N. (2008) Structure-activity relationship study of flowering-inducer FN against *Lemna paucicostata*. *Tetrahedron*, 64, 6760-6769.