

Additional file 3: Table S1. Comparative analyses of orthologous identified in biological pathways. Orthologous in RNAi pathway (A), immune responses (B), dauer/diapause pathway (C) and the synthesis, transportation and metabolism of neurotransmitters (D) were compared among *C. elegans*, *H. avenae* and other 6 nematodes with published genomic data. Cel, *Caenorhabditis elegans*; CCN, Cereal cyst nematode *H. avenae*; Gpa, *Globodera pallida*; Mha, *Meloidogyne hapla*; Min, *Meloidogyne incognita*; Bxy, *Bursaphelenchus xylophilus*; Asu, *Ascaris suum*; Ppa, *Pristionchus pacificus*.

Table S1A. Comparison of orthologous identified in RNAi pathway among *C. elegans*, *H. avenae* and other 6 nematodes with published genomic data.

Gene function	Cel	CCN	Gpa	Mha	Min	Bxy	Asu	Ppa
Small RNA biosynthetic proteins	<i>drh-3</i>	+	+	+	+	+	+	+
	<i>drsh-1</i>	+	+	+	+	+	+	+
	<i>xpo-1</i>	+	+	+	+	+	+	+
	<i>xpo-2</i>	+	+	+	+	+	+	+
	<i>dcr-1</i>	+	+	+	+	+	+	+
	<i>drh-1</i>			+	+	+	+	+
	<i>pash-1</i>	+	+	+	+	+	+	+
	<i>rde-4</i>							
	<i>xpo-3</i>	+					+	
	<i>smg-2</i>	+	+	+	+	+	+	+
Amplification	<i>smg-6</i>	+	+	+	+	+	+	+
	<i>ego-1</i>	+	+	+	+	+	+	+
	<i>rrf-3</i>					+	+	+
	<i>rrf-1</i>							
	<i>smg-5</i>							
Spreading	<i>rsd-2</i>							
	<i>rsd-3</i>	+	+	+	+	+	+	+
	<i>sid-1</i>						+	
	<i>rsd-6</i>							
	<i>sid-2</i>							
Argonautes	<i>alg-1</i>	+	+	+	+	+	+	+
	<i>wago-1</i>	+	+	+	+	+	+	+
	C04F12.1	+	+	+	+		+	+
	<i>wago-4</i>	+	+	+	+	+	+	+
	<i>alg-4</i>					+	+	+
	<i>rde-1</i>	+	+	+	+	+	+	+
	<i>hrde-1</i>	+	+	+	+	+	+	+
	<i>ppw-1</i>							
	<i>csr-1</i>	+	+	+	+	+		
	<i>ppw-2</i>	+	+	+	+	+	+	+
	<i>sago-1</i>							
	<i>alg-3</i>					+	+	+
	<i>wago-10</i>	+	+	+	+	+	+	+
	<i>alg-2</i>							
	<i>ergo-1</i>							
	<i>prg-1</i>							+
Other RISC components	<i>wago-2</i>	+	+	+	+	+	+	+
	<i>hpo-24</i>							
	<i>nrde-3</i>	+	+	+	+	+	+	+
	<i>sago-2</i>							
	T23B3.2						+	+
	<i>wago-11</i>	+	+	+	+	+	+	+
	<i>wago-5</i>	+	+	+	+	+	+	+
	<i>prg-2</i>							+
	<i>tsn-1</i>	+	+	+	+	+	+	+
	<i>ain-1</i>	+		+	+	+	+	
<i>vig-1</i>						+	+	+
	<i>ain-2</i>	+	+					

	<i>eri-1</i>	+	+	+	+	+	+
	<i>xrn-2</i>	+	+	+	+	+	+
	<i>adr-2</i>					+	+
	<i>xrn-1</i>					+	+
	<i>adr-1</i>					+	+
RNAi inhibitors		<i>lin-15b</i>					
	<i>eri-5</i>						
	<i>eri-6</i>						
	<i>eri-7</i>	+				+	+
	<i>eri-3</i>						
Nuclear RNAi effectors		<i>mut-7</i>	+	+	+	+	+
	<i>cld-1</i>	+	+	+	+	+	+
	<i>ekl-1</i>	+	+	+	+	+	+
	<i>gfl-1</i>	+	+		+	+	+
	<i>mes-2</i>	+	+	+	+	+	+
	<i>ekl-4</i>	+		+	+	+	+
	<i>mes-6</i>	+	+	+		+	+
	<i>rha-1</i>	+	+	+	+	+	+
	<i>ekl-6</i>					+	+
	<i>zfp-1</i>					+	+
	<i>mut-2</i>						+
	<i>ekl-5</i>						
	<i>mes-3</i>						
	<i>mut-16</i>						
	<i>rde-2</i>						

Cel, *Caenorhabditis elegans*; CCN, Cereal cyst nematode *H. avenae*; Gpa, *Globodera pallida*; Mha, *Meloidogyne hapla*; Min, *Meloidogyne incognita*; Bxy, *Bursaphelenchus xylophilus*; Asu, *Ascaris suum*; Ppa, *Pristionchus pacificus*.

Table S1B. Comparison of orthologous identified in immune responses among *C. elegans*, *H. avenae* and other 6 nematodes with published genomic data.

Immune signalling	Cel	CCN	Gpa	Mha	Min	Bxy	Asu	Ppa
DBL-1/TGF-beta signalling pathway	<i>dbl-1</i>	+	+	+	+	+	+	+
	<i>sma-2</i>	+	+	+	+	+	+	+
	<i>sma-3</i>	+	+	+	+	+	+	
	<i>sma-4</i>	+	+	+	+	+	+	+
ERK MAPK signalling pathway	<i>lin-45</i>	+	+	+	+	+	+	+
	<i>mak-2</i>	+	+	+	+	+	+	+
	<i>mpk-1</i>	+	+	+	+	+	+	
P38 MAPK signalling pathway	<i>nsy-1</i>	+	+	+	+	+	+	+
	<i>pmk-1</i>	+	+	+		+	+	+
	<i>sek-1</i>	+	+	+	+	+	+	
	<i>tir-1</i>	+	+	+	+	+	+	+
Toll signalling pathway	<i>tol-1</i>	+	+	+	+	+	+	
	<i>trf-1</i>	+				+	+	+
	<i>ikb-1</i>						+	+
	<i>plk-1</i>	+	+	+	+	+	+	
Insulin (DAF-2/DAF-16) signaling pathway	<i>age-1</i>	+		+	+	+	+	+
	<i>pdk-1</i>	+	+	+	+	+	+	+
	<i>akt-1</i>	+	+	+	+	+	+	+
	<i>akt-2</i>	+	+	+	+	+	+	+
	<i>daf-16</i>	+	+	+		+	+	
	<i>daf-2</i>	+	+	+	+	+	+	+

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Table S1C. Comparison of orthologous identified in dauer/diapause pathway among *C. elegans*, *H. avenae* and *G. pallida*.

Dauer pathway	Protein	Cel	CCN	Gpa
Guanylyl cyclase pathway	Transmembrane guanylate cyclase	<i>daf-11</i>	+	+
	cGMP-gated channel	<i>tax-2</i>	+	+
	cGMP-gated channel	<i>tax-4</i>	+	+
TGF β -like	TGF- β type I receptor	<i>daf-1</i>	+	+
	SMAD transcription factor	<i>daf-3</i>	+	+
	TGF β type II receptor	<i>daf-4</i>	+	+
	Proline rich protein	<i>daf-5</i>		
	BMP/TGF- β	<i>daf-7</i>	+	+
	SMAD transcription factor	<i>daf-8</i>		
	SMAD transcription factor	<i>daf-14</i>		
	Glutamine rich protein	<i>scd-1</i>	+	+
	Tyrosine kinase	<i>scd-2</i>	+	+
	Zn-finger protein	<i>bra-1</i>	+	
Insulin/IGF	Tyrosine kinase	<i>cam-1</i>	+	+
	cGMP-dependent protein kinase	<i>egl-4</i>	+	+
	Insulin receptor	<i>daf-2</i>	+	+
	Ortholog RAPTOR protein	<i>daf-15</i>	+	+
	FOXO transcription factor	<i>daf-16</i>	+	+
	Phosphoinositide 3-phosphatase PTEN	<i>daf-18</i>	+	
	β -insulin	<i>daf-28</i>		
Steroid hormone pathway	Phosphoinositide 3-kinase	<i>age-1</i>	+	
	3-phosphoinositide-dependent kinase	<i>pdk-1</i>	+	+
	Serine/threonine kinase	<i>akt-1</i>	+	+
	Serine/threonine kinase	<i>akt-2</i>	+	+
	Serine/threonine kinase	<i>sgk-1</i>	+	
	Cytochrome P450	<i>daf-9</i>	+	+
Other processes	Nuclear receptor	<i>daf-12</i>	+	+
	Rieske oxygenase, hormone pathway	<i>daf-36</i>	+	
	amphid morphology	<i>daf-6</i>		+
	WD-WAA rep	<i>daf-10</i>	+	+
	RFX transcription factor	<i>daf-19</i>	+	+
	HSP-90	<i>daf-21</i>		+

Cel, *Caenorhabditis elegans*; CCN, Cereal cyst nematode *H. avenae*; Gpa, *Globodera pallida*.

Table S1D. Comparison of orthologous identified in the synthesis, transportation and metabolism of neurotransmitters among *C. elegans*, *H. avenae* and other 4 plant-parasitic nematodes.

Neurotransmitter	Gene function	Cel	CCN	Gpa	Mha	Min	Bxy
Acetylcholine	Choline acetyltransferase	<i>cha-1</i>	+	+	+	+	+
	Synaptic acetylcholine transporter	<i>unc-17</i>	+	+	+		+
	Choline transporter	<i>cho-1</i>	+	+	+		+
	Post-synaptic transporter	<i>snf-6</i>	+	+	+	+	+
	Acetylcholinesterase	<i>ace-1</i>	+	+	+		+
	Acetylcholinesterase	<i>ace-2</i>	+	+		+	+
	Acetylcholinesterase	<i>ace-3</i>	+	+	+	+	+
	Acetylcholinesterase	<i>ace-4</i>					
Serotonin	Tryptophan hydroxylase	<i>tph-1</i>		+	+		+
	GTP-cyclohydrolase I	<i>cat-4</i>	+	+	+		+
	Aromatic AA decarboxylase	<i>bas-1</i>	+	+	+		+
	Vesicular monoamine transporter	<i>cat-1</i>	+	+	+		+
	Serotonin reuptake transporter	<i>mod-5</i>	+	+	+	+	+
	Monoamine oxidase	<i>amx-1</i>					+
	Monoamine oxidase	<i>amx-2</i>					
	Monoamine oxidase	<i>amx-3</i>					
Dopamine	Tyrosine hydroxylase	<i>cat-2</i>					+
	Dopamine reuptake transporter	<i>dat-1</i>	+	+	+		+
Tyramine	Tyrosine decarboxylase	<i>tdc-1</i>	+	+	+	+	+
Octopamine	Tyramine β -hydroxylase	<i>tbh-1</i>	+	+	+	+	+
Glutamate	Vesicular glutamate transporter	<i>eat-4</i>	+	+	+	+	+
	Plasma membrane glutamate transporter	<i>glt-1</i>	+	+	+	+	+
GABA	Glutamate decarboxylase	<i>unc-25</i>	+	+	+	+	+
	Esicular GABA transporter	<i>unc-47</i>	+	+	+	+	+
	GABA transporter	<i>snf-11</i>	+	+	+	+	+
	GABA transaminase	<i>gta-1</i>	+	+	+	+	+

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