Root-Knot Nematodes Exhibit Strain-Specific Clumping Behavior That Is Inherited as a Simple Genetic Trait

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Abstract
Root-knot nematodes are obligate parasites of a wide range of plant species and can feed only on the cytoplasm of living plant cells. In the absence of a suitable plant host, infective juveniles of strain VW9 of the Northern root-knot nematode, *Meloidogyne hapla*, when dispersed in Pluronic F-127 gel, aggregate into tight, spherical clumps containing thousands of worms. Aggregation or clumping behavior has been observed in diverse genera in the phylum Nematoda spanning free-living species such as *Caenorhabditis elegans* as well as both plant and animal parasites. Clumping behavior differs between strains of *M. hapla* and occurs with other species within this genus where strain-specific differences in clumping ability are also apparent. Exposure of *M. hapla* juveniles to a gradient formed using low levels of cyanide promotes formation of clumps at a preferred cyanide level. Analysis of F2 lines from a cross of *M. hapla* strains that differ in clump-forming behavior reveals that the behavior segregates as a single, major locus that can be positioned on the genetic map of this nematode. Clumping behavior may be a survival strategy whose importance and function depend on the niche of the nematode strain or species.

Introduction
Root-knot nematodes (*Meloidogyne* spp.) are obligate endoparasites of a wide range of plant species and globally cause large crop yield losses [1,2]. The infective stage, which is the second stage juvenile (J2), hatches from an egg in the soil and must find a host and establish a feeding site in order to survive. The J2 typically enters the host root in the zone of elongation then migrates to the vascular cylinder to establish a feeding site and complete its life cycle. Little is known about the behavior and survival of the infective juvenile. Many nematodes, including most plant parasitic species, inhabit the soil for at least part of their life cycle. Natural nematode behavior has been particularly difficult to study in this opeque, non-uniform environment [3–5]. As an attempt to gain insights into the behavior of pre-infective J2s, we developed a Petri dish assay using Pluronic F-127 (PF-127), a clear, non-toxic co-polymer that as a 23% aqueous solution is liquid at 15°C and solidifies into a soft gel at room temperature [6]. Nematodes suspended in PF-127 gel, rather than being limited to the surface as with agar plates, can respond to signals perceived in three dimensions. Movement of nematodes to root tips can be monitored as well as changes in behavior as they approach the zone of elongation where they penetrate the root. Because stable chemical gradients form in PF-127, we were able to show that root-knot nematode J2 accumulate at pH between 4.5 and 5.5, consistent with the low pH of the zone of elongation relative to other regions of the root [7].

A particularly interesting finding was that, when no seedling was included, J2 of *M. hapla* strain VW9, *M. javanica* strain VW4, and *M. incognita* strain 557R aggregate into tightly packed spherical clumps by two days after being uniformly suspended in the PF-127 gel [6]. The clumping behavior was accelerated by increasing the nematode concentration or by putting a coverslip on the gel surface, suggesting that low oxygen or a volatile signal from the nematode was involved. Interestingly, we found that clumps dispersed when a root of an appropriate host was added.

The obligate parasitism and long life cycle of plant parasitic nematodes compared to *C. elegans*, have hindered research approaches aimed at understanding the genes responsible for behavior and parasitism. However, recent advances have increased the feasibility of these goals. Genome sequences are now available for the root-knot nematode species *M. incognita* and *M. hapla* [8,9]. While several of the agriculturally damaging root-knot nematode species, including *M. incognita* and *M. javanica*, reproduce by mitotic parthenogenesis precluding any genetic analysis, many *M. hapla* strains reproduce by a novel mechanism that allows both outcrossing and selling [2,10]. This reproduction mechanism has facilitated the production of F2 lines and the generation of a genetic map based on molecular polymorphisms between the parental strains [9,10]. As a consequence of their novel reproductive mechanism, these F2 lines are homozygous for most markers and thus resemble recombinant inbred lines [10]. Due to the availability of the genome sequence, a genetic map,
and phenotypic variability within species, *M. hapla* is emerging as a model plant parasitic nematode. Here we investigate the inheritance in F2 lines of the ability to form clumps and present the first demonstration of the utility of these lines for mapping a biologically significant trait.

**Results**

**Clumping behavior differs among root-knot nematode strains and species**

We previously found that infective J2 of three different root-knot nematode species (*M. hapla* strain VW9, *M. javanica* strain VW4, and *M. incognita* strain 557R) aggregated into tightly packed spherical clumps by two days after being uniformly suspended in PF-127 [6]. To expand this analysis, 6 inbred *M. hapla* strains that originated from diverse geographic locations and hosts were compared for this behavior (Table 1). Four of these strains (VW9, VW10, VW11, and VW12) formed clumps by the second day. In contrast, the *M. hapla* strain NCS aggregated to some extent but never formed tight clumps, and strain LM did not aggregate at all. Analysis of cultures of closely related *M. javanica* strains VW4 and VW5 that differ in their ability to infect tomato with the resistance gene Mi-1 found that both formed clumps equally well. However, five *M. incognita* strains initially obtained from different locations and hosts tested differed dramatically in clumping behavior (Table 1). *Meloidogyne incognita* strain VW6 formed multiple clumps in one day, whereas strains Beltran and W1 formed 1–3 clumps, and strain Harmony formed none even after 48 h. As previously noted [6], when a coverslip was placed on the surface of the gel, the clumping was accelerated for each clump-forming strain and was visible under the coverslip within 24 hr (not shown). However, those strains that did not clump were not induced to do so by the addition of a coverslip.

Exposure to cyanide gradients promotes clumping behavior

In preliminary chemical screens we noted that low concentrations of potassium cyanide (KCN) appeared to attract *M. hapla* VW9 J2 and cause them to form clumps. To investigate this further, chemical dispensers (modified pipet tips; see reference 7) filled with 2 or 5 mM KCN were placed into Petri dishes containing J2 of VW9 dispersed in PF-127 gel. Based on our previous work with pH gradients [7], we anticipated that KCN would form a radial concentration gradient centered on the openings of the dispensers. By 5 h after assay initiation, nematodes could be seen to accumulate at both openings of the chemical dispensers with 2 mM KCN (Figure 1A), and in a region of about 5 mm diameter around the small opening of the dispenser with 5 mM KCN (Figure 1C). By 24 hr, nematodes had aggregated into a compact ball at the small opening of the dispenser (Figure 1B). For the dispenser with 5 mM KCN, the clumps formed a ring centered on the opening (Figure 1D). A circle of clumps was also seen around the large openings of the dispensers with 2 mM or 5 mM KCN with a larger diameter at the latter concentration (Figure 1F-H). For those cases with a ring of clumps, nematodes also accumulated within the circle, but the area closest to the opening of the dispenser was depleted of nematodes (Figure 1D). This novel distribution of J2 was reproducible in multiple replicates with independently hatched nematodes. In controls in which the dispenser contained only PF-127 gel, nematodes remained randomly distributed in the plate (Figure 1E).

| Table 1. Comparison of Meloidogyne strain clump formation and attraction to cyanide. |
|----------------------------------|---------------------------------|-----------------|-----------------|-----------------|
| *Species and strains* | *Nematode isolate origin: Host and geographic location* | *Clumps in PF-127 by 48 h* | *2 mM KCN: # J2 within 5 mm at 3 h* | *2 mM KCN: clump formation by 24 h* |
| *M. hapla* | | | | |
| VW9 | Tomato, CA, USA | Yes | 116.8a | Yes |
| VW10 | Hemp, The Netherlands | Yes | 65.4b | Yes |
| VW11 | Everlasting flowers, The Netherlands | Yes | 74.7b | Yes |
| VW12 | Tomato, CA, USA | Yes | 65.7b | Yes |
| LM | Unknown host, France | No | 50.0c | No |
| NCS | Unknown host, North Carolina, USA | No | 48.0c | No |
| *M. javanica* | | | | |
| VW4 | Unknown, CA, USA | Yes | 46.8c | Yes |
| VW5 | Unknown, CA, USA | Yes | 50.8c | Yes |
| *M. incognita* | | | | |
| VW6 | Cotton, CA, USA | Yes | 20.3d | No |
| Beltran | Lima bean, CA, USA | Yes | 31.3d | No |
| Harmony | Grape, CA, USA | No | 9.6e | No |
| 557R | Tomato, NC, USA | Yes | 30.3d | No |
| W1 | Tomato, CA, USA | Yes | 23.6d | No |

*Petri dishes contained 300 J2 per ml PF-127 gel.

1Average number of nematodes within 5 mm of the small opening of the dispenser minus the average number of nematodes in the corresponding region with a dispenser containing PF-127 gel alone. Significant differences are indicated by different letters (P<0.05). doi:10.1371/journal.pone.0015148.t001
were rigid at cyanide concentrations of 40 mM. Most nematodes were straight and did not move. All nematodes tested, including 110 mM potassium ions or elevated pH, was responsible for the attraction.

Because nematodes have been widely reported to be sensitive to cyanide, we examined the response of *M. hapla* VW9 J2 to a series of KCN concentrations in 10 μM increments from 0–110 μM in a 6-well-plate containing 1500 J2s in 5 ml water in each well. Nematodes were observed under microscope after 30 minutes. Nematodes in water displayed typical sinusoidal movement. In 10 and 20 μM KCN movement was slower and nematodes were curled, that is, head and tail were in close proximity resembling the conformation that we have previously observed as nematodes reach the root surface [6]. In 30 μM KCN, most nematodes were straight and did not move. All nematodes were rigid at cyanide concentrations of 40 μM or higher. By 24 h, nematodes had regained motility in all concentrations of cyanide. Many of the nematodes in the leading region at 24 h were observed to be curled in confirmation, rather than sinusoidal. The cyanide concentration in this region was estimated to be between 12 and 25 μM. In control assay tubes without the cyanide gradient, most J2 remained in sinusoidal shape, dispersed throughout the tube, and formed neither bands nor clumps.

**Meloidogyne strains differ in response to cyanide**

We compared the responses of different root-knot nematode species and strains to KCN gradients in PF-127 gels (Table 1). At 5 hours, VW9 had the greatest accumulation near the dispenser opening of the *M. hapla* strains tested, and strains LM and NCS were significantly less responsive than VW10, 11, and 12. The closely related *M. javanica* strains VW4 and VW5 were significantly less responsive than *M. hapla* strains VW9, 10, 11, or 12. The five *M. incognita* strains tested showed less accumulation than the other species, with *M. incognita* strain Harmony showing the least accumulation. By 24 h, the numbers of nematodes gathered near the dispenser opening were too high to count (>100), and some strains had formed clumps around the opening (Table 1). For *M. hapla* and *M. javanica* strains, the presence or absence of clumps correlated with that seen in plates with PF-127 alone at 48 h. However, no clumps were seen for any of the five *M. incognita* strains tested even though some of these formed clumps in PF-127 at 48 h.

**Inheritance of clumping trait in Meloidogyne hapla**

We assessed J2 from greenhouse cultures of 87 F2 lines from a cross of *M. hapla* strains VW8 and VW9 for clump formation in response to cyanide gradients produced around openings of chemical dispensers containing 2 mM KCN in PF-127 gel. Of the 87 lines, 40 were scored as non-clumping, 45 as clumping
Two lines produced ambiguous results giving small clumps in some replicates, but none in others. In addition, the degree of clumping varied to some extent between replicates in that some consistently produced large clumps whereas other produced only small ones. Nevertheless, using the 85 lines that we were able to score, the Clumping trait (Clm) could be assigned to Linkage Group 8 of the genetic map using JoinMap® 4 (LOD 10.0), and its relative position on the linkage group assigned (Figure 3B).

**Discussion**

All assays that attempt to examine natural behavior of microscopic organisms in the laboratory have limitations. However, such assays can lead to discovery of novel, relevant behaviors. PF-127 gel is highly transparent and provides sufficient traction to allow nematodes to move easily resolving difficulties with assays on agar surfaces where nematodes can be trapped in water films. We have found that the formation of tight clumps by
root-knot nematodes initially dispersed in PF-127 gel is a remarkably reproducible behavior. We have observed similar clumps on agar plates and in liquid, although in these two media the clumps were noted less consistently. Thus, while clumping is likely a natural phenomenon, the properties of the PF-127 gel, especially its ability to reduce or eliminate convection, may stimulate clumping behavior and stabilize the clumps. We observe three general conformations, sinusoidal, curled (head and tail in close proximity), and straight for nematodes in Pluronic gel. The sinusoidal shape was observed for nematodes undergoing directed movement, whereas the curled shape has been seen previously for nematodes in close proximity to their favored entry sites on host roots [6] and, in the current work, for nematodes in response to cyanide. Nematodes in the curled conformation change position in the gel more slowly than sinusoidal worms. The straight conformation has been observed for nematodes paralyzed by cyanide or other chemicals as well as for dead worms.

Aggregation, clumping and related behaviors have been observed in diverse nematode species, including free-living species and both plant and animal parasites [14–18]. For root-knot nematodes, the observations that the rate of clump formation is positively correlated with nematode population density and is accelerated by placement of a coverslip on the gel surface are consistent with the hypotheses that reduced oxygen, increased concentration of a volatile pheromone or contact with a solid surface can initiate or stabilize the clumping [6]. Clumping can also be seen when high numbers of infective J2 reach the zone of elongation of a healthy root, the targeted entry site for this nematode. In this case, signals from the root or from J2 may be signaling the clumping behavior [6].

Perhaps the most studied example of nematode aggregation is as a component of a set of related behaviors including social feeding (aggregation on a bacterial lawn), bordering (accumulation at the edges of bacterial lawns), and agitated movement in Caenorhabditis elegans. This group of behaviors is pronounced in many natural isolates of C. elegans, but is relatively weak in other strains, including the standard laboratory strain N2, due to a variant allele of the G-protein coupled receptor encoded by npr-1 [19]. Several other genes modulating this response have been identified and include genes whose products mediate oxygen sensation, suggesting that social feeding may be a response to hypoxic environments [20]. In contrast to social feeding in C. elegans, the clumping of root-knot nematode juveniles is observed when the nematodes are not feeding. In C. elegans, the phenomenon is manifested as loose aggregates with dozens of actively moving L4 or adult animals on an agar plate, whereas the clumps formed by root-knot nematodes contain more individuals and are formed by infective juveniles, an earlier and more uniform developmental stage. A more appropriate comparison may be with the desiccation-induced clumping behavior of certain entomopathogenic nematodes. Infective juveniles of some Heterorhabditis species, but not others, aggregate into large clumps when desiccated at high relative humidity, and this behavior is thought to confer desiccation tolerance [17]. Some plant parasitic nematodes, for example Aphelenchus avenae and Ditylenchus dipsaci, are full anhydrobiotes, aggregate into large clumps, classically referred to as “celworm wool,” and can survive for many years in a dehydrated state [15,21]. While root-knot nematodes and Heterorhabditis species do not have the capability to survive complete drying, clumping may help them to survive moderate levels of desiccation. This, together with our previous observation that clumps disappear when a seedling is introduced nearby [6], suggests that clumping may be a mechanism to increase survival against desiccation or other adverse condition as the J2s await the detection of a suitable host.

Our finding that KCN gradients attract RKNs was initially surprising as cyanide is toxic to many life forms. In fact, cyanide poisoning is responsible for the killing of C. elegans by some strains of Pseudomonas aeruginosa [11] and has been implicated as responsible for the biocontrol activity of some rhizosphere bacteria against soil pests and pathogens including nematodes [12,22]. Plant cyanogenesis, that is, release of HCN from damaged tissues, is considered to be a defense against herbivores including nematodes [23], and cyanogenic plant material has also been used for control of nematodes [13]. Potassium cyanide hydrolyzes slowly in a pH-dependent fashion to release HCN, which is highly volatile, resulting in concentration levels that are dynamic and difficult to measure accurately. By our estimates using two different assays, accumulation and clump formation for M. hapla VW9, the most responsive strain that we tested, occurs at about 15 μM KCN. We found that movement of VW9 was slowed, but worms were not paralyzed at this concentration. While they were paralyzed by higher concentrations, the paralysis was reversible, even for 110 μM KCN, presumably due to loss of cyanide by volatilization.

Figure 3. Segregation of clumping behavior in M. hapla F2 lines. A. Response to dispensers with 2 mM KCN in PF-127 gel 24 h after initiation of assay of F2 lines A16, A26, A54 (non-clumping), and lines A31, A58, and C61 (clumping). Petri dish contained 300 J2 per ml in 20 ml PF-127 gel for each assay. B. Map of linkage group 8 with the positions of DNA markers and the Clumping locus (Clm). Size bar in A is 1 mm.

doii:10.1371/journal.pone.0015148.g003
A possible explanation for the accumulation of J2 at specific HCN concentrations is that HCN interferes with oxygen perception. Caenorhabditis elegans accumulation at particular O2 levels is modulated by guanylate cyclases that are expressed in different sensory neurons and balance attraction and repulsion to specific oxygen levels [24,25]. These soluble guanylate cyclases bind O2 or other gases through associated heme groups, and cyanide, which can bind to heme groups, could compete or otherwise interfere with this signaling. Alternatively, concentration gradients of cyanide may be a natural signal for nematodes as there are many sources of cyanide in the rhizosphere. The formation of clumps at a specific concentration may simply be a consequence of localized accumulation of nematodes as high concentrations of nematodes promote clumping. In fact, other chemicals that cause accumulation, such as weak acids, also result in clump formation when initial concentrations of nematodes suspended in the PF-127 gel are high [7] (Williamson, unpublished).

The lower accumulation of M. incognita compared to M. hapla in response to cyanide may partially explain why M. incognita does not form clumps in the presence of CN. This is not due to a generally lower movement rate of M. incognita as this species accumulates more rapidly at root tips and at low pH than does M. hapla [6,7]. However it is possible that M. incognita is less attracted to cyanide gradients or is more sensitive to paralysis by cyanide.

While clumping behavior appears to be widespread in the phylum Nematoda, closely related species and even strains of a single species differ in their propensity to form clumps [17,19]. Perhaps each response has an advantage under particular environmental conditions. It may be relevant, for example, that the non-clumping M. incognita strain Harmony was isolated from a perennial crop (grape) whereas the other isolates examined were from annual crops. For C. elegans, it has been suggested that social feeding and solitary nematodes favor different food dispersal strategies [26]. While variant alleles of npr-1 explain much of the difference between isolates of C. elegans, genetic studies have uncovered several other genes that can modulate the response, and it may be that variant alleles of these genes modulate clumping behavior in other species [25,27,28].

The segregation of clumping behavior that we find for our M. hapla F2 lines is consistent with the possibility that in this species, as in C. elegans, a variant allele of a single gene accounts for much of the variation. However, we do see differences in degree of clumping among the F2 lines suggesting that modifiers of the phenotype may also be segregating in our population. The clumping trait in M. hapla is the first phenotype to be genetically mapped in this obligate parasitic species. While the current genetic and physical maps are so far insufficiently developed to identify the responsible genes, positional cloning should become feasible as a greater density map and more F2 lines become available. While transformation is not possible so far for root-knot nematode species, gene silencing by RNAi is an available strategy to assess candidate genes [29]. Identification of genes involved in chemical signaling or responsible for survival in the soil and location and modification of host roots will increase our understanding of this highly evolved parasite and may facilitate the development of novel strategies for control of these damaging pests.

Materials and Methods

Nematode cultures

Nematode sources and culture methods are as previously described [7,30]. Eggs were collected and surface-sterile second stage juveniles were produced as previously described [7].
The Spectroquant 14800 colorimetric HCN assay (EM Science 14000, Gibbstown, NJ) was used according to the manufacturer’s instructions with the following modifications: 1) total reaction volume was reduced to 0.5 ml; 2) solution CN-1A was kept on ice then added to cold cuvettes; 3) cuvettes were kept at room temperature for 5 min before adding CN-2 and CN-3 reagents; 4) readings were taken after 30 min, and 5) reaction mixtures were diluted 1:1 with H₂O before absorbance was read at 765 nm with a DU 640 Spectrophotometer (Beckman Coulter Inc., Fullerton, CA). Experimental samples were measured against the average of 3 controls containing PF-127 gel in 10 mM sodium phosphate, pH 7.0. A standard curve was generated using 0.25 to 50.00 µM KCN in PF-127 gel in 10 mM sodium phosphate, pH 7.0. Absorbance measurements were fit to a linear correlation model (R² = 0.99).

Mapping clumping trait in Meloidogyne hapla F2 lines

Juveniles of F2 lines previously developed from a cross of M. hapla strains VW8 and VW9 [10] were hatched from eggs harvested from greenhouse cultures then surface sterilized. Two or more Petri plates with 300 J2/ml in 23% PF-127 each with two dispensers filled with 2 mM KCN in 23% PF-127 were assessed for each F2 line. At 24 h after assay initiation, the presence or absence of clumps at the opening of the dispenser was scored. The linkage analysis of the clumping trait and polymorphic DNA markers was performed with the software JoinMap 4 [31] using the Kosambi mapping function. Polymorphic DNA markers include AFLP markers [9] and single nucleotide polymorphisms (SNPs). SNP polymorphisms are annotated by the letter ‘C’ followed by the contig number and the nucleotide position of the polymorphism [9] (http://www.pnlgg.org).

Acknowledgments

We thank George Bruning for comments on the manuscript and Mohit Bansal for help with experiments.

Author Contributions

Conceived and designed the experiments: CW SL VMW. Performed the experiments: CW SL VMW. Analyzed the data: CW SL VMW. Contributed reagents/materials/analysis tools: VMW. Wrote the paper: CW SL VMW.

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