

# Distribution of entomopathogenic nematodes (Steinernematidae and Heterorhabditidae) in natural habitats in California, USA

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Abstract. A total of 270 soil samples from 30 different habitats in 10 geographic regions of California were evaluated for the presence of rhabditid entomopathogenic nematodes. Nematodes were isolated from 26.3% of the samples. The recovered isolates were identified as *Steinernema carpocapsae*, *S. feltiae*, *S. kraussei*, *S. longicaudum*, *S. oregonense*, *Heterorhabditis marelatus* and *H. bacteriophora*. Among the steinernematids, *S. kraussei* and *S. feltiae* were the most commonly encountered species, generally occurring in acidic soils high in organic matter. Among the heterorhabditids, *H. bacteriophora* was isolated along the southern coast, whereas *H. marelatus* was recovered along the northern coast of California. Steinernematids were recovered from coniferous forests, oak woodlands and grasslands whereas heterorhabditids were isolated from coastal marshes.

Key words: California, insect-pathogenic nematodes, natural habitats, survey

#### Introduction

Entomopathogenic nematodes (EPN) in the families Steinernematidae and Heterorhabditidae are obligate parasites of soil-inhabiting insects and have great potential as biological control agents of many insect pests. They are found in a variety of soil habitats, and the various species and isolates exhibit considerable variation in terms of host range, reproduction, infectivity and conditions for survival (i.e. temperature, soil moisture, etc.) (Bedding et al. 1983; Bedding 1990). To further advance the use of EPN as biological control agents, locally-adapted species or isolates from native non-disturbed habitats need to be identified and their unique characteristics documented.

Surveys for EPN have been conducted in temperate, subtropical and tropical regions (Hominick et al. 1996). In the US, surveys have been conducted in Puerto Rico (Roman and Beavers 1983), Florida (Beavers et al. 1983), North Carolina (Akhurst and Brooks 1984), Hawaii (Hara et al. 1991), New Jersey (Gaugler et al. 1992), Tennessee (Rueda et al. 1993) and Oregon (Liu and Berry



1995). In California, several steinernematid and heterorhabditid nematodes have been recovered from agricultural areas and golf courses (Poinar 1990; Stock et al. 1996), however, no systematic survey has been conducted to document the occurrence of these nematodes from natural or non-disturbed habitats. In contrast to human modified areas, natural habitats are more likely uncontaminated by introduced nematodes and offer a better chance for finding native species. California is unique in the diversity of habitats found, and may contribute to an equally diverse distribution of EPN. The objective of this study was to survey EPN diversity in a variety of undisturbed (non-agricultural) habitats of California and identify the recovered isolates using classical and molecular taxonomy.

#### Methods

#### Collection and isolation of nematodes

Soil samples were collected from 10 landscape regions of California (Durrenberger 1959) between March and June 1996. Within each region, 1–4 of the predominant habitats (vegetation types) were selected for sampling (Table 1). Three sites were selected within each habitat and 3 samples were collected at each site. Each soil sample (approximately 1 kg) was a composite of 5 random subsamples taken at a depth of 10–20 cm in an area of 20 m<sup>2</sup>. Samples were taken at least 100 m apart at each site. Samples were placed in polyethylene bags to prevent water loss and were kept in coolers (ca. 15 °C) during transit to the laboratory.

EPN were recovered from soil samples by the insect baiting technique (Bedding and Akhurst 1975). Five *Galleria mellonella* (L.) larvae were placed in 250 ml plastic containers (5 containers/sample) with moistened soil obtained from each sample. Containers were covered with a lid, turned upside down and kept at room temperature ( $20 \pm 3 \text{ °C}$ ). After 7–8 days, all insects were recovered, and parasitized cadavers were individually placed in White traps to allow the emergence of the infective-stage juveniles. Emerging nematodes were pooled for each sample and used to infect fresh *G. mellonella* larvae to produce nematodes for identification and establishment of cultures. Soil samples with negative nematode recovery were baited a second time to confirm results of the first test.

For each sampling site, a portion of the soil from each of the 3 samples (ca. 100 g) was combined and analyzed for the following soil characteristics: pH, organic matter, sand, silt and clay content. Soil analysis was performed by the Division of Agricultural and Natural Resources (DANR) Analytical Laboratory at the University of California, Davis. Positive samples (EPN present) were correlated with landscape region, habitat and soil characteristics (Table 2).

Table 1. Landscape regions, natural habitats and sampling time.

Landscape regions	Habitats	Sampling time
R1. Klamath-Siskiyou	Mixed coniferous forest	5/96
	Grassland	6/96
	Chaparral	6/96
R2. Northeast volcanic	Mixed coniferous forest	5/96
	Grassland	5/96
	Chaparral	6/96
R3. Sierra Nevada foothills and low coastal mountains	Mixed coniferous forest	5/96
	Grassland	6/96
	Chaparral	6/96
	Oak woodland	4/96
R4. Great Valley	Oak woodland	5/96
	Grassland	5/96
	Wetland	6/96
R5. Sierra Nevada	Mixed coniferous forest (> 5000 ft.)	6/96
	Grassland (meadow) (> 5000 ft.)	6/96
	Lodgepole pine forest (>9000 ft.)	6/96
R6. Southwest mountains and valleys	Mixed coniferous forest	3/96
	Oak woodland	3/96
	Chaparral	3/96
	Grassland	3/96
R7. Desert and desert mountains	Pinion-juniper forest	3/96
	Joshua tree forest	3/96
	Desert scrub	3/96
	Grassland	3/96
R8. Northern coastal strip	Grassland	5/96
	Coastal chaparral	5/96
	Coastal marsh	5/96
R9. Southern coastal strip	Grassland	3/96
	Coastal chaparral	3/96
	Coastal marsh	3/96
R10. Redwoods	Redwood forest	5/96

## Taxonomic studies

Nematode isolates recovered were identified by morphological criteria using standard light microscopy and scanning electron microscopy (SEM) according to Kaya and Stock (1997). For biometric studies, approximately 30 adult males of first and second generation and 30 infective juveniles from selected representatives of each isolate were fixed in triethanolamine formalin (TAF) and processed to glycerin (Kaya and Stock 1997). Morphometrical analysis was done with the aid of Jandel Java software<sup>TM</sup> with a high resolution video

R1: Klamath-SiskyouMCF $6.3$ $7.13$ Sandy loam $3/3$ $5.$ carpoorR2: NE VolcanicGL $6.3$ $4.31$ Sandy loam $3/3$ $5.$ fattaceR3: Sierra foothills andOW $5.5$ $6.46$ Sandy loam $3/3$ $5.$ fattaceIow coastal mountainsOW $6.3$ $5.01$ Loam $3/3$ $5.$ fattaceNo coastal mountainsOW $6.3$ $5.01$ Loam $3/3$ $5.$ fattaceNo coastal mountainsOW $6.2$ $2.81$ Sandy loam $3/3$ $5.$ fattaceNo coastal mountainsOW $6.2$ $2.81$ Sandy loam $3/3$ $5.$ fattaceNo coastal mountainsOW $6.2$ $2.81$ Sandy loam $3/3$ $5.$ fattaceNo copyOW $6.2$ $5.96$ Sandy loam $3/3$ $5.$ fattaceR5: Sierra NevadaMCF $6.7$ $5.24$ Sandy loam $3/3$ $5.$ carporaNCF $6.7$ $5.24$ Sandy loam $3/3$ $5.$ longicatNCF $6.7$ $5.24$ Sandy loam $3/3$ $5.$ longicatNCF $5.5$ $5.49$ Sandy loam $3/3$ $5.$ latrasceNCF $5.6$ $5.49$ Sandy loam $3/3$ $5.$ latrasceNoMCF $5.6$ $5.49$ Sandy loam $3/3$ $5.$ latrasceNCN $5.5$ $5.49$ Sandy loam $3/3$ $5.$ latrasceNN $5.5$ $5.13$ Sandy loam $3/3$ $5.$	Region	Habitat	Hq	MO %	Soil type	No. positive samples /Total no. samples	Nematode species
R2: NE Volcanic         GL         6.3         4.31         Sandy loam         3/3         S. <i>fetture</i> R3: Sterra foothills and         0W         5.5         6.46         Sandy loam         3/3         S. <i>krausse</i> low coastal mountains         0W         6.3         5.01         Loam         3/3         S. <i>krausse</i> low coastal mountains         0W         6.3         5.01         Loam         3/3         S. <i>krausse</i> R4: Great Valley         0W         6.2         2.81         Sindy loam         3/3         S. <i>krausse</i> R4: Great Valley         0W         6.2         5.96         Sandy loam         3/3         S. <i>krausse</i> R5: Sierra Nevada         0CF         6.8         6.46         Sandy loam         3/3         S. <i>carpoca</i> R5: Sierra Nevada         MCF         6.7         5.24         Sandy loam         3/3         S. <i>carpoca</i> R6: SW Mountains and valleys         MC         5.73         Sandy loam         3/3         S. <i>krausse</i> MCF         6.1         4.67         Sandy loam         3/3         S. <i>krausse</i> S. <i>krausse</i> R6: SW Mountains and valleys         MC         5.5	R1: Klamath-Siskiyou	MCF	6.3	7.13	Sandy loam	3/3	S. carpocapsae
R3: Sierra foothils and low coastal mountainsOW5.56.46Sandy loam2/3S. krausselow coastal mountainsOW6.22.81Sandy loam3/3S. krausseR4: Great ValleyOW6.22.81Sandy loam3/3S. krausseR4: Great ValleyOW6.22.81Sandy loam3/3S. krausseR5: Sierra NevadaOW6.22.81Sandy loam3/3S. carpocaR5: Sierra NevadaMCF6.86.46Sandy loam3/3S. carpocaR5: Sierra NevadaMCF6.14.67Sandy loam3/3S. carpocaR6: SW Mountains and valleysMCF5.49Sandy loam3/3S. krausseMCF6.14.67Sandy loam3/3S. krausseMCF5.55.30Sandy loam3/3S. krausseMCF5.55.30Sandy loam3/3S. krausseMCF5.55.30Sandy loam3/3S. krausseMCF5.55.30Sandy loam3/3S. krausseNCF5.65.13Sandy loam3/3S. krausseNCF5.13Sandy loam3/3S. krausseNCF5.65.39Sandy loam3/3S. krausseNCF5.05.39Sandy loam3/3S. krausseNCF5.15.4Sandy loam3/3S. krausseNCF5.15.3Sandy loam3/3S. krausse<	R2: NE Volcanic	GL	6.3	4.31	Sandy loam	3/3	S. feltiae
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R4: Great Valley         OW $6.3$ 5.01         Loam $3/3$ S. krausse           R4: Great Valley         OW $6.2$ 2.81         Sandy loam $3/3$ S. krausse           OW $6.2$ 2.81         Sandy loam $3/3$ S. carpoca         S. krausse           R3: Sierra Nevada         OW $6.2$ 5.96         Sandy loam $3/3$ S. carpoca           R5: Sierra Nevada         MCF $6.8$ $6.46$ Sandy loam $3/3$ S. carpoca           R5: Sierra Nevada         MCF $6.7$ $5.24$ Sandy loam $3/3$ S. carpoca           R6: SW Mountains and valleys         MCF $6.1$ $4.67$ Sandy loam $3/3$ S. <i>cargoca</i> MCF $5.5$ $5.30$ Sandy loam $3/3$ S. <i>cargoca</i> MCF $5.6$ $5.13$ Sandy loam $3/3$ S. <i>cargoca</i> MCF $5.6$ $5.30$ Sandy loam $3/3$ S. <i>trausse</i> MCF $5.6$ $5.31$ Sandy loam $3/3$ S. <i>tr</i>	low coastal mountains						
R4: Great Valley         OW         6.2         2.81         Sandy loam         1/3         S. krausse           R4: Great Valley         OW         7.0         2.83         Sitt loam         3/3         S. carpoca           R5: Sierra Valley         OW         5.4         2.41         Sandy loam         3/3         S. carpoca           R5: Sierra Vevada         MCF         6.8         6.46         Sandy loam         3/3         S. carpoca           R5: Sierra Nevada         MCF         6.1         4.67         Sandy loam         3/3         S. carpoca           MCF         6.1         4.67         Sandy loam         3/3         S. longica           LPF         6.1         4.67         Sandy loam         3/3         S. longica           M         5.5         5.30         Sandy loam         3/3         S. longica           M         5.5         5.13         Sandy loam         3/3         S. longica           MCF         5.6         5.13         Sandy loam         3/3         S. longica           MCF         5.6         5.13         Sandy loam         3/3         S. longica           MCF         5.6         5.13         Sandy loam         3/3 </td <td></td> <td>OW</td> <td>6.3</td> <td>5.01</td> <td>Loam</td> <td>3/3</td> <td>S. kraussei</td>		OW	6.3	5.01	Loam	3/3	S. kraussei
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OW         5.4         2.41         Sandy loam         3/3         5. carpoca           R5: Sierra Nevada         MCF         6.8         6.46         Sandy loam         3/3         5. carpoca           MCF         6.2         5.96         Sandy loam         3/3         5. longica           MCF         6.1         4.67         Sandy loam         3/3         5. longica           LPF         6.1         4.67         Sandy loam         3/3         5. longica           LPF         6.1         4.67         Sandy loam         3/3         5. longica           MCF         5.2         5.49         Sandy loam         3/3         5. krause           MCF         5.6         5.13         Sandy loam         3/3         5. krause           MCF         5.6         5.13         Sandy loam         3/3         5. krause           MCF         5.0         5.39         Sandy loam         3/3         5. krause           MCF         5.0         5.34         Sandy loam         3/3         5. krause           MCF         5.0         5.33         Sandy loam         3/3         5. krause           OW         5.5         5.35         Sandy loam<	R4: Great Valley	OW	7.0	2.83	Silt loam	3/3	S. carpocapsae
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LPF       6.2       5.49       Sandy loam $3/3$ S. krausse         M       5.5       5.30       Sandy loam $3/3$ S. krausse         M       5.5       5.30       Sandy loam $3/3$ S. krausse         MCF       5.6       5.13       Sandy loam $3/3$ S. vrausse         MCF       5.0       5.39       Sandy loam $3/3$ S. oregone         MCF       6.1       4.81       Sandy loam $3/3$ S. oregone         OW       5.5       5.35       Sandy loam $3/3$ S. krausse         OW       5.2       2.44       Sandy loam $3/3$ S. krausse         OW       5.2       2.44       Sandy loam $3/3$ S. krausse         R8: North Coast       CM       6.3       2.01       Loam $3/3$ H. marela         R9: South Coast       CM <td< td=""><td></td><td>LPF</td><td>6.1</td><td>4.67</td><td>Sandy loam</td><td>3/3</td><td>S. kraussei</td></td<>		LPF	6.1	4.67	Sandy loam	3/3	S. kraussei
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R6: SW Mountains and valleys       MCF       5.6       5.13       Sandy loam       3/3       S. oregone         MCF       5.0       5.39       Sandy loam       3/3       S. oregone         MCF       6.1       4.81       Sandy loam       3/3       S. oregone         OW       5.5       5.35       Sandy loam       3/3       S. krausse         OW       5.2       2.44       Sandy loam       3/3       S. krausse         OW       5.2       2.44       Sandy loam       3/3       S. krausse         OW       6.3       2.81       Sandy loam       3/3       H. marela         R8: North Coast       CM       6.3       2.10       Loam       3/3       H. marela         R8: South Coast       CM       6.3       2.10       Loamy sand       3/3       H. marela         R9: South Coast       CM       7.1       1.27       Sand       3/3       H. bacteria		М	5.5	5.30	Sandy loam	3/3	S. kraussei
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OW       5.5       5.35       Sandy loam       3/3       S. krausse         OW       5.2       2.44       Sandy loam       3/3       S. krausse         OW       5.2       2.44       Sandy loam       3/3       S. krausse         OW       6.3       2.81       Sandy loam       3/3       S. krausse         OW       6.3       2.81       Sandy loam       3/3       H. marela         CM       7.2       2.77       Loam       3/3       H. marela         CM       6.4       3.96       Sandy loam       3/3       H. marela         CM       6.3       2.10       Loamy sand       3/3       H. marela         R9: South Coast       CM       6.3       1.10       Loamy sand       3/3       H. bacteria         CM       6.3       1.10       Loamy sand       3/3       H. bacteria		MCF	6.1	4.81	Sandy loam	3/3	S. oregonense
$ \begin{array}{cccccccccccccccccccccccccccccccccccc$		OW	5.5	5.35	Sandy loam	3/3	S. kraussei
OW         6.3         2.81         Sandy loam         3/3         S. krausse           R8: North Coast         CM         7.2         2.77         Loam         3/3         H. marela           CM         7.2         2.77         Loam         3/3         H. marela           CM         6.4         3.96         Sandy loam         3/3         H. marela           CM         6.3         2.10         Loamy sand         3/3         H. marela           R9: South Coast         CM         6.3         1.10         Loamy sand         3/3         H. bacteria           CM         6.3         1.10         Loamy sand         3/3         H. bacteria		OW	5.2	2.44	Sandy loam	3/3	S. kraussei
R8: North Coast         CM         7.2         2.77         Loam         3/3         H. marela           CM         6.4         3.96         Sandy loam         3/3         H. marela           CM         6.4         3.96         Sandy loam         3/3         H. marela           CM         6.3         2.10         Loamy sand         3/3         H. marela           R9: South Coast         CM         6.3         1.10         Loamy sand         3/3         H. bacteria           CM         6.3         1.10         Loamy sand         3/3         H. bacteria		OW	6.3	2.81	Sandy loam	3/3	S. kraussei
CM $6.4$ $3.96$ Sandy loam $3/3$ $H.$ marela           CM $6.4$ $3.96$ Sandy loam $3/3$ $H.$ marela           CM $6.3$ $2.10$ Loamy sand $3/3$ $H.$ marela           R9: South Coast         CM $6.3$ $1.10$ Loamy sand $3/3$ $H.$ bacteria           CM $6.3$ $1.10$ Loamy sand $3/3$ $H.$ bacteria	R8: North Coast	CM	7.2	2.77	Loam	3/3	H. marelatus
CM $6.3$ $2.10$ Loamy sand $3/3$ $H.$ marelaR9: South CoastCM $7.1$ $1.27$ Sand $3/3$ $H.$ bacteriaCM $6.3$ $1.10$ Loamy sand $3/3$ $H.$ bacteria		CM	6.4	3.96	Sandy loam	3/3	H. marelatus
R9: South Coast         CM         7.1         1.27         Sand         3/3         H. bacteri           CM         6.3         1.10         Loamy sand         3/3         H. bacteri		CM	6.3	2.10	Loamy sand	3/3	H. marelatus
CM 6.3 1.10 Loamy sand $3/3$ H. bacteri	R9: South Coast	CM	7.1	1.27	Sand	3/3	H. bacteriophora
		CM	6.3	1.10	Loamy sand	3/3	H. bacteriophora
GL 5.3 5.56 Loam 2/3 S. fetrae		GL	5.3	5.56	Loam	2/3	S. feltiae

Table 2. EPN-positive sites within California landscape regions.

camera. In general, and for all the nematode stages (first and second generation adults and infective juveniles), the following characters were analyzed: total length, greatest width, distance from anterior end to excretory pore, distance from anterior end to nerve ring, distance from anterior end to base of esophagus, tail length, width at anus/cloaca, and values of ratio a, b, c, d (expressed as D%) and e (expressed as E%). In particular, within the males, testis reflexion, length and width of the spicules, length and width of gubernaculum and number and arrangement of genital papillae were also considered. For first and second generation females, morphology of the vulva, V% and shape of tail were also examined. First generation males and infective juveniles of selected representatives from each group were examined by SEM for confirmation of observations done with light microscopy. According to their morphology, all isolates were placed into different species groups using taxonomic criteria suggested by Stock and Kaya (1996) and Stock et al. (1997).

To further confirm the identity of the *Steinernema* isolates recovered, crossbreeding tests were conducted according to Kaya and Stock (1997). Representative isolates of each species group were cross-bred with each other and with known isolates of the species group they were assigned and were more closely related (Table 3). With the exemption of *S. longicaudum* (no 'type isolate' was available), the following species were considered to confirm identity of the species: *S. feltiae* (SN strain), *S. carpocapsae* (ALL strain), *S. kraussei* (Westphalia and Nash strains), *S. oregonense* (OS10 strain). Similarly, each of the *Heterorhabditis* isolates recovered were cross-bred with each other and with reference isolates OH10 (*H. marelatus*) and Chino Hill (*H. bacteriophora*) using the lipid agar method of Dix et al. (1992). In both cases, controls consisted of crosses between the same isolate/species.

In addition to morphological and cross-breeding studies, two randomly selected isolates of each identified *Steinernema* species were sent for molecular characterization to A. P. Reid (International Institute of Parasitology, St. Albans, UK) (Figures 2–3) (Table 3). These isolates were subjected to RFLP analysis (Reid and Hominick 1992) and the resulting band patterns were compared to an RFLP library representing of over 35 *Steinernema* species.

## Results

EPN were recovered from 71 (26.3%) of the 270 soil samples collected from 12 distinct habitats in 8 out of the 10 surveyed regions (Figure 1). No nematodes were recovered from soil samples collected from the desert and redwood regions. With the exception of two sites in Region 3 (Sierra Nevada foothills and

Table 3. Hybridization to	ests o	of sel	lecte	ЯE	ΓN	isola	tes fi	rom	Cali	forni	a.																			
Isolates/species	1	7	Э	4	5	9	7	8	6	10	11	12	13	14	15	16	17	18	9 2	0 2	1 22	23	24	25	26	27	28	29 3	0 31	32
S. carpocapsae isolates																														
1. R1-mcf21 <sup>a</sup>	+																													
2. R4-ow12 <sup>a</sup>	+	+				Ι			I																					
3. R4-ow21	+	+	+			Ι				I																				
4. R4-ow33	+	+	+	+				Ι																						
5. S. c. <sup>1</sup> (All)	+	+	+	+	+	1				I						I				I	I							I	1	
S. feltiae isolates						-																								
0. K2-g152 7. R9-g111 <sup>a</sup>					I	+ +	+		I													I								
8. S. f. <sup>2</sup> (SN)	Ι	Ι				+	+	+		Ι		Ι		I	I				I										Ι	
S. kraussei isolates																														
9. R3-ow11						I				+ -	-																			
10. R3-0w24 11. R3-0w31 <sup>a</sup>						I				+ +	+ +	+																		
12. R5-lpf11			Ι			Ι				+	+	+																		
13. R5-lpf23 <sup>a</sup>	I	I							+		+	+																		
14. R5-g112		I	Ι			Ι			+		+	+	+					+												
15. R6-ow12						Ι			+		+									Ι	Ι	I								
16. R6-ow23	I	Ι				Ι				+	+				+		+			Ι	Ι									
17. R6-ow31	Ι	I				Ι				+	+	+				+	+			I	I									
18. S. k <sup>3</sup> (Westphalia)			Ι	Ι	Ι	Ι			+	+	+	+	+	+	+	+	+	+	+		I	Ι			I			1		
19. S. k <sup>3</sup> (Nash)			Ι	I	I	I			+	+	+	+	+	+	+	+	+	+	I		Ι	Ι			I			1		
S. longicaudum isolates 20. R5-mcf12 <sup>a</sup>																				Т										
21. R5-mcf23 (B2) <sup>a</sup>																				+										
S. oregonense isolates	I	I		I		I			I											- 1	+									
23. R6-mcf22			Ι							I			I						1		+	+								

(Continued)
Table 3.

Isolates/species	1	7	ŝ	4	5	5	3 2	6	1(	11	12	13	14	15	16	17 1.	8 15	9 20	21	22	23	24 2	25 2	6 2	7 28	29	30	31	32
24. R6-mcf32 <sup>a</sup> 25. S. o. <sup>4</sup> (OS10)		I	I	1	,				I	I			I					I		+ +	+ +	+ +	+						
<ul> <li><i>H. bacteriophora</i> isolates</li> <li>26. R9-cm12</li> <li>27. R9-cm22</li> <li>28. H. b.<sup>5</sup> (Chino Hills)</li> </ul>																							1 1 1	+ + +	+	1 1 1			1 1
<i>H. marelatus</i> isolates 29. R8-cm11 30. R8-cm22 31. R8-cm31 32. H. m. <sup>6</sup> (OH10)																										+ + + +	+ + +	+ +	+

Abbreviations: <sup>1</sup> S. carpocapsae; <sup>2</sup> S. feltiae; <sup>3</sup> S. kraussei; <sup>4</sup> S. oregonense; <sup>3</sup> H. bacteriophora; <sup>6</sup> H. marelatus. +: progeny; -: no progeny; in blank: no crosses. <sup>a</sup>Submitted for molecular characterization (RFLP analysis).



*Figure 1.* Geographic regions of California showing sampling sites. 1. Klamath-Siskiyou; 2. Northeast Volcanic; 3. Sierra foothills and low coastal mountains; 4. Great Valley; Sierra Nevada; 6. Southwest Mountains and Valleys; 7. Desert and desert mountains; 8. North coastal strip; 9. South coastal strip; 10. Redwoods. Positive sampling sites: ( $\Box$ ) *S. carpocapsae*; ( $\triangle$ ) *S. feltiae*; ( $\blacklozenge$ ) *S. kraussei*; ( $\blacksquare$ ) *S. oregonense*; ( $\bigcirc$ ) *S. longicaudum*; ( $\blacktriangle$ ) *H. bacteriophora*; ( $\blacklozenge$ ) *H. marelatus*; ( $\bigstar$ ) negative sampling sites.

low coastal mountains) and one site in Region 9 (southern coastal strip), all positive sampling sites yielded 3 out of 3 EPN-positive samples (Table 2). Of the isolates recovered, 80% were steinernematids and 20% were hetero-rhabditids. Nematodes were identified as: *Steinernema feltiae* (Filipjev 1934), *S. kraussei* Steiner 1923 (Figure 2), *S. carpocapsae* (Weiser 1934), *S. long-icaudum* Shen and Wang 1992 (Figure 3), *S. oregonense* Liu and Berry 1996a, *Heterorhabditis bacteriophora* Poinar 1975 and *H. marelatus* Liu and Berry 1996b.

Steinernematids were more ubiquitous than the heterorhabditids, having been isolated from 7 out of 10 regions. *S. kraussei* was the most abundant (32.8% of the total positive samples) and widely distributed species. It was found in 3 different landscape provinces – the Sierra Nevada foothills and



1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 M

Figure 2. PCR amplified products from the internal transcribed spacer (ITS) of S. kraussei digested with 17 different enzymes. Fragments were separated with ethidium bromide stained 1.5% (W/V) agarose gel. A. Westphalia isolate; B. R3-OW24 isolate (California). Lane 1 is a digest of S. feltiae (UK, site 76) with Alu I. Lanes 2-18 are individual digests of S. kraussei with the following restriction enzymes: 2. Alu I; 3. Bst 0 I; 4. Dde I; 5. ECOR I; 6. Hae III; 7. Hha I; 8. Hind III; H inf I; 10. Hpa I; 11. Kpm I; 12. Pst I; 13. Pvu I; 14. Rsa I; 15. Sal I; 16. Sau A I; 17. Sau 96; 18. Xba I; 19. molecular weight marker and the band sizes are shown in base pairs.

coastal mountains, Sierra Nevada and SW mountains and valleys. S. feltiae was recovered in the northeast volcanic and southern coast regions. S. carpocapsae was isolated in the Klamath-Siskiyou and Great Valley. S. oregonense



*Figure 3.* PCR amplified products from the internal transcribed spacer (ITS) of *S. longicaudum* digested with 17 different enzymes. Fragments were separated with ethidium bromide stained 1.5% (W/V) agarose gel. A. Chinese isolate; B. R5-mcf23 (B2) isolate (California). Lane 1 is a digest of *S. feltiae* (UK, site 76) with *Alu* I. Lanes 2–18 are individual digests of *S. longicaudum* with the following restriction enzymes: 2. *Alu* I; 3. *Bst* 0 I; 4. *Dde* I; 5. *ECOR* I; 6. *Hae* III; 7. *Hha* I; 8. *Hind* III; H inf I; 10. *Hpa* I; 11. *Kpm* I; 12. *Pst* I; 13. *Pvu* I; 14. *Rsa* I; 15. *Sal* I; 16. *Sau* A I; 17. *Sau* 96; 18. *Xba* I; 19. molecular weight marker and the band sizes are shown in base pairs.

and *S. longicaudum* were isolated from the SW mountains and valleys and the Sierra Nevada, respectively (Table 1).

The *Heterorhabditis* species were restricted to the Northern and Southern coastal regions, but with a distinct geographic delimitation between the two species (Table 1). *H. marelatus* was exclusively isolated along the northern

coastal strip, and *H. bacteriophora* was only recovered along the south coastal strip.

## Habitats

The majority of the positive samples (48 of 71) were recovered from woodlands (coniferous forest (33.8%) and oak forests (33.8%)). Coastal marshes and grasslands yielded the remaining positive samples (23.9% and 8.5%, respectively). No nematodes were recovered from soil samples taken from chaparral habitats, redwood forest, or any of the desert habitats (scrub, chaparral and Joshua tree forest).

In terms of species diversity, the coniferous forests were the richest habitats, yielding 4 out of the 5 species of *Steinernema* recovered. Next were the coastal marshes, where two *Heterorhabditis* species (*H. marelatus*, *H. bacteriophora*) and one steinernematid (*S. feltiae*) were recovered. Oak woodlands were characterized by the presence of *S. kraussei* and *S. carpocapsae*, and the grasslands were represented by *S. kraussei* and *S. feltiae*.

In terms of EPN recovery, several habitats had significantly higher recovery frequencies than others. Most notable were the oak woodlands in which 100% of the sites sampled were positive in all regions where they were sampled. Also showing 100% recovery rate were the mixed coniferous forests of the southwest mountains and the north coastal marshes. South coastal marshes, Sierra Nevada mixed coniferous forests, and Sierra Nevada lodge-pole pine forests had recovery frequencies of 67%.

## Soil type

Soil types classified from sand to clay with a pH that ranged from acidic (4.2) to slightly alkaline (7.2). In general, steinernematids were found in sandy loam soils that ranged from acidic (pH 5.0) to neutral (Table 2). The organic matter content of the nematode-positive soil samples varied from 2.4% to 7.1%. As for the *Heterorhabditis* species, *H. bacteriophora* was recovered from sand to loamy sand soils, with a pH ranging from slightly acidic to slightly alkaline (6.3–7.1). The organic matter content of these soils was the lowest recorded from all the positive soil samples (Table 2). The soil types where *H. marelatus* was found ranged from loam to sandy loam. The pH of the soils was similar to those for *H. bacteriophora* (6.3–7.2). The organic matter content for these soils was higher than that for *H. bacteriophora* (2.1–3.9% vs. 1.1–1.2%).

## Discussion

The present study has shown for the first time the occurrence of EPN in nondisturbed habitats in California. Moreover, an accurate identification (to the species level) of all the isolates recovered was accomplished.

In this survey, 26.3% of the samples taken were nematode-positive with a predominance of steinernematids (80%) over heterorhabditids (20%). Most of the *Steinernema* species were recovered from coniferous and oak forests. Even though no nematodes were recovered from soil samples taken from chaparral habitats, redwood forest, or any of the desert habitats, a more intensive survey of these habitats would be necessary to fully confirm these results.

Among the steinernematids, *S. kraussei* was the most commonly encountered species in California. It was isolated above 7000 ft from coniferous forests and a grassland in the Sierra Nevada Mountains; at 3000 ft in oak woodlands in the Valley SW Mountains and Valleys and near sea level in oak woodlands in the Sierra Foothills and Coastal Mountains. This is in agreement with European studies in which this species has been found at both high (coniferous forests) and low elevations (meadows) (Mracek et al. 1992; Steiner 1996). Although this is the first report of *S. kraussei* in California, recent studies have confirmed the presence of this species in US (New York) and Canada (Alberta and British Columbia) (Stock et al. unpublished).

An unexpected finding from this survey was the recovery of *S. longicaudum* from coniferous forests (lodgepole pine) in the Sierra Nevada mountains. This species was originally isolated from an orchard in China (Shen 1992), and has subsequently been isolated from Australia (Hominick et al. 1996). This is the first report of this species in North America and expands its known geographic range.

*S. oregonense* was originally isolated from a grassland in Oregon (Liu and Berry 1996). In our survey, this species was found in mixed coniferous forests in the Southwest Mountains and Valleys, expanding the habitat range and natural distribution of this nematode to Southern California.

*S. carpocapsae* was recovered from oak woodlands in the Great Valley and mixed coniferous forests in the Klamath-Siskiyou region. However, this species has also been reported from an apple orchard (Poinar 1990) and golf courses in the foothills of the Sierra Nevada (Kaya and Koppenhofer pers. comm.), which suggests this species may be widely distributed in California.

*S. feltiae* was recorded from a grassland and a coastal marsh in Southern California, and from grasslands in the Northeast volcanic region. This is in agreement with results from surveys conducted in other parts of the world, where grasslands seem to be the habitat of preference for this species (Boag et al. 1992; Hominick et al. 1995; Miduturi et al. 1996; Steiner 1996).

In this survey, *Heterorhabditis* species, were isolated in coastal marshes from both North and South coastal strips. This is in agreement with other studies

where heterorhabditids have been found to be more common in sandy coastal soils (Griffin et al. 1991; Hara et al. 1991; Liu and Berry 1995). However, previous studies have reported the presence of *H. bacteriophora* from orchards and golf courses in California (Stock et al. 1996) indicating this species is not restricted to the coastal strips.

Like other surveys (Hara et al. 1991, Griffin et al. 1994), our study showed that sandy loam soils seem to be the preferred soil types for EPN. This is supported by the fact that mobility and survival of EPN are favored in soils with a high sand content, whereas soils with high clay content restrict nematode movements (Molyneux and Bedding 1984; Kung et al. 1990). In terms of pH tolerance, EPN were found in soils with a pH ranging from acidic (pH 5) to slightly alkaline (pH 7.2). This agrees with other studies where the pH of nematode-positive soil samples varied from 4.6 to 8 (Hara et al. 1991; Griffin et al. 1994).

Our results extend the geographic range of several species. The presence in California of *S. longicaudum* indicates this species has a more extended geographic distribution. According to Hominick et al. (1996) this species has been isolated from two distant locations China and Australia. In view of this, the Californian isolate of *S. longicaudum*, as well as the Chinese and Australian, may represent a species with a Pacific Rim distribution or relic populations of a previously widespread species. The recovery of *S. oregonense* in California extends the known geographic range of this species further south than previous reports (Liu and Berry 1996a, 1996b; Stock et al. 1996). The finding of *S. carpocapsae*, *S. feltiae* and *H. bacteriophora* in California supports the notion sustained by Hominick et al. (1996) that these three species are ubiquitous.

An important finding of this survey is that at all the EPN-positive sites, only one single nematode species was recovered. Furthermore, at over 90% of the EPN-positive sites, all three samples collected contained nematodes. This information should be considered in future surveys in California so that emphasis can be done on sampling more sites rather than multiple samples per site.

Finally, the richness of species encountered in this survey, reflects the diversity of habitats found in California and suggests the adaptability of EPN to a wide range of habitats. This is a very encouraging aspect since the nematodes isolated during this survey may contribute to the expanded use of native isolates of entomopathogenic nematode in future biological control programs in California and other areas with similar climates.

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Westphalia isolate of. S. kraussei and OS10 isolate of S. oregonense, respectively.

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