

# A Comparative Study of Life History Traits in *C. briggsae* and *C. elegans*

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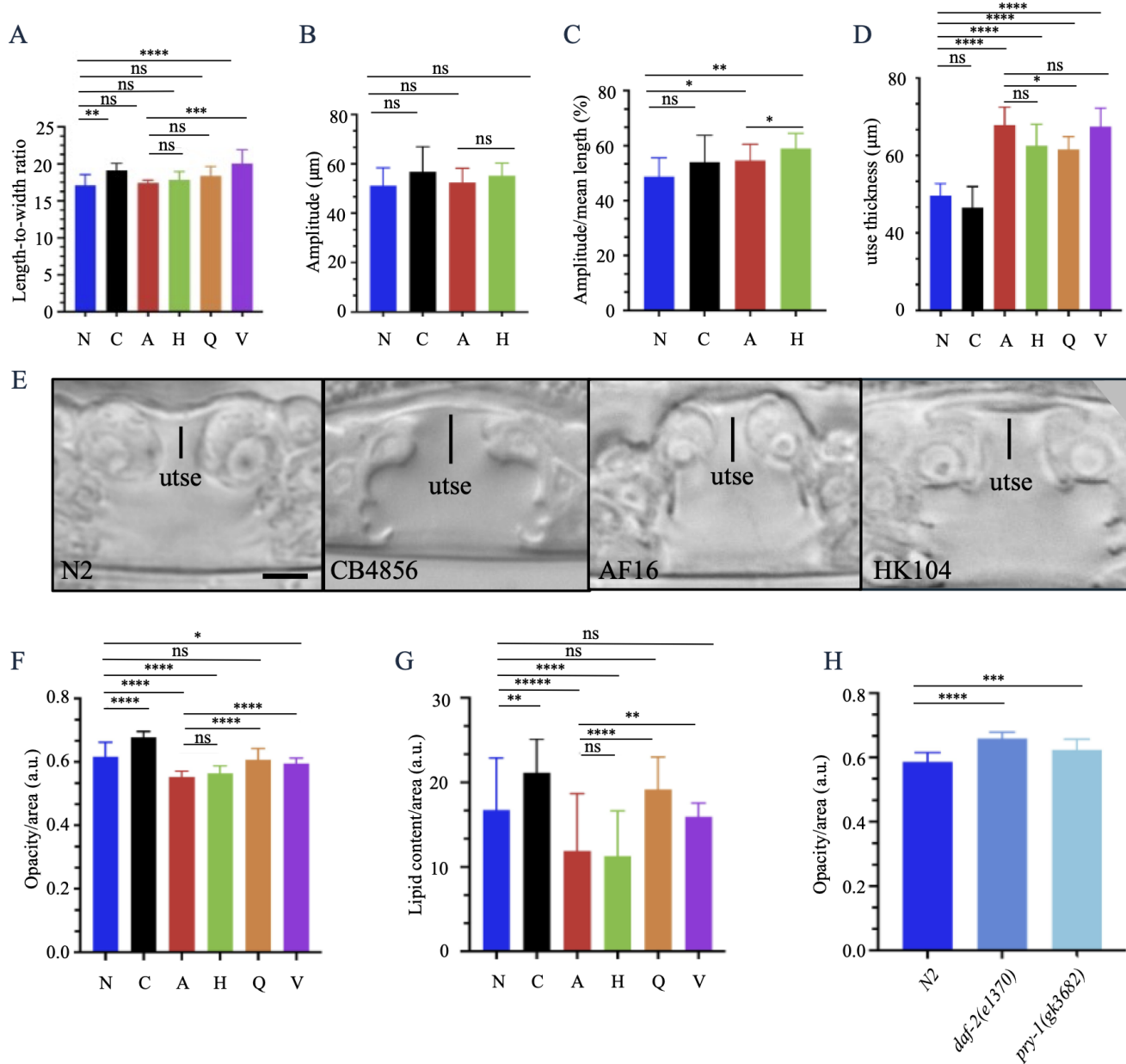
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## Abstract

The nematodes *C. elegans* and *C. briggsae* are key models for genetic studies. Despite their overall similar morphology, these two species exhibit notable differences. We used the isolates from tropical ([AF16](#) and [QX1410](#)) and temperate ([HK104](#) and [VX34](#)) regions to characterize the life history traits of *C. briggsae*. Our findings reveal significant variations in body dimensions, movement patterns, utse morphology, and lipid contents across isolates, highlighting species-level distinctions that further establish *C. briggsae* as a valuable comparative model for genetic research.



**Figure 1. Characterization of various traits in *C. briggsae* and *C. elegans* wild isolates:**

Strains are abbreviated as N (N2), C (CB4856), A (AF16), H (HK104), Q (QX1410), and V (VX34). **A.** Length-to-width ratios of day-1 adult hermaphrodites. Overall, N2 has the smallest ratio and VX34 the largest. Within *C. briggsae*, AF16 has the smallest ratio. The numbers of animals and other details are provided in Table 1. **B., C.** Movement analysis of day-1 adult hermaphrodites. *C. briggsae* strains have amplitudes comparable to *C. elegans*, however the amplitude per unit length shows some differences.  $n = 8-10$  worms for each strain in three or more batches. **D., E.** utse thickness in hermaphrodites at the L4 larval stage. *C. briggsae* isolates have thicker utse than *C. elegans*.  $n = 8 - 12$  worms for each strain, combined from two-three batches. Scale bar 5 μm. **F.** Opacity (measured as pixel brightness) of different isolates, measured in triplicates in day-1 adult hermaphrodites.  $n = 20$  to 30 worms for each strain. **G.** Oil Red O staining of day-1 adult hermaphrodites, done in triplicates with  $n = 20$  to 30 worms for each strain. **H.** Opacity of N2, *daf-2(e1370)* and *pry-1(gk3682)* day-1 adult hermaphrodites. Mutants are darker than N2.  $n = 17$  to 27 animals in a total of 3 batches for each strain. The units are arbitrary (a.u.) in panels F-H. In all graphs, data are shown as mean  $\pm$  SD. Statistical analyses were performed using one-way ANOVA with Dunnett's multiple comparisons test for interspecies comparisons in panels A-D, F, G. Student's unpaired *t*-test was used for intraspecies comparison in panels A-D, F, G, and also to compare the mutants to N2 in panel H. Statistically significant values are indicated by star (\*): \*  $p < 0.05$ ; \*\*  $p < 0.01$ ; \*\*\*  $p < 0.001$ ; \*\*\*\*  $p < 0.0001$ ; ns, not significant.



<a href="#">N2</a> hermaphrodites	1051.3 +/- 81.4	-	61.5 +/- 2.6	-	17.1+/-1.5	-	18
<a href="#">CB4856</a> hermaphrodites	1002.6 +/- 31.4	ns ( <a href="#">N2</a> h)	52.7 +/- 3.8	**** ( <a href="#">N2</a> h)	19.1+/- 1.0	** ( <a href="#">N2</a> h)	15
<a href="#">AF16</a> hermaphrodites	961.4 +/- 21.4	** ( <a href="#">N2</a> h)	55.2 +/- 1.4	* ( <a href="#">N2</a> h)	17.4+/-0.4	ns ( <a href="#">N2</a> h)	10
<a href="#">HK104</a> hermaphrodites	934.6 +/- 47.6	*** ( <a href="#">N2</a> h), ns ( <a href="#">AF16</a> h)	52.7 +/- 3.6	**** ( <a href="#">N2</a> h), ns ( <a href="#">AF16</a> h)	17.8+/-1.2	ns ( <a href="#">N2</a> h), ns ( <a href="#">AF16</a> h)	10
<a href="#">VX34</a> hermaphrodites	1076.0 +/- 93.3	ns ( <a href="#">N2</a> h), ** ( <a href="#">AF16</a> h)	54.6 +/- 2.2	** ( <a href="#">N2</a> h), ns ( <a href="#">AF16</a> h)	20.0+/- 1.9	**** ( <a href="#">N2</a> h), *** ( <a href="#">AF16</a> h)	13
<a href="#">QX1410</a> hermaphrodites	1023.0 +/- 94.2	ns ( <a href="#">N2</a> h), ns ( <a href="#">AF16</a> h)	55.7 +/- 3.6	* ( <a href="#">N2</a> h), ns ( <a href="#">AF16</a> h)	18.4 +/- 1.3	ns ( <a href="#">N2</a> h), ns ( <a href="#">AF16</a> h)	16
<a href="#">N2</a> males	968.2 +/- 20.3	-	49.9 +/- 3.0	-	-	-	10
<a href="#">AF16</a> males	795.7 +/- 40.6	**** ( <a href="#">N2</a> m)	38.5 +/- 2.6	**** ( <a href="#">N2</a> m)	-	-	10
<a href="#">HK104</a> males	842.2 +/- 43.9	**** ( <a href="#">N2</a> m), * ( <a href="#">AF16</a> m)	41.2 +/- 1.9	**** ( <a href="#">N2</a> m), * ( <a href="#">AF16</a> m)	-	-	10

## Methods

Worms were cultured on NG-Agar plates using standard methods (Brenner, 1974). Cultures were maintained at 20°C, which is an optimum temperature for growth, fecundity, and other characteristics of *C. elegans* and *C. briggsae*. Plates were seeded with *E. coli* [OP50](#) as the bacterial food source (Stiernagle, 2006). For Nomarski differential interference contrast (DIC) imaging, live animals were anesthetized with 1 mM sodium azide and mounted on 5% agar pads on glass slides. The slides were examined using Nikon Eclipse 80i and Zeiss Apotome microscopes. Images were captured using Nikon and Zeiss Zen 3.0 software. For each assay, multiple biological replicates of isolates were processed on different days.

Day-1 adult hermaphrodites were measured using Zeiss Zen 3.0 software attached to a Zeiss Nomarski microscope. L4-staged worms were picked 24 hours prior to analysis and incubated overnight at 20°C on OP50-seeded plates. Measurements of body length and width were performed on young adult hermaphrodites the following morning.

To quantify the amplitude of sinusoidal movement, individual worms were allowed to move freely on NG-Agar plates seeded with an overnight-grown [OP50](#) bacterial lawn. The distance between the peak and trough of the sine wave produced by the worm's movement was measured. The amplitude was calculated as half of this distance. At least one sine wave per worm was analyzed. Additional details on sample sizes are provided in the figure legend.

utse thickness was measured in L4 larvae of hermaphrodites, with the width determined at the center of the hymen region. Opacity (optical density) was measured in day-1 adult hermaphrodites using Nomarski microscopy on anesthetized animals. Lipid content was quantified following fixation and Oil Red O staining of day-1 adults, according to a protocol published

earlier (Ranawade et al., 2018). ImageJ (<https://imagej.net/>) software was used for image analysis. Worm outlines were traced, and pixel intensities and areas were measured to assess opacity and lipid levels.

## Reagents

Strain	Genotype	Source
<a href="#">N2</a>	Wild-type <i>C. elegans</i>	<i>Caenorhabditis</i> Genetics Center
<a href="#">CB4856</a>	Wild-type <i>C. elegans</i>	Sternberg lab
<a href="#">CB1370</a>	<i>daf-2(e1370)</i>	<i>Caenorhabditis</i> Genetics Center
<a href="#">VC3710</a>	<i>pry-1(gk3682)</i>	Gupta lab
<a href="#">AF16</a>	Wild-type <i>C. briggsae</i>	<i>Caenorhabditis</i> Genetics Center
<a href="#">HK104</a>	Wild-type <i>C. briggsae</i>	<i>Caenorhabditis</i> Genetics Center
<a href="#">VX34</a>	Wild-type <i>C. briggsae</i>	Andersen lab
<a href="#">QX1410</a>	Wild-type <i>C. briggsae</i>	Andersen lab

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## References

- Brenner S. 1974. The genetics of *Caenorhabditis elegans*. *Genetics* 77: 71-94. DOI: [10.1093/genetics/77.1.71](https://doi.org/10.1093/genetics/77.1.71)
- Delattre M, Félix MA. 2001. Polymorphism and evolution of vulval precursor cell lineages within two nematode genera, *Caenorhabditis* and *Oscheius*. *Current Biology* 11: 631-643. DOI: [10.1016/s0960-9822\(01\)00202-0](https://doi.org/10.1016/s0960-9822(01)00202-0)
- Félix MA, Ashe A, Piffaretti Jp, Wu G, Nuez I, BÉlicard T, et al., Wang. 2011. Natural and Experimental Infection of *Caenorhabditis* Nematodes by Novel Viruses Related to Nodaviruses. *PLoS Biology* 9: e1000586. DOI: [10.1371/journal.pbio.1000586](https://doi.org/10.1371/journal.pbio.1000586)
- Fitch DHA. 1997. Evolution of Male Tail Development in *Rhabditid* Nematodes Related to *Caenorhabditis elegans*. *Systematic Biology* 46: 145-179. DOI: [10.1093/sysbio/46.1.145](https://doi.org/10.1093/sysbio/46.1.145)
- Fitch DHA, Emmons SW. 1995. Variable Cell Positions and Cell Contacts Underlie Morphological Evolution of the Rays in the Male Tails of Nematodes Related to *Caenorhabditis elegans*. *Developmental Biology* 170: 564-582. DOI: [10.1006/dbio.1995.1237](https://doi.org/10.1006/dbio.1995.1237)
- Flemming AJ, Shen ZZ, Cunha A, Emmons SW, Leroi AM. 2000. Somatic polyploidization and cellular proliferation drive body size evolution in nematodes. *Proceedings of the National Academy of Sciences* 97: 5285-5290. DOI: [10.1073/pnas.97.10.5285](https://doi.org/10.1073/pnas.97.10.5285)
- Franz CJ, Zhao G, Félix MA, Wang D. 2012. Complete Genome Sequence of Le Blanc Virus, a Third *Caenorhabditis* Nematode-Infecting Virus. *Journal of Virology* 86: 11940-11940. DOI: [10.1128/JVI.02025-12](https://doi.org/10.1128/JVI.02025-12)
- Frézal L, Jung H, Tahan S, Wang D, Félix MA. 2019. Noda-Like RNA Viruses Infecting *Caenorhabditis* Nematodes: Sympatry, Diversity, and Reassortment. *Journal of Virology* 93: 10.1128/jvi.01170-19. DOI: [10.1128/JVI.01170-19](https://doi.org/10.1128/JVI.01170-19)
- Gumienny TL. 2013. TGF- $\beta$  signaling in *C. elegans*. *WormBook* : 1-34. DOI: [10.1895/wormbook.1.22.2](https://doi.org/10.1895/wormbook.1.22.2)

- Gupta BP, Sternberg PW. 2003. The draft genome sequence of the nematode *Caenorhabditis briggsae*, a companion to *C. elegans*. *Genome Biology* 4: 238. DOI: [10.1186/gb-2003-4-12-238](https://doi.org/10.1186/gb-2003-4-12-238)
- Hammerschmith EW, Woodruff GC, Moser KA, Johnson E, Phillips PC. 2022. Opposing directions of stage-specific body shape change in a close relative of *C. elegans*. *BMC Zoology* 7: 10.1186/s40850-022-00131-y. DOI: [10.1186/s40850-022-00131-y](https://doi.org/10.1186/s40850-022-00131-y)
- Inoue T, Ailion M, Poon S, Kim HK, Thomas JH, Sternberg PW. 2007. Genetic Analysis of Dauer Formation in *Caenorhabditis briggsae*. *Genetics* 177: 809-818. DOI: [10.1534/genetics.107.078857](https://doi.org/10.1534/genetics.107.078857)
- Kammenga JE, Doroszuk A, Riksen JAG, Hazendonk E, Spiridon L, Petrescu AJ, et al., Bakker. 2007. A *Caenorhabditis elegans* Wild Type Defies the Temperature-Size Rule Owing to a Single Nucleotide Polymorphism in *tra-3*. *PLoS Genetics* 3: e34. DOI: [10.1371/journal.pgen.0030034](https://doi.org/10.1371/journal.pgen.0030034)
- Maulana MI, Riksen JAG, Snoek BL, Kammenga JE, Sterken MG. 2022. The genetic architecture underlying body-size traits plasticity over different temperatures and developmental stages in *Caenorhabditis elegans*. *Heredity* 128: 313-324. DOI: [10.1038/s41437-022-00528-y](https://doi.org/10.1038/s41437-022-00528-y)
- Nyaanga J, Andersen EC. 2022. Linkage mapping reveals loci that underlie differences in *Caenorhabditis elegans* growth. *G3 Genes|Genomes|Genetics* 12: 10.1093/g3journal/jkac207. DOI: [10.1093/g3journal/jkac207](https://doi.org/10.1093/g3journal/jkac207)
- O'Rourke EJ, Soukas AA, Carr CE, Ruvkun G. 2009. *C. elegans* Major Fats Are Stored in Vesicles Distinct from Lysosome-Related Organelles. *Cell Metabolism* 10: 430-435. DOI: [10.1016/j.cmet.2009.10.002](https://doi.org/10.1016/j.cmet.2009.10.002)
- Ranawade A, Mallick A, Gupta BP. 2018. PRY-1/Axin signaling regulates lipid metabolism in *Caenorhabditis elegans*. *PLOS ONE* 13: e0206540. DOI: [10.1371/journal.pone.0206540](https://doi.org/10.1371/journal.pone.0206540)
- Rezai P, Salam S, Selvaganapathy PR, Gupta BP. 2011. Effect of pulse direct current signals on electrotactic movement of nematodes *Caenorhabditis elegans* and *Caenorhabditis briggsae*. *Biomicrofluidics* 5: 10.1063/1.3665224. DOI: [10.1063/1.3665224](https://doi.org/10.1063/1.3665224)
- Stiernagle T. 2006. Maintenance of *C. elegans*. *WormBook* : 10.1895/wormbook.1.101.1. DOI: [10.1895/wormbook.1.101.1](https://doi.org/10.1895/wormbook.1.101.1)
- Van Voorhies WA. 1996. Bergmann Size Clines: A Simple Explanation for Their Occurrence in Ectotherms. *Evolution* 50: 1259-1264. DOI: [10.1111/j.1558-5646.1996.tb02366.x](https://doi.org/10.1111/j.1558-5646.1996.tb02366.x)
- Wang X, Chamberlin HM. 2002. Multiple regulatory changes contribute to the evolution of the *Caenorhabditis lin-48* ovo gene. *Genes & Development* 16: 2345-2349. DOI: [10.1101/gad.996302](https://doi.org/10.1101/gad.996302)
- Winston WM, Sutherland M, Wright AJ, Feinberg EH, Hunter CP. 2007. *Caenorhabditis elegans* SID-2 is required for environmental RNA interference. *Proceedings of the National Academy of Sciences* 104: 10565-10570. DOI: [10.1073/pnas.0611282104](https://doi.org/10.1073/pnas.0611282104)
- Yen K, Le TT, Bansal A, Narasimhan SD, Cheng JX, Tissenbaum HA. 2010. A Comparative Study of Fat Storage Quantitation in Nematode *Caenorhabditis elegans* Using Label and Label-Free Methods. *PLoS ONE* 5: e12810. DOI: [10.1371/journal.pone.0012810](https://doi.org/10.1371/journal.pone.0012810)

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