



WHY WORMS MATTER: THE IMPORTANCE OF NEMATODE DIVERSITY FOR STUDYING BIOLOGY AND EVOLUTION

DR ERIK ANDERSEN, OF NORTHWESTERN UNIVERSITY IN THE USA, IS A MAN WITH A MISSION – TO CAPTURE THE GENETIC DIVERSITY OF *CAENORHABDITIS ELEGANS*. THIS TINY WORM IS USED BY GENETICISTS AROUND THE WORLD TO IMPROVE OUR UNDERSTANDING OF HUMAN BIOLOGY. SO FAR, SIX NOBEL PRIZES HAVE BEEN AWARDED FOR DISCOVERIES USING *C. ELEGANS*. EVEN THOUGH THIS LITTLE WORM IS ONLY THE SIZE OF A COMMA, IT HAS HAD A HUGE IMPACT ON OUR UNDERSTANDING OF BIOLOGY. MOST STUDIES USE ONLY A SINGLE STRAIN OF *C. ELEGANS*, WHICH IS LIKE STUDYING A SINGLE HUMAN TO LEARN ABOUT OUR ENTIRE SPECIES. CLEARLY, WE KNOW THAT DIVERSITY EXISTS IN OUR SPECIES, SO INCORPORATING *C. ELEGANS* DIVERSITY INTO THIS POWERFUL MODEL ORGANISM CAN ENABLE EVEN MORE DISCOVERIES. FIND OUT HOW YOU CAN HELP DR ANDERSEN ACCOMPLISH HIS MISSION AND CONTRIBUTE TO ADVANCING THE FIELD OF GENETICS

TALK LIKE A GENETICIST

NEMATODE – a diverse range of roundworms that inhabit a wide range of environments

PARASITE – an organism that lives off another organism (the host), sometimes causing damage to the host

MODEL ORGANISM – a species that is easy to grow in the laboratory and helps scientists increase our understanding of biological processes shared with humans

CAENORHABDITIS ELEGANS (*C. ELEGANS*) – a non-parasitic nematode, which is used as a model organism for genetic experiments

GENETIC DIVERSITY – the variation in the genes of individuals among a population or species

GENOME – all the genetic material belonging to an individual organism

ANTHELMINTIC DRUGS – a group of drugs that kill parasitic worms

Roundworms are nematodes, a diverse group of organisms that have adapted to nearly every ecosystem, with over 25,000 known species and likely up to a million different species found across the world. Some of these species are parasites that infect plants, livestock, pets, and even humans. Although these infections are often not deadly, nematodes do cause illness and decrease agricultural productivity.

One species of nematode, *Caenorhabditis elegans*, is one of the most genetically studied organisms on the planet because it is used as a model organism. Model organisms are non-human species that scientists can study to increase their understanding of biological processes shared with humans. *C. elegans* shares many genes with humans and can easily be grown in a lab, making it a desirable target for studying genetic traits that are relevant to humans. The importance of *C. elegans* in genetic research is exemplified by its key role in six Nobel Prize winning studies.

However, most of the research conducted using *C. elegans* is limited to a single laboratory-adapted strain that was isolated from the wild in the 1950s. This means almost all biological and genetic research from

C. elegans involves identical genetic information from a single individual organism. “This is like trying to study the whole of human biology by just looking at a single person!” explains Dr Erik Andersen.

Dr Andersen is an Associate Professor at Northwestern University, where he is building a collection that contains the genetic information of wild *C. elegans* strains. By understanding the natural variation within *C. elegans* populations, studies using *C. elegans* as a model organism can be improved, advancing our understanding of human biology.

WHY IS *C. ELEGANS* USED AS A MODEL ORGANISM?

C. elegans is a powerful model organism used to study genetics, development, cell biology, and neurobiology. “We know the locations of every cell and the connections of every neuron in *C. elegans* nematodes,” explains Dr Andersen. “Importantly, these cells and neurons are the same in every single *C. elegans* animal.”

The life cycle of *C. elegans* is only 3.5 days and they can be stored frozen then brought back to life. The worms are about 1 millimetre in size and their genome is 1/30 the size of the human



genome and so is much easier to study. All these features mean that *C. elegans* is a powerful model organism. Additionally, *C. elegans* is also a good model to study parasitic nematodes. Unlike parasitic nematodes, *C. elegans* does not require a host organism to grow, meaning it can easily be grown and experimented on in the laboratory. “Unsurprisingly, not many people will volunteer to be a host for a parasitic nematode!” says Dr Andersen.

COLLECTING WILD NEMATODES

To determine the genetic diversity of wild populations of *C. elegans*, Dr Andersen and his team collect broad and unbiased samples of nematodes from natural environments. Dr Andersen needed a location that represented the largest range of genetic diversity and so extensive fieldwork was carried out on the Hawaiian Islands. “We believe that Hawaiian *C. elegans* have more genetic diversity than the rest of the world,” explains Dr Andersen. The Pacific region, where Hawaii is located, is likely the source of *C. elegans* before they spread around the globe. This means they have a large stable population that has allowed for greater genetic variation over time.

Dr Andersen and his team collect nematodes in Hawaii every few months, allowing them to discover diverse strains (or genetically unique individuals) that may only exist in certain climates at certain times of the year. “On each collection trip, we go on hikes and pick up rotting fruits, tubers, nuts, fungus or leaf litter every 100 steps,” explains Dr Andersen. “With each sample collection, we take a picture, record the substrate temperature, ambient temperature, ambient humidity, and a variety of other environmental parameters.” The samples are mailed back to the laboratory, where the nematodes crawl off the vegetation and are collected by scientists, who place them in Petri dishes and allow them to grow.

After a couple of days, the species of each strain can be determined. Any *C. elegans*

strains will have their entire genome sequenced, allowing genetic differences in the wild strains to be identified. Each new strain is added to the genetic dataset, building a resource of global *C. elegans* genetic diversity.

A ‘CYCLE OF DISCOVERY’

Dr Andersen’s laboratory is using this resource to conduct a range of genetic research. The discoveries from *C. elegans* can inform studies in parasitic nematodes, which in turn can inform new studies on *C. elegans* in a ‘cycle of discovery’. This cycle of discovery is a powerful method because both *C. elegans* and parasitic nematode species have distinct advantages and disadvantages for their use in laboratory studies.

Dr Andersen has used a cycle of discovery to look at the genetic factors responsible for resistance to anthelmintic drugs. “Discoveries that broadly apply to nematode drug resistance would not be possible without this interplay between the two species,” says Dr Andersen. “We use the advantages of *C. elegans* genetics to discover specific genes and molecular mechanisms for drug resistance.” And by determining how natural diversity in nematodes enables them to develop resistance to drugs, Dr Andersen and his team are identifying new anthelmintic resistance genes using genetic mapping experiments.

Dr Andersen has also investigated the sensitivity of some *C. elegans* strains to chemotherapy drugs, showing that their natural genetic diversity influences the molecular mechanisms that cause differences in the responses to chemotherapy in humans. These discoveries show that genetic research with nematodes can be transformed to knowledge in human medicine.

HOW CAN YOU HELP DR ANDERSEN?

Dr Andersen’s resource of genetic information is essential for ensuring that research using *C. elegans* as a model organism reflects the



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FIELD OF RESEARCH

Genetics, Genomics, Evolutionary
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RESEARCH PROJECT

Collecting wild strains of the nematode
C. elegans and building a resource of
natural genetic diversity

FUNDERS

National Institutes of Health, National
Science Foundation, Human Frontiers
Science Program, American Cancer
Society, The Pew Charitable Trusts,
March of Dimes

true genetic diversity of the species. He hopes to increase the number of *C. elegans* strains available to the research community, saying, “We collect, organize, and disseminate these wild *C. elegans* strains and their genome data to the world so that these resources can impact the broader community.” Other researchers who work with *C. elegans* can visit Dr Andersen’s strain resource and order the strain they want to work with. Dr Andersen’s lab will send frozen nematodes of this strain so that the researcher can thaw and experiment on these strains in their own laboratory.

And you can help Dr Andersen by collecting more nematodes for his resource! When you are next on a hike, pick up some rotting material and send it to Dr Andersen. This will almost always have nematodes within it, which Dr Andersen will analyse and add to his collection. “You could add crucial new information to our natural diversity resources,” says Dr Andersen. “You might even find a new species!”

ABOUT GENETICS

Model organisms are incredibly important in biology, allowing scientists to learn about shared processes in humans. However, for model organisms to be accurate and useful, they must reflect the true genetic diversity of the population.

This is why Dr Andersen is highlighting the importance of using the natural diversity in wild populations of *C. elegans*. It is important to understand that every individual of a species has slightly different genetic information. Researchers should study this range of genetic information from wild nematodes, not only a single laboratory-adapted strain.

WHAT DOES DR ANDERSEN FIND MOST REWARDING ABOUT HIS FIELD?

Dr Andersen is excited about bringing more

genetic diversity to his field. He wants to stop the focus of experiments on a single *C. elegans* strain and for researchers to embrace the genetic diversity that is all around us. "I am excited to bring *C. elegans* natural diversity to the community that focuses on only a single strain," he says. "We have so many more discoveries to make once we incorporate these new strains."

WHAT ISSUES ARE FACING THE NEXT GENERATION OF GENETICISTS?

Dr Andersen believes the biggest challenge facing the next generation of scientists is the recent obsession by funding agencies to focus on 'translational' research, where scientific advances in the laboratory are applied to human health issues, such as for developing new methods to diagnose or treat disease. This makes funding 'basic' research, where scientists

try to answer fundamental scientific questions, much more difficult. However, Dr Andersen highlights that basic research often advances human medicine much more than more applied approaches. "From signalling pathways in cancer to genome editing to new mRNA vaccines, no one predicted that basic research would have had these huge impacts," says Dr Andersen.

PATHWAY FROM SCHOOL TO GENETICIST

- Genetics is a branch of biology, so take biology classes at school. Some universities may offer undergraduate degrees in genetics, but a general biology degree will allow you to specialise in genetics as you progress through your studies.

- Genetics covers a broad range of topics, from molecular to population to quantitative to developmental genetics. "Take as many of these courses as you can," says Dr Andersen. "Each one will teach you new aspects of inheritance and how to make discoveries using genetics."

- Computer programming and statistics skills are incredibly important. "Programming skills in R, python, and the UNIX command line will give you a huge advantage over standard biologists," says Dr Andersen.

EXPLORE A CAREER IN GENETICS

- You can visit the website for Dr Andersen's lab (www.andersenlab.org) and also visit his website where the genetic collection of wild *C. elegans* strains is organised (www.elegansvariation.org).
- Find a lab that works on a topic that excites you. Contact the lab and ask about working as a volunteer. It's never too early to start laboratory research!
- Most faculty at Dr Andersen's institution, Northwestern University, and at other universities, host high school students, undergraduates, and community volunteers to do independent research.

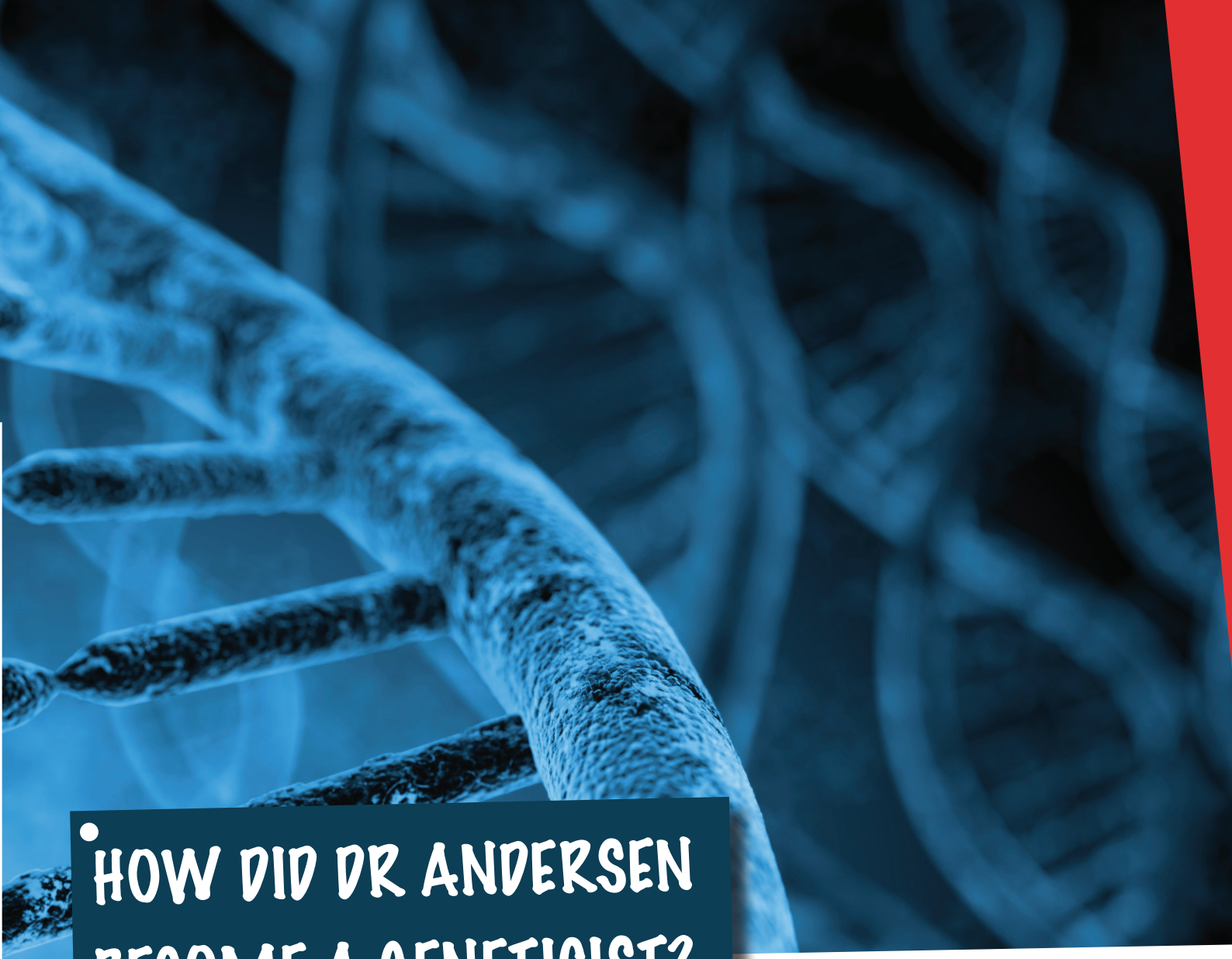
DR ANDERSEN'S TOP TIPS

01 Stay curious. Never trust what you read on the internet or in the literature. Come to your own conclusions after experiments and lots of thinking.

02 Work hard. Throughout history, advances are made by people who do not give up and commit to a problem more than other people.



Dr Andersen showing plates of nematodes in his laboratory storage room



• HOW DID DR ANDERSEN BECOME A GENETICIST?

WHAT WERE YOUR INTERESTS WHEN YOU WERE YOUNGER?

From a young age, I was interested in nature and being outdoors. I grew up around lakes and forests, which defined much of my childhood interests. Additionally, I am an avid swimmer. This interest eventually was enhanced by SCUBA diving, providing me with another way to experience nature.

WHAT INSPIRED YOU TO BECOME A SCIENTIST?

In sixth grade, my science class read small packets of current science topics weekly. In one of these classes, we read about genetically engineered corn where genes that encode a crystal toxin from bacteria were added to corn to naturally fight off pest infections. I was amazed that you could take genes from one species and put them into the other. That was the point I realised I wanted to be a geneticist. My early interest was amplified by short lab experiences working with *Drosophila* (fruit flies) and

molecular genetics at my local college while I was in high school. I chose my undergraduate school because I knew I could start doing genetics research in my first year.

WHAT ATTRIBUTES HAVE MADE YOU SUCCESSFUL AS A SCIENTIST?

My mother was an Italian immigrant who inspired me to work hard. I get up early and go to bed late, putting everything I can into my research and the people in my laboratory. Additionally, it's important for a scientist to be resilient. I am not slowed by the many difficulties and failures that I often encounter in science and in life. I strongly believe that hard work on many different topics can always yield new discoveries and make every day a learning experience.

WHAT DO YOU ENJOY OUTSIDE OF WORK?

Outside of my research, I spend time with my family, cook, swim, SCUBA, hike, and read.

WHAT ARE YOUR PROUDEST CAREER ACHIEVEMENTS SO FAR?

I am most proud of two things. First, the creation of the *C. elegans* natural diversity resource – we have brought natural diversity to the broader *C. elegans* community. I look forward to expanding this resource to other *Caenorhabditis* species. Second, the application of natural diversity to anthelmintic resistance in parasitic nematodes. We have made significant impacts on the discoveries of new resistance loci.

WHAT ARE YOUR AMBITIONS FOR THE FUTURE?

I look forward to expanding our discoveries to new genetic approaches in parasitic nematodes. With current advances in genomics and genetics, many new species can be made into model organisms relatively quickly and easily. I am excited to see what discoveries and new biology these 'models' will bring us.