

Prevalence of nematodes in wildlife and livestock animals and their control by medicinal plants

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Abstract

All animal species are infected with parasitic nematodes, which are extremely widespread and diverse. Their significance in natural ecosystems is becoming increasingly clear. Host movement has long been known to have a significant role in deciding the genetic makeup of parasitic nematode populations. More recently, studies have emphasized the significance of nematode life histories, environmental factors, and other host ecological components. Parasitic infections by nematodes continue to be a major global health concern, particularly in developing nations, despite advancements in contemporary human and veterinary treatment. Research on alternative treatments for nematode infections has increased due to mounting evidence of these nematodes' multidrug resistance and the adverse effects of currently available synthetic pharmaceuticals. In this situation, investigating possible botanical antiparasitics which are widely available and priced might be a practical substitute. In this review the prevalence of nematodes is summarized and how the infection of nematodes can be treated through plant-based drugs instead of conventional approaches.

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1. Introduction

Nematodes, cestodes, and trematodes belong to helminths that cause parasitic infections in domestic and wildlife species including small and large ruminants. Among all helminths, nematodes have significant importance due to their host-parasite interaction and their effect on the host immunity (King and Li 2018). Nematodes and other protozoans cause a complex immunological response. The infections due to nematodes are much more severe and most of the time they develop resistance due to their genetic modifications. Nematodes also have zoonotic importance because they cause many human infections but most importantly, they target domestic and wildlife species (Thompson 2023). Several wildlife species are affected by nematode infections which cause huge economic and species losses. In most cases, light infections will drop the growth and development of the animals, and heavy infections cause the death of the animals (Stear et al. 2003). Nematode infections play a significant role in determining wildlife health and biodiversity. These parasitic worms are common in many wildlife species and can affect population dynamics, affecting both individual animals and ecosystem stability. Wildlife infected with nematodes may experience reduced health, reproductive output, and

increased vulnerability to predators, which can lead to population declines, particularly in species with already small populations (Stear et al. 2023). Infection with nematodes can result in severe health consequences, including organ damage, nutritional depletion, and compromised immune function. For example, gastrointestinal nematodes in ruminants and lungworms in some wildlife species can lead to malnutrition and respiratory issues, which make animals more susceptible to other diseases and environmental stressors (Pybus et al. 2023). In ecosystems where predator-prey relationships are tightly interwoven, such health impacts can also alter predation patterns by weakening prey species or reducing their populations, which in turn affects predator species and other members of the ecosystem (Postema 2023).

The significance of nematode infections extends beyond individual hosts to influence biodiversity by potentially reducing genetic diversity within affected populations. When infection levels are high, selective pressure can lead to an increased frequency of resistant individuals, potentially limiting genetic variability. Additionally, nematode infections can create competition between host species, as some hosts become reservoirs for these parasites, spreading them to other species, thus altering interspecies dynamics (Elhady et al. 2024).

Moreover, nematodes can have indirect impacts on biodiversity. Some nematode species are zoonotic, meaning they can be transmitted to humans, especially in regions where wildlife and human populations overlap. The presence of zoonotic nematodes requires careful management to protect public health while preserving biodiversity, which adds complexity to conservation efforts. The dynamic interplay between host species, parasite load, and environmental factors underscores the importance of understanding and managing nematode infections within wildlife populations to safeguard both ecosystem health and biodiversity (Otranto and Deplazes 2019). This review is focused on various nematode species present in different animals in different geographical areas and their transmission to other animals of wildlife importance. The second part of the study described the role of medicinal plants in controlling nematode infections because they are present in every part of the world and more specifically their prevalence is higher in North America, Europe, the Middle East, and some countries of Asia.

2. Prevalence of Eustrongylides nematodes

2.1. North America

USA and Mexico were two countries where firstly the Eustrongylides nematodes were reported. The prevalence of infection in *Neogobius melanostomus* ranged from 0.5% in Michigan Lake to 95% in *Perca flavescens* in Lake Huron (Muzzall 1999). Among other fishes such as *Fundulus heteroclitus*, Eustrongylidosis was also reported. The intermediate hosts of the infection due to Eustrongylides in the waters of Mexico were reported among fishes such as *Ictalurus balsanus* and *Allophororus robustus* and the prevalence of the infection ranged from 5.8% to 19% (Quiroz-Martinez and Salgado-Maldonado 2012). The fish-eating birds of the orders Ciconiformes, *Egretta thula*, *Ardea Herodias*, and *E. caerulea* are the definitive hosts of Eustrongylides (Ziegler et al. 2000). In some snakes *Thamnophis sirtalis parietalis* the group of reptiles Eustrongylidosis was also reported. Some investigators described on biting activity of Eustrongylides nematodes in humans in the states of New York, California, New Jersey, and Maryland, but they reported no data regarding the prevalence of infection (Narr et al. 1996).

2.2. Europe

Norway, Bulgaria, the UK, Romania, Italy, Lithuania, Serbia, Poland, Ukraine, Moldova, and Russia were among the most prominent countries in Europe where the prevalence of nematodes was reported. In fish that are intermediate hosts, the prevalence of the infection varied from 0.13% (Sokolov and Moshu 2014) to about 100% in the members of *Esox tunica*, *Silurus glanis*, and *Perca fluviatilis* (Branciari et al. 2016). In Ukraine (12-100%), Moldova (up to 100%), and Russia (33.1 to 62.2%) highest prevalence of Eustrongylides was recorded in Eastern Europe. In fishes of *Acipenseridae*, *Gobiidae*, *Salmonidae*, *Percidae*, and *Cyprinidae* the Eustrongylides were most commonly found (Sloboda et al. 2010). The statements also stated the incursion of definitive hosts, for instance, Waterfowl (*Phalacrocorax carbo*) with the pervasiveness of disease from 7.14% (Svazas et al. 2011) and wild ducks up to 3.71% (Kavetska et al. 2012). Reptiles and mammals were unintended hosts for Eustrongylides (Sloboda et al. 2010).

2.3. The Middle East

Turkey and Iran were the prominent countries in the Middle East where Eustrongylides was reported. The pervasiveness of disease varies in intermediate hosts such as fish ranging from 3.5% in *Rutilus rutilus*

species to near about 100% in some species of the Sander genus such as *Sander lucioperca* (Metin et al. 2014). Among many other fishes such as *Perca fluviatilis*, *Abramis brama*, and *Esox Lucius*, Eustrongylides were found (Fallah et al. 2015). Among many accidental hosts from Amphibian, *Rana ridibunda* and *Pelophylax ridibundus* were reported (Leon Regagnon 2019). Although in the Middle East prevalence of nematodes was high there was no mentioned case of human infection in the study.

2.4. Asia

Japan, China, India, and Java were the lands where Eustrongylides were found (Honcharov et al. 2022). The infection rate varies among fishes from intermediate hosts fluctuating from 6.3% in *Hypomesus transpacificus* to 29.3% in *Synbranchus bengalensis* (Subekti et al. 2020). Recently prevalence of Eustrongylides nematodes known as *Gnathostome spinigerum* were reported in *Monopterus albus* (Asian swamp eel) (Zhang et al. 2021). Another study revealed that Eustrongylides spp. Nematodes are prevalent in various species in China, as evidenced by molecular analysis of their intra- and interspecific evolutionary variations using the COI genes and ITS rDNA regions (Xiong et al. 2013). Similarly, Eustrongylides were collected from the *Channa punctata* found in polluted rivers in India along with other ectoparasites (Biswas et al. 2023). Other than Eustrongylides various gastrointestinal nematodes were detected in wildlife species in various regions of Asia including India, Pakistan, Bangladesh, Sri Lanka, and China (Otranto and Deplazes 2019; Chandrakar et al. 2020; Ahmad et al. 2024).

3. Nematodes of terrestrial vertebrates of wildlife Animals

3.1. Parasitic nematodes of marsupials

The parasitic nematodes that transmit disease to macropod marsupials are often the topic of population genetic research. A single species of nematodes can infect multiple host species, reflecting its adaptability in host-parasite interactions and its potential impact on biodiversity and disease dynamics (Heppert et al. 2022) of different families. This is due to the variation in the genetic structure of the nematodes that they adapt according to the situation. It is not always true because some studies confirm no genetic variation or variation of some genes that does not express much, but it may be due to a smaller number of studies on parasites and their less interaction with the host. Table 1 shows different species of nematodes and their respective hosts as well as their genetic makeup.

3.2. Nematodes of terrestrial carnivores

There are various nematodes including the genera of *Ancylostoma*, *Capillaria*, *Strongyloides*, *Toxocara*, and *Trichinella* that affect the carnivorous vertebrates but the most important are *Trichinella* spp. They have wide geographic and host species ranges. Certain population genetic traits, such as differentiation between genetic variations in different populations, are shared by *Trichinella* spp. (La Rosa et al. 2012). For instance, genetic divergence was discovered between populations of *Trichinella pseudospiralis* from Australia, North America, Europe, and Asia, as well as between populations of *Trichinella spiralis* from Asia and Europe. *Trichinella* spp. frequently exhibit different genetic structures in different population that also vary with the change in geographical area (Dunams-Morel et al. 2012). Other species of *Trichinella* such as *T. nelsoni* prevalent in Kenya and Tanzania are different from the species found in South Africa (Di Cesare et al. 2014). Similar to other worms that

Table 1 Marsupial parasitic Strongylid nematodes

Nematode species	Definitive host species	Genetic markers	Reference
<i>Cloacina caenis</i>	<i>Petrogale assimilis</i> , <i>P. inornate</i>	ITS1 and ITS2	(Chilton et al. 2017)
<i>C. robertsi</i>	<i>P. assimilis</i> , <i>P. purpureicollis</i> , <i>P. persephone</i>	ITS1 & ITS2	(Song et al. 2024)
<i>C. ernabella</i>	<i>P. purpureicollis</i>	ITS1 & ITS2	(Peacock et al. 2022)
<i>Globocephaloides trifidospicularis</i>	<i>Macropus rufogriseus</i> , <i>M. giganteus</i>	24 isozymes	(Jabbar et al. 2013)
<i>G. macropodis</i>	<i>M. agilis</i> , <i>M. dorsalis</i>	ITS1 & ITS2	(Juillard and Ramp 2022)
<i>Labiostrongylus bancrofti</i>	<i>M. dorsalis</i> , <i>M. parryi</i>	18 isozymes	(Chilton et al. 2011)
<i>L. unicus</i>	<i>M. dorsalis</i> , <i>M. parryi</i>	17 isozymes	(Palomba et al. 2021)
<i>Macrostrongylus baylisi</i>	<i>M. giganteus</i> , <i>M. erubescens</i> , <i>M. robustus robustus</i> , <i>M. parryi</i>	27 isozymes	(Beveridge and Gasser 2014)
<i>Papillostrongylus labiatus</i>	<i>M. dorsalis</i> , <i>M. rufus</i>	ITS2	(Cole and Viney 2018)
<i>Paramacropostrongylus typicus</i>	<i>M. giganteus</i> , <i>M. fuliginosus</i>	37 isozymes	(Sukee et al. 2018)
<i>Rugopharynx australis</i>	<i>M. eugenii</i> , <i>Wallabia bicolor</i> , <i>Thylogale billardierii</i> , <i>M. robustus</i> , <i>M. rufus</i> , <i>M. giganteus</i> , <i>M. fuliginosus</i>	17 isozymes	(Cole and Viney 2018)
<i>R. omega</i>	<i>T. stigmatica</i> , <i>M. rufogriseus</i>	23 isozymes	(Flores et al. 2019)
<i>R. zeta</i>	<i>P. assimilis</i> , <i>M. dorsalis</i>	21 isozymes	(Akhoundi et al. 2017)
<i>Zoniolaimus mawsonae</i>	<i>M. rufus</i>	ITS2	(Coley and Viney 2018)

parasitize big carnivorous animals, *Trichinella* spp. appear to lack population genetic heterogeneity within countries and even in continents. This is due to the long-distance dispersion of hosts, which encourages substantial gene flow of parasites. Smaller hosts (like rats and foxes) may also encourage gene flow for *Trichinella* spp., allowing parasites to spread across otherwise dispersed populations of highly mobile hosts (Rothmann and de Waal 2017).

3.3. Parasitic nematodes of rodents

According to mitochondrial sequence analysis of nematode species in rodents, populations of the parasite *Heligmosomoides polygyrus*, which infects the European wood mouse (*Apodemus sylvaticus*), show significant population genetic structure throughout the host species' range. The mitochondrial mutation rate and generation time of *H. polygyrus* are quicker than those of its host. This suggests that mitochondrial genetic drift occurs more quickly in *H. polygyrus* than in *A. sylvaticus*, which contributes to the relatively stronger population genetic structure of *H. polygyrus* (Nieberding et al. 2008). Both *Trichuris* species are widespread in Europe. *T. muris* infects rats and mice, including *A. sylvaticus*, and *T. arvicolae* infects arvicoline rodents, such as lemmings and voles. Analysis of both nuclear and mitochondrial loci reveals that *T. muris* and *T. arvicolae* exhibit widespread population genetic structure throughout their geographic distribution, similar to *H. polygyrus* (Wasimuddin et al. 2016). *H. polygyrus* and both *Trichuris* species showed mostly identical patterns of population genetic structure, with a distinction between eastern and western populations and more variety in southern populations than in northern ones. The rodent hosts' range extension from southern refugia during the last ice age, at least 12,000 years ago, may be reflected in these patterns (Callejon et al. 2012).

3.4. Amphibian and reptilian parasitic nematodes

Anole lizards (*Anolis* spp.) are infected by *Spauligodon anolis*, whilst a variety of lizards and snakes are infected by the species complex *Parapharyngodon cubensis* (*P. cubensis* A, *P. cubensis* B, and *P. cubensis* C). According to a study on the population genetics of these nematodes,

which were obtained from a variety of Caribbean *Anolis* spp. hosts, genetic variation was divided across and between islands (Falk and Perkins 2013). Nevertheless, populations of *S. anolis* exhibited more genetic differentiation than populations of *P. cubensis* A or *P. cubensis* B. This is probably because *S. anolis* has a limited host species range consisting of poor dispersers. Nevertheless, populations of *S. anolis* exhibited more genetic differentiation than populations of *P. cubensis* A or *P. cubensis* B. This is probably due to the fact that *S. anolis* has a limited host species range consisting of poor dispersers (Calsbeek 2009). However, each of the *P. cubensis* complex's species uses a greater variety of hosts, some of which may be more mobile. However, even though their host range varied, cryptic species of *Spauligodon atlanticus*, parasites of *Gallotia* spp. lizards, all displayed robust genetic structures within and across the Canary Isles Islands (Jorge et al. 2011). This may be because the geographical ranges of the host species of *S. atlanticus* do not overlap, precluding nematode gene flow between them.

3.5. Parasitic nematodes of ungulates

Since ungulate (hoofed mammal) individuals travel far longer distances than rodents, their parasitic nematodes may be able to spread their genes more readily. Similar to *Teladorsagia boreoarcticus* in muskoxen (*Ovibos moschatus*), *Ostertagia gruehneri* and *Marshallagia marshalli*, parasites of reindeer (*Rangifer tarandus*), exhibit a lack of population genetic organization (Cole and Viney 2018). On the other hand, populations of the parasite *Mazamastrongylus odocoilei*, which affects white-tailed deer (*Odocoileus virginianus*), exhibited genetic structure spread to almost all continents (Long et al. 2008). A parasite of many deer species (*Cervus* spp. and *Dama* spp.) is *Dictyocaulus eckerti*. While *D. capreolus* displayed relatively lesser genetic diversity and more highly genetically structured populations when sampled sympatrically.

3.6. Parasitic nematodes of marine mammals and birds

Among the enopleane vertebrate nematodes, two families named Trichinelloidea and Dioctophymatoidea are present in coastal birds, while the remaining group of enopleane nematodes is present in aquatic organisms. Nematodes of these two families are mostly present

in every organ of the bird but most of the nematodes are found in the gastrointestinal tract, gizzard, and proventriculus of the birds. They may be found in other organs such as the heart and dermal layers but their number is not more than the gastrointestinal parasites. Even every nematode has its predilection site. The species of different systematic groups generally have their specific predilection sites. Prior to reaching the final host, the majority of nematode infections that affect marine mammals and birds spread tropically among intermediate species. Determinate host individuals will sample broadly from the parasite population since hosts of each trophic level are likely to eat several infected hosts at the lower trophic levels, causing hosts to mass parasites from a range of sources. This might result in genetically varied parasite intrapopulation that prevents inbreeding and encourages elevated N_e levels (Cole and Viney 2018). The gene flow of their parasitic nematode populations is predicted to be strong due to the long travel lengths of many marine fish, mammal, and bird hosts; this implies that these worm populations will show minimal genetic structure. A large number of parasitic nematodes that infect marine animals indeed have minimal genetic organization within their populations. A group of cryptic species with distinct geographic and definite host ranges make up the *Anisakis simplex* complex (Pontes et al. 2005).

4. Impact of nematodes on wildlife animals

In parasite ecology, one of the most common features is the aggregation of parasites within the groups of the host and a number of parasites are found in a small percentage of hosts. Individual hosts often differ in their visible levels of infection in natural populations. A wide range of variables, both internal to the host (host immune system or metabolic activities) and related to host-specific "intrinsic" (some characters that become specie specific due to acquired or hereditary factors) characteristics, contribute to this host heterogeneity in infection. The literature study revealed that different parasitic infections, particularly nematode infections, have a relationship with the age, sex, and reproduction of host animals. These have been studied in various wildlife species of deer, sheep, primates, rodents, and birds (Sol et al. 2003). It has been seen that nematodes mostly affect females compared to males but in most of the studies, it was investigated that both have equal chances of parasitic susceptibility (Lynsdale et al. 2020). The chances of nematode infections in females are more due to their estrous cycle phase, during pregnancy when immunity is decreased, and due to the lactation phase, when females bear a deficiency of micro and macronutrients. The parasitic burden of nematodes affects the overall growth and development of the host, hence leading to diseases. Similarly, young animals are more prone to infections as compared to adults. In young animals, the level of immunity is not much to combat parasitic infections and if there is any bacterial or viral infection then there will be chances of mortality of the host. During heavy infections, various chemical drugs have been used but due to their frequent use nematodes have developed resistance and their control is very necessary. Various control strategies have been used for this purpose and some of these are discussed below.

5. Conventional strategies to control parasitic nematodes of wildlife

5.1. Environmental and habitat management

Managing habitats to reduce nematode exposure is the main goal of environmental management strategies. Changing these factors (habitat

characteristics, wildlife source, climatic conditions) can interfere with the growth and transmission of many wildlife nematodes because they have intricate life cycles that call for intermediate hosts or certain environmental conditions. For instance, depending on the species, nematode larvae might live in plants, water, or soil. Rotational grazing is one practice that helps polluted land recover and prevents excessive parasite burdens. As many nematode larvae prefer damp conditions, eliminating feces and controlling soil moisture levels can work well (Bricarello et al. 2023). According to a recent study, wildlife managers can lower nematode infection rates by making changes to the environments where species exist (Mukherjee et al. 2023). For instance, exposure to certain nematodes is reduced when possible intermediate hosts, such as rodents, are removed from grazing grounds. Certain wildlife populations' parasite burden can also be decreased in wooded regions by habitat changes like vegetation removal or controlled burning (Cable et al. 2017). According to studies, environmental management practices reduce the chances of nematodal infection that lessens the need for chemical treatments and promotes ecological balance (Bouchtaoui et al. 2024).

5.2. Chemical anthelmintic treatments

One of the main strategies for controlling nematode infections in wildlife is still chemical management. Adult and larval nematodes are the targets of anthelmintics, which include ivermectin and benzimidazoles. For instance, in controlled environments, these compounds are often used in feeding or through baiting programs for wild ungulates. However, there are drawbacks to using these medications, most notably the possibility that misuse or incorrect dosage can cause parasite populations to become resistant (Gianechini 2024). Anthelmintic resistance is a growing problem in wildlife parasitology, according to scientific investigations due to accumulation of chemicals in the body and body become immune to it. To minimize misuse of these treatments and to create resistance profiles for various nematode species, checking systems have been put in place in a number of national parks (Kapinder and Verma 2023). Based on diagnostic evaluations, researchers support selective therapy strategies in which only animals with severe infections are treated. This keeps the animal population at a low infection threshold while lowering the chance of resistance (Mukherjee et al. 2023).

5.3. Biological control using predatory fungi

Natural enemies of nematodes, such as nematophagous fungus, are used in biological control techniques to capture and introduce worm larvae in the soil. After adhering to and penetrating the nematode's cuticle, the fungus release enzymes that break down the nematode's body. This method has proven effective in controlled settings, especially with cattle, and new studies are looking into how it may be used to manage wildlife (Berhanu et al. 2024). According to field research, adding these fungus spores to soil or water sources that animals use will lessen nematode infestations. Researchers are looking at using the fungus in natural wildlife habitats since it has been effective in lowering nematode numbers in livestock fields (Gianechini 2024). Nevertheless, there are still difficulties in using these fungi extensively in natural settings, and further research is required to comprehend the ecological effects on creatures that are not the intended targets (Panayotova-Pencheva 2024).

5.4. Immunization and genetic resistance

Research on immunization has grown in attention as a strategy to

manage parasitic nematodes in animals. Certain nematode vaccines have been created for household animals and have the potential to be used in wildlife. Vaccines that target *Haemonchus contortus* in sheep, for example, have been shown to drastically lower parasite burdens; similar vaccines are currently being investigated for wild ruminants (Adduci et al. 2022). The broad application of this technique is, however, constrained by practical issues, such as delivering vaccinations to wildlife that is free to roam (Rupprecht et al. 2023). For captive or semi-wild populations, selective breeding initiatives to improve genetic resistance to nematodes offer a workable alternative. According to studies, certain species are inherently resistant to particular nematodes; populations can become more resilient by selectively breeding individuals with these qualities. For instance, selective breeding has improved the general health of controlled populations of several deer species by lowering nematode loads (Panayotova-Pencheva 2024).

5.5. Integrated pest management (IPM) approach

To manage parasites sustainably, an Integrated Pest Management (IPM) strategy integrates several control techniques. This approach figures out infection levels through diagnostic monitoring and adjusts control measures accordingly. By combining biological agents, selective chemical treatments, and environmental management, IPM reduces dependency on any one technique and aids in keeping parasite populations below harmful thresholds (Singh et al. 2023). IPM techniques for animal protection include habitat management, treatment rotation, and the sporadic use of biological agents. In addition to aiding in nematode population management, this strategy supports ecological stability and biodiversity. Research has demonstrated that IPM can successfully lower the incidence of nematodes in animals without resulting in the environmental damage linked to heavy pesticide use (Abd-Elgawad 2024).

6. Problems with conventional control methods of nematodes

Nematode infestations in agriculture and wildlife can be handled using a variety of techniques, each having exclusive hazards and effects on the ecosystem. Normally used chemical anthelmintic therapies, such as ivermectin and benzimidazoles, can become the reason for environmental pollution, parasite population resistance, and detrimental effects on beneficial microbes and soil biodiversity (Navrátilová 2024). Rotational grazing and controlled burning are two environmental administration techniques that decline nematode habitats but have the potential to upset local ecosystems by affecting food webs and biodiversity. Although large-scale application may unintentionally harm non-target creatures, biological control, which targets nematodes without leaving chemical residues, provides an environmentally acceptable option by utilizing nematophagous fungus like *Duddingtonia flagrans* (Jones 2020). These techniques are used in Integrated Pest Management (IPM), which takes a more sustainable approach by employing focused tactics to lower chemical reliance, boost biodiversity, and manage parasite numbers (Deguine et al. 2021). For animal conservation initiatives, IPM's intricate coordination in natural environments is still difficult.

7. Control of nematodes by medicinal plants

Gastrointestinal (GI) parasite infections continue to be a major global health concern, particularly in impoverished nations, despite

advancements in contemporary human and veterinary treatment. Research on alternative treatments for parasitic diseases has grown as a result of mounting evidence of these parasites' multidrug resistance and the adverse effects of currently available synthetic pharmaceuticals (Harhay et al. 2010). In this situation, investigating possible botanical antiparasitics that are widely available and reasonably priced might be a viable substitute (Ranasinghe et al. 2023). Many ancient medicinal systems across the world, including Chinese (Ayurveda), Arabic (Unani), and Indian (Ayurvedic), have their roots in plants. These ancient medical systems' treatments are grounded on thousands of years' worth of empirical research, some of which has been thoroughly documented (Patwardhan 2014). Similarly, a nematode known as *Hemonchus contortus* causes severe infections in almost all wildlife ruminants and in most of the cases it causes death of the animals. Many other nematodes are acknowledged to cause severe loss to wildlife, and many control methods are suggested but with their limitations (Emery et al. 2016).

8. Use of various plants to treat nematode infections

Various botanicals of therapeutic and medicinal effects have been used over the years to treat different infections and diseases in humans and animals. They have significant importance in the veterinary field to treat bacterial, viral, fungal, and parasitic infections. These medicinal values of the plants are due to the presence of various phytochemicals in them. Various laboratory examinations have confirmed the presence of active chemical components such as polyphenols, saponins, monocarboxylic acids, alcohols, flavonoids, terpenes, sesquiterpenes etc. However, control of nematodal infections by these phytochemicals is eco-friendly and cost saving. Various studies have been carried out to check the effect of different plants on the parasites of animals and some are listed below in the Table 2.

9. Compounds of plant origin with anthelmintic action

The molecules that have been shown to have nematocidal action are members of a broad family of compounds known as secondary metabolites, which are derived from the allelopathic interactions that occur between plants and their surroundings. As the name implies, these compounds are not connected to the main metabolism of plants, hence they are not essential to their growth (Santos et al. 1999). They are created by alternate cell metabolic processes that use amino acids and shikimic acid and are stored in sections like trichomes (epidermal appendages) or cellular vacuoles, depending on their chemical makeup, frequently in cells and organs far from the site of synthesis (Wink 2008). In response to diverse stimuli, plants simultaneously create a variety of chemicals. In certain situations, the same stimulus or agent might cause the production of many substances (KhokharVoytas et al. 2023). Every part of the plant, including the leaves, stems, roots, flowers, and seeds, can create secondary metabolites, and the species, stage of development, and geographical and climatic factors all influence concentration (Luz et al. 2010).

10. Mode of action

Secondary metabolites are ideal candidates for phytotherapeutic treatments because they can be linked to defense mechanisms against infection and other plants vying for resources, and solar protection. Additionally, they can serve to distribute seeds and draw pollinators to blooms and to encourage the fixation of nitrogen (Acamovic and Brooker 2005). These chemicals can interact with many molecules in

Table 2 Different medicinal plants and their target parasites

Plant species	Effective plant part	Parasites and associated diseases	Reference
<i>Acokanthera oppositifolia</i>	Leaves	Gastrointestinal nematodes	(Sanhokwe et al. 2016)
<i>Allium cepa</i>	Cloves	G.I nematodes	(Cheraghipour et al. 2019)
<i>Allium sativum</i>	Cloves	G.I nematodes	(Soliman et al. 2023)
<i>Azadirachta indica</i>	Leaves	nematode infestation and toxemia	(Ranasinghe et al. 2023)
<i>Canabis sativa</i>	Resin	GIT nematodes	(Guerra et al. 2023)
<i>Dature metel</i>	Garden fresh fruits	GIT nematodes	(Chatterjee et al. 2022)
<i>Mentha spicata</i>	Entire plant	GIT nematodes	(El Menyiy et al. 2022)
<i>Musa paradisiaca</i>	Leaves	GIT nematodes	(Oguntibeju 2019)
<i>Nicotiana tabacum</i>	Leaves	Ectoparasites and GIT nematodes	(Nouri et al. 2016)
<i>Piper nigrum</i>	Seeds	IDyspepsia, diarrhea, flatulence, and poisoning	(Saad et al. 2022)
<i>Senna italica</i>	Entire plant	G.I parasite and ectoparasites	(Yongwa et al. 2020)
<i>Trachyspermum ammi</i>	Seeds	Ectoparasites and endoparasites	(Abbas et al. 2019)
<i>Trianthema portulacastrum</i>	Entire plant	G.I parasite	(Hussain et al. 2011)
<i>Veronia anthelmintic</i>	seeds	G.I parasite	(Dogra et al. 2020)

mammals, such as hormone receptors, neurotransmitters, and enzymes, hence it is uncommon for a cell to location that does not have any active secondary metabolites (Rodrigues and Guedes 2006). Secondary metabolites include tannins, non-protein amino acids, alkaloids, saponins, lignin, glycosides, and other polyphenols. Among these, anthelmintic properties are most closely linked to tannins (Rodrigues and Guedes 2006). It is generally recognized that some proteins can be linked to polyphenol chemicals, particularly tannins. Tannins are divided into condensed and hydrolysable categories based on their molecular structures. Tropical forage grasses often contain these compounds, which are further classified into different groups, highlighting the potent anthelmintic properties of procyanidins and prodelfinidins (Baxter et al. 1997). The rumen of ruminants is where binding of procyanidins and prodelfinidins dietary proteins, particularly those high in proline, takes place. In the abomasum, the protein/tannin macromolecular complex is broken down due to the low pH, and the proteins are broken down and taken up in the digestive system. As a result, a diet high in protein strengthens the immune system's defense against dangerous substances, including nematodes, which is an indirect effect of these metabolites on helminth

infections (Waterman et al. 2010). Because of their propensity to attach to the parasite's proteins, tannins have a direct action that alters the cuticle structure and degenerates the intestinal and muscular cells. Because of the metabolic changes brought on by the cuticle's structural breaking, these injuries may cause the worm to become less motile (Brunet et al. 2011). Destructuring the reproductive appendage of females might also hinder their ability to deliver eggs, and abnormalities in the front end can influence the parasite's feeding. The interaction of these metabolites with the L3 larvae's sheath results in another impact of tannins, which hinders their ability to penetrate the host's gastrointestinal tract by blocking their exsheathment (Hoste et al. 2012).

11. In vitro trials of the anthelmintic effectiveness of phytotherapy substances

Conducting in vitro studies of plant extracts, excluding host and environmental factors, and focusing solely on the target parasite and candidate plant, is the first stage in the process of confirming phytotherapeutic compounds. Numerous plant species have been assessed for their anthelmintic properties in this regard (Table 3). Much little research has been done on other livestock species, with the vast bulk of studies examining the impact of phytotherapeutic compounds on parasites concentrating on sheep. This study bias can be attributed to two factors: the higher pathogenicity of the parasites and the increasing frequency of reports of resistance (Borges and Borges 2016).

12. Limitations in use of plant-based medicines and future prospective

The use of plant-based medications to treat and prevent parasitic illnesses shows great promise. However, bioavailability can restrict their use. The main phytochemicals, such as flavonoids, glycosides, and tannins, are poorly soluble in water and lipids, which restricts their capacity to pass through biological membranes and causes inadequate absorption (Gao and Hu 2010). To get bioactive components, plants are also put through a variety of processes, including fermentation, distillation, purification, concentration, and extraction. The stability of active ingredients is questioned because they are subjected to oxidation

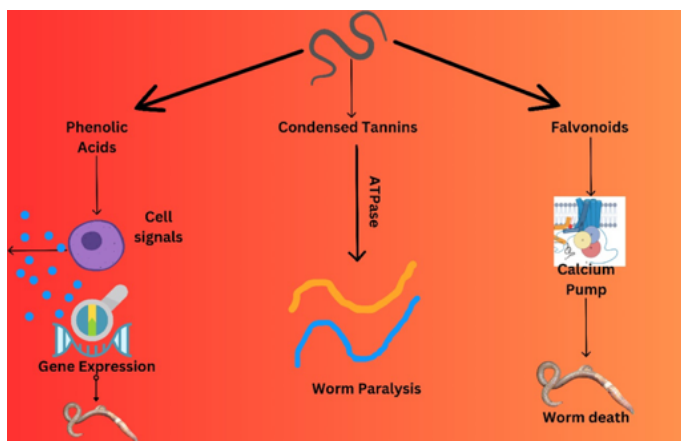


Fig. 1: Schematic representation of the mechanism of action and different pharmacological targets of plant extracts/compounds

Table 3 Different plant species and their target parasitic life stage

Plant species	Target parasite	Life stage of parasite	Reference
<i>Acacia baileyana</i>	Cyathostominae sp.	Larvae	(Payne et al. 2013)
<i>A. holorecicea</i>	Cooperi sp. & <i>Haemonchus contortus</i>	L3 larvae	(Moreno et al. 2010)
<i>A. meanrsnii</i>	<i>H. contortus</i>	Eggs and larvae	(Yoshihara et al. 2014)
<i>A. Melanoxydon</i>	Cyathostominae sp.	Larvae	(Payne et al. 2013)
<i>A. podalyrrifolia</i>	Cyathostominae sp.	Larvae	(Waterman et al. 2010)
<i>A. polycacantha</i>	<i>Caenorhabditis elegans</i>	Juveniles	(Payne et al. 2013)
<i>A. salicina</i>	<i>Cooperi</i> sp. and <i>H. placei</i>	L3 larvae	(Moreno et al. 2010)
<i>Alectryon oleifolius</i>	Cyathostominae sp.	Larvae	(Elghandour et al. 2023)
<i>Allocasuarina torulosa</i>	<i>H. placei</i>	L3 larvae	(Riley 2019)
<i>Anogeissus lelocarpus</i>	<i>C. elegans</i>	Juveniles	(Maroyi 2017)
<i>Artemisia annua</i>	<i>Bunostomum</i> sp.	Eggs and larvae	(Ekanem and Brisibe 2010)
<i>Bridelia micranta</i>	<i>C. elegans</i>	Juveniles	(Kevin et al. 2023)
<i>B. micranta</i>	<i>C. elegans</i>	Juveniles	(Waterman et al. 2010)
<i>Cambretum nigricans</i>	<i>C. elegans</i>	Juveniles	(Ignagli et al. 2024)
<i>Casuarina cunninghamiana</i>	<i>H. placei</i>	L3 larvae	(Moreno et al. 2010)
<i>Citrus sinensis</i>	<i>H. contortus</i>	Eggs and larvae	(Githiori et al. 2002)
<i>Duboisia hopwood</i>	Cyathostominae sp.	Larvae	(Payne et al. 2013)
<i>Eucalyptus gomphocephala</i>	Cyathostominae sp.	Larvae	(Ishaq 2014)
<i>Flemingia vestita</i>	Trematode and Cestoda	Adults	(Tandon et al. 1997)
<i>Grewia bicolor</i>	<i>C. elegans</i>	Juveniles	(Waterman et al. 2010)
<i>Jatropha curcas</i>	<i>H. contortus</i>	Eggs and larvae	(Monteiro et al. 2011)
<i>Markhamia obtusifolia</i>	<i>Trichostrongylus colubriformis</i>	Eggs	(Nchu et al. 2011)
<i>Melaleuca quinquenervia</i>	<i>H. contortus</i>	Eggs and larvae	(Githiori et al. 2002)
<i>Melia azedarach</i>	<i>H. contortus</i>	Eggs and larvae	(Kamaraj et al. 2010)
<i>Petoporum africanum</i>	<i>T. colubriformis</i>	Eggs and L1, L2 and L3 larvae	(Bizimenyera et al. 2006)
<i>Phillyrea latifolia</i>	<i>T. circumcincta</i>	Eggs and larvae	(Azaizeh et al. 2013)
<i>Pistacia lentiscus</i>	<i>Teladorsagia circumcincta</i>	Larvae	(Azaizeh et al. 2013)
<i>Santalum spicatum</i>	Cyathostominae sp.	Larvae	(Payne et al. 2013)
<i>Strychnos spinosa</i>	<i>C. elegans</i>	Juveniles	(Waterman et al. 2010)
<i>Tabernaemontana citrifolia</i>	<i>H. contortus</i>	Eggs, larvae, and adults	(Marie-Magdeleine et al. 2010)
<i>Ziziphus mucronate</i>	<i>C. elegans</i>	Juveniles	(Waterman et al. 2010)

and hydrolysis during these procedures (Rangari 2009). Additionally, plant products frequently deteriorate, especially when stored, which results in the loss of active ingredients and the creation of inactive metabolites (Thakur et al. 2011). Concerns about the safety of plant-based medications are becoming more prevalent as their use grows worldwide. Despite their widespread use and enticing potential, many plants have not yet been confirmed safe or poisonous. This results in a lack of awareness regarding their possible side effects and makes it challenging to determine the safest and most efficient treatments (Gao and Hu 2010).

The research discussed here supports more research into plants or plant derivatives as sources of innovative treatments for nematodal parasite infections. The studies do, however, also highlight a number of areas that need more research (Liu 2020). Future study is encouraged by the majority of the studies in literature to find the ideal dosage for maximizing the efficiency of the plants under investigation. It is also

crucial to translate the findings of in vitro research into in vivo experiments. Furthermore, to prove the new compounds' safety and effectiveness, clinical trials involving successful animal experiments using either the compounds alone or in combination with well-established antiparasitic medications are necessary.

13. Conclusion

In order to find new medications and lead compounds, this study focused on research that assessed plants and plant products as nematode antiparasitic medication. Although there are few studies on some species of nematodes affecting animals, plants or their isolates have been tested against all prevalent parasites. Given the circumstances, the outcomes of this review offer insightful data about genetic structures in Table 1 that can guide the development of future research concerning procedures, dosages, and experimental setups. According to this review, nematode parasites are significantly affected

by plants and chemicals resulting from plants both in vitro and in vivo. Broad-spectrum antiparasitic medications and many plant extracts have shown comparable benefits. Although this element needs to be examined, the traditional use of plants offers vital evidence for finding and creating synergistic medications.

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