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USE OF NON-MAMMALIAN ALTERNATIVE MODELS FOR NEUROTOXICOLOGICAL STUDY

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Abstract

The field of neurotoxicology needs to satisfy two opposing demands: the testing of a growing list of chemicals, and resource limitations and ethical concerns associated with testing using traditional mammalian species. National and international government agencies have defined a need to reduce, refine or replace mammalian species in toxicological testing with alternative testing methods and non-mammalian models. Toxicological assays using alternative animal models may relieve some of this pressure by allowing testing of more compounds while reducing expense and using fewer mammals. Recent advances in genetic technologies and the strong conservation between human and non-mammalian genomes allows for the dissection of the molecular pathways involved in neurotoxicological responses and neurological diseases using genetically tractable organisms. In this review, applications of four non-mammalian species, Zebrafish, cockroach, *Drosophila*, and *Caenorhabditis elegans*, in the investigation of neurotoxicology and neurological diseases are presented.

Introduction

There was a time when non-mammals were thought to be far from ideal materials for the study of biomedical sciences because they are phylogenically too distant from humans. However, it has now become abundantly clear that some non-mammals are not only convenient materials but also are endowed with physiological and pharmacological properties common to humans. Thus, several such species have become very popular alternative organisms and are being used extensively as models. Here we would like to present a few such examples: *Drosophila*, *Caenorhabditis elegans*, cockroach, and zebrafish. Each of them is now being used not only for genetics, biochemistry, physiology and pharmacology of the nervous system, but also for neurotoxicology. This article summarizes the symposium on “Use of Non-Mammals for

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Neurotoxicological Study” which was held as part of the 11th Meeting of the International Neurotoxicological Association in 2007.

Zebrafish are amenable to high-throughput screening in small molecule discovery and cardiac toxicology. Zebrafish small molecule screening takes advantage of the small size, chemical permeability, and optical transparency of the zebrafish embryo. Transgenic lines expressing fluorescent proteins in specific neuronal subpopulations have also been developed, which could facilitate screening. Cardiotoxicity is perhaps the most thoroughly tested zebrafish toxicity to date. Zebrafish screens have also been used to discover novel compounds that suppress the effects of genetic vascular defects.

The nematode *C. elegans*, another useful neurotoxicological model, has been used to study of Parkinson’s disease and manganism. The nematode’s nervous system is highly conserved with mammals, and contains almost all of the known signaling and neurotransmitter systems found in vertebrates. In addition, the means to screen potential neurological and developmental toxicants using *C. elegans* have been developed in a medium-throughput setting. Assays are designed to assess chemical sensitivity to specific endpoints including growth, reproduction, movement, and feeding. Several additional toxicological assays are currently under development including green fluorescent protein-based, stress-responsive transgenic *C. elegans*.

Mammalian Na⁺ channels consist of a large pore-forming α -subunit and several small auxiliary β -subunits in various tissues and cell types. In *Drosophila melanogaster*, however, the *para* appears to be the only gene that encodes a functional sodium channel. Insects employ extensive alternative splicing and RNA editing to generate many functionally diverse sodium channel variants from a single sodium channel gene. Most of these alternative splice sites are conserved in *D. virilis*, the house fly *Vssc1*, and the cockroach *BgNa_v*. The cockroach sodium channel gene *BgNa_v* undergoes extensive alternative splicing and RNA editing to produce functionally distinct sodium channel variants. Interestingly, variants BgNa_v1-1 and BgNa_v2-1 showed different sensitivities to pyrethroids. BgNa_v2-1 channel variant is 100-fold less sensitive to deltamethrin than the BgNa_v1-1 variant. This is the first example of involvement of alternative splicing of a sodium channel gene in differential sensitivity to neurotoxins. In many cases, insects and mammals have the same type of target site for an insecticide, but with differential sensitivity. It has become increasingly clear that invertebrates including insects and *C. elegans* have inhibitory glutamate-gated chloride channels (GluCl_s) which are not present in mammals and which are highly sensitive to insecticides. Because of the presence only in invertebrates, GluCl_s are a potentially important target of insecticides. At least three types of currents were recorded in response to 100 μ M glutamate: a fast-desensitizing current, a slow-desensitizing current, and a mixed type of current. Methods have recently been developed for recording them differentially. Slow-desensitizing currents could be inhibited selectively by trypsin, whereas fast-desensitizing currents were blocked selectively by soybean trypsin inhibitor or polyvinylpyrrolidone. The slow-desensitizing GluCl_s were much more sensitive to the blocking action of fipronil than the fast-desensitizing GluCl_s with IC₅₀s of 10 nM and 800 nM, respectively. Fipronil is known to be degraded to fipronil sulfone via biotic/abiotic oxidation and to a desulfinyl photoproduct via photolysis. Fipronil sulfone blocked both slow- and fast-desensitizing GluCl_s, the former being slightly more sensitive than the latter.

Potential applications for zebrafish in neurotoxicology (R.T.P.)

The zebrafish, a favorite model organism for developmental geneticists, has been shown to be amenable to high-throughput screening in applications including small molecule discovery and cardiac toxicology (Zon and Peterson, 2005) (Fig. 1). Recent developments in zebrafish small

molecule screening were reviewed to explore the possibility that the approach could be applied to neurotoxicity testing.

Zebrafish small molecule screening takes advantage of the small size, chemical permeability, and optical transparency of the zebrafish. Embryonic and larval stages of the zebrafish can be grown in 96- or 384-well assay plates, exposed to small molecules by adding the compounds to the water in the wells, and the effects can be observed in the transparent embryos using microscopy. The first small molecule screens performed employed wild-type zebrafish and visual screening to identify obvious morphological defects (Peterson et al., 2000; Sternson et al., 2001; Spring et al., 2002; Shafizadeh et al., 2004; Moon et al., 2002; Khersonsky et al., 2003). These screens identified defects in numerous organ systems including the central nervous system (CNS). Phenotypes identified in this way were generally severe, and for the CNS ranged from loss or expansion of brain ventricles to truncation of the telencephalon to severe neuronal necrosis. While these studies demonstrate the ability of this approach to identify small molecules that cause severe developmental neurotoxicities, it is doubtful that such screens could reliably identify subtle neurotoxicities that aren't manifest in obvious morphological changes. More sophisticated assays will likely be necessary. Vital dyes like acridine orange have been reported to stain apoptotic cells in zebrafish and may help detect subtle neurotoxicities (Parg et al., 2004). Transgenic lines expressing fluorescent proteins in specific neuronal subpopulations have also been developed, which could facilitate screening. Numerous functional and behavioral assays, including assays of vision, hearing, touch responsiveness, memory, anxiety, and startle habituation have been developed and could also be useful for identifying neurotoxicants that do not cause obvious developmental phenotypes (Brockerhoff et al., 1995; Bang et al., 2002; Fetcho et al., 1998; Peitsaro et al., 2003). It is possible that a panel of several high-throughput morphological and functional assays could be used to screen broadly for neurotoxicants.

Increasing the number and sophistication of high-throughput neuronal assays for zebrafish will be of little value if zebrafish and human neurotoxicities do not correlate. Much work remains to be done to determine the extent to which zebrafish toxicities are predictive, but initial data from other organ systems are encouraging. Cardiotoxicity is perhaps the most thoroughly tested zebrafish toxicity to date. In an assay for drug-induced bradycardia, 22 of 23 compounds known to cause human QT prolongation were detected among 100 tested compounds, suggesting a high degree of correlation between zebrafish and human cardiotoxicity (Milan et al., 2003). Similar types of studies focused on neurotoxicity would be very useful but have not been reported. However, some individual compounds have been reported to have predictable neurotoxicities in zebrafish, including ethanol, 6-hydroxydopamine, acrylamide, MPTP, and pentylenetetrazole (Parg et al., 2007; McKinley et al., 2005; Baraban et al., 2005).

Beyond screening for neurotoxicants, might the zebrafish high-throughput platform be useful for identifying neuroprotectants? Zebrafish screens have been used to discover novel compounds that suppress the effects of a genetic vascular defect (Peterson et al., 2004; Hong et al., 2006). Similar screens have discovered a small molecule that suppresses the effects of a mutation that causes a cell cycle defect in zebrafish (Stern et al., 2005). This approach could be applied to neuroprotection by exposing thousands of zebrafish en masse to a neurotoxicant, then screening in high-throughput for novel small molecules that block the neurotoxic effects of the toxicant. As preliminary evidence that such an approach may be feasible, several known neuroprotectants have been shown to protect zebrafish from L-hydroxyglutamic acid neurotoxicity (Parg et al., 2006), and in a separate study, l-deprenyl and nomifensine were shown to protect zebrafish from MPTP-induced neurotoxicity (McKinley et al., 2005).

Conclusions

In the search for alternative models for neurotoxicity testing, the zebrafish offers much: low cost, high throughput, an almost limitless range of morphological and functional assays, and an apparently high degree of correlation with mammalian systems. What is missing is history. Unlike many mammalian models that have been used for decades, the zebrafish cannot benefit from a large reservoir of historical data establishing the system's validity and limitations. If the zebrafish is to become useful, it will require a commitment to accumulating and sharing that reservoir, a process that hopefully could be accelerated by the ability to acquire data rapidly in zebrafish. Although such an effort would be formidable, the competing pressures for additional testing and reduced use of mammals may indicate that an investment in zebrafish is a sound one.

Toxicogenetic analysis in a novel *C. elegans* model of Parkinson's disease and manganism (R.N.)

The identification of the molecular components and mechanisms of neurodegenerative diseases have often been inhibited by the complexities of the vertebrate brain and the difficulties of modeling the diseases in cell cultures. The recent advances in genetic technologies and the high sequence similarity between human and invertebrate genomes allows for the dissection of the molecular pathways involved in neurological diseases using model organisms. Here we briefly summarize how we are using the nematode *C. elegans* to model Parkinson's disease and manganism, and describe how this worm has the potential for identifying novel proteins and compounds that may be involved in, or protect against, these neurodegenerative disorders.

C. elegans is a powerful genetic model for exploring the molecular mechanisms of neuron function and human disease (Riddle et al., 1997; Nass et al., 2001). The worm's nervous system contains almost all of the known signaling and neurotransmitter systems found in the vertebrates (Bargmann, 1998; Nass and Blakely, 2003). Its genome has been sequenced and contains approximately 20,000 genes, over 90% of the human genome. Its small size (1 mm long), large brood size, quick generation time of 3.5 days, and ease of maintenance in the laboratory allows for inexpensive production and rapid (Riddle et al., 1997). The animals can also easily be propagated in liquid medium in standard 96-well microtiter plates, allowing for high throughput screening (HTS) of animals with particular behavioral phenotypes or optical properties (Link et al., 2000; Kaletta and Hengartner, 2006). The worms are anatomically simple and well-characterized with approximately one-third of the over 1000 cells identified as neurons. The nematode is transparent and the ease of making transgenic animals containing reporter constructs greatly facilitates examination of neuron protein expression and localization (Mello and Fire, 1995; Thomas and Lockery, 1999). Primary cultures, first developed by Laird Bloom in 1993 during his MIT Ph.D thesis, differentiate and appear to retain many of their *in vivo* cellular properties (Bloom, 1993; Buechner et al. 1999).

Significant advantages of using *C. elegans* for genetic analysis is the opportunity to incorporate forward and reverse genetic screens to relatively quickly identify proteins and molecular pathways that are involved in particular cellular processes. Forward genetic screens using chemical mutagens have the advantage that no *a priori* knowledge is needed to determine a gene's function in order to determine whether it plays a role in the particular phenotype. Mutations can be identified within as little as a week (Wicks et al., 2001), and since the worms can easily be mated with each other, strains containing several mutations can be generated within days (Wicks et al., 2001). *C. elegans* is also amenable to reverse genetics that allows for the identification of molecular pathways in which a particular gene acts upon. RNA interference, or RNAi, where the introduction of dsRNA induces sequence specific degradation of homologous mRNAs and subsequent protein expression, is one example of a reverse genetic

approach. The 2006 Nobel Prize in Physiology or Medicine was awarded to Andrew Fire and Craig Mello for their RNAi efforts.

Parkinson's disease (PD) is the second most prevalent neurodegenerative disease and results from the loss of at least 80% of the DA neurons within the substantia nigra pars compacta (SNpc). A hallmark of the disorder is the formation of protein aggregates termed Lewy bodies (LBs) in a significant number of the surviving neurons. Although the molecular basis of this disorder has not been identified, etiological and pathological data suggest that there is both a genetic and environmental component that causes and contributes to the DA neuron cell death (Jenner, 1998). Environmental exposure to manganese, as well as several other metals, has been associated with the development of PD. Mn-induced Parkinsonism, also called manganism, has been associated with Mn mining and welding (Calne et al., 1994; Chia et al., 1993). Acute Mn²⁺ toxicity results in symptoms similar to those seen in patients with PD, including rigidity, tremors, and bradykinesia (Stredrick et al., 2004).

One of the most common mechanisms to model Parkinson's disease in vertebrates is through exposure of the animals to a DA neuron neurotoxin. Within weeks following toxin exposure, most of the animals display a Parkinsonian-like syndrome. The most common neurotoxins used are 6-hydroxydopamine (6-OHDA), 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine (MPTP), 1-methyl-4-phenylpyridinium ion (MPP⁺ [the active metabolite of MPTP]). 6-OHDA and MPP⁺ are transported into the cell by the high affinity dopamine transporter, DAT, which is also the target of drugs of abuse such as cocaine and amphetamine (Reading, 1994).

We generated the first *C. elegans* PD model by developing a *C. elegans* transgenic line that expresses GFP in all eight DA neurons within the hermaphrodite (Nass et al., 2002). When we briefly exposed the animals to 6-OHDA (0.5–1 hr), we find a time and concentration-dependent loss of DA neuron integrity. Also, this affect can be blocked by co-incubation with DAT agonist (e.g. amphetamine) or antagonist (e.g. cocaine), consistent with these vertebrate models of PD that DAT is required (Nass et al., 2002). These exciting results mimic several significant aspects of the vertebrate PD model, and should allow us to utilize the power of invertebrate genetics to explore *in vivo* the function of genes previously identified to play a role in PD, as well as identify potential environmental components such as Mn²⁺ exposure and novel endogenous molecules that could contribute to the DA neuron cell death.

For example, epidemiological studies indicate that the etiology of PD likely involves specific molecular pathways involved in protein aggregation and degradation. To date, at least seven genes have been independently identified that are associated with rare, familial forms of PD, including α -synuclein, parkin, DJ-1, UCH-1, NURR1, PINK-1, and LRRK2. Most of these proteins have strong homologues in the worm. α -synuclein, the first gene identified, is a presynaptic protein that appears to interact with synaptic vesicles and could be involved in the regulation of both dopamine biosynthesis and dopamine transporter function, and is also a major component of the PD associated protein aggregates LBs. α -synuclein aggregation is also dramatically accelerated *in vitro* in the presence of the transition metal Mn²⁺ (Uversky et al., 2001). Mn²⁺ exposure to DA and α -synuclein expressing mammalian cells also cause greater cell death than those not exposed to Mn²⁺ (Pifl et al., 2004). These examples suggest that Mn²⁺ exposure could be an aggravating factor for triggering DA neuron cell death in PD.

We have also generated transgenic animals overexpressing either human wild-type (WT) or mutant A53T α -synuclein and gfp in the DA neurons within the worm. We have found that both WT and A53T expression confer DA neuron cell death in the worm even in the absence 6-OHDA (Lakso et al., 2003). Motor deficits were also observed when α -synuclein was expressed behind the pan-neuronal promoter, and α -synuclein containing inclusion bodies are seen in some of the DA neurons. Our preliminary results also indicate that the DA neurons are

sensitive to a number of heavy metals, and that some of these metals amplify the effects of 6-OHDA and α -synuclein *in vivo*. These observations suggest that *C.elegans* could be a useful model for α -synuclein-induced pathologies, and help to identify the relationship between α -synuclein, heavy metals, and PD-associated proteins.

The establishment and initial characterization of our *C. elegans* PD model also provides an opportunity to utilize genetic screens to identify novel genes involved in DA neuron cell death. For example, in one forward genetic screen we could utilize our DAT-1::GFP reporter line to identify molecules involved in Mn^{2+} -induced neuronal death. We would first mutagenize the genome of the parental animals and then isolate the second generation hermaphrodites (to homozygose the mutation; Nass et al., 2003) in which GFP is still retained in the DA neurons following exposure to Mn^{2+} . Animals that have DA neurons that are insensitive to Mn^{2+} could have mutations within Mn^{2+} transport proteins or proteins involved in DA neuron viability or cell death. We have implemented a similar genetic screen with 6-OHDA and have identified a number of mutants that have varying degrees of DA neuron 6-OHDA insensitivity (Nass et al., 2005). Three of these mutants contain mutations within DAT that render them completely resistant to 6-OHDA. The identification of these mutants provides proof-of-concept that we should be able to isolate genes involved in toxin-induced cell death.

With our worm PD model we should also be able to identify novel genes involved in Mn^{2+} -induced DA neuron sensitivity by utilizing the worm friendly methods of RNA interference (RNAi). The Medical Research Council (Cambridge) has generated a remarkable library of bacteria that express RNAi molecules against roughly 90% of the known worm genes (~18,000 genes). Since feeding the bacteria expressing the dsRNA to the worms is an efficient way to knockdown the expression of a gene (Timmons et al., 2001), we could envision feeding our gfp reporter line to the bacteria also on medium containing Mn^{2+} . We would then select for animals in which the DA neurons are not affected by the Mn^{2+} , and therefore could quickly map and identify genes are involved in Mn^{2+} -induced DA neuron sensitivity.

Conclusions

C. elegans provides remarkable opportunities to identify and characterize genes and proteins involved in Parkinson's disease and manganism *in vivo*. The high similarities on the molecular level between the worm and humans suggest that the paradigms discovered using this system are highly relevant to these devastating diseases. *C.elegans* can also be utilized to screen and identify environmental agents such as heavy metals that could cause or contribute to susceptibility to disease.

Toxicological studies of environmental agents using *C. elegans* (W.A.B. and J.H.F.)

The National Toxicology Program (NTP) is responsible for the development of sound scientific tests designed to estimate the effects of chemicals on human health. National and international government agencies, such as the NTP and the Environmental Protection Agency (EPA), have defined a need to reduce, refine or replace vertebrate animals in toxicological testing with alternative testing methods and models (Becker et al. 2006). Toxicological assays using invertebrate species are more rapid and less expensive than traditional mammalian-based tests due in part to shorter life spans and the ability to assay in multi-well plate formats. Invertebrate species are also ideal model organisms because of the lack of animal welfare concerns.

C. elegans has been recognized as an attractive biological and genetic model organism for some time (Brenner 1974). Recently, these advantages have led to a rise in the use of *C. elegans* as a toxicity test organism. Short life cycles, easy and inexpensive maintenance, and

detailed biological knowledge allow for the development of rapid, low-cost tests that readily lend themselves to mechanistic studies of toxicant action. There is also a high degree of conservation in the molecular toxicological responses between *C. elegans* and mammals. For example, many signal transduction pathways involved in general stress responses are well conserved (National Research Council, 2000). Several studies have also demonstrated the predictive potential of *C. elegans* lethality and changes in locomotion for mammalian toxicity (Cole et al. 2004; Tataru et al. 1998). With advances in technology, the assessment of phenotypes of thousands of nematodes can now be quantified in a high-throughput fashion, rather than by direct observation of only a few organisms.

Our group has developed the means to screen potential neurological and developmental toxicants using *C. elegans*. To this point, sublethal toxicological assays have been automated using liquid handling robotic workstations, a Complex Object Parametric Analyzer and Sorter (COPAS) Biosort (Pulak 2006), which is used to dispense and analyze nematode length and fluorescence, and an imaging workstation for motion tracking and multidimensional image analyses. To optimize the toxicological assays, a 96-well plate format is used for sample preparation, dispensing of nematodes, and quantification of specific toxicological endpoints. As nematodes are dispensed, the COPAS Biosort measures the time of flight (TOF) or length, extinction (EXT) or optical density, and green and red fluorescence of each nematode.

C. elegans matures from fertilized egg to adult through four distinct larval stages, termed L1-L4, in approximately 3.5 days and has an average life span of 10 days. Fig. 2 illustrates the size distributions of *C. elegans* as they developed from the first larval stage (L1) to adults over 72 h, as measured by the COPAS Biosort. After 48 h, the nematode population was observed by microscopy to be mostly L4s. After 72 h, nematodes were either gravid adults, with the highest TOF and EXT, or the second generation of L1s and eggs, with the lowest TOF and EXT. As the nematodes grow and develop, they increase in length and optical density.

Three toxicological assays have been developed in order to assess chemical sensitivity at specific developmental stages: L1s for growth, L4s for reproduction, and adults for movement and feeding. In experiments similar to the one shown in Fig. 2, the *C. elegans* growth assay was used to measure the TOF and EXT from L1s at the beginning of the assay and then after a 72 h exposure to the test toxicant. The *C. elegans* reproduction assay was developed to assess the effects of chemicals at the L4 stage, just before nematodes are reproductive. After 48 h exposures, the COPAS Biosort was used to count the number of offspring. The *C. elegans* feeding assay was developed to monitor neurotoxicity in adult nematodes exposed to toxicants for 24 h (Boyd et al., 2007). After exposures, adult nematodes are allowed to feed on 0.5 μm red fluorescent microspheres and the level of red fluorescence in individual nematodes is measured using the COPAS Biosort. Feeding rates of exposed and non-exposed control *C. elegans* can then be calculated. Fig. 3 presents a typical dose-response curve showing the effects on feeding after exposing *C. elegans* to increasing concentrations of toxicant. In this study, nematodes were exposed to the organophosphate pesticide chlorpyrifos. In all assays, dose-response curves are calculated to estimate toxicological endpoints such as EC_{50} s and benchmark doses. Our group has validated the growth, reproduction, and feeding assays by screening almost 60 chemicals, which include metals, pesticides, mutagens, and non-toxic agents (Table 1).

In order to increase the throughput of the assays, the reproduction assay has been modified and used to screen a library of 1408 chemicals selected by the NTP. Results from the *C. elegans* reproduction assay will then be compared to those obtained by the NIH Chemical Genomics Center, which performs cell-based high-throughput screens (Feng et al. 2007). All of the chemicals that cause a significant effect on reproduction will be entered into a second tier of testing. In the second tier, detailed concentration responses will be assayed.

Several additional toxicological assays are currently under development. These include the development of green fluorescent protein-based, stress-responsive transgenic *C. elegans*, which will be used to improve the sensitivity and specificity of toxicity screens and semi-automated motion tracking (Fig. 4). Transgenic strains are being constructed that will be used to monitor several toxicological processes including metal response, biotransformation, apoptosis, and DNA damage response. After creating the transgenic strains, the effects of toxicant exposure on GFP expression will be measured using a COPAS Biosort or a plate-reading fluorometer. Semiautomated motion tracking is being developed to monitor the effects of potential neurotoxicants on *C. elegans* locomotion. Many parameters are simultaneously calculated from individual nematode tracks including rate of movement, number of reversals, and sinuosity.

In addition to the technological deployment of medium and high throughput *C. elegans* toxicological assays, our group is continuing to develop statistical tools that will allow us to properly analyze the large volume of data that is produced in each of these assays. Several of the tools currently under development include multi-dimensional (e.g. TOF, EXT, GFP, etc.) modeling to assess changes in *C. elegans* populations after exposure to chemicals. Statistical algorithms that can be used to classify nematodes into discrete growth stages and to characterize the statistical properties of nematode populations at different developmental stages are also being developed. These new statistical tools will be used to quantify changes in *C. elegans* biology in response to toxicant exposure.

Conclusions

In response to the need to develop inexpensive and rapid toxicological assays, and to decrease the numbers of vertebrate animals used in the current assays, our group has explored the use of *C. elegans* as a test organism and developed several toxicological assays. We have found that *C. elegans* will make a valid alternative test organism for the screening of toxic agents.

Differential sensitivity of insect sodium channel variants to pyrethroid insecticides (K.D.)

Sodium channels are responsible for the initiation and propagation of action potentials in almost all excitable cells. Because of their crucial role in membrane excitability, sodium channels are targeted by a great variety of naturally occurring neurotoxins, such as tetrodotoxin, scorpion toxins, and batrachotoxin, which are produced by plants and animals for defense or predation (Cestele and Catterall, 2000; Wang and Wang, 2003). Sodium channels are also the primary target of the synthetic insecticides pyrethroids, which are structural derivatives of pyrethrins found in extracts of the flowers of *Chrysanthemum* species (Narahashi, 1988). Different classes of neurotoxins bind to different sites on the sodium channels and alter channel functions. For example, pyrethroids cause prolonged opening of sodium channels primarily by inhibiting channel deactivation and inactivation and stabilizing the open configuration of the sodium channel (for review see Narahashi, 1988; 1996; 2000; Soderlund and Bloomquist, 1989; Raymond-Delpech et al., 2005).

Mammalian sodium channels, encoded by nine distinct genes, are known to exhibit different sensitivities to pyrethroid insecticides. For example, tetrodotoxin (TTX)-sensitive sodium channels in the rat dorsal root ganglion neurons are less sensitive to pyrethroids than TTX-resistant sodium channels in the same neurons (Ginsburg and Narahashi, 1993; Tatebayashi and Narahashi, 1994; Song and Narahashi, 1996; Tabarean and Narahashi, 1998). Rat Nav1.2 and Na1.4 channels are resistant to pyrethroids (Warmke et al., 1997; Vais et al, 1997; Smith and Soderlund, 1998; Wang et al., 2001), whereas rat Nav1.8 sodium channel is highly sensitive

(Choi and Soderlund et al., 2006). However, the molecular basis of the differential sensitivities to pyrethroids remains unknown.

In contrast to multiple sodium channel genes in mammals, *Drosophila melanogaster* and other insects appears to have only a single sodium channel gene, *para* in *Drosophila* and *BgNa_v* in *Blattella germanica* (Warmke et al., 1997; Dong, 2007). However, electrophysiological studies showed that the sensitivity of the insect nervous system to pyrethroids varies greatly depending on nerve preparations, suggesting the presence of distinct subtypes of sodium channels (References in Dong, 2007). For example, permethrin affects the insect sensory neurons more profoundly than the neuromuscular synapses (Osborne and Hart, 1979). However, the molecular basis of the different sensitivities of insect sodium channels to pyrethroids was not known until recently. We found that the cockroach sodium channel gene *BgNa_v*, undergoes extensive alternative splicing and RNA editing to produce at least twenty functionally distinct sodium channel variants (Tan et al., 2002; Song et al., 2004). This observation is consistent with what was found in the *D. melanogaster para* and housefly *Vssc1* sodium channel transcripts. Most of these alternative splice sites are conserved in *D. virilis* (Thackeray and Ganetzky, 1995), the house fly *Vssc1* gene (Lee et al, 2002), and the cockroach *BgNa_v* gene (Tan et al., 2002; Song et al., 2004). Interestingly, cockroach variants *BgNa_v1-1* and *BgNa_v2-1* showed greatly different sensitivities to pyrethroids: The *BgNa_v2-1* channel variant is 100-fold less sensitive to deltamethrin than the *BgNa_v1-1* variant (Tan et al., 2002).

Like mammalian α -subunits, the insect sodium channel contains four repeated homologous domains (I–IV), each having six membrane-spanning segments (S1–6) connected by intracellular or extracellular loops of amino acid sequences. As shown in Fig. 5, two mutually exclusive exons G1/G2 encode IIIS3-IIIS4. The *BgNa_v2-1* channel contains exon G2, whereas the *BgNa_v1-1* channel contains exon G1. When the G1/G2 exons were swapped between *BgNa_v1-1* and *BgNa_v2-1*, recombinant *BgNa_v1-1* channel carrying exon G2 (*BgNa_v1-1^{G2}*) was 10-fold more resistant to deltamethrin than *BgNa_v1-1*; whereas recombinant *BgNa_v2-1* channel carrying exon G1 (*BgNa_v2-1^{G1}*) was 10-fold more sensitive to deltamethrin than *BgNa_v2-1* (Tan et al., 2002). Therefore, the different pyrethroid sensitivities between *BgNa_v1-1* and *BgNa_v2-1* channels can be partially attributed to the presence of exon G1 or G2. There are 14 amino acid differences between exons G1 and G2 (Fig. 5). To identify the residue(s) in exons G1/G2 that are responsible for the different deltamethrin sensitivities, we made single, double, or triple amino acid substitutions (depending on whether the amino acid differences are adjacent to each other or not) in *BgNa_v1-1* or *BgNa_v2-1*. We found that a V to A change at the amino acid position 1356 (V1356A) in exon G2 was responsible for the exon G2-associated low sensitivity to pyrethroids (Du et al., 2006). This is the first example of involvement of alternative splicing of a sodium channel gene in differential sensitivity to neurotoxins. The G1/G2 exons are also conserved in *D. melanogaster*, which allows us to use powerful tools and resources for further genetic manipulation.

Another excellent example of using insect sodium channels to study the molecular action of pyrethroids on sodium channels is the study of naturally occurring sodium channel mutations that confer knockdown resistant (*kdr*) to pyrethroids (Soderlund, 2005; Dong, 2007). Specifically, a large number of single amino acid mutations were found in the sodium channels of *kdr* mutant insects, many of which were later confirmed to confer reduced pyrethroid-sensitivity to sodium channels expressed in *Xenopus* oocytes. Because these mutations also confer insecticide resistance at the whole insect level, the biological relevance can be established beyond *in vitro* or cell culture systems. A similar combination of approaches should prove useful in using insects as models for the study of other classes of neurotoxins that target sodium channels.

Conclusions

Insect models have proven to be excellent in advancing our understanding of the molecular action of pyrethroids, which act on both mammalian and insect sodium channels. The pyrethroid-sodium channel research illustrates the great potential of using naturally occurring neurotoxin-resistant insects or target variants as excellent tools for understanding the mechanisms underlying mammalian neurotoxicological processes involving the same or similar insect targets.

Glutamate-activated chloride channel: unique chemical target present in insects but not in mammals (T.N.)

Most insecticides are more toxic to insects than to mammals, and this is one of the important characteristics for a chemical to become a useful insecticide. In many cases, insects and mammals have the same type of target site for an insecticide, but with differential sensitivity. For example, the major target site of pyrethroids is the voltage-gated sodium channels in both insects and mammals, but the insect sodium channel is much more sensitive to pyrethroids than the mammalian sodium channel with a difference in EC_{50} of ~1000-fold (Ginsburg and Narahashi, 1993; Narahashi, 2001; Narahashi et al., 2007). However, in some other cases, an insecticide affects the same insect and mammalian target site almost equally. An example is endosulfan which blocks the GABA receptors of both cockroach neurons and rat dorsal root ganglion neurons with almost the equal potency, with IC_{50} s of 5 nM and 10 nM, respectively (Zhao et al., 2007).

It has become increasingly clear that invertebrates including insects and *C. elegans* have inhibitory glutamate-gated chloride channels (GluCl_s) which are not present in mammals and which are highly sensitive to insecticides. Whereas glutamatergic synaptic transmission in vertebrate is excitatory mediated by glutamate-activated cation channels, glutamate serves as both an excitatory (Gration et al., 1979; Patlak et al., 1979; Ultsch et al., 1992) and an inhibitory (Cleland, 1996) transmitter in invertebrates. Inhibitory glutamatergic synaptic transmission is mediated by glutamate-gated chloride channels (Cleland, 1996; Raymond and Sattelle, 2002) that are present only in invertebrates.

Because of the presence only in invertebrates, GluCl_s are a potentially important target of insecticides. Many studies of GluCl_s were performed with *C. elegans*, particularly as a target site of the anthelmintic/insecticide ivermectin (Hejmadi et al., 2000; Burkhart, 2000; Arena et al., 1995). Studies of GluCl_s using insects in connection with ivermectin were limited (Kane et al., 2000), but increasing attentions are recently focused on insect GluCl_s (Ikeda et al., 2003; Zhao et al., 2004a,b; Ihara et al., 2005; Eguchi et al., 2006; Janssen et al., 2007).

We present the physiological and toxicological characteristics of GluCl_s. Whole-cell patch clamp experiments were performed using cockroach thoracic ganglion neurons. Neurons were isolated as described in our previous paper (Zhao et al., 2004a).

At least three types of currents could be recorded in response to glutamate application (Zhao et al., 2004b; Narahashi et al., 2007). One was a fast-desensitizing current (Fig. 6A). Some neurons generated a slow-desensitizing current (Fig. 6B). Other neurons produced a mixed type of current (Fig. 6C). As will be shown later, since fast- and slow-desensitizing currents exhibit different characteristics and different pharmacological responses, it is often necessary to analyze them separately. We have recently developed methods for recording them differentially (Zhao et al., 2007). Slow-desensitizing currents could be inhibited selectively by trypsin, whereas fast-desensitizing currents were blocked selectively by soybean trypsin inhibitor or polyvinylpyrrolidone. These selective inhibitors now make separate recordings of two types of currents much easier than before.

These glutamate-induced currents have been shown to be carried by chloride ions (Zhao et al., 2004b). With symmetrical chloride concentrations across the membrane, the reversal potential was estimated to be +2.5 mV for both fast- and slow-desensitizing currents. This value is very close to the calculated chloride equilibrium potential. When the external chloride concentration was reduced to a quarter, the reversal potential was shifted to +37.5 mV. This amount of shift agreed with the calculated shift of reversal potential. Thus, it is concluded that GluCl_s are carried by chloride ions.

Since cockroach GABA receptors and glutamate receptors are both accompanied by chloride channels, a question arises as to whether these two receptors are different entities. Several types of experiments have proven that this is indeed the case (Zhao et al., 2004b).

Fipronil is known to block GABA receptors thereby causing hyperexcitation of insects and mammals (Ikeda et al., 2001; Zhao et al., 2003). Fipronil has also been found to be a potent blocker of GluCl_s (Zhao et al., 2004a, b; 2005). The slow-desensitizing GluCl_s were much more sensitive to the blocking action of fipronil than the fast-desensitizing GluCl_s with IC₅₀s of 10 nM and 800 nM, respectively (Fig. 7) (Zhao et al., 2004a; Narahashi et al., 2007).

Fipronil is known to be degraded to fipronil sulfone via biotic/abiotic oxidation and to a desulfinyl photoproduct via photolysis (Bobe et al., 1998; Ngim et al., 2000). The biological conversion of fipronil to fipronil sulfone was blocked by piperonyl butoxide (Hainzl et al., 1998; Caboni et al., 2003). However, it was unclear whether the conversion of fipronil to its sulfone represented detoxication in mammals and insects. Fipronil sulfone blocked both slow- and fast-desensitizing GluCl_s (Zhao et al., 2005). The IC₅₀ for slow- and fast-desensitizing currents were calculated to be 8.5 ± 0.4 nM (n = 4) and 25 ± 2 nM (n = 4), respectively. Thus, fipronil sulfone plays an important role in the toxicity caused by fipronil.

The blocking action of fipronil in cockroach neuron GluCl_s is much more potent than that in mammalian neurons. The IC₅₀ values are: 10 nM in cockroach slow GluCl_s, 800 nM in cockroach fast GluCl_s, 30 nM in cockroach GABAergic chloride channels, and 1600 nM in rat GABAergic chloride channels. Thus, the high sensitivity of cockroach GluCl_s and GABAergic chloride channels plays a crucial role in selective fipronil toxicity in insects over mammals. Insect GluCl_s will become an excellent target site for development of new insecticides with a high degree of selectivity over mammals.

Conclusions

Glutamate-activated chloride channels, which are present in insects but not in mammals, are crucial target site responsible for the selective toxicity to insects over mammals for certain insecticides. Therefore, GluCl_s are a unique and important site for developments of new insecticides. This represents an excellent example in which non-mammals are irreplaceable materials for the study of neurotoxicology.

Overall Conclusions

Certain non-mammalian preparations have been shown to be excellent materials for the study of neurotoxicology. Recent advances in genetic techniques and the strong conservation between human and non-mammalian genomes make it possible to dissect out the molecular pathways involved in neurotoxicological responses using genetically tractable organisms. In many cases, toxicant's target receptors are similar between mammals and non-mammals. However, in some other cases, non-mammals have a target receptor that is not present in mammals; this makes the receptor important for the selective toxicity of insecticides. Low costs and easiness for maintaining non-mammals in the laboratory are another great advantage.

Zebrafish offers several advantages: low costs, a variety of morphological and functional assays, and a high degree of correlation with mammalian systems. *C. elegans* provides remarkable opportunities to identify and characterize genes and proteins involved in Parkinson's disease. It can also be utilized to screen and identify environmental agents and to develop several toxicological assays. An example is also given in which molecular target sites of sodium channels for pyrethroid insecticides can be analyzed using *Drosophila* and cockroach. Cockroach neurons also represent a unique material for the electrophysiological analysis of selective toxicity as they are endowed with glutamate-activated chloride channels which are highly sensitive targets of fipronil and which are present in insects but not in mammals. It is expected that use of non-mammals for a variety of neurotoxicological studies will become more important in the coming

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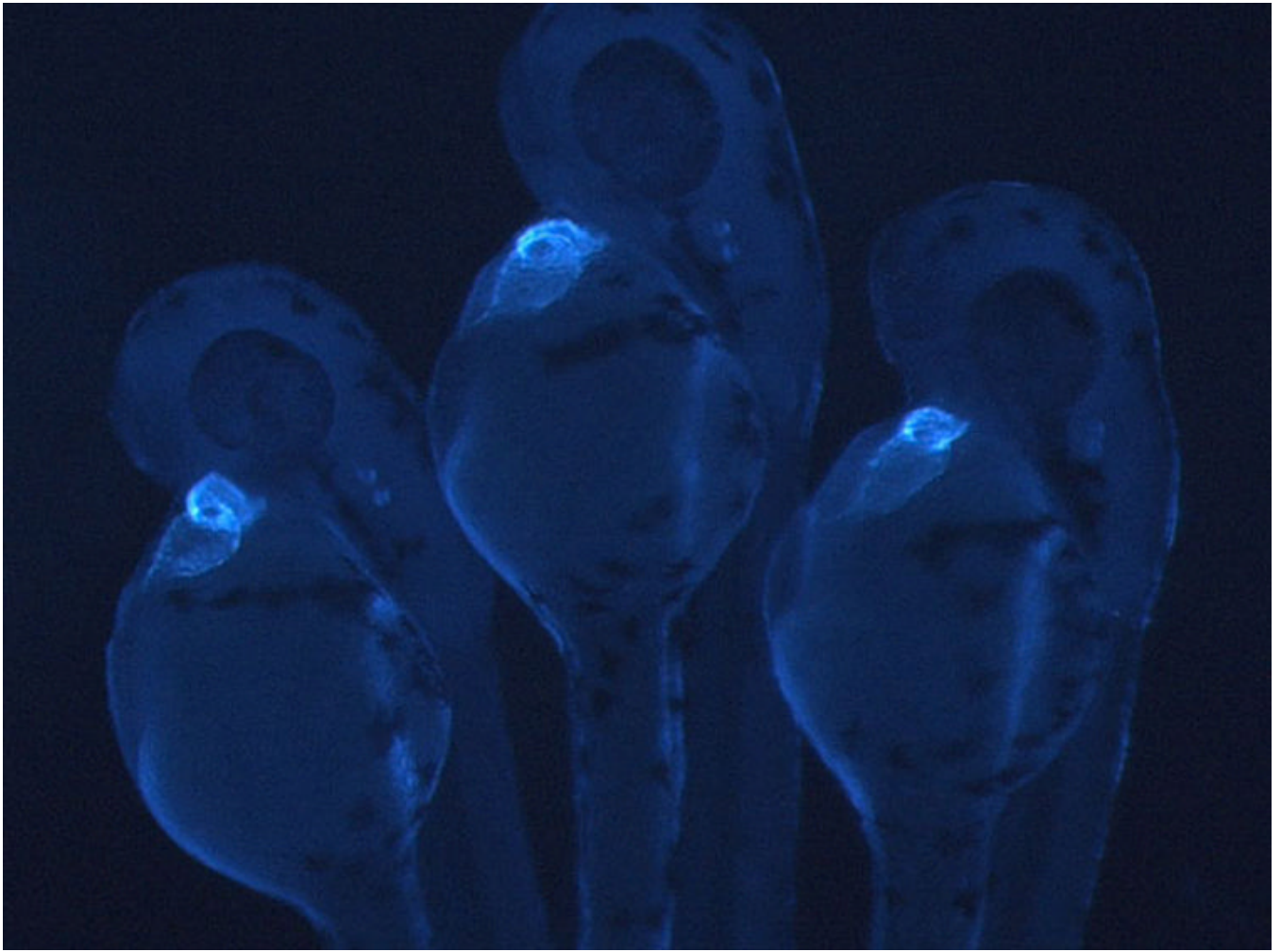


Figure 1. Transgenic zebrafish expressing a blue fluorescent protein from the cardiac myosin light chain 2 promotor. Image courtesy of Peter Schlueter.

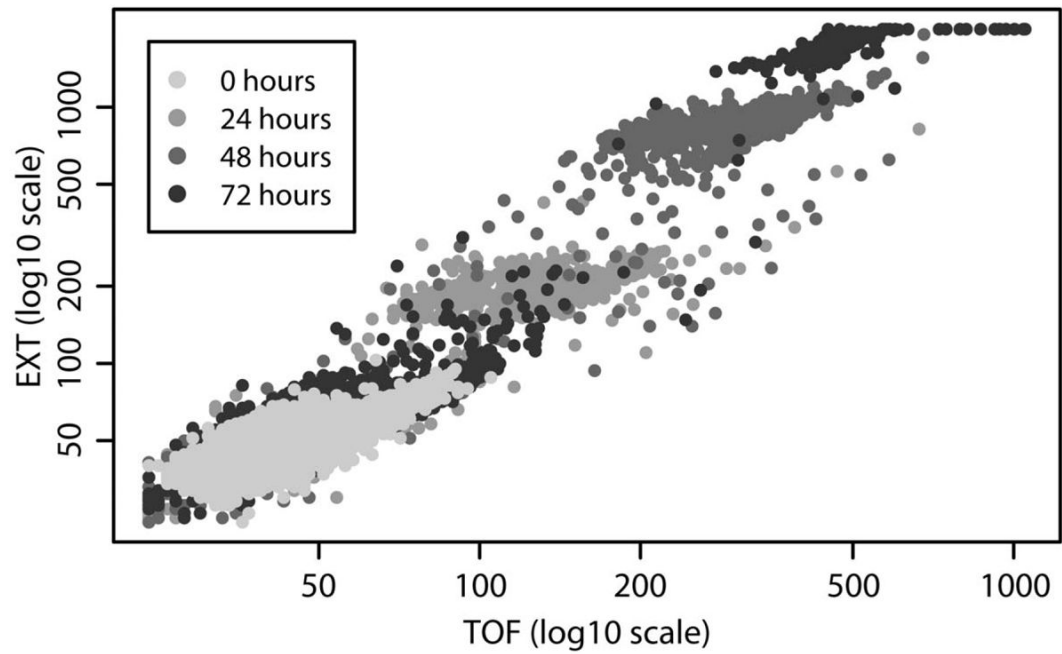


Figure 2. Development of *C. elegans* from larvae to adult over 72 h. Extinction (optical density) versus time of flight (length) for nematodes incubated at 20°C for 0, 24, 48, or 72 h. Each point corresponds to a single nematode.

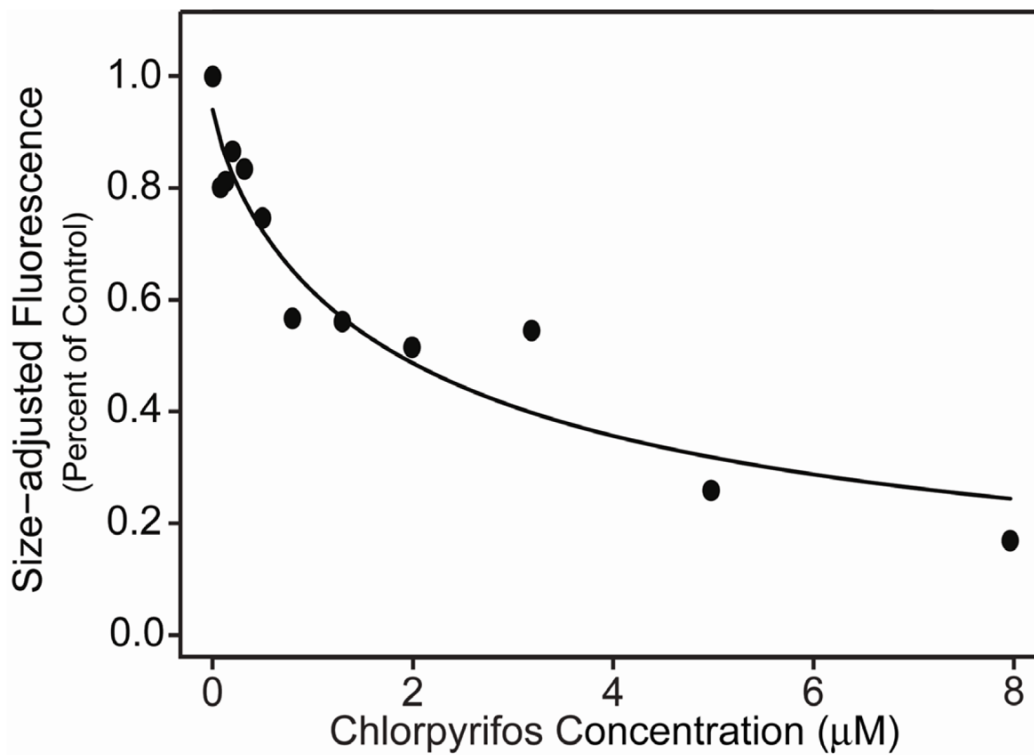


Figure 3. Neurotoxic effects of chlorpyrifos on adult *C. elegans* feeding. Fitted concentration-response curve of a representative experiment based on observed mean size-adjusted fluorescence measurements as a percent of the control for chlorpyrifos. Each point represents approximately 120 nematodes on average, with counts ranging from 104 to 141 (For details, see Boyd et al., 2007).



Figure 4. Effect of cadmium-exposure on *mtl-1::GFP* transgenic *C. elegans*. Transgenic nematode expressing GFP under the control of the *C. elegans* metallothionein promoter (*mtl-1*) were grown in the absence (upper panel) or presence (lower panel) of 100 μ M cadmium for 24 h. Constitutive *mtl-1* transcription is observed in the pharynx of the nematodes, while metal-inducible transcription occurs in the nematode intestine.

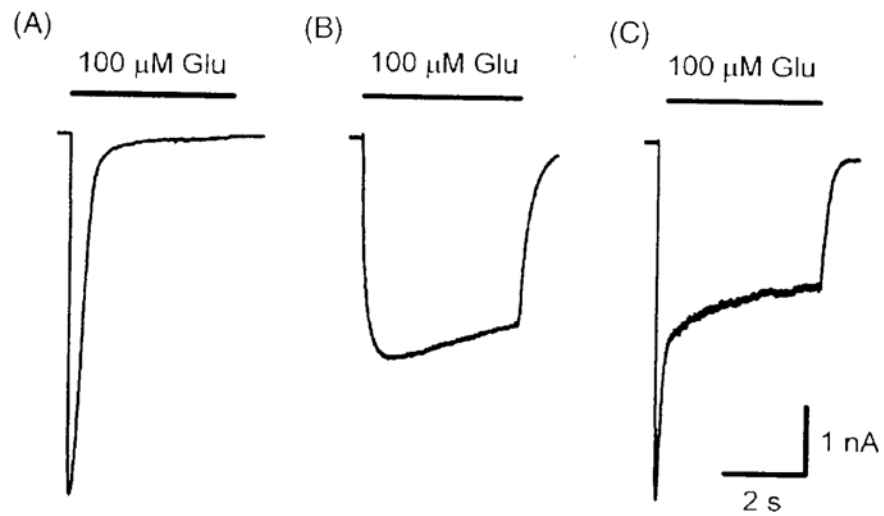


Figure 6. Glutamate-activated currents in cockroach thoracic ganglion neurons recorded by the whole-cell patch clamp method. Three types of currents were evoked following the U-tube applications of 100 μ M glutamate for 8 sec at a holding potential of -60 mV, with the symmetrical chloride concentrations between internal and external solutions (Zhao et al., 2004b).

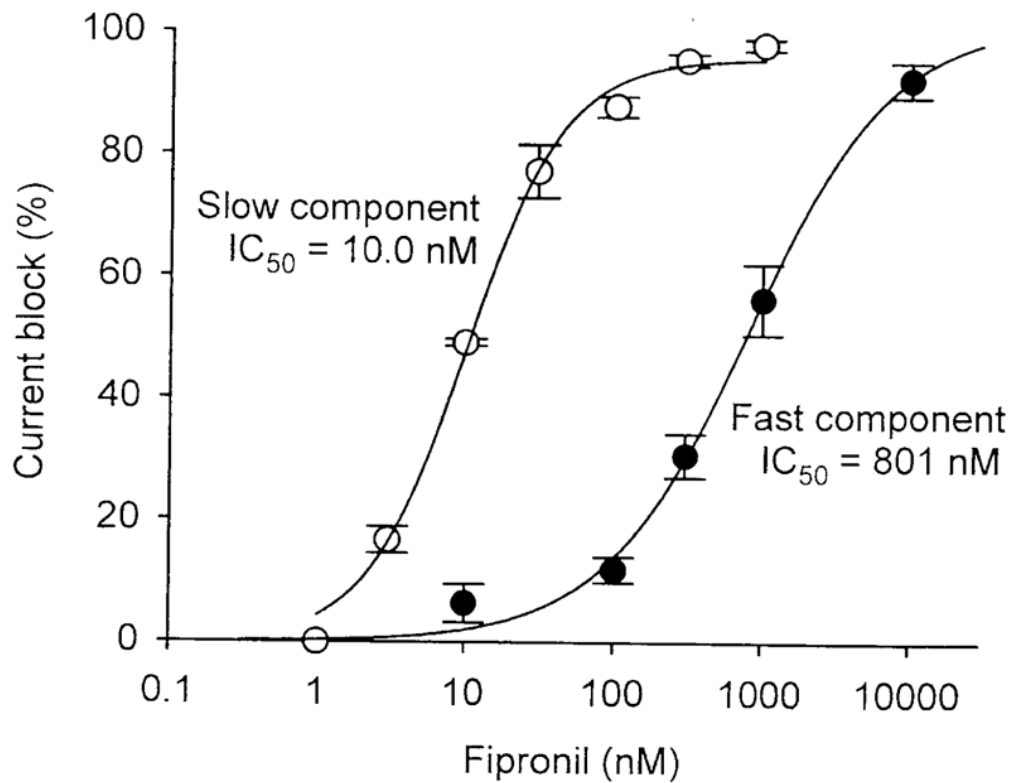


Figure 7. Dose-response relationships of fipronil block of slow- desensitizing and fast-desensitizing GluCl_s of cockroach neurons. The currents were evoked by 20-sec U-tube and bath application of 100 μ M glutamate and various concentrations of fipronil at a holding potential of -60 mV. The maximum peak current was measured for the fast-desensitizing current and the steady-state current was measured for the slow-desensitizing current (Zhao et al., 2004a).

Table 1Chemicals tested in *C. elegans* medium-throughput growth, reproduction and feeding assays.

Metals		Organics	
Aluminum Chloride	Acetaminophen	EtOH	Pyridine
Cadmium Chloride	α -Cyclodextrin	Fumonisin	Sodium Metam
Chromium Oxide	All-trans Retinoic Acid	Glyphosate	Tamoxifen
Cobalt Chloride	Ascorbic acid	Lindane	Tebuconazol
Copper Sulfate	β -Cyclodextrin	Methadone	Valproic acid
Lead Acetate	β -Cyclodextrin hydrate	Methanol	
Lead Nitrate	Caffeine	Methyl cellulose	
Manganese Oxide	Carbaryl	Methyl Parathion	
Mercuric Chloride	Chlorpyrifos	Methylisothiocyanate	
Methyl Mercury	Demeton-S-methylsulfone	MMS	
Nickel Sulfate	Dichlorvos	MNNG	
Silver Nitrate	Diphenylhydantoin	Monocrotophos	
Sodium Arsenite	Diquat	Nicotine	
Sodium Selenite	DMSO	Paraquat	
Thimerosal	EMS	Parathion	
Vanadium Oxide	ENU	PCB mixture	
Zinc Sulfate	Ethephon	PEG-60	