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Animal Models of Endocrine Disruption

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Abstract

Endocrine disrupting chemicals (EDCs) are compounds that alter the structure and function of the endocrine system and may be contributing to disorders of the reproductive, metabolic, neuroendocrine and other complex systems. Typically, these outcomes cannot be modeled in cell-based or other simple systems necessitating the use of animal testing. Appropriate animal model selection is required to effectively recapitulate the human experience, including relevant dosing and windows of exposure, and ensure translational utility and reproducibility. While classical toxicology heavily relies on inbred rats and mice, and focuses on apical endpoints such as tumor formation or birth defects, EDC researchers have used a greater diversity of species to effectively model more subtle but significant outcomes such as changes in pubertal timing, mammary gland development, and social behaviors. Advances in genomics, neuroimaging and other tools are making a wider range of animal models more widely available to EDC researchers.

Keywords

zebrafish; vole; peromyscus; toxicology; neurosciences; collaborative cross; PFOA

Introduction

An endocrine disrupting chemical is an “exogenous chemical, or mixture of chemicals, that interferes with any aspect of hormone action” [1, 2]. Although classical toxicology heavily

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relies on rodent models, especially rats, the EDC field owes much of its origins to studies in wild animal species. Its core principles and concepts were derived from studies conducted in a wide range of taxa both terrestrial and aquatic, and species diversity remains a core element of ongoing EDC research [3]. Despite comprehensive experimental and epidemiological data for some of the most notorious chemicals, regulatory action on their manufacture and use has been glacially slow to nonresponsive, leading to frustration among families, communities, activists, and scientists. Difficulty translating effects in animals to humans has been cited as a core obstacle, thus experimental animal model selection is critically important. There is growing consensus that classical, apical endpoint-based toxicity testing is not being conducted at human-relevant doses, during the appropriate life stages, or in appropriately susceptible test models to identify or predict endocrine-related disorders such as endometriosis or premature puberty, or to fully assess complex neurodevelopmental disorders that do not have clear pathology, such as schizophrenia and autism. Thus, although there is growing pressure to move away from animal-based toxicity testing, whole organism-based studies remain a critical tool for EDC research because they allow for the interrogation of chemical influences at the phenotypic, physiological, behavioral, and molecular levels and are particularly useful to assess outcomes that are difficult to model in simpler *in vitro* or organotypic systems.

EDCs have since garnered considerable attention and rapidly compounding evidence reveals that exposure, particularly during critical windows of organ development, is likely contributing to rising rates of multiple disorders and chronic diseases in humans including premature female puberty, compromised fertility, obesity, cardiovascular disease risk, and disorders of neurodevelopment [1, 2]. The incidence of these diseases/disorders has increased faster than can be explained by genetics alone, and is now thought to be heavily attributable to environmental factors, including EDCs [4]. Although, historically, the field has focused primarily on the estrogen-disrupting effects of EDCs, and effects on reproductive development and function [3], it is now recognized that EDCs can also act via other mechanisms and have impacts on non-reproductive physiology.

Common mechanisms of EDC action include hormone agonism, hormone antagonism, modulation of hormone receptor expression, and disruption of hormone production and/or clearance. While most work still heavily focuses on estrogen, androgen and thyroid disruption via their respective receptors, non-steroidal hormone disruption has repeatedly been shown [5]. For example, studies in multiple species of birds and mammals have revealed that kisspeptin, gonadotropin releasing hormone (GnRH), and oxytocin (OT)/ vasopressin (AVP) pathways (vasotocin (AVT) in non-mammalian species) are vulnerable [5–9]. Some of these effects involve steroid hormone receptor dependent mechanisms, but others do not. There is also growing interest in possible epigenetic, and immune mechanisms of disruption [2, 10, 11].

There are an estimated 90,000+ anthropogenic chemicals in the wild and built environment, although an accurate accounting has proved nearly impossible to obtain, even for regulators such as the US Environmental Protection Agency (EPA) charged with monitoring their distribution and potential toxicity (<http://cen.acs.org/articles/95/i9/chemicals-use-today.html>). The vast majority have not been tested for any form of toxicity at all, let alone

endocrine disruption, so information on their potential health risks is patchy and often contested. A subset of at least a few hundred (estimates vary) are categorized as endocrine disrupting compounds (examples shown in Table 1) with dozens in our bodies at any given time [12]. This complex exposure landscape is largely unavoidable illustrating the critically importance of understanding how EDCs affect human health. Although there is considerable interest and pressure to develop high throughput screening assays and other tools which do not use whole animals to more efficiently and rapidly accomplish the goal of “predictive toxicology [13, 14],” the development and acceptance of effective approaches remains controversial and primarily focused on estrogen, androgen and thyroid activity [15–17]. Additionally, the inherent biological complexity of the whole organism has not been adequately replicated or modeled in simpler systems and complex behavioral phenotypes and processes such as pubertal onset, affiliative interactions, and learning can only be observed in whole animals. Animal models, with organ and endocrine systems modeling those in humans, allow for the evaluation of chemical influences at many levels and are particularly useful to investigate mechanisms of action, critical windows of susceptibility, sex and age specific effects, and dose responses.

Animal Models in Toxicology: Lessons Learned

Justification and utility of animal-based work rests on several key assumptions, the most fundamental of which is that other organisms can serve as accurate predictive models of toxicity in humans. Thus, selection of an appropriately sensitive animal model is key for accurately guiding health decisions. Animal models are simply that, models, all of which have strengths and weaknesses. Understanding the advantages and limitations of any particular model is essential to maximizing the translational value of the data collected. Although this may seem obvious in principle, there are a number of species and strain-specific limitations that are unique to some toxicants, and can lead to erroneous conclusions about human risk.

Thalidomide

The thalidomide tragedy is a seminal example (reviewed in [18]). Introduced in 1957 by a German pharmaceutical company as a nonaddictive, nonbarbiturate sedative, and then as an anti-emetic for morning sickness in pregnant women, it was widely marketed and distributed in 46 countries until the early 1960s (not the US, despite intense pressure on the FDA to approve its use). Unfortunately, thalidomide caused infant mortality rates as high as 40% and severe birth defects in over 10,000 children including, most commonly, phocomelia, but also malformations of the face, eyes, ears, genitalia, and multiple internal organs, including heart, kidney, and gastrointestinal tract. Although the details remain disputed, the drug appears to have been tested primarily in mice, which was standard practice for the era, but clearly insufficient and later found to have failed for several reasons: (1) species differences in metabolism, (2) species differences in susceptibility, (3) species differences in the nature of the resulting adverse outcomes and, (4) a very short embryonic window of exposure susceptibility. Consequently, the thalidomide tragedy completely revolutionized how drugs were tested for safety and efficacy, including how animal models are selected and used. For the first time, there was recognition that critical species differences exist in drug reaction/

response. It is now appreciated that mice, traditionally used to screen for drug action, are less sensitive to thalidomide than other species, and the primary outcome is fetal resorption not phocomelia. This species-specific resilience is not unusual, and differences in sensitivity can also exist across rodent strains. For example, vulnerability to the potent toxicant 2,3,7,8-tetrachlorodibenzo-*p*-dioxin (TCDD) varies considerably across species with some strains being completely resistant to this and other chemical carcinogens, while others succumb to even very low doses [19, 20].

In humans, Thalidomide acts via several mechanisms [21], including interference with the Wnt/beta-catenin pathway, thereby increasing apoptosis in the developing limbs, eyes, brain and other organs. The same occurs in chick and rabbit embryos, but not mice [22]. Mice also metabolize Thalidomide differently and much faster than humans and other rodent species. Notably, the active metabolites are responsible for much of the drug's activity, but these metabolites are species specific, explaining, at least in part, why thalidomide toxicity is species dependent. Additional work over the intervening decades has shown thalidomide damage in non-human primates, rabbits, armadillos, *Xenopus*, hamster, chicken, zebrafish, marine fish, and even hydra and bacteria. This wide range of vulnerable taxa is a powerful reminder to the biomedical community not to rely so heavily on mice, but also that effects in wildlife and other “non-model organisms” can produce informative, human-relevant data for toxicants.

The thalidomide tragedy also demonstrated the importance of critical windows. The severity and location of the deformities exquisitely depend on when fetal exposure occurs, with the most sensitive window being between 34 and 49 days after the last menstrual period [21]. Exposure on the early end of that critical window results in central brain damage, with risk then shifting to the eyes, ears, face, arms and legs on subsequent days. Critically, teratogenicity is not seen if taken beyond these early days of gestation. Thus, a mouse study with exposure in the later stages of pregnancy would yield the deceptively false conclusion that thalidomide is safe.

Diethylstilbestrol (DES)

A similar story unfolded for the notorious EDC, diethylstilbestrol (DES). Administered to as many as 10 million pregnant women and newborns (to promote weight gain) in the USA between 1938 and 1971, this potent estrogen agonist produced no obvious teratogenic effects in newborns but rather more insidious, and less obvious outcomes which only became apparent years after exposure. An estimated 90–95% of DES daughters suffer from unusual cancers of the reproductive tract, and/or reproductive problems including reproductive tract malformations, endometriosis, infertility, and more complicated and unsuccessful pregnancies [23]. DES exposure is also associated with increased risk of psychiatric disorders including depression, anorexia, phobias and learning disabilities [24]. DES sons also have adverse outcomes including elevated rates of urogenital malformations, undescended testes, urogenital inflammation, low sperm density/quality, and testicular cancer [23]. As with thalidomide, outcome depends on timing of exposure [25, 26]. Most of the reproductive outcomes following fetal exposure to DES were recapitulated in mouse models [27] suggesting that appropriate animal studies could have theoretically caught the

effects. At the time, however, the focus was primarily on overtly teratogenic endpoints and not more subtle, long-term outcomes such as compromised fertility and pregnancy complications. With EDCs, latency between exposure and outcome can be years to decades long with prenatal exposure heightening risk to later in life cancers, such as breast or prostate cancer, or fertility-related deficits such as low sperm count or poor implantation.

Perfluorinated Compounds: A Contemporary Example

There are also contemporary examples where metabolism of the EDC varies between rodent species and even between strains within a species. Having some information on the pharmacokinetics in humans of the test chemical is imperative in selection of a model species/strain for further testing. For example, the link between perfluorooctanoic acid (PFOA), an industrial surfactant and common water pollutant, and altered breast development timing and function was first uncovered by studies in animal models. PFOA delivered to pregnant mice caused a lack of normal lactational competency in the exposed dam [28] and led to increased offspring mortality [28, 29]. Gestational PFOA exposure resulted in delays in mammary gland epithelial cell development and ductal elongation, and reductions in ductal branching and terminal end bud appearance in CD-1 female mouse offspring [28, 30]. Subsequent studies using lower doses and exposures spanning full or late gestation to focus on the time of mammary bud development, were sufficient to stunt mouse mammary gland development [31] and alter gene expression required for normal gland development in C57Bl/6 mice [32]. PFOA exposure during the peripubertal period can also induce altered estrous cyclicity, decreased ovarian steroid hormonal synthetic enzymes, and reduced gene expression of steroid-induced mammary growth factors but effects vary significantly by strain [33]. A recent paper highlighting the strain differences in metabolism or excretion of PFOA, reported that when Sv/129 mice received doses 3 to 300 times higher than those that affected CD-1 mice, Sv/129 mice had no mammary effects and lower than expected PFOA blood concentrations [34]. Enhanced excretion of PFOA in C57Bl/6 mice was also the basis for species differences in low dose effects in mammary tissue [32]. Female rats cannot be used to investigate these human relevant PFOA exposures because they exhibit a sex-specific increase in elimination rates compared to male rats; a difference that is not found in mice or humans.

Human Chemical Catastrophe and the Need to Model Mechanisms of Toxicity

Environmental accidents provide unfortunate and tragic evidence of human susceptibility to chemical exposures and critical windows of exposure. The 1976 pesticide plant explosion in Seveso, Italy [35], revealed a relationship between dioxin exposure and significantly increased cancer rates in women [36], increased metabolic disease in women who were 12 or younger at the time of the explosion [37], permanently reduced sperm quality in men who were breastfed as infants just after the explosion [38], and a dose-related association between serum dioxin levels and time to pregnancy and infertility in women [39]. Other pollution events have also been correlated with health effects in large residential cohorts. Specific examples include Agent Orange exposure to servicemen in south Vietnam [40], contamination of drinking water sources with a myriad of volatile organic chemicals at the

Camp Lejeune Marine base, Jacksonville, NC, from the 1950s-mid 1980s [41, 42], and widespread PFOA contamination of the Little Hocking River and surrounding areas of Northern Kentucky and Ohio from as early as the 1950s to the present [43]. In all of these instances, exposure was linked to numerous documented health effects, but their mechanisms of action were unclear, making it difficult to predict possible future health effects and predict risk in other exposed populations. When correlations are made between health outcomes in humans and a particular chemical, the confirmation of cause-and-effect and the elucidation of a mechanism or mode of action must be derived from experimental studies, usually animal models.

For any toxicological or EDC study, a number of criteria for choosing the animal model must be met (Table 2). Some (of the many) reasons that a given animal model may be inappropriate include differences in metabolic fate, differences in the critical window(s) of sensitivity, lack of a sensitive or homologous target (the organism fails to model the human-relevant pathway), and the presence of an irrelevant target (the drug acts via a mechanism not relevant to humans). Practicalities such as cost and housing availability may also be important considerations. Housing challenges, for example, have limited use of classic EDC models such as sheep [44] and quail [45, 46]. Other species sometimes used in toxicology such as the mini pig, dogs, ferrets, rabbits, hamsters and non-human primates are rarely used for EDC research for similar reasons. *Daphnia Magna* have been used for decades by some groups in the context of chemical screening, albeit with some concerns about sensitivity and chemical specificity [47, 48]. Wildlife species including fish, birds, and reptiles such as alligators also remain critically valuable sentinels of organism and ecosystem health but pose their own logistical challenges [3, 49]. For some endpoints, particular model organisms have unique features that make them more advantageous. The guinea pig, for example has a more human-typical placental structure, but are more expensive per unit than rats and mice. A list of some commonly used EDC models with their unique strengths and weaknesses is listed in Table 3.

Novel Animal Models in EDC Research

Although classic toxicology still heavily relies on inbred lines of rats and mice, powerful new options in rodents and other species created to leverage significant advances in gene editing, genomics, and neuroimaging hold the potential to significantly advance EDC research. Although some fields, particularly genetics and the neurosciences, have made significant discoveries with lower order species such as the nematode *Caenorhabditis elegans*, and the fruit fly *Drosophila melanogaster*, there is reticence to use them for toxicological testing because of concerns about data translation to humans. However, a greater diversity of vertebrate models is gaining in popularity including transgenic mouse (and rat) lines, zebrafish *Danio rerio*, and monogamous rodents such as the prairie vole *Microtus ochrogaster*. Perhaps most exciting are powerful new mouse lines for interrogating population-level effects, and gene by environment interactions.

Zebrafish

The zebrafish model is rapidly being adopted as an option for EDC research, particularly chemical screening, because of several critical advantages [50]. Compared to mammalian models zebrafish are easy to rear and relatively inexpensive to maintain, even for small labs, and can be bred and studied at high speed, and used in high numbers. Less than a few millimeters in size, the larvae are small enough to be accommodated in multiwell plates, making their large scale use in semi-automated systems feasible [50–52]. The translucent body of zebrafish embryos combined with advances in high dimensional imaging tools facilitates non-invasive visualization of organs and biological processes *in vivo*, including neurodevelopment and the fate/transport of putative EDCs. The relative ease by which the zebrafish genome can be manipulated is also a unique advantage. The zebrafish genome is comprehensively annotated, a large number of transgenic lines are readily available, and alteration of gene expression is easily achievable using antisense morpholinos or gene-editing techniques, such as TALENs or CRISPR/Cas9. There are also robust genomic tools and consortium-level projects enhancing their utility. Launched in 2011, the Zebrafish Mutation Project (ZMP) is an initiative to create a knockout allele in every protein-coding gene in the zebrafish genome. The data and resources are readily available on the ZMP website and via the Zebrafish International Resource Center (ZIRC). The mammalian and zebrafish genomes display reasonably high homology with an overall conservation of over 70%, and an estimated 80% of human genes expressed in zebrafish [53]. They also effectively model some of the most basic modes of endocrine disruption. Many basic physiological, sensory, and anatomical and signal-transduction mechanisms are also homologous to those of mammals [54].

Significant advances in instrumentation and –omics technology have made it possible to use early life stage zebrafish for EDC screening. For example, several studies have used such systems to screen the EPA’s ToxCast chemicals for endocrine disrupting and other toxicity outcomes, and to evaluate the validity of comparable *in vitro* and *in vivo* toxicity data [50, 52, 55]. Zebrafish were also used to identify a critical pathway by which Thalidomide inhibits limb development [56].

As with all model systems, zebrafish have noteworthy limitations. Most obvious is that EDC exposure is typically via absorption from tank water (dermally and, as the gills mature, via circulation), meaning that chemical uptake and metabolism can be quite different from the human experience where exposure is typically oral or inhaled. Tissue concentrations are also poorly understood and rarely reported, which can make it difficult to extrapolate concentration-response results in zebrafish to dose-response studies in mammals. By extension, although zebrafish metabolism is similar to other vertebrates, there are subtle differences that could yield inaccurate conclusions. As with any animal model, some pathway components may not necessarily be highly conserved and some cellular systems may bear little homology. For example zebrafish lack brown adipose fat so exploring pathways activated by non-shivering thermogenesis, such as the β -adrenergic system, will be limited. Finally, as with rodents, inter-laboratory replication difficulties have occurred, most likely due to strain, housing and other differences, but are not well documented or understood. As in mammals, lab-reared colonies are vastly different in many respects from

wild-derived strains, and allelic variations between individuals can result in marked differences in chemical susceptibility [57]. Accounting for this in future study designs could enhance an already uniquely advantageous EDC screening tool.

Transgenic Rodents

Developed in the 1970s and now commonly used in virtually every biomedical field, transgenic rodents are surprisingly seldom used in toxicology. Literally thousands of different lines are now available and have proved to be powerful tools for modeling human disease and exploring the roles of specific genes in biological pathways and systems. Although the older, more traditional techniques for generating rodent transgenics only reliably worked in mice, newer tools such as CRISPR/Cas9 are rapidly making it possible to create transgenic lines in rats and other species [58]. Using *Cre-Lox*, Flp-Frt, or similar approaches, it is also now possible to make site-specific conditional transgenics where genes can be manipulated, for example, in specific cell populations, or under certain conditions [59]. Using a combination of approaches transgenics can be used to perform gene targeting experiments to achieve genotypic/phenotypic “rescue,” spatio-temporally inactivate genes in specific cells, and perform cell fate mapping and gene/protein expression profiling. For example, transgenic techniques have been used in combination with viral vectors and other tools to introduce visual (such as GFP) and other reporters to manipulate and visualize entire neural circuits and pathways [60].

In classical toxicology transgenics are grossly underutilized. To date, they have really only been used for hazard characterization of mutagens, and even then only rarely [61]. They have, however, been used by EDC researchers to probe mode of action and generate humanized mouse models. For example a variety of ER-mutated and knockout mouse lines are now available [62, 63] and have been used to explore non-canonical signaling mechanisms [64] as well as mechanisms of estrogenic endocrine disruption [65, 66]. Examples of models developed to overcome known species differences in toxicological vulnerability and outcomes include mutated and humanized aryl hydrocarbon receptor (AhR) [67] and PPAR α -humanized mice [68]. Use of knockout and other transgenic models have also been used to probe EDC conjugation and metabolism [69].

While powerful tools to elucidate mechanisms of endocrine disruption and possible gene by environment interactions, transgenic animals have limitations, most significantly the high cost and time required to generate them. There is also the potential for incomplete knockout of the target gene, off-target expression, biological compensation, and incomplete recapitulation of the expected disease phenotype (many “Alzheimer’s mice” have this problem). Numerous Cre lines have phenotypes that are not obvious or fully characterized, and Cre activity in off-target tissues is proving to be fairly common. In some strains, Cre mosaicism is also a known limitation and Cre recombinase activity can be less robust when paternally inherited. Researchers should be sure proper controls are used to validate and maintain the models.

Population-Based Genetic Resources

Natural genetic variation plays an important role in nearly all biological processes and traits, including response to EDCs [70]. Population-based mouse resources capture individual differences in EDC susceptibility and may better model diverse human populations. Multi-parent populations such as the Collaborative Cross (CC) and Diversity Outbred (DO) mice are designed to maximize genetic variation while controlling the relationships between individuals, so that they are highly amenable to genetic analysis such as quantitative trait locus (QTL) mapping [71–73]. The CC population consists of 83 inbred mouse strains that are all descended from eight existing strains. These eight founder strains selected are a combination of common strains that were historically important in biomedical research and wild-derived strains that were included to maximize genetic diversity. The recombinant inbred line breeding design means that each strain is equally related to the others, and eliminates population structure that can limit QTL mapping ability or create false positive QTL. Each CC line is an inbred mouse strain, and can be reproduced at will in multiple environmental contexts. DO mice are descended from the same eight founder strains as the CC, and therefore both populations have the same sets of alleles. In contrast to the CC, DO mice are outbred and each individual is unique; hundreds of individual DO mice can be used in a single mapping study. These large populations combined with the outbred genomes make the DO uniquely powerful for detecting genetic associations. While each DO population has similar properties in terms of genotype frequencies, individual DO mice are not reproducible.

Both of these mouse resources allow replicable populations to be studied across a range of environmental contexts [74] and reveal gene by environment (GxE) interactions. Population studies can both measure the range of response to a specific toxicant and to identify specific genes driving susceptibility. In one example, DO mice exposed to benzene showed an order of magnitude more variation in chromosomal damage than isogenic B6C3F1 mice. QTL mapping in the population revealed a novel role for the sulfotransferase *Sult3a1* in benzene response [75, 76]. A separate study found that trichloroethylene metabolism varied in CC populations due to multiple interacting factors, and individual CC strains showed distinct dose-response trajectories [77]. Other studies using these resources with environmental factors have found new candidate genes involved in the asthma response [78, 79], differential toxicity to chemotherapeutics [80, 81], and sensitivity to a popular food additive [82]. Thoughtful application of these resources to EDC research will directly inform human studies regarding population-level variation and mechanisms of susceptibility.

Monogamous Rodents

Rapidly and inexplicably rising rates of autism spectrum disorder (ASD), attention deficit activity disorder (ADHD) and other developmental disorders of non-reproductive behaviors including social interaction have spurred widespread interest in understanding how EDCs might impact the social brain. Genetic factors contribute, at best, only an estimated 30–40% of ASD heritability [83, 84], indicating that it and other disorders of sociality have a significant environmental component. Although it is widely speculated that chemical exposures are contributory, there is actually very little *direct evidence* linking any specific chemical, including EDCs, to adverse effects on the social brain or social impairments in

humans [84–89]. Additionally, relatively little is known about the mechanisms by which the social brain could be vulnerable to chemical exposures, or which chemical classes are most likely to be detrimental [90–92].

Because classical rodent models are limited in terms of what human-relevant social behaviors they can recapitulate, animal model suitability has been cited as a significant experimental barrier. For example, common ASD mouse models often have some sort of neuropathology not seen in human patients (ex. the BTBR mouse completely lacks a corpus callosum and has a severely reduced hippocampal commissure) and none display partner attachment or paternal care; defining elements of human sociality. In that regard, monogamous rodent species could prove uniquely valuable. Although no animal model can fully recapitulate the sophisticated complexity of human social behavior, the neuroendocrine pathways coordinating social traits are highly conserved [93, 94] including the sexually differentiating influence of steroid hormones, and coordinating roles of the neuropeptides vasopressin (AVP) and oxytocin (OT).

Much of what is known about the evolution and manifestation of pro-social traits came from groundbreaking work in microtine voles. Prairie and pine voles (*Microtus ochrogaster* and *Microtus pinetorum* respectively) are socially monogamous while montane and meadow voles (*Microtus montanus* and *Microtus pennsylvanicus* respectively) are socially promiscuous and solitary providing an ideal comparative system for exploring species-level differences in social brain structure and development. Decades of transformative work in these and other microtine species has linked pro-social traits to the OT/AVP system and its interactions with mesolimbic dopamine pathways [95–97]. The translational importance of the vole model has now been demonstrated in humans (reviewed in [97]). Most significantly, intranasal OT administration is being used therapeutically for ASDs. Because OT/AVP and the downstream dopaminergic pathways they project to are heavily influenced by sex steroids across the lifespan [98–101], it is highly plausible that their sexually dimorphic ontogeny and function may be susceptible to endocrine disruption. With the prairie vole genome now sequenced, it is feasible to do the types of gene by environment experiments needed to identify possible mechanisms by which EDCs might contribute to social decrements relevant to ASD and other behavioral disorders [96, 102].

Only a handful of studies have used voles to explore endocrine disruption of the social brain. The first linked perinatal methoxychlor exposure to reduced affiliative behavior by exposed females, and reduced oxytocin receptor binding in the cingulate [103], a region thought to play a role in stress responses and emotional processing [104]. More recent studies have reported that BPA can alter aspects of anxiety-related behaviors, eliminate or reverse sexually dimorphic social behaviors, alter OT/AVP and dopaminergic neuron numbers in multiple brain regions critical for sociality, and disrupt microglial colonization [105, 106].

Similar pro-social models include *Peromyscus* mice (often collectively referred to as deer mice). As in *Microtus*, some species are socially monogamous (most notably *Peromyscus californicus* but also *Peromyscus polionotus*) while others are not. The mechanisms underlying monogamous behavior in this genus, however, may differ from *Microtus*, and are not as well understood [107]. As with zebrafish, *Peromyscus* are a well-established model

for studying the evolution and genomics of complex traits and have been backed by consortium-level projects to advance their use. Most significantly, various “wild type” and mutant lines have been created and maintained by the *Peromyscus* Genetic Stock Center at the University of South Carolina.

Like *Microtus* they can be more challenging to breed and maintain the laboratory environment than rats or mice, but have key advantages including their genetic diversity and the spontaneous display of human-relevant social traits such as pair bonding and paternal care. Notably, deer mice have been utilized to monitor real world exposures by comparing exposed wild animals with laboratory-bred controls [108, 109]. *Peromyscus* have also been used to explore the possible effects of BPA on exploratory and socio-sexual behaviors [110, 111]. Because of their limited sources, the cost may be higher than other rodent models, but their environmental relevance may outweigh the disadvantages for some studies.

Summary

In conclusion, numerous considerations must be made when designing studies to address disease conditions that occur in human populations as the result of EDC exposure. Although cell models may be useful in identifying some components of signaling pathways, they lack feedback loops, endocrine systems, and initiation and progression of disease processes that allow for translation of EDC effects. They are also incapable of modeling or recapitulating complex behavior. Rodent and other relevant animal models are critical in testing the effects of EDCs on developmental exposures, systems biology, immunity, reproductive and neurobehavioral outcomes. Emerging models such as zebrafish, monogamous rodents, and population-based mouse resources hold tremendous promise for addressing key challenges in EDC research including differences in individual and population-level susceptibility, effects on complex behaviors, including social behaviors, the capacity for rapid EDC screening, and the ability to explore gene by environment interactions.

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Practice Points

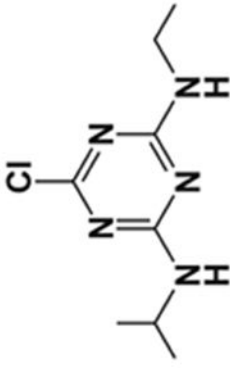
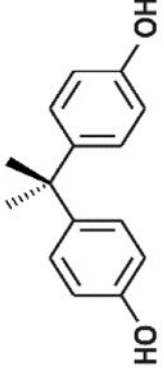
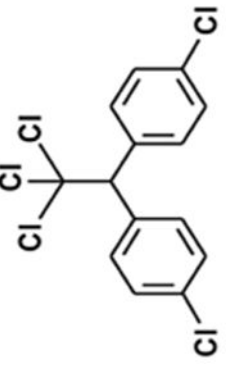
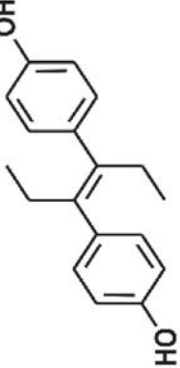
- EDCs are widespread and likely contributing to a myriad of human diseases, thus greater physician and patient awareness of these chemicals and their sources is needed.
- Human exposure is typically low, life long, and to complex mixtures with exposures highest in children.
- Some exposures can be mitigated by simple lifestyle changes such as using unscented products and glass instead of plastic.

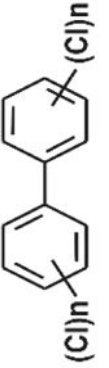
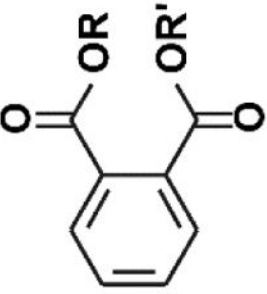
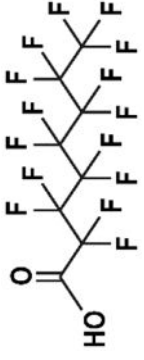
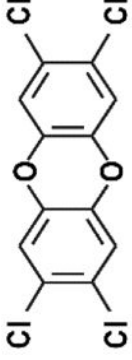
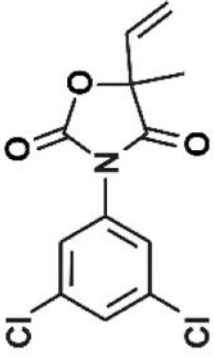
Research Agenda

- Research has typically focused on reproductive and thyroid targets but exciting new research is identifying immune, cardiovascular, metabolic, epigenetic and other endpoints.
- An overreliance on one or only a few strains of mice can limit the type of risk and biology that can be effectively modeled.
- Zebrafish are emerging as cost effective tool for EDC screening and should be incorporated into regulatory testing strategies to improve screening capacity and accuracy.
- New rodent models including a wide range of transgenic strains, monogamous species, and those specifically designed to examine population-level effects are rapidly emerging to address significant limitations in our current capacity to probe complex issues like inter-individual variability and gene by environment interactions.

Table 1

Summary of Common EDCs and their Histories, Uses, Sources and Effects*

EDC	General chemical structure	Group	Introduction date	Restrictions	Route of exposure	Sources	Half-life	Effects/animal model notes
Atrazine		Chlorotriazine herbicide	1959	European Union ban 2004	Ingestion, inhalation	Pesticide/herbicide, water and soil contaminant	10–12 hours	Endocrine, respiratory and nervous system targets, liver damage
BPA		Bisphenols	1960s	Varies by country. In the United States, voluntarily restricted in baby products 2012	Ingestion, inhalation, dermal absorption	Polycarbonate plastics, epoxy resins, plastic toys and bottles, lining of food cans	4–5 hours	Estrogenic, obesogenic, neurological effects, reproductive and developmental effects
DDT		Organochloride	1940s	Widely banned 1972 but still used in some countries.	Ingestion, inhalation, dermal absorption	Contaminated water, soil crops, fish	6–10 years	Estrogenic, anti-androgenic, reproductive effects, carcinogen, central nervous system, kidney, liver and peripheral nervous system effects
DES		Non-steroidal synthetic estrogen	1941–1947	Restricted 1971–1975	Ingestion, injection, vaginal suppository	Pharmaceutical for humans and livestock	2–3 days	Transplacental carcinogen, teratogen

EDC	General chemical structure	Group	Introduction date	Restrictions	Route of exposure	Sources	Half-life	Effects/animal model notes
PCBs		Organochloride	1927	Banned 1979	Ingestion, inhalation, dermal absorption	Contaminated air and food, skin contact with old electrical equipment	12 days to 16 years	Carcinogen, stomach and liver damage, reproductive and nervous system effects including IQ loss, thyroid injury
Phthalates		Plasticizers	1920s	Restricted 2009	Ingestion, inhalation, dermal absorption	Contaminated food, PVC plastics and flooring, personal care products, fragrance, medical devices and tubing	~12 hours	Antiandrogenic activity, carcinogen, liver damage, reproductive and developmental effects, asthma, obesogen, possible neuroendocrine disruptor; Rat testis is vulnerable, mouse and human fetal testis xenografts are not.
PFOA		Fluorosurfactant	1940s	United States 2015 voluntary production restriction	Ingestion, inhalation	Contaminated food and water, dust, floor waxes, fire fighting foam, electrical wiring, lining of food wrappers, stain resistant carpeting	2-4 years	Liver, developmental, and immune system toxicant, carcinogen; Rats an inappropriate model
TCDD		Polychlorinated dibenzo-p-dioxin	Synthesized 1872		Ingestion, inhalation	By-product of chlorinated herbicide production, smelting, chlorine bleaching of paper; can be naturally occurring	7-11 years	Liver damage, weight loss, atrophy of thymus gland, immunosuppression, reproductive effects and cancer; susceptibility varies widely across species and strains
Vinclozolin		Dicarboximide fungicide	1981		Ingestion, inhalation, dermal absorption	Diet and occupational	Aerobic soil 28 days, plasma 20 hours	Antiandrogenic activity, male reproductive and neurological effects, transgenerational reproductive effects, potential carcinogen

BPA=bisphenol A; DDT=dichlorodiphenyltrichloroethane; DES=diethylstilbestrol; MXC=methoxychlor; PCBs=polychlorinated biphenyls; PFOA=perfluorooctanoic acid; TCDD= 2,3,7,8-tetrachlorodibenzodioxin

* Adapted from Gore, A. C. et al. EDC-2: The Endocrine Society's Second Scientific Statement on Endocrine-Disrupting Chemicals. Endocr. Rev. 36, E1-E150, doi:10.1210/er.2015-1010 (2015).

Table 2

Considerations in Animal Model Selection and Use for EDC Research

<ul style="list-style-type: none">• Account for possible species differences in metabolism and generation of biologically active metabolites.
<ul style="list-style-type: none">• Select the most sensitive species possible.
<ul style="list-style-type: none">• Ensure the presence of a relevant target (a similar mechanism of action).
<ul style="list-style-type: none">• Ensure the outcome is relevant for human disease and not unique to that species (some types of cancers in rodents are not seen in humans).
<ul style="list-style-type: none">• Use both sexes – unless only one is relevant (prostate cancer).
<ul style="list-style-type: none">• Make sure dosing occurs over the appropriate critical period for that species.
<ul style="list-style-type: none">• Be aware that for EDCs latency between exposure and effect could be long and stretch into advanced adulthood.
<ul style="list-style-type: none">• Make sure enough individuals are used to have sufficient statistical power. For mammalian developmental toxicology studies the litter should be the statistical unit, not the individual pup.
<ul style="list-style-type: none">• Methodological reporting should use ARRIVE guidelines or similar to ensure comprehensive and transparent reporting (https://www.nc3rs.org.uk/arrive-guidelines)

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Table 3

Strengths and Weaknesses of Common EDC Animal Models

Species	Strain	Unique Characteristics
Mouse	CD1	Large litters, excellent maternal behavior, sensitive to estrogens, robust data on historical control disease incidence, outbred strain so higher variability in experiments
	C57BL/6J	Inbred strain particularly sensitive to immune challenges, good embryo donor and transplant recipient
	BTBR	Used as a model of autism but lacks a corpus callosum and has a severely reduced hippocampal commissure.
	B6C3F1	Moderate sized litters, cross of 2 inbred strains, large dataset on controls, longevity
	Collaborative Cross	Genetic diversity with reproducibility
	Diversity outbred	Genetic diversity with maximum QTL mapping power and resolution
	Rat	Wistar Han
Long Evans		Outbred strain, not albino, spontaneously displays more ethologically relevant behaviors
Sprague		Vendor strains are not congruous; Harlan or Taconic are
Dawley		preferred source. Background incidence of cardiomyopathy can be as high as 100%, CRL demonstrate high rate of mammary galactocoeles, poor longevity, and low estrogen sensitivity.
F344/N		High spontaneous tumor rate in testes and mammary gland due to prolactin sensitivity
Voles (<i>Microtus</i>)	Some strains are socially monogamous, display paternal care, alloparental care and other pro-social traits; biological basis for pro-sociality well understood	
Deer mice (<i>Peromyscus</i>)	Some strains are socially monogamous, display paternal care and other pro-social traits; have been used to characterize “real world” exposures from contaminated sites.	
Zebrafish	Transparent; relatively easy and inexpensive to house, breed and maintain; rapid development	
Guinea Pig	Human-relevant placental structure	
Sheep	Human-relevant placental structure	