

Reframing sexual differentiation of the brain

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In the twentieth century, the dominant model of sexual differentiation stated that genetic sex (XX versus XY) causes differentiation of the gonads, which then secrete gonadal hormones that act directly on tissues to induce sex differences in function. This serial model of sexual differentiation was simple, unifying and seductive. Recent evidence, however, indicates that the linear model is incorrect and that sex differences arise in response to diverse sex-specific signals originating from inherent differences in the genome and involve cellular mechanisms that are specific to individual tissues or brain regions. Moreover, sex-specific effects of the environment reciprocally affect biology, sometimes profoundly, and must therefore be integrated into a realistic model of sexual differentiation. A more appropriate model is a parallel-interactive model that encompasses the roles of multiple molecular signals and pathways that differentiate males and females, including synergistic and compensatory interactions among pathways and an important role for the environment.

The value of understanding sex differences in the brain is both self-evident and underappreciated. The effects of sex on neural phenotypes are often as large as the effects of other important variables, and conclusions based on the study of one sex have not always been found to hold in the other^{1,2}. Moreover, susceptibility to disease or dysfunction and the effect of injury can be as much as 2–5-fold greater in one sex. These include higher rates of neuropsychiatric and learning disorders with developmental origins in males and higher rates of aging-related neurodegenerative diseases and mental health dysfunctions in females^{3–5}. Heuristically, contrasting males and females has revealed previously unknown mechanisms of neural development that were not otherwise accessible^{6–8}. However, most studies of the brain and other tissues continue to focus on one sex, usually males, or fail to report the sex of the animals⁹. Thus, the still widespread assumption that the influence of sex is negligible retards progress in our field¹⁰. Even more paradoxical is that factors present in one sex sometimes counteract other sex-specific factors to eliminate sex differences in phenotype^{11,12}. Thus, sexual equality of phenotype does not imply sexual equality of physiology or development and, more importantly, sex differences are more pervasive than can be realized just from considering traits in which males differ from females.

The study of sexual differentiation of the brain has long focused on a few model cases of brain function (for example, sex behavior, control of ovulation) and brain regions that are predominantly involved in reproduction and therefore show large sex differences that are amenable to study. The repeated investigation of a relatively small number of sexual dimorphisms may have contributed to the false impression that a few discrete male or female circuits sit in an otherwise sexually monomorphic brain. The notion that for specific behaviors there is a discrete male neural circuit versus a discrete

female neural circuit remains widely held despite a lack of empirical evidence of the existence of either. Preconceived notions, stemming from the hormonally controlled elimination or retention of female (Müllerian) and male (Wolffian) reproductive tracts, may have contributed to the view of a similar system in the brain. Moreover, studies of only a few robustly dimorphic brain structures have contributed to the perception that sex differences in brain function are controlled by a unitary program: genetic sex determines gonadal sex and gonadal hormones determine brain sex. Gonadal steroids were believed to act via common mechanisms on a restricted group of brain regions to cause sex differences, promoting the formation of male circuits in males and female circuits in females. This traditional view, seductive in its simplicity, must now be replaced. Sufficient new evidence has accumulated to warrant a shift away from the old serial model and toward a more complex and nuanced model in which numerous sex-specific factors, hormonal, genetic and epigenetic, act in parallel to cause or eliminate sex differences in the brain and other tissues, by mechanisms that frequently are region specific and heterogeneous in terms of their intracellular mechanisms and mode of cell-to-cell communication. The modern view emphasizes a diversity of proximate mechanisms and an interaction of multiple sex-specific factors in many brain regions.

Biological theories of sexual differentiation have largely underemphasized or even excluded the differential effect of sex-specific environments. The environment has far-reaching influences on self-concept and gendered behavior of humans and is poorly modeled by studies of rodents. Sex differences in the environment likely have major effects on brain biology, as has been suggested by recent studies of the importance of environmentally triggered epigenetic changes in the brain¹³. The effect of environment is rarely controlled for or empirically tested, but environmental and biological factors likely interact in complex ways to sculpt the female and male phenotype. The goal of this review is to present the historic serial model of sexual differentiation of the brain and propose its replacement by a parallel model that incorporates all of the variables relevant to brain development in the sexes, including the environment.

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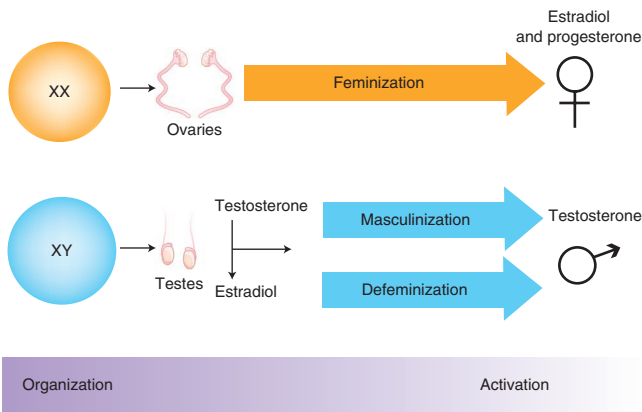


Figure 1 Twentieth-century linear view of sexual differentiation. For the past 50 years, the prevailing view of sexual differentiation of the brain has been a linear model in which chromosomal sex determines gonadal sex, which determines brain sex. Feminization of the brain is the default process that occurs in the absence of high levels of gonadal steroids during a perinatal sensitive period. Masculinization and defeminization are separate hormonally driven processes that organize the neural substrate to promote male-typical behaviors while suppressing female-typical behaviors. The organized neural substrate is activated by adult gonadal steroids and required for sex-typical behaviors to be expressed. This iconic model based on the organizational/activational hypothesis¹⁴ has proved a sturdy framework for elucidating some, but not all, of the aspects of sexual differentiation of the brain.

Classic model: hormone-mediated organization/activation

The concept that sex differences in adult brain and behavior are sexually differentiated during development by the action of gonadal hormones was first illustrated by the finding that female guinea pigs exposed to testosterone as fetuses have a permanent tendency to copulate like males rather than females¹⁴. This iconic study provided the conceptual framework for discriminating two types of sex-specific action of gonadal steroid hormones: organizational and activational¹⁵ (Fig. 1). The embryonic testes of mammalian species synthesize and release testosterone, which acts throughout the body to masculinize the genitalia, sperm ducts, brain and other tissues. These organizational effects are differentiating in the classic sense of developmental biology, in that the tissues permanently and irreversibly adopt a restricted fate (in this case, a sex-specific fate) with concomitant loss of cellular pluripotency. Contrasted with these are the reversible activational effects of gonadal hormones, which can occur at any time of life, but are predominantly studied in adults. As adults the male brain is exposed to testicular hormones and the female brain to ovarian hormones, resulting in sex differences that are abolished by gonadectomy of adults. In most instances, such as hormonal control of sexual behavior, activational effects of steroids are constrained by earlier organizational effects, including additional organization that occurs at puberty¹⁶. Thus, the dominant theory of mammalian sexual differentiation, championed even as recently as 2010 (ref. 17), is that the embryo is sexually indifferent until differentiation of the gonads and that all sex differences arise afterwards by differential action of gonadal hormones.

Parallel model: multiple sex-specific signals and pathways

All sex differences must ultimately stem from the inherent imbalance of genes encoded by the sex chromosomes, which are the only factors thought to differ in the male and female zygote. By 1959, the mammalian Y chromosome was found to contain a dominant testis-determining gene that was later identified as *Sry*¹⁸, which initiates

testis differentiation and is often placed at the top of the molecular cascade that differentiates testicular from ovarian development. The differentiation of gonads sets up sex differences in the level of gonadal hormones, which cause differences in male and female cells. Each XX and XY cell, however, inherently has a different complement of sex chromosomes. In brain and other tissues, XX and XY mice show differences in phenotype that are explained by differences in expression of X and Y genes. For example, *Sry* is expressed in the dopamine-containing cells of substantia nigra pars compacta (SNpc) that project to the striatum and has direct male-specific effects¹⁹. These neurons are the targets of Parkinson’s disease, which shows a 1.5-fold higher incidence in men. When the expression of *Sry* is temporarily reduced in adult rats, the expression of tyrosine hydroxylase in the SNpc and striatum are markedly reduced and motor performance declines. These results constitute one of the first demonstrations of a direct sex-specific effect on brain of an identified sex chromosome gene. One paradox is that, despite the sex differences in Parkinson’s disease, striatal function and SNpc neuron number^{19,20}, the SNpc and striatum function quite well in both sexes so that a male-specific dynamic role for *Sry* is unexpected. The paradox raises the question of whether a female-specific factor maintains tyrosine hydroxylase expression in females and if another male-specific factor (testosterone?) has an undesirable effect in the SNpc that is counteracted by *Sry* expression. This case raises a common theme in the investigation of sex chromosome effects, that sex-specific factors do not always make males and females different, but sometimes counteract each other to make the sexes more similar^{11,12,21}.

Another gene, *Xist*, is also encoded by the sex chromosomes and has sex-specific effects. *Xist* is expressed from one of the two X chromosomes in non-germline XX cells of eutherian mammals and initiates inactivation of that chromosome²². The ultimate effect of *Xist* is that only one X chromosome is transcriptionally active in females, and *Xist* is typically viewed as a factor that makes females more similar to males. As a result, *Xist* is rarely included on lists of genes that cause sexual differentiation. Its role in causing mosaicism of X gene effects in females, but not males, has, however, been emphasized^{23,24}. But clearly, XX cells are different from XY cells precisely because they express *Xist* and engage a major epigenetic machinery that is not active in XY cells. However, we know little about the differentiating effects of *Xist*, perhaps because its role in compensating for sex differences has been emphasized.

The direct roles of sex chromosome genes on sex differences are not well studied (and therefore probably are underestimated), not only because of the dominance of the hormonal theory of sexual differentiation, but also because few animal models have allowed manipulation of the sex chromosomes without also causing large changes in hormone levels²⁵. To date, most models for studying direct sex chromosome effects are available in mice in which the sex chromosomes can be engineered²⁶. The best studied is the four core genotypes (FCG) model, in which the complement of sex chromosomes (XX versus XY) is made independent of gonadal sex^{12,27,28}. This is possible because the *Sry* gene is ‘moved’ to an autosome and the Y chromosome is therefore no longer testis-determining. The model produces four types of offspring in a 2x2 pattern: they are either XX or XY and they either do (producing gonadal males, XXM or XYM) or do not (producing gonadal females, XXF or XYF) possess the *Sry* transgene. The model allows simultaneous appreciation of the effects of gonadal sex (comparing the phenotypes of gonadal males and females) and of sex chromosome complement (comparing XX and XY mice of either gonadal sex; Fig. 2). The results available to date from analysis of FCG mice confirm the importance of hormones, but



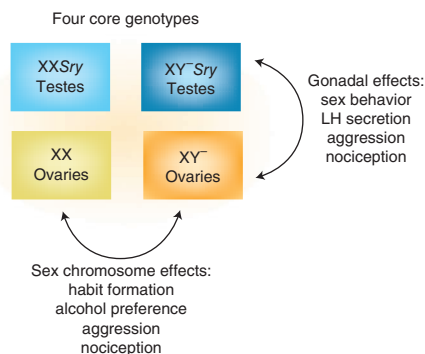


Figure 2 Genetics matter. The ability to distinguish the contributing role of genes versus gonads was markedly advanced by the development of the four core genotypes model of mice. These mice bear a Y chromosome from which the *Sry* gene has been deleted (denoted Y⁻) and carry *Sry* on an autosome, allowing the development of XX individuals with testes and XY individuals with ovaries. Analysis of this model supports the view that sexual differentiation of reproductive endpoints is largely driven by the testicular hormone testosterone or estradiol synthesized in the developing nervous system from this testosterone, consistent with the organizational/activational hypothesis. Conversely, many nonreproductive endpoints involve direct genetic contributions to variability between males and females.

undermine their hegemony. Numerous sex differences in neural and behavioral phenotypes, such as sex behavior and the neural underpinnings of ovulation, are largely, if not entirely, sexually differentiated by perinatal gonadal steroid hormones. However, a number of variables that differ in males and females are robustly influenced by sex chromosome complement, with genetic sex exerting influences sometimes as large as those exerted by gonadal hormones. These include sex differences in vasopressin innervation of the lateral septum²⁷, aggressive and parenting behavior²⁹, nociception^{30,31}, formation of habits³², alcohol abuse³³, susceptibility to neural disease^{34,35}, social behaviors^{36,37}, and gene expression^{38–40}. Notably, some sex differences stem from direct effects of X genes that are present in two copies in females and one copy in males^{34,38}, which result in constitutive sex differences in the dose of X genes or their parent of origin^{12,15,41}. Thus at the genetic level, *Sry* is not the only gene causing sex differences in brain phenotype (although this gene acts both indirectly by virtue of its effects on hormone levels and directly because of its expression in brain cells). Rather, various genes encoded on the X or Y chromosome have sex-specific effects, including *Sry* and *Xist* and other yet to be identified X and/or Y genes. Often the phenotypes differentiated by direct sex chromosome effects are also influenced by gonadal hormones, so there is a great need to understand the interaction of sex-specific hormonal and sex chromosome effects. Moreover, these results support the conclusion that every cell in the brain of males may differ from those in females, by virtue of differences in their sex chromosome complement, as well as in response to the important hormonal effects discussed next. Thus, sex differences are likely pervasive in the brain and not limited to a few sexually dimorphic regions. This view is further supported by recent evidence for a substantial sex-specific parental bias in gene expression across brain regions⁴², although the ramifications and importance of sex differences in imprinting are not yet understood.

Parallel hormonal effects on diverse pathways and circuits

The permanent differentiating (organizational) effects of testosterone in the pre- or postnatal brain are often caused by estradiol, a major brain metabolite of testosterone (for a review, see ref. 43). Estrogens act on estrogen receptors to masculinize (enhance behaviors and

functions typical of males) and defeminize (suppress behaviors and functions typical of females). Advances in understanding of steroid-mediated brain differentiation are occurring on two fronts: elucidation of cellular mechanisms of steroid action and downstream effects, and characterization of behavioral and neuronal phenotypes of genetically modified mice. For instance, loss of the estrogen receptor alpha (*Esr1*) results in males with greatly reduced sex behavior, although they retain simple mounting behavior⁴⁴. Knockout of estrogen receptor beta (*Esr2*) alone has no effect on male sex behavior, but when both estrogen receptors are dysfunctional male sex behavior is lost completely⁴⁵. *Esr2* is specifically implicated in the suppression of female sex behavior (defeminization) in males⁴⁶. The importance of estradiol, as opposed to the estrogen receptor, for masculinizing sex behavior is confirmed by the disruption of the gene coding for aromatase, the enzyme required for estradiol synthesis from testosterone⁴⁷. Notably, experiments in aromatase knockout mice show that there is a requirement for estradiol in normal female brain development⁴⁸. Estradiol is unlikely to masculinize males and feminize females at the same site and time; thus, sexual differentiation involves temporally and/or spatially separate estradiol-induced patterns in the two sexes⁴⁹. Disrupting the androgen receptor also predictably impairs male sexual behavior^{50,51}. Thus, the study of mice bearing null mutations for steroid receptors mediating sexual differentiation largely confirms the major conclusions of earlier studies using manipulations of steroid levels or steroid receptors⁵², but also reveals multiple molecular pathways responding to estrogens and androgens in males and females.

The study of mice has been highly informative in parsing out the mechanisms of a major contributor to sex differences in the brain, differential cell death. Many regions and subnuclei in the brain are larger in one sex than in the other. In mammalian males, the spinal cord nucleus, SNB, which contains motoneurons controlling striated muscles of the penis, is larger in males, as are several nuclei in or directly associated with the medial preoptic area of the hypothalamus (MPOA), a major brain region controlling male sexual behavior⁵³. In each instance this is a result of a greater number of neurons surviving through the perinatal period of hormonal sensitivity in males as opposed to high rates of cell death in females. Treatment of females with estrogens or androgens during the sensitive period will rescue the cells from death and result in a permanently masculinized, larger nucleus. Examination of mice lacking the cell death gene, *Bax*, confirms that the higher rate of cell death in several brain nuclei in females results from apoptosis⁵⁴. Conversely, a preoptic ventral forebrain nucleus, the AVPV, is larger in females than in males and is a critical node in the neural circuit controlling ovulation. In this case, estradiol actively promotes cell death in males via a complex mix of classic caspase-3-mediated cell death of the tyrosine hydroxylase-expressing population⁵⁵, combined with an independent program of death in the GABAergic cells mediated by downregulation of TNF α ⁵⁶. This orchestrated killing of discrete populations of neurons in the male AVPV is mediated by the estrogen receptor⁵⁷ and appears to occur in response to estradiol only in cells expressing estrogen receptor. These studies indicate that the sex-specific cellular responses to the same gonadal hormones are different in specific CNS regions (SNB versus AVPV) and involve different molecular pathways in specific cell populations of a single brain region (AVPV).

Studies beyond steroid receptor null mutant mice are needed to elucidate the cellular events downstream of the nuclear steroid receptors, which involve active organization of neural elements and the neuropil. This is illustrated in part by emerging evidence of sex differences in cell proliferation in the rat hippocampus, which is in marked

contrast to the well-documented sex differences in cell death seen in reproductively relevant brain regions. Measures of cell birth indicate that twice as many new cells are born in the male rat hippocampus during the perinatal sensitive period than in the female⁵⁸, and this sex difference is a product of higher endogenous estradiol action in males stimulating neurogenesis⁵⁹. The majority of cells born in the first few days of life will endure until at least the juvenile stage and differentiate into neurons. The overall hippocampal volume is only modestly larger in male rats than in females⁶⁰, suggesting that the enhanced neurogenesis in males may serve purposes other than contributing to increased volume.

In marked contrast to the male bias in neurogenesis in the hippocampus, more new cells are born in the developing amygdala of female rats and, in this instance, those that survive to adulthood largely differentiate into astrocytes. Moreover, sex differences in endocannabinoids mediate the higher rate of cell genesis in females⁶¹. These provocative findings of a role for differential cell birth in brain regions outside those directly involved in reproduction further emphasize the potential for widespread, but region-specific, sex differences.

A third strategy for increasing the size of specific brain region is differentiation of more cells into the phenotype by which that region is defined. The male bed nucleus of the stria terminalis contains more vasopressin-expressing cells than the female nucleus and is therefore larger. Null mutation of *Bax* and overexpression of the anti-cell death gene, *Bcl2*, reveal that, although cell death regulates the overall number of vasopressin neurons in this nucleus, it does not regulate the sex difference. Taken together, these observations suggest, but do not prove, that sex differences in the rate of phenotypic differentiation is the most likely underlying mechanism⁶².

Diverse mechanisms also mediate the estradiol-induced sexual differentiation of synaptic patterning in specific brain regions (Fig. 3). The higher level of estradiol during the perinatal period in males organizes the number and/or density of dendritic spine versus axosomatic synapses, resulting in 2–3-fold differences between males and females in specific nuclei. These are particularly well characterized in preoptic and hypothalamic nuclei such as the MPOA, ventromedial nucleus of the hypothalamus (VMN) and arcuate nucleus⁵². The regional specificity of response to estradiol is evident by comparing the MPOA, where male preoptic neurons have a 2–3-fold greater density of dendritic spine synapses than females⁶, the hypothalamic arcuate nucleus, where female neurons have twice the density of dendritic spine synapses as males⁶³, and the VMN, where males have more overall dendritic spine synapses secondary to the longer and more highly branched dendrites of male VMN neurons⁶⁴. The cellular mechanism organizing the synaptic pattern is also distinct for each nucleus, with estrogen receptor activation invoking unique strategies in each case. In the MPOA, estradiol upregulates the cyclooxygenases genes *COX1* and *COX2* to increase prostaglandin synthesis, activating EP2 and EP4 receptors linked to adenyl cyclase, and activating PKA, which promotes stabilization and membrane insertion of AMPA glutamate receptors^{65,66}. In the arcuate nucleus, estradiol upregulates GAD and GABA production, which act on and differentiate neighboring astrocytes⁶⁷, and estradiol rapidly and nongenomically activates PI3 kinase in the VMN and promotes presynaptic glutamate release, activating postsynaptic AMPA and NMDA receptors^{7,68}.

Although these mechanisms downstream of estradiol have no observable overlap, a common theme is that releasable factors and cell-to-cell communication are important. Estradiol-induced glutamate release alters the synaptic profile of the downstream neuron, with no requirement for estrogen receptor in that neuron. Similarly, GABA released from neurons permanently alters the morphology of

neighboring astrocytes with no requirement for estrogen receptor in those cells. The consequence of emancipation from expression of estrogen receptor to be organized by estrogen receptor is that many more cells are affected both locally and in a domino fashion as one brain region projects to and alters the sexual differentiation of others. Similar cross-cellular effects are seen in the song control circuit in birds, where implants of estradiol near nucleus HVC masculinize nucleus RA downstream of HVC, and lesions of HVC block estradiol's masculinizing effect on nucleus RA^{69,70}. Put more simply, steroid-mediated sexual differentiation of neural circuits is not limited to direct targets of the hormone. Just as every brain cell has a genetic sex, many cell types in specific regions are organized during development by virtue of interactions with other cells in its milieu, so that any information coming into that region is integrated in the context of its sex. This concept argues against the idea that a few steroid-response neurons sit in an otherwise sexually monomorphic brain.

The idea that there are sex-specific circuits may stem in part from the existence of numerous robust and reliable anatomical sex differences in brain regions critical for the expression of sex behavior^{43,53,57}. But consider the nature of the differences. One of the most celebrated, the sexually dimorphic nucleus of the preoptic area (POA), is 3–5-fold larger in males, but is nevertheless present in females, the only difference being that it is smaller than the sexually dimorphic nucleus in males. Similarly, males have 2–3-fold more dendritic spines on POA and VMN neurons, but again, females have plenty of dendritic spines and attendant synapses, just not as many as males. Thus, instead of two distinct neural circuits, it is equally likely that there is only one neural network and that it is differentially weighted toward sex-specific responses as a function of early organization and adult context and hormonal activation.

Recent studies have also found previously unappreciated diversity in the cellular mechanisms of steroid hormone action in the brain. According to the serial model of sexual differentiation of the last century, steroids were viewed as being synthesized by the gonads and acting at sites far from their origin. Steroid receptors such as androgen receptor, estrogen receptor and progesterone receptor are nuclear transcription factor receptors that are capable of directly interacting

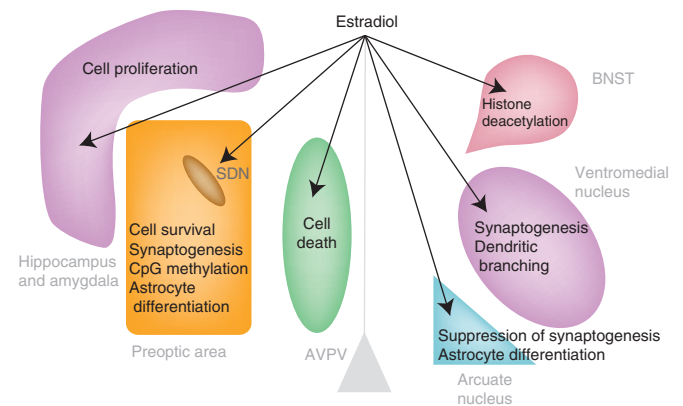


Figure 3 Multiple mechanisms of estradiol-induced differentiation. In the rodent, estradiol is a masculinizing hormone, but it exerts multiple region-specific effects via distinct cellular mechanisms. Thus, during a perinatal sensitive period, the same hormone, estradiol, promotes cell survival, cell death and cell proliferation in separate brain regions. Estradiol also promotes the formation of new dendritic spine synapses in some brain regions while suppressing them in others. The enduring consequences of the organizational effects of estradiol may be mediated in part via epigenetic changes to the DNA and chromatin in processes that are region-specific, but are still poorly understood.

with DNA at specific response elements and modulating gene transcription^{71,72}. Induction of transcription, translation and the construction of new biologically active proteins require time. Combined, these characteristics contributed to the view of steroids as slow mediators of distant cellular processes with long onset and offset. Although that view still has validity, it is equally true that steroids act rapidly on membrane bound receptors, activating signal transduction pathways associated with dynamic changes in cell physiology, including excitability⁷³. Moreover, we now know that steroids, including estradiol, can be synthesized locally, quickly and on demand by neural cells. This is a shift from the concept of steroids as humoral signaling molecules and has led to speculation that estrogens can function in a manner akin to neurotransmitters^{74,75}. The fact that steroids are not stored distinguishes them from neurotransmitters, but places them in a category similar to that of endocannabinoids and gaseous messengers, such as nitrous oxide, that are synthesized on demand⁷⁶. Finely tuned changes in the rate of synthesis and degradation of these signaling molecules regulates their 'tone' and thereby their effect on neural functioning. Responses to external stimuli that induce internal changes, such as activating the stress axis, often involve changes in neuromodulatory tone⁷⁷ and this potential may also exist for steroids. However, the process of steroid-mediated permanent sexual differentiation of the brain was not expected to be modulated by rapidly initiated signaling cascades. That expectation was proved wrong by the discovery of rapid membrane effects of estradiol leading to permanent organization of dendritic morphology in the mediobasal hypothalamus⁷. The precise nature of membrane receptors for estradiol is still being debated, but their distribution both in and between cells appears to be far greater than that of the classic nuclear transcription factor, in part because estradiol can be synthesized in glia and likely has effects on non-neuronal cells that interact with neurons^{78–80}. These findings further

underscore the diversity of cellular responses to steroids and, because of the possibility that these effects are more distributed than effects mediated by nuclear steroid receptors, belie the notion of a limited set of hormonally responsive neurons that are dedicated to the control of sex-specific functions. Nevertheless, there are quite likely some nodes in the brain that are more critical than others in the regulation of sexually dimorphic physiology and behavior, such as the POA for sexual behavior and the AVPV for control of gonadotropin secretion⁵⁷.

Experience matters

The biological sex of a child immediately influences its social and physical environment, even before birth. Our gendered place in society strongly conditions our life history, concept of self, and reaction to social and nonsocial events⁸¹. Parents and teachers create robust sex-specific expectations for children, fostering gendered behavioral development⁸². The different environments for boys and girls contribute to strong sex differences in choice of occupation and other gender-specific environmental stratification⁸³, no doubt contributing to life-long sex differences in social roles, stress and disease. The sex-specific effects of these differentiated social environments, no doubt pervasive and profound, have long been the purview of social psychology and not a topic integral to the study of brain sexual differentiation. In part, this deficit stems from the difficulty of modeling human social environments in animal models. Studies on humans are also problematic because of confounding sex-specific biological and environmental factors, which makes it impossible to disentangle the effects of the two.

This situation is likely to change soon as a result of the emerging discovery of epigenetic modifications of the genome caused by specific environments, which can be measured in model animals. A salient example is the rat dams' differential response to male and

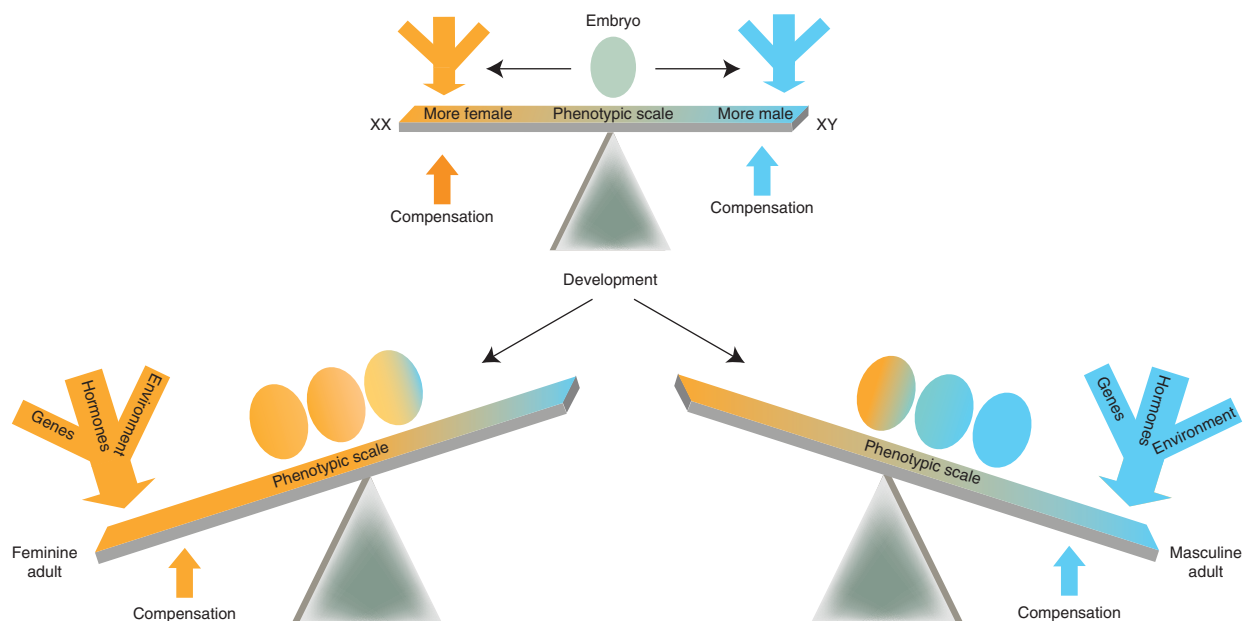


Figure 4 Redefining sexual differentiation. In a twenty-first-century view of sexual differentiation of the brain, the importance of genetics and environment are incorporated along with the effects of hormones to provide a more nuanced portrayal of the types of variables that cause sex differences. Included in this view are the principles that hormones, sex chromosome genes and sex-specific environments have independent parallel differentiating effects that can interact with each other, often synergistically, to cause sex differences in the brain. However, there are also compensatory sex-specific variables that act to reduce sex differences rather than induce them. The result is that some aspects of male and female brain, behavior and physiology are unique from each other, whereas others are highly similar. Two important aspects of the redefined view are not illustrated here: sex differences are pervasive throughout the brain and not restricted to reproductively relevant neural circuits, and variability in the degree to which brain regions are masculinized or feminized in one individual results in a mosaic of relative maleness or femaleness and thereby greatly increases the variance between individuals of the same sex in a population.

female pups. The anogenital grooming of newborn pups is a critical component of maternal care, and variation in the amount of attention a dam gives her offspring has enduring consequences for adult behavior, an effect that is mediated, at least in part, by epigenetic changes. Lasting changes in the responsiveness of the stress axis and behavioral strategies for coping with fear and novelty are correlated with the degree of CpG methylation in the promoter region of the genes coding the glucocorticoid receptor in the hippocampus and estrogen receptor in the POA⁸⁴. Dams distinguish between their male and female pups by providing more anogenital grooming of males than is required for their survival. This not only provides critical somatosensory stimulation needed for normal development of the nerves controlling the penis^{85,86}, but also produces a sex difference in the percentage of CpG methylation of the *Esr1* promoter in the preoptic area, which is correlated with a sex difference in *Esr1* expression later in development⁸⁷.

Another example of sex-specific epigenetic programming is that estradiol aromatized from testicular androgens during the perinatal sensitive period influences the degree of CpG methylation of the *Esr1* promoter⁸⁸, as well as that of *Esr2* and *Pgr*⁸⁹, revealing a potential for steroid hormone feedback on its own sensitivity throughout the lifespan. Sex differences in the epigenetic modulation of histones in specific brain regions⁹⁰ and evidence for an epigenetic underpinning to the differential cell death observed in at least one brain region, the bed nucleus of the stria terminalis⁹¹, further emphasizes the potential for enduring and widespread effects of early hormone exposure via epigenetic modulation. These recent findings foreshadow extensive future work to measure the epigenetic effects of sex-specific environments and biological signals, which likely interact with the hormonal and sex chromosomal control of sexual differentiation. We are only beginning to be able to frame this question and little is known in this area⁹². The combination of experience, steroid hormonal milieu and epigenetic changes represents a convergence point of hormonal and genetic influences on sex differences in the brain that are further modified by individual experience.

Summary: reframed view of brain sexual differentiation

This review presents a new framework for integrating the multiple factors affecting developing brains of males and females (Fig. 4) and highlights several important conclusions drawn from recent studies that alter our concept of sexual differentiation of the brain. First, the sex chromosomes, both X and Y, harbor multiple genes, not just *Sry*, that initiate sexual differentiation. At the genetic level, these are the factors that are the root cause of all sex differences in phenotype. Second, the proximate signals that act directly on brain cells to cause sexual differentiation are not just gonadal hormones, but include other factors, such as those encoded by the sex chromosomes and non-gonadal gene products downstream from sex chromosome genes. Third, different brain regions have different programs of response to the sex-specific signals, involving regional cell type-specific responses, cell-to-cell communication, effects mediated by membrane and nuclear hormone receptors, local steroid synthesis, and compensatory sex-specific effects that antagonize each other and reduce sex differences in phenotype. Changes in gonadal hormone levels over the lifespan and other dynamic changes likely condition sex differences and enhance or suppress them over time. Finally, sex differences in the environment have an enormous effect on gender in humans and are arguably more potent in sculpting the gender-based social phenotype of humans. There is virtually nothing known about the biological basis for these environmental effects, but the rodent literature (epigenetic effects of early stress, differences in maternal

behavior) hints at epigenetic mechanisms that could mediate environmental effects on the brain, throughout the lifespan. We conclude that the long established and mature field of study of sex differences in the brain is as vibrant and dynamic as ever, with many valuable lessons left to be learned.

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