Neural Tube Defects

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Keywords

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ABSTRACT

Neural tube defects (NTDs), including spina bifida and anencephaly are severe birth defects of the central nervous system that originate during embryonic development if the neural fails to completely close. Human NTDs are multifactorial, with contribution of both genetic and environmental factors. The genetic basis is not yet well understood but several non-genetic risk factors have been identified as well as the possibility for prevention by maternal folic acid supplementation. Mechanisms underlying neural tube closure and NTDs may be inferred from experimental models, which have revealed numerous genes whose loss of function causes NTDs, as well as details of critical cellular and morphological events whose regulation is essential for closure. Such models also provide an opportunity to investigate potential risk factors and to develop novel preventive therapies.

1. INTRODUCTION

Neural tube defects (NTDs) are severe birth defects of the central nervous system that originate during embryogenesis and result from failure of the morphogenetic process of neural tube closure. In higher vertebrates the neural tube is generated by shaping, bending and fusion of the neural plate and fusion in the dorsal midline progressively seals the neural tube as it forms. If closure is not completed, neuroepithelium remains exposed to the environment and consequently subject to degeneration and neuronal deficit. The type and severity of these 'open' NTDs varies with the level of the body axis which is affected. Thus, failure of closure in the prospective brain or spinal cord result in anencephaly and open spina bifida (myelomeningocele), respectively.

While the unifying feature of open NTDs is the failure of completion of neural tube closure, there are many different possible causes, both genetic and environmental. In humans, it appears that most NTDs are multifactorial in causation, resulting from an additive contribution of several risk factors which are each individually insufficient to disrupt neural tube closure (the multifactorial threshold model)(Harris & Juriloff 2007). The challenge of identifying the primary cause of NTDs in individual patients is highlighted by the numerous candidate genes and environmental factors indicated by epidemiological studies and experimental models. Moreover, the potential for gene-gene and gene-environment interactions introduces further potential complexity.

2. UNDERSTANDING THE EMBRYONIC BASIS OF NTDS - NEURAL TUBE CLOSURE

Determination of the specific causes of NTDs is best achieved in the context of an understanding of the mechanisms underlying neural tube closure (reviewed by (Copp & Greene 2013, Greene & Copp 2009). Given the inaccessibility of the neurulation-stage human embryo, our knowledge of the key principles of neural tube closure comes mainly from analysis of experimental models, particularly other mammals, amphibians and birds, in which primary neural tube closure is achieved through folding and fusion of the neuroepithelium.

2.1 Primary neurulation; sub-types of NTDs relate to stages of closure

In the prospective brain and most of the spinal cord, neural tube formation essentially involves the bending of the neuroepithelium in the midline to generate neural folds that elevate, meet and fuse in the dorsal midline (primary neurulation). Rather than simultaneously rolling up along the extent of the rostro-caudal axis, neural tube closure is discontinuous with distinct sites of initiation located at characteristic axial levels. Moreover, the morphological and molecular requirements for closure vary along the body axis, such that an individual NTD usually only affects a portion of the neural tube.

NTDs can thus be attributed to failure of particular initiation events or disruption of the progression of closure between these sites.

In mice, closure is first achieved on embryonic day 8.5 at the level of the hindbrain/cervical boundary (Closure 1) and failure of this event leads to craniorachischisis (Copp et al. 2003). Closure initiates at a second site on embryonic day 9, Closure 2, in the caudal forebrain or forebrain/midbrain boundary. Once initial contact and fusion have been established between the tips of the neural folds, closure spreads bi-directionally from the sites of Closure 1 and 2 and in a caudal direction from the rostral end of the neural tube (Closure 3). The open regions of neural folds, termed neuropores, gradually shorten leading to complete closure of the anterior neuropore (between Closures 2 and 3) on embryonic day 9, and the hindbrain neuropore (between Closures 1 and 2) a few hours later. Cranial NTDs (anencephaly) result from failure of Closure 2, or incomplete 'zippering' between Closures 1 and 2, which closes the midbrain and hindbrain. If fusion does not progress from the anterior end of the neural plate (Closure 3), the resultant phenotype is a 'split face' usually accompanied by forebrain anencephaly.

Unlike the cranial region where closure proceeds bidirectionally, spinal neurulation is entirely caudally directed as the embryo continues to grow. Primary neurulation completes with final closure of the posterior neuropore on embryonic day 10. Impaired progression of closure, and consequently the presence of a persistently open posterior neuropore, results in spina bifida and the size of the ensuing lesion relates directly to the axial level at which closure stops.

2.2 Primary neurulation in humans

Examination of human embryos suggests that initiation of closure is discontinuous, as in the mouse (Nakatsu et al. 2000, O'Rahilly & Müller 2002). Bending of the neural plate begins at around 17-18 days after fertilisation, with an equivalent event to Closure 1 at around 22 days and completion of closure at the posterior neuropore by 26-28 days post-fertilisation. It appears that closure of the forebrain and midbrain in human embryos may be achieved by progression between the site of Closure 1 and the rostral end of the neural plate without an intervening initiation site analogous to Closure 2 (O'Rahilly & Müller 2002, Sulik et al. 1998).

2.3 Secondary neurulation

In mice and humans, the neural tube caudal to the mid-sacral region is continuous with the caudal end of the primary neural tube but forms by a distinct process, termed secondary neuralation

(Schoenwolf 1984, Copp & Brook 1989). This process involves condensation of a population of tail bud-derived cells bud to form an epithelial rod that undergoes canalisation to form the lumen of the tube in the lower sacral and coccygeal regions. Malformations resulting from disturbance of secondary neurulation are 'closed' (skin covered) and often involve tethering of the spinal cord, with associated ectopic lipomatous material (Lew & Kothbauer 2007).

3. MECHANISMS UNDERLYING NEURAL TUBE CLOSURE

Studies of neurulation-stage embryos, both normal and developing NTDs, provide insights into key molecular and cellular pathways underlying the morphological tissue movements of neural tube closure (Copp & Greene 2010). In addition to ubiquitous requirements, the occurrence of isolated NTDs at cranial or caudal levels in humans and different mouse models suggests the likely involvement of region-specific mechanisms, dependent on different gene products.

3.1 Shaping of the neural plate – convergent extension is required for initiation of closure

Concomitant with the onset of neural tube closure, the neural plate undergoes narrowing in the medio-lateral axis (convergence) and elongation in the rostro-caudal axis (extension), owing to intercalation of cells at the midline (Keller 2002). Convergent extension depends on activity of a non-canonical Wnt signalling pathway, homologous to the planar cell polarity (PCP) pathway first described in *Drosophila* as regulating cell polarity in the plane of epithelia (Goodrich & Strutt 2011). Signalling occurs via a Frizzled (Fzd) membrane receptor and cytoplasmic Dishevelled (Dvl), but without stabilisation of beta-catenin.

Functional disruption of PCP mediators prevents convergent extension and the neural plate remains broad in *Xenopus* (Wallingford & Harland 2001, Wallingford & Harland 2002) and mouse embryos (Greene et al. 1998, Ybot-Gonzalez et al. 2007). Hence, closure 1 fails, leading to craniorachischisis, in mice homozygous for mutations in 'core PCP' genes including *Vangl2*, *Celsr1*, or double mutants for *Dvl-1* and *-2*, or *Fzd-3* and *-6* (Juriloff & Harris 2012). Craniorachischisis also results from mutation of the PCP-related genes *Scrb1* (Murdoch et al. 2001) and *Ptk7* (Lu et al. 2004) or genes encoding accessory proteins, such as *Sec24b* which affects Vangl2 transport (Merte et al. 2010). Ultimately, failure of closure initiation in PCP-mutant embryos is thought to result from insufficient proximity of the neural folds owing to the broadened midline.

Failure of closure 1 in the majority of 'core' PCP mutant embryos precludes analysis of a requirement for convergent extension at later stages of neurulation. However, spina bifida occurs in some *loop*-

tail heterozygotes (*Vangl2*^{Lp/+})(Copp et al. 1994) and in compound heterogotes of *Vangl2*^{Lp/+} with mutations of *Ptk7*, *Sec24b* or *Sdc4* (Lu et al. 2004, Merte et al. 2010, Escobedo et al. 2013). Moreover, non-canonical Wnt signalling is compromised in *Lrp6* null embryos that develop spina bifida (Gray et al. 2013). These observations suggest a likely continued requirement for PCP signalling as spinal neurulation proceeds.

Despite the entirely open spinal neural tube in *Vangl2^{Lp/Lp}* embryos with craniorachischisis, closure does occur in the forebrain and much of the midbrain implying that PCP-dependent convergent extension is not required throughout the cranial region. Nonetheless, exencephaly is observed in digenic combinations of *Vangl2^{Lp/+}* with some Wnt pathway genes (e.g. *Dvl3^{+/-}*, *Fzd1^{+/-}* and *Fzd2^{+/-}* (Etheridge et al. 2008, Yu et al. 2010). Exencephaly also develops in mutants for the PCP 'effector' genes *Fuz* or *Intu* but the role of these genes in cilium-dependent hedgehog signalling seems more likely to explain their loss-of-function effect on cranial neural tube closure than a role in regulating convergent extension (Gray et al. 2009, Zeng et al. 2010, Heydeck & Liu 2011) (see Section 3.2). Thus, components of PCP signalling potentially impact on neural tube closure via multiple cellular mechanisms.

3.2 Bending of the neural folds – regulation by Shh and BMP signalling

In order to achieve closure, the neuroepithelium must bend to bring the tips of the neural folds into apposition. Bending occurs in a stereotypical manner at 'hinge points'; a median hinge point (MHP) in the midline and paired dorsolateral hinge points (DLHPs) that arise laterally (Shum & Copp 1996). The morphology varies along the body axis with differing modes in the upper (MHP only), mid- (MHP and DLHPs) and caudal (DLHPs only) regions of the primary neural tube.

The mechanisms underlying neuroepithelial bending are not fully understood, but one notable feature of the MHP is the predominance of wedge-shaped cells (wider basally than apically) compared to non-bending regions (Schoenwolf & Smith 1990). At neural plate stages the neuroepithelium is a pseudostratified epithelium in which nuclei move to the basal pole during Sphase owing to inter-kinetic nuclear migration. Prolongation of Sphase at the MHP provides a possible means by which regulation of the cell cycle may contribute to cell wedging and hence MHP-formation (Schoenwolf & Smith 1990).

Bending is regulated by signals emanating from non-neural tissues dorsal and ventral to the neural folds (reviewed by (Greene & Copp 2009). The MHP is induced by signals from the notochord, located immediately ventral to the midline of the neuroepithelium (Smith & Schoenwolf 1989, Ybot-

Gonzalez et al. 2002). At the molecular level, notochord-derived Shh induces the floor plate of the neural tube at the site of the MHP (Placzek & Briscoe 2005, Chiang et al. 1996). However, this is not essential for spinal neural tube closure which completes in the absence of a floor plate in mouse embryos lacking Shh or Fox A2 (Chiang et al. 1996, Ang & Rossant 1994). Thus, the MHP may be functionally important in floor plate development but is not essential for neural tube closure.

In contrast to the MHP, DLHPs appear essential for closure of the neural tube in the low spinal region. For example, *Zic2* mutant embryos, in which DLHPs are absent, develop severe spina bifida (Ybot-Gonzalez et al. 2007). The formation of DLHPs is actively regulated, with interplay of inhibitory and inductive signals determining their appearance at different axial levels (Copp & Greene 2013). These include inhibitory effects of Shh signalling from the notochord and BMP signalling from the surface ectoderm at the dorsal tips of the neural folds. These signals are opposed by the BMP antagonist noggin whose expression in the dorsal neural folds is sufficient to induce DLHP (Ybot-Gonzalez et al. 2002, Ybot-Gonzalez et al. 2007).

In contrast to absence of Shh signalling, NTDs do result from mutations which enhance Shh signalling, for example through deficient function of inhibitory or cilia-related genes such as *Gli3*, *Rab23*, *Fkbp8*, *Tulp3* and *Ift40* (Murdoch & Copp 2010, Miller et al. 2013). Mutants involving increased Shh signalling display NTDs at cranial and/or spinal levels. While spina bifida appears to be associated with suppression of dorsolateral bending of the neural folds (Murdoch & Copp 2010), the mechanism underlying cranial NTDs is not clear.

3.3 Cranial neurulation – additional complexity and sensitivity to disruption

The neural folds in the cranial region bend in the midline and dorsolaterally as in the spinal region but the closure process appears morphologically more complex. The folds are initially biconvex, with the tips facing away from the midline, and then switch to a biconcave shape allowing the tips to approach in the midline. The additional complexity of cranial compared with spinal neurulation appears to be reflected in a more extensive genetic underpinning and a greater sensitivity to disruption, at least in rodents. Exencephaly occurs in approximately three times as many knockout mouse models as spina bifida and is the NTD type most commonly induced by teratogens (Copp et al. 1990, Harris & Juriloff 2010).

Cranial neurulation may rely on specific contributory factors that are not involved in the spinal region such as expansion of the mesenchyme underlying the neural folds (Greene & Copp 2009,

Zohn & Sarkar 2012). Moreover, disruption of the actin cytoskeleton prevents closure in the cranial but not the spinal region (Morriss-Kay & Tuckett 1985, Ybot-Gonzalez & Copp 1999). Similarly, exencephaly is observed but spinal neurulation completes successfully in null mutants for several cytoskeletal components (e.g. *n-cofilin*, *vinculin*) (Gurniak et al. 2005, Xu et al. 1998). Nevertheless, apically-located actin microfilaments are present throughout the neuroepithelium (Sadler et al. 1982), while functional disruption of the cytoskeleton—associated proteins *MARCKS-related protein* or *Shroom3* cause both spinal and cranial NTDs (Hildebrand & Soriano 1999, Xu et al. 1998), suggesting that regulation of the acto-myosin cytoskeleton plays a role in closure in both regions. Shroom proteins appear to play a key role: expression of Shroom in *Xenopus* is sufficient to induce apical constriction of epithelial cells while functional disruption inhibits neural fold bending and suppresses closure (Haigo et al. 2003).

3.4 Adhesion and fusion of the neural folds

Once the neural folds meet at the dorsal midline, a process of adhesion, fusion and remodelling gives rise to two discrete epithelial layers, with the nascent neural tube overlain by an intact surface ectoderm (Pai et al. 2012). At the closure site the neural fold tips are composed of neuroepithelium continuous with the non-neural surface ectoderm. The cell type that adheres first may differ at varying axial levels (Geelen & Langman 1979, Ray & Niswander 2012). Nevertheless at all levels, initial contact appears to involve sub-cellular protrusions, resembling lamellipodia and filopodia, observed by electron microscopy (Geelen & Langman 1979) and in live embryos (Pyrgaki et al. 2010). The molecular basis of adhesion is not well characterised, perhaps due to functional redundancy among the proteins involved. However, a role for interaction of cell surface ephrin receptors with Eph ligands is suggested by the occurrence of cranial NTDs in mice lacking ephrin-A5 or EphA7 (Holmberg et al. 2000), and delayed spinal closure in embryos exposed to peptides that block ephrinA/EphA interactions (Abdul-Aziz et al. 2009).

Knockout of protease-activated receptors (PAR1 and PAR2) in the surface ectoderm also causes cranial NTDs, implicating a role for signalling via these G-protein coupled receptors in closure (Camerer et al. 2010). Further evidence for the function of the non-neural ectoderm is provided by *Grhl2* null mutants which fail in closure throughout the cranial region and exhibit spina bifida (Rifat et al. 2010, Werth et al. 2010, Brouns et al. 2011). *Grhl2* is expressed in the surface ectoderm overlying the neural folds and regulates expression of several components of the apical adhesion junction complex, including E-cadherin (Werth et al. 2010, Pyrgaki et al. 2011).

3.5 Regulation of cell proliferation and cell death

During neurulation the embryo grows rapidly. Cell cycle exit and neuronal differentiation begin in the neuroepithelium shortly after closure and maintenance of adequate proliferation in the neuroepithelium appears crucial for closure, particularly in the cranial region. Thus, in mice NTDs can be caused by exposure to anti-mitotic agents (Copp et al. 1990) or mutation of genes encoding proteins associated with cell-cycle progression (e.g. *neurofibromin 1, nucleoporin*) or prevention of neuronal differentiation (e.g. Notch pathway genes *Hes1*, *Hes3*, *RBP-J\kappa*)(Harris & Juriloff 2010, Harris & Juriloff 2007). Conversely, excessive cell proliferation is also associated with NTDs in several mouse models, such as *Phactr4* mutants (Kim et al. 2007).

Characteristic patterns of apoptotic cell death occur in the neural folds and the midline of the closed neural tube (Geelen & Langman 1979, Massa et al. 2009, Yamaguchi et al. 2011). Increased cell death could hypothetically inhibit closure through compromising the functional and/or mechanical integrity of the neuroepithelium. It is associated with NTDs in a number of teratogen-induced and genetic models, although only rarely has a direct causal link been definitively established (Copp & Greene 2013, Fukuda et al. 2011). The occurrence of exencephaly in mice lacking apoptosis-related genes such as *caspase3* or *Apaf1* suggests a requirement for apoptosis in closure (Harris & Juriloff 2010). However, forebrain and spinal closure occurs normally in these models and pharmacological suppression of apoptosis does not cause NTDs, suggesting that it is dispensable for completion of closure (Massa et al. 2009).

4. CLINICAL FEATURES OF NEURAL TUBE DEFECTS

4.1 Open NTDs and associated conditions

Open NTDs can result from failure of closure at a *de novo* initiation site or incomplete progression of closure following successful initiation. Where embryos are available for examination, as in experimental models, NTDs can be recognised during or immediately after neurulation stages owing to the persistently open neural folds. However, at later embryonic and fetal stages the morphological appearance varies considerably owing to secondary changes and degeneration.

In cranial NTDs, the open neural folds undergo growth and differentiation and typically appear to bulge from the developing brain, termed *exencephaly*. Inability to form the skull vault over the open region leads to degeneration of the exposed neural tissue and the characteristic appearance of

anencephaly, observed later in human or rodent pregnancy (Wood & Smith 1984, Seller 1995). Both anencephaly and craniorachischisis (~10% of NTDs) are lethal conditions at or shortly after birth.

Open neural folds in the spinal region prevent the sclerotome-derived vertebral arches from covering the neuroepithelium, the consequent opening in the vertebral column giving rise to the term *spina bifida* (Copp et al. 2013). The neural tissues may be contained within a meninges covered sac that protrudes through the open vertebrae (myelomeningocele; spina bifida cystica), or exposed directly to the amniotic fluid (myelocele). Babies born with open spina bifida usually survive with appropriate medical care, but suffer neurological impairment whose severity depends on the level of the lesion. Associated conditions include hydrocephalus, Chiari type II malformation and vertebral abnormalities as well as genitourinary and gastrointestinal disorders.

4.2 Diagnosis, treatment and maternal-fetal surgery

NTDs can be diagnosed prenatally by ultrasound (Cameron & Moran 2009). However, where prenatal diagnosis is not routinely available and/or therapeutic abortion not an option, many babies with NTDs are born. Post-natal medical care for babies born with open spina bifida usually involves surgery to close and cover the lesion. Multiple subsequent surgeries are commonly required to alleviate tethering of the spinal cord, treat hydrocephalus and/or address orthopaedic and urological problems.

As open NTDs arise early during pregnancy, there is a prolonged period during which secondary neurological damage may occur owing to exposure of nervous tissue to the amniotic fluid environment. These considerations provided impetus for development of *in utero* fetal surgery for spina bifida which may improve neurological outcome compared with post-natal repair, although with fetal and maternal risks (Adzick et al. 1998, Adzick et al. 2011). Experimental models of spina bifida are being used to investigate the possible combination of surgical intervention with additional therapy, intended to remediate neural damage. Examples include the implantation of biodegradable scaffolds to promote neural regeneration and/or neural stem cells to populate the damaged spinal cord (Saadai et al. 2011, Saadai et al. 2013).

4.3 Disorders of the closed neural tube

This review focuses on open NTDs, characterised by failure of neural tube closure. Various other conditions are also associated with abnormalities of the closed spinal cord and are often categorised as NTDs under a broader definition. There is also a less well-defined group of closed spinal NTDs in

which the vertebral arches are malformed but covered by skin. These conditions, including spina bifida occulta and 'spinal dysraphisms', vary widely in clinical presentation. The more severe subtypes are associated with various abnormalities of the spinal cord, lipoma and/or anorectal abnormalities. The embryonic origin of closed spina bifida is not well defined but is hypothesised to involve abnormalities of secondary neurulation (Copp et al. 2013).

Abnormal development of the vertebrae or cranium may also allow herniation of the closed neural tube through the affected region in the rare form of spina bifida, meningocele (spina bifida cystica) or encephalocele, respectively.

5. CAUSES OF NTDs

NTDs are among the most common birth defects worldwide with a prevalence that varies from 0.5 to more than 10 per 1,000 pregnancies. This likely reflects differing contributions from risk factors such as nutritional status, prevalence of obesity and diabetes, usage of folic acid supplementation and/or fortification, the presence of environmental toxicants and differing genetic predisposition between ethnic groups. In most populations there is also a striking gender bias with a higher prevalence of anencephaly among females than males. Many NTD mouse strains also show a female preponderance among cranial NTDs, apparently reflecting a fundamental higher sensitivity of cranial neural tube closure to disturbance in female embryos (Juriloff & Harris 2012). Overall, although a number of risk factors have been identified these may account for less than half of NTDs, suggesting that additional genetic and non-genetic factors remain to be identified (Agopian et al. 2013).

5.1 Environment factors

Various teratogenic agents induce NTDs in rodent models (Copp et al. 1990, Copp & Greene 2010). In humans, teratogens that have been associated with NTDs include the anti-convulsant drug valproic acid (Wlodarczyk et al. 2012), and the fungal product fumonisin (Missmer et al. 2006). Other non-genetic risk factors include maternal fever and excessive use of hot tubs (Moretti et al. 2005), consistent with the induction of NTDs by hypothermia in rodent models.

Maternal obesity or diabetes are well-recognised risk factors for NTDs (Correa et al. 2003). Determination of the cause of diabetes-related NTDs is hampered by the complexity of the diabetic milieu, although hyperglycemia alone is sufficient to cause NTDs in cultured rodent embryos. It has been proposed that NTDs may result from increased oxidative stress, altered expression of genes such as *Pax3*, and neuroepithelial cell apoptosis (Fine et al. 1999, Reece 2012). Recent findings

suggest that activation of apoptosis signal-regulating kinase 1 (ASK1) in hyperglycaemic conditions, leads to activation of the apoptosis mediator caspase 8 via stimulation of the FoxO3a transcription factor (Yang et al. 2013).

Nutritional factors and folate

The historical link between lower socioeconomic status and higher risk of birth defects led to examination of the possible involvement of nutritional factors in NTDs. Lower levels of the B-vitamin folate were observed in mothers of NTD fetuses (Smithells et al. 1976), prompting an intervention trial of a folic acid-containing multivitamin supplement for prevention of NTD recurrence (Smithells et al. 1981, Schorah 2008). A multi-centre randomised controlled trial confirmed that maternal folic acid supplementation (at 4 mg/day) significantly reduces the recurrence risk (Wald et al. 1991). Additional clinical trials provided evidence for reduction of occurrence risk (Czeizel & Dudás 1992, Berry et al. 1999, Czeizel et al. 2011).

Questions remain over the mechanism by which folic acid prevents NTDs (Blom et al. 2006, Copp et al. 2013). Although maternal folate status is a risk factor, in most cases, maternal folate levels are within the 'normal' range and rarely clinically deficient. Nonetheless, there is an inverse relationship between blood folate concentration and risk of an affected pregnancy (Daly et al. 1995). It has been suggested that sub-optimal folate levels may contribute to development of NTDs in individuals who are genetically susceptible. Such a gene-environment interaction has been demonstrated in mice, where folate deficiency does not cause NTDs, unless present in combination with mutation of a predisposing gene, such as *Pax3* (Burren et al. 2008).

Folate one-carbon metabolism comprises a complex network of inter-linked reactions that mediate transfer of one-carbon groups for a number of biosynthetic processes (Stover 2009). Among these, attention has particularly focussed on the requirement for nucleotide biosynthesis and methylation reactions in neural tube closure. Abnormal thymidylate and purine biosynthesis have been identified in mouse NTD models (Fleming & Copp 1998, Beaudin et al. 2011) and in a proportion of NTD cases (Dunlevy et al. 2007), while deficient methylation may also be implicated in NTDs (Section 5.3).

5.2 Genetics of NTDs

Most NTDs occur sporadically, with a relative scarcity of multi-generational families. Nevertheless, there is strong evidence for a genetic component in the etiology of NTDs and the pattern of inheritance favours a multifactorial polygenic or oligogenic model, as opposed to an effect of single

genes with partial penetrance (Harris & Juriloff 2007). Most studies of NTD genetics have focussed on one or more candidate genes (reviewed by (Boyles et al. 2005, Greene et al. 2009, Harris & Juriloff 2010). In general these have been: (i) human orthologues of genes whose mutation causes NTDs in mice, of which there are more than 200 examples; or (ii) genes related to environmental risk factors, particularly folate metabolism.

Case-control association studies have implicated several genes while mutation screening by sequencing has identified putative pathogenic mutations. However, the definitive assignment of a gene variant as causative is complicated by the apparent multigenic nature of NTDs, and the large number of possible candidate genes, modifier genes, epigenetic factors and environmental influences. Moreover, where putative mutations have been identified in specific genes, each has only been involved in a small proportion of NTD patients, suggesting that there is considerable heterogeneity underlying the genetic basis of NTDs. Thus, although the morphological and cellular basis of neural tube closure has become increasingly well understood, the genetic basis of NTDs in individual cases remains largely unclear.

Gene-gene interactions and effect of modifier genes

Mouse studies suggest three broad mechanisms by which genetic interactions may result in NTDs. (1) In some instances functional redundancy makes it necessary for mutation of two orthologous genes, (e.g. *Dvl1-Dvl2* (Hamblet et al. 2002), *Cdx1-Cdx2* double knockouts (Savory et al. 2011)), in order to reveal a requirement in neural tube closure. (2) Additive effects of heterozygous mutations may result in NTDs that resemble those of individual homozygotes (e.g. *Dvl3* with *Vangl2^{Lp}* (Etheridge et al. 2008). (3) Variation in the penetrance and expressivity of NTD phenotypes between inbred strains of mice is widely reported and thought to reflect variants in modifier genes. For example, the rate of exencephaly resulting from *Cecr1* mutation is strongly affected by FVB/N strain background (Davidson et al. 2007). While the identity of modifier genes for NTDs has rarely been determined, a variant in *Lmnb1* is present in some mouse strains and significantly increases the frequency of NTDs in *curly tail* (*Grhl3^{ct}*) embryos (de Castro et al. 2012).

Genes implicated through experimental models

In mice, mutation of genes encoding components of the PCP pathway causes NTDs (Section 3.1). Sequencing of PCP genes in humans has identified putative mutations in *CELSR1*, *VANGL1*, *VANGL2*, *FZD6*, *SCRIB1* and *DVL2* in a proportion of patients with craniorachischisis, spina bifida, anencephaly and closed forms of spina bifida (Kibar et al. 2007, Lei et al. 2010, Robinson et al. 2012, De Marco et

al. 2013, Chandler et al. 2012, Lei et al. 2013) and reviewed by (Juriloff & Harris 2012). As in mice, heterozygous human PCP mutations may hypothetically interact with other genetic NTD risk factors in a digenic or polygenic fashion, to cause a range of NTD types. This could potentially involve summation of multiple variants in PCP genes. For example, a putative mutation in *DVL2* was identified in a spina bifida patient in combination with a second, previously identified missense variant in *VANGL2* (De Marco et al. 2013).

Among other genes implicated in NTDs from mouse models, association studies have not provided evidence for a major contribution to risk and few positive results have emerged from sequencing-based mutation screens. As data begins to emerge from large-scale exome sequencing studies of NTD patients, it will become possible to evaluate the contribution of multiple genes in the same patient cohorts and the mutational load associated with individual risk.

Analysis of genes related to environmental risk factors

The identification of environmental factors such as maternal diabetes and folate status as risk factors for NTDs provides impetus for analysis of related genes in affected families. Risk could potentially be associated with maternal genotype, if genetic variation alters maternal metabolism and secondarily affects the developing embryo. The inheritance of maternal alleles by the embryo complicates interpretation of such effects. Alternatively, a genetically determined abnormality in the embryo itself could influence risk of NTDs; potentially through interaction with a predisposing environmental factor. For example, it may be informative to analyse genetic data on folate-related genes in the context of maternal folate status (Etheredge et al. 2012).

Association with risk of spina bifida has been reported for several genes implicated in diabetes, obesity, glucose metabolism and oxidative stress, including *GLUT1*, *SOD1* and *SOD2* (Davidson et al. 2008, Kase et al. 2012). Maternal variants in the obesity-related genes *FTO*, *LEP* and *TCF7L2* are also associated with NTDs, consistent with maternal obesity being a risk factor (Lupo et al. 2012).

Genes related to folate one-carbon metabolism have been perhaps the most intensively group of candidates for NTDs (reviewed by (Blom et al. 2006, Greene et al. 2009, Shaw et al. 2009)). The C677T polymorphism of *MTHFR*, which encodes an alanine to valine substitution, has been associated with NTDs. The TT genotype is found at higher frequency among cases than controls in some populations (e.g. Irish) but not others (e.g. Hispanics) (Botto & Yang 2000). Several studies

indicate positive associations with other folate-related genes, including *MTRR*, although these have generally not been not observed in all study populations.

In mice, mutations in folate-metabolising enzymes (e.g. *Mthfd1*) are sometimes lethal before the stage of neural tube closure (e.g. (MacFarlane et al. 2009, Christensen et al. 2013) while others do not disrupt closures (eg. (Chen et al. 2001, Di Pietro et al. 2002). Null embryos for the folate receptor, *Folr1* die pre-neurulation but develop NTDs when supplemented with sufficient folic acid to prevent early lethality (Piedrahita et al. 1999). NTDs are also observed in *Shmt1* knockouts, under folate-deficient conditions (Beaudin et al. 2011). In contrast, NTDs occur 'spontaneously' in mice carrying loss-of-function alleles of *Amt* (Narisawa et al. 2012) or *Mthfd11* (Momb et al. 2013), both of which encode enzymes of mitochondrial folate metabolism (Tibbetts & Appling 2010). Interestingly, the homologous genes in humans has also been linked to NTDs. Missense mutations have been identified in NTD patients in *AMT*, as well as *GLDC* which encodes its partner enzyme in the glycine cleavage system (Narisawa et al. 2012). Genetic associations with NTDs have been reported for *MTHFD1L* (Parle-McDermott et al. 2009) and *SLC23A32* (*MFTC*), encoding a mitochondrial folate transporter (Pangilinan et al. 2012). Altogether, these findings suggest that NTD risk is influenced by function of mitochondrial folate metabolism, a major source of one-carbon units to the cytoplasm.

5.3 Gene-regulatory mechanisms and NTDs

In addition to the potential multigenic nature of NTDs, identification of causative genes may be complicated by the potential involvement of aberrant gene expression, perhaps resulting from mutations in regulatory elements. For example, mutations resulting in insufficient expression of *Grhl3* or excess expression of *Grhl2* cause NTDs in mice in the absence of coding mutations (Gustavsson et al. 2007, Brouns et al. 2011). Further complexity may be added by the potential for regulation by epigenetic modifications such as DNA methylation, histone modification or chromatin remodelling, each of which has been associated with NTDs in mice and in some cases in humans (reviewed by (Harris & Juriloff 2010, Greene et al. 2011)). For example, methylation of LINE-1 genomic elements was lower than normal in DNA of anencephalic but not spina bifida fetuses (Wang et al. 2010).

A simple model predicts a positive correlation between folate status and methylation. However, data from human pregnancy suggests the relationship is not straightforward (Crider et al. 2012). A recent study found an inverse correlation of LINE-1 methylation with maternal and cord blood folate, while different imprinted genes showed positive or negative associations (Haggarty et al.

2013). Somewhat counter-intuitively use of folic acid supplements was associated with reduced LINE-1 methylation.

A requirement for DNA methylation in mouse neural tube closure is suggested by the occurrence of NTDs in knockouts of *Dnmt3b*, encoding a DNA methyltransferase, and in embryos cultured with 5-azacytidine (Okano et al. 1999, Matsuda & Yasutomi 1992). Similarly, inhibition of the methylation cycle reduces DNA methylation and causes NTDs in cultured mouse embryos (Dunlevy et al. 2006, Burren et al. 2008). However, *Mthfr* null embryos do not develop NTDs despite a significant reduction in global DNA methylation (Chen et al. 2001), nor is there an exacerbating effect of *Mthfr* loss-of-function on *Pax3* or *curly tail* mutants, although both show increased rates of NTDs under folate-deficient conditions (Pickell et al. 2009, Burren et al. 2008, de Castro et al. 2010). Thus, questions remain over the relationship between folate status, DNA methylation and risk of NTDs.

Other epigenetic mechanisms include various modifications of histone proteins, which potentially miss-regulate genes that influence neurulation. NTDs occur in mice carrying mutations in the histone demethylases *Jarid2* (Takeuchi et al. 1999) and *Fbxl10* (Fukuda et al. 2011). Similarly, histone acetylases and deacetylases, which regulate the equilibrium of histone acetylation, are implicated in NTDs. An acetylase-specific knock-in mutation of *Gcn5* causes cranial NTDs (Bu et al. 2007), as does loss of function of another histone acetylase, p300 (Yao et al. 1998). Increased acetylation is also associated with NTDs. For example, cranial NTDs occur in mice carrying mutations in histone deacetylases *Sirt1* or *Hdac4* (Cheng et al. 2003, Vega et al. 2004). The teratogenic effects of valproic acid and trichostin A may also be mediated through their inhibition of histone deacetylases (Finnell et al. 2002).

6. PRIMARY PREVENTION OF NTDs

6.1 Folic acid supplementation and fortification

The reduction in risk of NTDs following maternal folic acid supplementation led to public health recommendations that women who may become pregnant should consume 0.4 mg of folic acid daily or 4 mg daily following a previous affected pregnancy (Czeizel et al. 2011). To ensure that additional folate was received, food fortification programmes were introduced in many countries. This approach has raised blood folate levels and been associated with lower frequency of NTDs (Crider et al. 2011). The magnitude of effect varies, with greatest reduction where pre-existing rates were higher (Blencowe et al. 2010, Rosenthal et al. 2013). Some countries have delayed decision on fortification owing to safety concerns (e.g. possible enhancement of bowel cancer) but a recent

meta-analysis found no evidence for increased cancer rates following folic acid supplementation (Vollset et al. 2013).

6.2 Folate-resistant NTDs

Folic acid supplementation in clinical trials has not approached 100% NTD prevention and an estimated one-third of NTDs may be folic acid-resisant (Blencowe et al. 2010). A study in the USA, where folate fortification of food is mandatory, found no apparent protective effect of folic acid supplements (Mosley et al. 2009), suggesting that increased dosage would not necessarily provide additional preventive effects.

Given the multifactorial causation of NTDs it seems reasonable to suppose that optimal prevention will require a combination of multiple interventions. Possible approaches may relate to folate one-carbon metabolism. For example, like folate there is a graded relationship between lower levels of circulating vitamin B_{12} and increasing risk of an NTD-affected pregnancy (Molloy et al. 2009). Perhaps use of B_{12} supplements would further reduce the frequency of NTDs, although this remains to be tested.

Another possibility is that folic acid cannot ameliorate some defects that result from abnormal folate metabolism, owing to defects in the intervening enzymes required to transfer one carbon units to key downstream metabolites. In this case supplementation with alternative folates, such as 5-methyl THF (Czeizel et al. 2011), or key downstream molecules could potentially be advantageous. For example, supplementation with formate prevented NTDs in *Mthfd1L* null mice (Momb et al. 2013), while combinations of thymidine and purine precursors prevented NTDs in *curly tail* mice, in which folic acid is not protective (Leung et al. 2013).

In addition to folate and vitamin B_{12} , lower maternal levels of other vitamins, including vitamin C, have been reported in NTDs (Smithells et al. 1976). Conversely, intake of several vitamins and maternal diet are associated with lower risk of NTDs suggesting that nutrients other than folic acid may be beneficial (Chandler et al. 2012, Sotres-Alvarez et al. 2013). Experimental analysis of individual vitamins found myo-inositol deficiency to cause NTDs in cultured rodent embryos (Cockroft 1988). Inositol supplementation significantly reduced the frequency of NTDs in $curly\ tail$ mice (Greene & Copp 1997) and in rodent models of diabetes (Reece et al. 1997).

7. FUTURE PERSPECTIVES

Experimental models provide systems for analysis of the developmental events of neural tube closure and fundamental cellular and morphological processes continue to be defined in more detail. In principal NTDs may result from insufficiency of one or more of the key driving forces (eg, cellular properties and/or morphological movements) that are necessary to achieve closure, for example, through mutation of a PCP gene. Alternatively, a genetic lesion or environmental insult may disrupt the closure process even where the underlying machinery is intact, for example through induction of aberrant cellular behaviour such as excess apoptosis. Experimental models require careful analysis to disentangle these possibilities. A key challenge will be to understand how the molecular and cellular determinants of neurulation relate to the biomechanical forces required to fold the neuroepithelium to achieve closure.

Advances in exome and whole genome sequencing offer the potential to begin to understand the genetic basis of NTDs in humans. The multifactorial complexity of NTDs means that analysis of data from such studies will present a major challenge. Moreover, there will be a need to integrate genetic data with information on epigenetic and environmental factors to obtain a more complete understanding of the cause of individual NTDs.

Folic acid supplementation provides a means to reduce NTD risk and represents a major public health advance. Nevertheless, the heterogeneity of NTDs suggests that primary prevention may be best achieved by multiple interventions and use of additional micronutrients alongside folic acid may provide an opportunity to further reduce risk.

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Figure 1

