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**Author Manuscript**

*Birth Defects Res A Clin Mol Teratol*. Author manuscript; available in PMC 2010 April 1.

### Published in final edited form as:

*Birth Defects Res A Clin Mol Teratol*. 2009 April ; 85(4): 314–321. doi:10.1002/bdra.20542.

## **Mechanistic insights into folate supplementation from** *Crooked tail* **and other NTD-prone mutant mice**

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## **Abstract**

Despite two decades of research since R. Smithells and colleagues began exploring its benefits, the mechanisms through which folic acid supplementation supports neural tube closure and early embryonic development are still unclear. The greatest progress toward a molecular genetic understanding of folate effects on neural tube defect (NTD) pathogenesis has come from animal models. The numbers of NTD-associated mouse mutants accumulated and studied over the past decade have illuminated the complexity of both genetic factors contributing to NTDs and also NTDgene interactions with folate metabolism. This article discusses insights gained from mouse models into how folate supplementation impacts neurulation. A case is made for renewed efforts to systematically screen the folate responsiveness of the scores of NTD-associated mouse mutations now identified. Designed after *Crooked tail*, supplementation studies of additional mouse mutants could build the molecular network maps that will ultimately enable tailoring of therapeutic regimens to individual families.

## **Keywords**

Folic acid; neural tube defects; folate metabolic pathway; mouse models; nucleotide biosynthesis; methylation

## **INTRODUCTION**

This special issue honors the estimable contribution of Richard Smithells and colleagues for recognizing as early as 1976 that deficiencies of folic acid, a simple water-soluble vitamin, might indicate increased risk of neural tube defects (NTDs) including anencephaly and spina bifida (Smithells and others, 1976). His subsequent landmark study showed that periconceptual folate supplementation could reduce the recurrence of neurulation defects, comparing control and supplemented cohorts of approximately 240 women each (Smithells and others, 1983). Larger therapeutic trials in the UK and eastern Europe established folate supplementation as the most effective birth defect prevention method ever identified, capable of reducing NTD occurrence rates by 70% or more (Czeizel and Dudas, 1992; MRC, 1991; Seller and Nevin, 1984). These and later supportive studies led to the current recommendation by the Center for Disease Control and the World Health Organization that all women of child bearing age receive folate supplementation of 0.4 mg daily and those women who have had a child with an NTD

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should receive 4 mg daily in the periconceptual period. Indeed, after voluntary folate supplementation rates indicated relative public inattention to this important measure, the US mandated fortification of food stuffs such as cereals to provide an additional 0.1 mg daily folate intake in the average US diet, implemented starting in 1996. A reduction in US NTD rates of 20–30% has been attributed to this fortification over the subsequent decade (Canfield and others, 2005; Honein and others, 2001; Pitkin, 2007; Wald and Oakley, 2007).

Surprisingly, more than 20 years after recognition that folic acid supplementation can reduce the occurrence of spina bifida and anencephaly, the mechanism of this rescue and why some families benefit while others do not is still largely unknown. The lack of present clarity is in no small measure due to the complexity of the interactions of genetics, metabolic status and environment that must converge to result in an NTD. Advances in this intriguing field have led to an appreciation of the intricacies of these interactions. Examining NTD mechanisms and their intersection with folate metabolism in mouse models, linked with investigation of human genetics and epidemiology, will likely unravel patterns of these gene-environment interactions. This level of understanding would certainly herald improved strategies for NTD prevention in individual families, including new options for the 30% or more who will not benefit from folate supplementation.

Here, we discuss insights from mouse models demonstrated to be protected from NTD by folate supplementation and highlight questions that these studies raise.

## **MOUSE MODELS as an INROAD to FOLATE ACTION in NTDs and PREVENTION**

Studies in mice provide numerous advantages as an adjunct to clinical investigation. The availability of inbred mouse strains and their shorter life cycles makes it possible to reduce the complexities of genetic background, environmental influences, and limited observations per family that are inherent in human studies. Experimental mouse models are more amenable to pharmacological manipulation of metabolism and detailed investigation of pathological outcomes in various tissues and fetuses. They also enable gene manipulation to perturb systems and probe gene-environment interactions. Animal studies have proven challenging, as no single mouse model appears to encompass the features of clinical experience. Nevertheless, a number of insights into folate-dependent influences on neurulation have been mined from mouse models. Evolving technical capabilities provide new options for the analysis of complex geneenvironment interactions that underlie folate supplementation effects.

#### **Folate Deficiency is not Enough (or may be too much)**

Several studies in mice suggest that folate deficiency alone can impair implantation and early embryogenesis, but it is insufficient in isolation to produce NTDs. Heid and colleagues compared the reproductive tracts of female Swiss-Webster mice 12 days postcoitum (dpc) that had been maintained on low-folate diets ranging from 45 nmol/kg chow to 2266 nmol/kg chow (Heid and others, 1992). They found that diets containing less than 181 nmol/kg prevented implantation and diets containing 221–453 nmol/kg folate produced 100% resorption rates. Dietary folate levels producing 75% resorptions did not produce NTDs in viable embryos. A decade later, Burgoon and colleagues examined ICR mice maintained on defined folate diets containing succinyl sulfathiazole, an antibiotic used to reduce gut flora to remove sources of absorbable folate not present in the diet (Burgoon and others, 2002). Those few implantations that were viable past dpc 11–12 did not display NTDs.

Nevertheless, genetic disruption of folate intracellular transport in folate binding protein 1 (*Folbp1*, also known as folate receptor 1*, Folr1*) or reduced folate carrier 1 (*RFC1*) knockout

mice (C57BL6 and SWV backgrounds, respectively) results in NTDs and this strongly argues that impairment of intracellular folate metabolism is deleterious to neurulation (Gelineau-van Waes and others, 2008; Piedrahita and others, 1999). These nullizygous embryos die periimplantation unless dams are heavily supplemented with folic acid. A detailed study of the RFC1 deficient embryos from dams receiving 50 mg/kg/day folic acid indicate the critical importance of folate transport in chorioallantoic fusion, hematopoesis and the development of neural tube, limbs, lungs, heart, and skin (Gelineau-van Waes and others, 2008).

These are prime examples of numerous investigations recently reviewed elsewhere (Beaudin and Stover, 2007; Blom and others, 2006; Harris and Juriloff, 2007). The ultimate message of these studies in mice is that the folate metabolic pathway is crucial for implantation and many aspects of embryonic development, and severe impairment of the pathway is incompatible with survival beyond early embryogenesis. However, intracellular folate utilization is an important aspect of neurulation. Therefore, it is likely that lesser alterations in the molecular pathways supported by folic acid metabolism would result in viable fetuses with defects more restricted to neural tube and its derivatives. In other words, a compilation of partial loss- or gain-offunction mutations must conspire to result in an NTD. This contention certainly parallels clinical experience. While this complexity has led to a degree of pessimism over how well studies in mice can emulate the human condition, one must also be heartened that the degree of complexity demonstrated in the mouse is up to the challenge of providing clinically relevant information. The field has only to be clever enough to take advantage of the opportunity.

#### **Mouse Mutants and NTD Prevention by Folate Supplementation**

Perhaps the most under utilized opportunity for investigation of folate actions in NTD prevention is the examination of folate supplementation in mice bearing a genetic predisposition to neurulation defects. Well over 190 naturally occurring and engineered mouse mutations are known to produce exencephaly (the mouse equivalent of anencephaly in humans), spina bifida or both, and over 155 of the responsible genes have been identified (Harris and Juriloff, 2007). Yet only a small handful has been tested for the ability of added folate to reduce the occurrence of NTDs in these mice and still fewer have been rigorously evaluated (Table 1). Studies in several mouse mutants, among them *Crooked tail*, suggest that screening of folate supplementation in genetic NTD models should be revisited to more broadly apply experimental paradigms that have proven informative.

**Crooked tail (Cd)—This naturally occurring mutation arose first in the A/J strain as a** semidominant allele on mouse chromosome 6 (Lane, 1972; Morgan, 1954). Heterozygous *Cd* mice display a crooked tail while homozygous mice express phenotypes including early embryonic lethality, exencephaly or viable neonates that are runted with more severe skeletal deformity confined to tail and thoracolumbar spinal column. These skeletal deformities are characteristic of a condition recognized as the Wiebel-Rippen syndrome, placing *Cd* in a group of mouse mutations including *Malformed vertebrae, pudgy* (mutated in *delta-like 3Dll3*, (Kusumi and others, 1998)), *Rachiterata, Rib fusions, Rib-vertebrae and Fused* (mutated in *Axin* (Zeng and others, 1997)) (Theiler, 1988). More recently, this group was expanded to include *Ringelschwanz* (*Lrp6* hypomorph) and *doubleridge* (*Dkk1* hypomorph) (Kokubu and others, 2004; MacDonald and others, 2004).

Mapping of *Cd* to a 0.2 cM region of chromosome 6 enabled the assessment of prenatal dietary folate supplementation effects on birth defect rates in this line, which proved to parallel human anencephaly responses in a number of respects (Carter and others, 1999). First, NTDs and other defects in *Cd* embryos were incompletely penetrant, as exencephaly occurred in roughly 30% of *Cd*/*Cd* embryos, and heterozygous offspring demonstrated variable rates of the dominant crooked tail phenotype, depending on the strain background carrying the *Cd* allele (Carter and

others, 1999). This was consistent with the clinical experience that recurrence risks in families with one NTD affected offspring almost never approach the Mendelian distribution of an autosomal recessive trait (Elwood and others, 1992). Second, NTD rates in *Cd*/*Cd* embryos displayed a gender bias, affecting more females by 3 to 1, similar to the gender bias observed in human studies (Elwood and others, 1992). Third, folate supplementation reduced NTD incidence by as much as 55%, lowering rates in females to equal those of males. Importantly, the rescue effect occurred even though *Cd* dams were not inherently folate deficient, since baseline and supplemented red blood cell folate levels were indistinguishable from wild type mice.

Among the controversies aired in the clinical literature, is the question of how added folate may reduce the incidence of birth defects--that is, could folate reduce the apparent incidence of defects by hastening fetal loss (Klein, 1996). Dietary folic acid supplementation of *Cd* mice, at near-physiological doses, affected a true rescue of NTDs (Fig. 1) (Carter and others, 1999). Not only was the NTD incidence lowered by folate supplementation, but also embryo viability was increased. Importantly, when all observations were scored and genotyped, a Mendelian distribution was obtained  $(1:2:1; +/+.Cd/cd)$ , indicating that pre-implantation loss of mutant embryos was not induced by folate supplementation (Fig. 1B).

Positional cloning identified *Cd* as a missense mutation, substituting an aspartate amino acid in place of a highly conserved glycine in the second extracellular propeller domain of *Lrp6* (Carter and others, 2005). A member of the low density lipoprotein receptor (LDLR) family, Lrp6, like its close cousin Lrp5, is a single-pass transmembrane protein and an obligate coreceptor with Frizzled that is required for canonical Wnt signaling (He and others, 2004; Logan and Nusse, 2004; Schweizer and Varmus, 2003). This connection of NTD with Lrp6 is of particular interest, since unlike other Wnt pathway genes associated with neurulation, Lrp5/6 are thought to be synonymous with the canonical pathway. Other Wnt-related mutations associated with NTD have been shown to impair neurulation through the non-canonical Wnt pathway and planar cell polarity (PCP) signaling. These include *Frizzled 3* and *6* (Wang and others, 2006b), and *Disheveled 2* (Wang and others, 2006a), which genetically interact with the *loop tail* gene, *Vangl 2*. Those NTD-associated Wnt pathway gene mutations that have yet to be mechanistically studied are known to have multiple roles, both in canonical and noncanonical Wnt pathways. The *Lrp6Cd* mutation indicates that either canonical Wnt signaling also has a role in neural tube closure or Lrp6 function also has an as yet unrecognized impact on the non-canonical Wnt pathway and this will be an important issue to resolve.

The *Lrp6Cd* allele provides a first link between folate metabolic status and a major developmental signaling pathway. However, the mechanism of folate rescue from NTD and other birth defects in *Cd* mice is not yet known. Array data analysis of folate gene expression and metabolite levels in tissues obtained from wild type and mutant *Cd* mice indicate an intracellular folate utilization defect due to the mutation (Ernest and others, 2006). One significant step toward mechanistic understanding will be the determination of folate effects on other *Lrp6* mutants. A gene trap mutation inactivating *Lrp6* is associated in null embryos with severe defects of limbs, truncation of the caudal body axis and NTDs including exencephaly and spina bifida (Pinson and others, 2000). In contrast, the *Ringelschwanz* mouse that carries a hypomorphic point mutation in *Lrp6* displays lumbosacral bone abnormalities and spina bifida, but apparently not exencephaly (Kokubu and others, 2004). If NTDs in other *Lrp6* mutants (*Lrp6* nulls or *Ringelschwanz* mice) are also sensitive to folate status, then the effect of the vitamin must be due to a direct interaction of folate metabolism with Lrp6 function and not to an idiosyncratic interaction with the *Cd* mutation. Similarly, establishing whether NTDs associated with the *Disheveled 1*/*2* (Hamblet and others, 2002) mouse lines respond to folate supplementation would help to determine whether the folate metabolic pathway impacts Lrp5/6 function specifically or Wnt signaling more globally, downstream of the receptor.

Regardless of whether the folate interaction is unique to *Cd* or relates to Lrp6 action, it will be important to investigate how the *Cd* mutation leads to impaired intracellular folate utilization.

**Axial defects (Axd)—NTD-prone mutants found to be unresponsive to folic acid may also** be highly informative. Spina bifida occurs in some 30% of *Axd*/+ mice and 100% of homozygotes. While folic acid or B12 supplementation did not affect occurrence rates, high doses of methionine given on dpc 8&9 reduced NTD rates 40–47% (Essien and Wannberg, 1993). An essential amino acid, methionine is required in the folate metabolic pathway for the production of the body's principal methyl donor, S-adenosyl methionine. The lack of response in *Axd* to folate, yet significant reduction in its NTD rate by methionine, implies the presence of a metabolic block in this mutant line that can be circumvented by high doses of methionine. Thus, there are likely other agents feeding into folate/homocysteine metabolism that will be effective in preventing NTDs, depending on the individual mutation and genetic background of the individual.

#### **Possible Mechanisms Affecting Folate-dependent NTD Prevention**

The folate metabolic pathway encompasses a multi-faceted series of reactions that regulate one carbon metabolism (OCM) through the formation and utilization of formate and tetrahydrofolate (THF), enzymatically converting THF to different oxidation states from formyl-to methylated versions (Beaudin and Stover, 2007; Stover, 2004). The two major arms of the pathway lead to nucleotide biosynthesis through thymidylate and to methylation reactions through the methyl-donor S-adenosyl methionine (Fig. 2). It is likely that the success of folic acid supplementation in the face of the complex genetic underpinnings of NTDs is due to the diversity of pathways to which folate contributes. Connections between NTDs and both arms of the folate metabolic pathway have been uncovered with the aid of mouse mutant models.

#### **Nucleotide Biosynthesis**

A principal arm of the folate metabolic pathway contributes to purine and thymidine biosynthesis. Reduction in nucleotide availability can impact DNA replication and repair and decrease mitotic rates during critical points in morphogenesis. (Keller-Peck and Mullen, 1997). Particularly important is the increased incorporation of Uracil into DNA in face of thymidylate deficiency, impacting DNA replication and repair, thereby reducing proliferation and promoting genomic instability (Andersen and others, 2005; Courtemanche and others, 2004; Mashiyama and others, 2004).

**Pax3/Splotch—**Mutations in the *Pax3* transcription factor lead to deficits in neural crestderived cells in heterozygous mice, giving them the white patches of the 'splotch' coat color, and to either exencephaly or spina bifida in homozygous *Splotch (Sp*/*Sp)* mice (Epstein and others, 1993; Vogan and others, 1993). The NTD phenotypes can be rescued by folate supplementation in culture and in utero (Fleming and Copp, 1998; Wlodarczyk and others, 2006). Using the deoxyuridine (dU) suppression assay, the ability of *Sp*/*Sp* embryos to synthesize thymidylate was shown to be impaired (Fleming and Copp, 1998). In addition, either folic acid or thymidine added to culture media could prevent NTD and relieve the metabolic block in homozygous embryos, but addition of methionine exacerbated the metabolic defect, inducing NTD in 50% of heterozygous *Sp*/+ embryos (Fleming and Copp, 1998). Thus, rescue of *SpPax3* mutants from NTDs is associated with that segment of the folate/homocysteine metabolic pathway that supports nucleotide biosynthesis, while in contrast, tipping the balance toward the methylation arm of the pathway exacerbates embryopathy.

Connection of *Pax3* loss of function with a folate metabolism defect that impacts nucleotide biosynthesis suggests that the rescue of NTD in this case may relate to folate effects on cell

proliferation. Indeed, proliferation defects have been observed in *Sp* mutant embryos (Keller-Peck and Mullen, 1997). A recent examination of the *Sp2H* mutant line has shown that folate deficiency increases NTD rates in mutant, but not in wild type, embryos (Burren and others, 2008). This is associated with reduced growth rates in the mutants, especially in female *Sp2H* embryos. At the same time, no changes in global DNA methylation were found in that study, leading those investigators to hypothesize that the cellular defect in *Sp2H* mice is at least in part compensated for by FA supplementation that restores a folate metabolic pathway block, normalizes nucleotide biosynthesis and supports intrauterine growth (Burren and others, 2008). Nevertheless, this study does not rule out the possibility of methylation effects on specific key genes that would not be detected in this experimental design.

Studies of *Splotch* mutants support the contention that there is a critical balance between the nucleotide biosynthesis and the methionine biosynthesis/methylation arms of OCM that must be maintained to promote successful neural tube closure (Beaudin and Stover, 2007; Stover, 2006). There is some evidence that these insights from mouse models can be translated into clinical relevance. Impaired thymidylate biosynthesis has recently been reported in a subset of human NTD-affected embryos, assessed using a dU suppression assay to screen primary fibroblast cells derived from human fetuses affected by NTD (Dunlevy and others, 2007). It will be of considerable interest whether dU suppression will prove abnormal in fibroblasts derived from mothers of those affected fetuses with thymidylate biosynthesis abnormalities.

An intriguing question with regard to folate rescue of NTD is whether a flaw or at least partial impairment within the folate metabolic pathway is necessary in order for FA supplementation to have a protective effect on neurulation. For example, might it be possible for FA to rescue neurulation by virtue of enhancing cell proliferation, in effect giving mutant embryos a "boost" without directly addressing the underlying genetic functional defect? While certainly possible, this has yet to be demonstrated. The observation that folate deficiency had no impact on intrauterine growth rate or neurulation in wild type embryos (Burren and others, 2008) suggests that added FA would have an impact via this mechanism only if a cell division defect were part of the neurulation failure. On the other hand, clearly not all NTD mutants expressing a proliferation defect will respond to folate. For example, the *ct* mutant suffers NTD attributed to reduced proliferation in the hindgut, notochord and caudal tailbud, but *ct* is insensitive to FA supplementation (van Straaten and Copp, 2001). At this early stage, the limited data available from NTD mutants like *SpPax3* , *CdLrp6* and *ctGrhl3* seems to support that neurulation sensitivity to supplementation requires some inefficiency of folate intracellular uptake or utilization in the embryo, or possibly a maternal component of OCM homeostasis.

#### **Methylation Cycle**

There are multiple indications that methylation—especially of DNA or histones—is a robust contributor to neurulation. Knockout in the mouse of the DNA methyltransferase, *Dnmt3b*, produces NTDs and distortions of the neural tube (Okano and others, 1999). In fact, no fewer than eight mutations in genes involved in DNA methylation or in regulating chromatin structure produce neurulation defects, including *Cerc2, Dnmt3b, Dnmt3l, Gtf2i, Hdac4, Sirt1, Smarca4* and *Smarcc1* (Cheng and others, 2003; Harris and Juriloff, 2007; Okano and others, 1999; Vega and others, 2004). Variant alleles of methylenetetrahydrofolate reductase (MTHFR) have been associated with increased NTD risk in some but not all clinical populations studied (Boyles and others, 2005). This enzyme is a hub in the pathway and diverts OCM toward methylation reactions at the expense of purine and thymidine biosynthesis. The MTHFR 677 T/T genotype has been associated with a global reduction in DNA methylation (Castro and others, 2004; Friso and others, 2002). Experiments in chick and mouse embryos indicate that exposure to homocysteine or inhibitors of the methylation cycle delayed neural tube closure in a dose-dependent manner (Afman and others, 2005; Dunlevy and others,

2006). There remains some controversy in the literature as to whether folic acid supplementation results in hyper- or hypomethylation of DNA globally, whether the most important effects for neurulation result from the methylation status of specific target genes, or whether the methylation of histone proteins to influence chromatin structure is the key event (Waterland, 2006). Certainly it would be of interest, in the eight or more mutants affecting DNA, chromatin or histone methylation, to determine whether their NTDs are prevented by folate supplementation. However, this may not be relevant if the ability to respond to folate by increasing DNA/histone methylation is substantially precluded by the mutation. The more pertinent question is whether the expression of methylation-sensitive genes contributing to NTDs will be favorably modulated by folate supplementation.

Although much attention has focused on the role of DNA/chromatin modification in NTDs, it must be recognized that methylation is an important modulator of other molecules including RNA, proteins, lipids and even neurotransmitters. Indeed, neurulation defects were induced in wild type rat embryos when cultured in reduced levels of methionine (Coelho and Klein, 1990). Methionine deficiency was associated with a failure of the neural folds to turn medially, suggesting a deficit in 'microfilaments' and diminished cytoskeletal contractility. The authors' hypothesis was supported by measured reductions in microfilament-associated methylated amino acids (Coelho and Klein, 1990). It is likely that one or more of the several recognized possible folate-related mechanisms contribute to NTD prevention in a particular genetic context.

A sobering insight regarding folate supplementation is that its effects on methylation can result in sustained alteration of gene expression, as demonstrated by the  $A<sup>VT</sup>$  mouse (Dolinoy and others, 2007; Dolinoy and others, 2006; Waterland, 2006; Wolff and others, 1998). Coat color in this inbred mouse line is determined by the agouti locus and hypomethylation of CpGs in its transcription unit shifts the adult coat toward yellow. Maternal supplementation with methyl donor precursors folate, choline and betaine leads to hypermethylation of the  $A<sup>VT</sup>$  locus and shifts coat color from yellow to pseudo-agouti (Wolff and others, 1998). Intrauterine exposure to bisphenol A (BPA) leads to hypomethylation at the  $A<sup>VY</sup>$  locus and this is prevented by either folate supplementation or treatment with the phytoestrogen, genistein. Importantly, these effects on CpG methylation at the  $A<sup>VY</sup>$  locus created a permanent alteration in the individual epigenome. Moreover, varying the methyl donor sources folate, methionine and choline in a post-weaning diet can permanently influence the imprinting of insulin like growth factor 2 (*Igf2*) (Waterland and others, 2006). Thus, effects of folate metabolism can be felt long after periods of supplementation, underscoring the importance of understanding the range of possible effects of folate, the dose dependence of those effects and likely outcome for an individual genotype.

## **CONCLUDING REMARKS**

The examples provided by *Cd, Axd, Folb1, RFC1, Pax3* and several other mutations in mice suggest that, with knowledge of the maternal-fetal genetic background, it is within our reach to target particular aspects of the folate metabolic pathway and so optimize NTD prevention for families using individually tailored supplement regimens. The gene expression and metabolite 'signatures' indicating NTD risk and likely effective prevention for a particular profile will be best identified in mouse models, followed by searches for parallels in human subjects. It may indeed now be time to revisit metabolic studies in mice to take advantage of the over 155 genetic mutants in hand, with more NTD-associated mutations being identified every day.

In such metabolic studies, it will be important to determine for a particular NTD-associated gene, not only whether the rate of NTDs is reduced by folate supplementation, but also by what

mechanism. That is, for a particular genetic profile, does folic acid affect a true rescue and improve embryo viability or enhance embryonic lethality? Does folate alter proliferation in the neural tube or in surrounding mesenchyme? Does it alter convergence-extension during neurulation or is there another agent that does a better job preventing NTDs due to defects in planar cell polarity (PCP) pathways? What does folate supplementation do to the gene expression pattern of that particular allele and genetic background? Does folate impact posttranslational regulation of protein function during neurulation? How much supplementation is enough—where is the boundary defining too much of a good thing? Indeed, there may be a level at which folate supplementation becomes counterproductive. For example, dietary supplementation of *Sp* mice with very high levels of folate (200 mg/kg chow, or 20 times higher folate content than in other dietary supplementation studies) resulted in a 30% reduction in NTDs in homozygous embryos, but also a six-fold increase in the number of resorptions (Wlodarczyk and others, 2006). However, the level of supplementation required is likely to vary according to the mutation involved. Certainly, very high folate levels are required to ameliorate the Folbp1 or RFC1 knockouts (Gelineau-van Waes and others, 2008). Only when we have these more complete datasets will therapeutically useful parallels be obtainable in clinical populations.

Over decades of study in animal models and in humans, it is now clear that folate deficiency has significant deleterious effects on embryonic and fetal development. There is also compelling evidence that folate supplementation can, in the right genetic setting, exert a decided rescue from neurulation failure and other serious birth defects. In addition, supplementation can bring measurable benefits even in the absence of folate deficiency. However, more must be understood regarding the intracellular folate levels that will be optimal for a particular genetic mutation and background, in order to avoid unintended consequences and achieve better individual therapeutic advantage over population-directed strategies for supplementation (Pitkin, 2007; Smith and others, 2008; Stover, 2004; Stover and Garza, 2002).

More than 20 years have elapsed since Richard Smithells and colleagues blazed the trail for folic acid supplementation to prevent NTDs, and significant reductions in incidence have been achieved. Nevertheless many questions remain and a new era is emerging in which focus will shift away from populations at risk for NTDs to individual couples. The field has made breathtaking advances over the last two decades, though much remains to be understood. With accurate assessment of the relative contributions from specific genes, expression patterns of genetic networks, and the role of epigenetic and environmental factors, individually tailored strategies for NTD prevention will become feasible. It all began with the connection between a small molecule and a devastating malformation. To even think that vitamin supplementation could reduce the occurrence of birth defects was a bold stroke, for which much is owed to Dr. Smithells.

## **Acknowledgments**

Supported by NRSA NS059562 to JDG and the WCMC-THMRI initiative to MER

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#### **Figure 1.**

Dietary folic acid supplementation produces a true rescue of *Cd*/*Cd* embryos. **(A)** Distribution of embryonic phenotypes as a percentage of all embryos scored at E12.5 from dams maintained on diets containing defined amounts of folic acid (FA) (parts per million, ppm = mg/kg chow). On a 0 ppm FA diet, homozygous phenotypes accounted for 22%, nearly all of the expected 25% for a Mendelian distribution of an autosomal recessive mutation. However on a 10 ppm FA diet, 7.5% displayed embryopathy (30% of the expected 25%). Compared to embryos on a 4 ppm FA diet, the exencephaly encountered on a 10 ppm FA diet was reduced more than half. **(B)** *Cd*/*Cd* embryos are not lost prior to implantation. The genotype distribution observed

for E12.5 embryos maintained on defined FA diets showed the expected Mendelian pattern of 1:2:1 for a heterozygous parental cross. Data from Carter et al., 1999



#### **Figure 2.**

Summary of folate-homocysteine metabolism. The primary outputs of the pathway are to **1. support proliferation** and DNA repair by the synthesis of nucleotides thymidine via thymidylate synthase (and purines from 10-Formyl THF); **2. support methylation** of DNA, RNA, proteins and lipids through the major methyl donor, S-adenosyl methionine (SAM). In this part of the cycle, Hcy is converted to Met by methionine synthase, gaining a methyl group from methyl-THF. \* In some tissues, Hcy can also be converted to Met from betaine through conversion of choline to betaine and transfer of a methyl group catalyzed by betainehomocysteine methyltransferase. DHF=dihydrofolate, THF=tetrahydrofolate, MTHF=methyltetrahydrofolate, Hcy=homocysteine, Met=methionine, SAH=S-adenosyl homocysteine, dUMP=deoxyuridine monophosphate, dTMP=deoxythymidine monophosphate, MTHFR=methyltetrahydrofolate reductase.

#### **Table 1**

#### Mouse NTD mutants tested for responsiveness to Folate



#### **Folate Unresponsive NTD Mutant Mice**



i.p., intraperitoneal; sc., subcutaneous; d, day; dpc, days postcoitum