



A REVIEW STUDY ON THE IMPACT OF MTHFR POLYMORPHISMS ON FOLATE RELATED DISEASES

Genetics

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ABSTRACT

Folate plays a crucial role in homocysteine metabolism, DNA and methionine synthesis, and other biochemical pathways. These are collectively known as one-carbon metabolism mediated by folates. Methylenetetrahydrofolate reductase (MTHFR) gene loci present on chromosome 1 encode an enzyme critical for folate metabolism, cellular metabolism, and methylation processes. MTHFR gene variations like A1298C and C677T, are responsible for hyperhomocysteinemia and folate deficiencies leading to various health-related adverse outcomes. The current review attempts to highlight the role of genomics in MTHFR expression and how gene-level variations impact MTHFR gene etiologies. Knowledge about such associations can help devise therapeutic approaches for folate-related conditions.

KEYWORDS

5,10-methylenetetrahydrofolate reductase, Disease risk, MTHFR gene, Polymorphism

INTRODUCTION

Folate is an essential micronutrient and cofactor in one-carbon metabolism. It is associated with the vitamin B group and regulates purine and pyrimidine synthesis, and amino acid metabolism¹. Folate act as methyl donors for the conversion of homocysteine, a sulfhydryl-containing amino acid, to methionine². Mammals are unable to synthesize folate and thereby rely on supplementation. Despite this, drug interactions and poor absorption are major reasons for folate deficiency³.

Folate deficiency leads to neural tube defects, cardiovascular diseases, congenital heart defects, cancers, migraines, dementia, and depression, impaired embryonic development, birth defects, morbidity and mortality in children, preterm and low birth weight^{4,5}.

The enzyme methylenetetrahydrofolate reductase (encoded by the MTHFR gene) regulates homocysteine metabolism, folate pathways, synthesis of methionine, cysteine, etc. The MTHFR gene is present on chromosome 1 at the edge of the short arm (1p36.6)⁶. It is involved in DNA methylation and RNA synthesis by catalyzing the conversion of 5,10-methylenetetrahydrofolate (5,10-MTHF) to 5-methyltetrahydrofolate (5-MTHF) as observed in figure 1. This is the prime folate found in the blood which regulates the homocysteine to methionine conversion. Cells use this type to synthesize glutathione, taurine, S-adenosylmethionine (SAM), and creatine for the functioning of normal bodily processes.

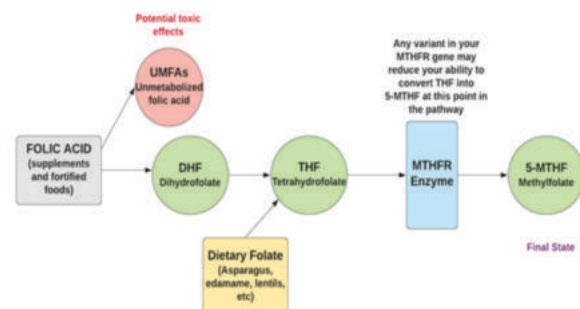


Figure 1: The folate pathway wherein supplemented folic acid is converted into 5-MTHF methyl folate via different intermediates

There are nearly 14 polymorphisms associating MTHFR with enzymatic deficiency⁶. Amongst all, C677T and A1298C are the most common ones, decreasing enzyme capacity to 70% and 40%, respectively. C677T (rs1801133) results in a valine-to-alanine conversion at codon 222, while A1298C (rs1801131) results in a transversion of adenosine to cytosine at nucleotide position 1298 resulting in glutamate-to-alanine substitution⁷. MTHFR variations

affect folate and homocysteine levels, thereby disrupting associated pathways as observed in figure 2, leading to diseases.

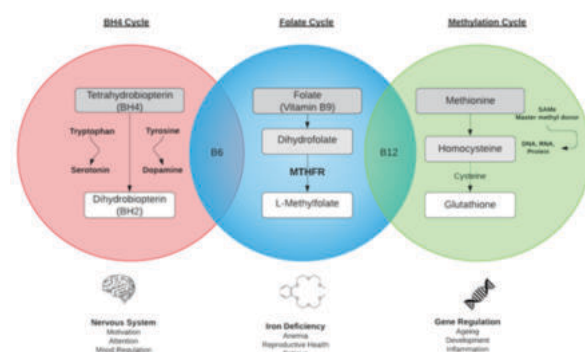


Figure 2: The MTHFR gene and its associated pathways

METHODOLOGY

A systematic literature review was conducted on MTHFR gene variations and associated conditions. Scientific databases such as PubMed, Google Scholar, and other resources were used to find the genotype-phenotype relationship between MTHFR gene and its clinical implications globally. These were obtained by using the following keywords: “MTHFR” or “methylenetetrahydrofolate reductase” or “polymorphism” or “variant”, “C677T” or “A1298C” and “diseases”.

Inclusion criteria:

- Well-defined meta-analysis, systematic review, case-control studies and review articles
- Publication year 2015 onwards

Exclusion criteria:

- Preclinical studies
- Articles in a different language than English
- Study with lower sample number

MTHFR gene variants and their role in fertility

Approximately 10%-15% of couples globally are affected by fertility complications, of which nearly half are attributed to males and about 20%-25% are ascribed to idiopathic male infertility⁸. Neural tube defects have an incidence of around 1 in 1,000⁹. Low folate and vitamin B12 levels are associated with infertility. MTHFR variants can cause homocysteine imbalance, thereby affecting DNA methylation and consequently affecting reproductive function. The C677T T allele expression is significantly higher in impaired spermatogenesis cases than in controls. Male infertility risk decreased following vitamin B9 and B12 intake, wherein a decreased total homocysteine level and development in semen parameters were observed¹⁰.

The 677TT genotype remarkably decreased vitamin D levels and increased homocysteine than its counterpart genotypes (CC & CT) in recurrent pregnancy losses (RPL)¹¹. A significant association was compared between MTHFR C677T and A1298C genetic variants and unexplained pregnancy loss (UPL) using samples of Korean origin¹². An Eastern Indian study suggested that the rs1801133 polymorphism was associated with an increased risk of neural tube defects (NTD) in mothers with NTD-containing fetuses¹³. Studies have proved a significant correlation between MTHFR polymorphisms and neural tube defect development, fertility problems, and recurrent pregnancy loss¹⁴. Summarily, evidence shows that MTHFR variations may cause poor pregnancy-related outcomes, hence, genetic screening is important to understand the causal variants and undergo supplementation if required.

MTHFR gene variants and their role in neurological and psychiatric diseases

MTHFR activity reduction or folate deficiency is associated with depression, autism, schizophrenia, mood disorders, bipolar disorder, and ADHD^{15,18}. Research suggests associations between C677T and Parkinson's disease, Alzheimer's disease, migraine, cerebral venous thrombosis and unipolar depressive disorder^{6,16}. It is also associated with hyperactivity-impulsivity among East Indians, indicating that folate-homocysteine pathway variants affect ADHD pathogenesis¹⁷. The MTHFR enzyme reduces gray matter volume in the hippocampus, indicating susceptibility hyperhomocysteinemia correlation with epilepsy or spastic paraparesis¹⁹ and antiepileptic-induced psychosis in a case with C677T polymorphism which may improve medical management for epileptic patients²⁰. MTHFR may also be associated with suicide risk in schizophrenic patients²¹. It is important to understand variants underlying such conditions for better management and prevention.

MTHFR gene variants and their role in cancer

DNA destabilization, leading to reduction of cytosine methylation due to folate deficiency, may trigger pro-oncogene expression. Since MTHFR is involved in folate metabolism, its variants affect cancer susceptibility⁷. Several research studies state the association of A1298C polymorphism with an increased cervical cancer risk and cervical intraepithelial neoplasia (CIN) development in patients and a decreased risk for lung cancer^{22, 23, 24}. Studies also demonstrated that C677T variant was significantly associated with lung cancer and adenocarcinoma among Chinese individuals, especially in females as well as decreased gastric cancer risk among elders^{24, 25, 26}. A Taiwan-based study found that the T allele of this variant is a predictive marker for colorectal cancer given its lower risk-conferring property than the wild-type C allele²⁷. Meta-analysis studies also showed that C677T increased breast and ovarian cancer risk, especially in Asians, and is associated with pancreatic cancer^{28, 29}. It is considered a therapeutic target for breast cancer³⁰ but research among Northern Sardinians CT genotype exerted a protective effect on its development³¹.

Various studies show associations between polymorphisms in folate-pathway genes, and pancreatic cancer development³². Hence it can be inferred that folate deficiency may increase but also be protective for various risks of cancer including thyroid, colorectal, ovarian, lung, prostate, cervical and breast^{33,34,35,36,37,38,39,40}. Considering that MTHFR variants play a substantial role in cancer development, screening to prevent disease onset is essential for well-being.

MTHFR gene variants and their role in vascular health

In a meta-analysis, C677T showed association with abdominal aortic aneurysm (AAA) risk, especially in non-smoking and younger patients⁴¹. In another study, C677T was more prevalent in women with abnormal triglyceride levels, linking it to inflammatory, vascular and metabolic dysfunction⁴². Evidence indicates associations between C677T, hyperhomocysteinemia and severe coronary lesions in acute coronary syndrome patients⁴³.

Another study highlighted a significant association between C677T and deep venous thrombosis, intracerebral hemorrhage^{44,45}, ischemic stroke⁴⁶, coronary artery disease⁴⁷ premature coronary artery disease⁴⁸, congenital heart defects⁴⁹, atherosclerosis⁵⁰, coronary atherosclerotic lesions⁵¹. C677T also demonstrates a protective role by negatively impacting left ventricular hypertrophy, with increased resistance of intrarenal arterial and hyperparathyroidism⁵². While A1298C polymorphism is a factor in the development of congenital heart disease⁵³ and conotruncal heart disease⁵⁴.

MTHFR variants are highly prevalent in non-valvular atrial fibrillation (NVAf) cardioembolic stroke subjects causing a higher risk of developing cardiovascular-related comorbidities with an enhanced inflammatory state, dyslipidemia, lacunar stroke or stroke recurrences. The detection of risk-related MTHFR variants may serve a major role in reinforcing recommendations towards nutrient intake and personalized management.

MTHFR gene variants and their role in diabetes and related complications

A study indicates that C677T confers type 2 diabetes mellitus risk, especially among Asians⁵⁵. Studies show its association with diabetic nephropathy⁵⁶, diabetic neuropathy^{57,58}, and diabetic retinopathy⁵⁷. It may also be associated with higher total homocysteine levels in type 2 diabetes patients with dyslipidemia⁵⁸. Additionally, C677T could impact folic acid supplementation efficacy in diabetic polyneuropathy⁵⁹.

The A1298C polymorphism significantly increased diabetic retinopathy, diabetic polyneuropathy⁵⁷, diabetic nephropathy⁶⁰, hypermethylation related to hyperglycemia, and dyslipidemia⁶¹ susceptibilities.

Table 1: Summary of the correlation between MTHFR gene polymorphisms and its clinical implications

Category	Disease	Variant	References (PMID)
Neurology and Psychiatric Diseases	Schizophrenia	MTHFR C677T A1298C	23318463 28822116
	Bipolar Disorder, Depression	MTHFR C677T A1298C	25744938, 28968218, 24751310
	Autism Spectrum Disorders	MTHFR C677T A1298C	31033224, 31920317
	Attention Deficit Hyperactivity Disorder	MTHFR C677T A1298C	29407547, 23227261 21618410
	Parkinson's Disease	MTHFR C677T A1298C	33574311, 29097250
	Alzheimer's disease	MTHFR C677T A1298C	21663380, 25359311, 26820674, 32804129
	Migraine	MTHFR C677T A1298C	31045246, 30451038, 19559392, 19925624,
	Epilepsy	MTHFR C677T A1298C	33096746, 31354331
	Cerebral Venous Thrombosis	MTHFR C677T A1298C	33375456, 26261166, 30466296
Vascular Diseases	Atherosclerosis	MTHFR C677T A1298C	32404177, 29087287
	Ischaemic Stroke	MTHFR C677T A1298C	31932513, 31456518, 32722170, 32098547
	Congenital Heart Defects	MTHFR C677T A1298C	33219830, 32418960, 30333252, 20664391
	Abdominal Aortic Aneurysm	MTHFR C677T	27603386, 24613192
	Coronary Artery Disease	MTHFR C677T A1298C	32682401, 30115070, 21567207, 29972410
	Acute Coronary Syndrome	MTHFR C677T	29245302, 27051002
	Dyslipidemia	MTHFR C677T A1298C	32795354, 28890888
	Embolism, Deep Vein Thrombosis	MTHFR C677T A1298C	33128086, 28500484, 26261166, 30466296

	Conotruncal Heart Defects	MTHFR C677T A1298C	22495907
Pregnancy and infertility	Male Fertility Problems/Infertility	MTHFR C677T A1298C	24265582, 33371103, 32912251, 30813130, 28081209, 27173242, 16888682
	Recurrent Implantation Failure	MTHFR C677T A1298C	27560137, 29073227
	Recurrent Pregnancy Loss	MTHFR C677T A1298C	29785531, 33751263
Developmental Anomalies	Down's Syndrome Child Birth	MTHFR C677T A1298C	27659321, 24668664, 20680153, 15834031
	Neural Tube Defects	MTHFR C677T A1298C	29222906, 31238314, 24737468, 23056169
Diabetes	Type 2 Diabetes	MTHFR C677T A1298C	31663297, 24452036
	Hypermethylation Profile	MTHFR A1298C	30675189
	Diabetic Nephropathy	MTHFR C677T A1298C	32871871, 22554825, 29227003
	Diabetic Neuropathy	MTHFR C677T A1298C	24452036, 28231061, 31557408
	Diabetic Retinopathy	MTHFR C677T A1298C	32582807, 24452036
	Gestational Diabetes	MTHFR C677T A1298C	32440179
Cancer	Breast	MTHFR C677T A1298C	32192442, 22291746
	Cervical	MTHFR C677T A1298C	33406167, 29740106
	Ovarian	MTHFR C677T A1298C	32639550, 28487897, 22524826
	Hepatocellular	MTHFR C677T A1298C	26662389, 31694048, 23457501, 27803768
	Prostate	MTHFR C677T A1298C	22296369, 22938427, 26782572, 15298954
	Lung	MTHFR C677T A1298C	24648965, 30323671, 27173216
	Brain	MTHFR C677T A1298C	25809864, 30583664, 18483342
	Thyroid	MTHFR C677T A1298C	24805831, 25070812, 27173331
	Gastric	MTHFR C677T A1298C	33920562, 25170232, 30884202, 20237899
	Colorectal	MTHFR C677T A1298C	28983337, 31740010, 20078613

CONCLUSION

MTHFR regulates protein synthesis and the folate pathway. It is majorly involved in the conversion of homocysteine to methionine, which is important for DNA methylation and gene regulation. Variations are associated with MTHFR deficiency, homocystinuria and hyperhomocysteinemia. This enhances risk of conditions related to fertility, cardiovascular, diabetes, cancers, neurodevelopmental, and psychiatric disorders.

Hyperhomocysteinemia is rarely attributed to two variants alone.

Patients should be evaluated for factors affecting homocysteine levels such as diet, age, thyroid disease, certain medications, hypercholesterolemia, or lifestyle-related factors. In the absence of such factors, identification of the genetic basis of high homocysteine levels is helpful.

Considering the clinical implications of polymorphisms, genetic counselors should focus on various queries on etiologies to understand the role of MTHFR in health. In this review, we documented a brief on the role of MTHFR in diseases as summarized in table I. Advancement in research and genomics is leading to a better understanding of folate, methylation, and epigenetics which can pave the way for advanced research in folate metabolism efficacy roles for therapeutic purposes.

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