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Introduction

Dependence to opiates is a chronic relapsing brain disease that can cause major health, social and economic problems [1]. Dependence vulnerability is affected by environmental factors, drug-induced factors

CT and TT genotypes are 3.7 and 13.9 times more likely to be linked to drug dependence than genotype CC; respectively. In addition, the presence of allele T is associated with increased likelihood of dependence compared to allele C (OR=4.1; CI=2.8-5.96) (Table 3).

(Table 4) shows that the CT and TT genotypes have a significantly longer duration of dependence, higher severity score, more relapse, and larger dose of daily Heroin intake and higher percent of positive family history compared to genotype CC. The same pattern was observed in

allele T compared to allele C. These data are not deviated from Hardy-Weinberg equilibrium. (Table 5) shows that homocysteine level is significantly higher in cases than control. Among control homocysteine level does not show significant variation according to genotype and allele type. On the other hand, in cases the homocysteine level is significantly higher in genotype CT and TT compared to CC and with allele type T compared to C.

Discussion

Present study revealed that cases and control were matched regarding their age, gender, religion, marital status, education and residence. However, they differ significantly in their working status which could be explained by the effect of substance use and its hazard on social and occupational function.

In the current study, MTHFR C677T genotype CC was more prevalent in the control group (75%) with low prevalence of CT and TT (21% and 3%, respectively) while CT represented the most common genotype in the cases group (42%) then CC (38%) and lastly TT (19%). (P=0.19 for the control group and 0.10 for cases group). There are statistically significant increases in CT & TT genotypes as well as T allele in the cases group when compared to the control group. These results confirm the idea recently emerged that opioid abuse is closely related to genetic polymorphism of some opioid receptor genes apart from MTHFR gene, [23]. For our knowledge, it is the first study to investigate the association of MTHFR C677T gene polymorphism and Heroin dependence. Previous studies showed that T allele and T carrier genotypes (TT & CT) were more prevalent in nicotine smokers [11] alcohol [12] and cocaine dependent patients [13] than in the control group. Yuferov et al., [1] found that there is a strong association between Heroin and cocaine addiction and polymorphism in genes. The similarity of the present study results and other drug dependence (cocaine, alcohol, and nicotine) could return back to the fact that these drugs share common neural neurotransmitter release (dopamine) in the same brain area (brain reward system) and many other genotype polymorphism [1,23].

The current study shows statistically significant increase in plasma homocysteine level in the cases group than the control group. Also, there is increase in the TT genotype of MTHFR C677T gene and T allele frequency in comparison to other genotypes (CT and CC) and C allele frequency in the cases group but not in the control group. These results are in agreement with Tomedi et al. [24]. Their study showed that the plasma homocysteine level is statistically higher in the opiate drug addict group even the treated subjects of them than that of their control group. The hyperhomocysteinemia associated with T allele and T

	Control (192) N (%)	Cases (178) N (%)	χ ² / P
Age: <30	49(25.5)	48(27.0)	F=0.23,
30-40 & more	97(50.5)	91(51.1)	P=0.88
Mean ± SD	46(24.0)	39(21.9)	t=1.1, P=0.3
Gender: Female	60(31.2)	55(30.0)	χ ² =0.01,
Male	132(68.8)	123(69.1)	P=0.9
Religion: Muslim	185(96.4)	171(96.1)	χ ² =0.02,
Christian	7(3.6)	7(3.9)	P=0.3
Marital status: Married	54(28.1)	48(27.0)	F=0.5, P=0.9
Divorced	9(4.7)	7(3.9)	
Single	117(60.9)	114(64.0)	
Widow	12(6.2)	9(5.1)	
Education *: literate	64(33.3)	50(28.1)	F=4.2, P=0.24
Primary-preparatory	46(24.0)	40(22.5)	
Secondary school	68(35.4)	80(44.9)	
Faculty education	14(7.3)	8(4.5)	
Work *: Not working	94(49.0)	66(37.1)	F=14.96, P=0.002
Student	71(37.0)	97(54.5)	
Semi-skilled	16(8.3)	13(7.3)	
Skilled worker	11(5.7)	2(1.1)	
Residence : Rural	93(48.4)	89(50.0)	χ ² =0.1, P=0.76
Urban	99(51.6)	89(50.0)	

	Genes			Alleles	
	CC Median (min-max)	CT Median (min-max)	TT Median (min-max)	C Median (min-max)	T Median (min-max)
Duration	14(12-26) ^{AB}	28(12-39) ^A	27.5(12-39) ^B	18(12-39)	28(12-39)
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and hydrogen peroxidase by auto-oxidation leading to oxidative stress and reduce the expression of glutathione peroxidase (an antioxidant enzyme) thereby causing the accumulation of its neurotoxic oxidative products [29]. Moreover, homocysteine was found to provoke neuronal cell damage and brain atrophy by stimulating N-methyl-D-aspartate (NMDA) receptors [30]. Therefore, it is not strange to find that elevated homocysteine (Hcy) levels associated with neuro-degeneration, vascular diseases and brain atrophy [31].

In this study, the dependence severity was more associated with the presence of a positive family history. This was consistent with previous studies by Pickens et al., [32] found that family history of substance dependence is a good predictor for more opioid dependence symptoms and more likely to be classified as severely dependent patient. Moreover, the current study revealed an association between the severity of dependence as measured by duration of illness, number of relapses and severity of dependence scale and the dose of the substance with genotype and alleles. All these measures are higher in CT & TT genotypes than CC genotype as well as T allele than C allele. A similar increase in severity of dependence with an increase in gene polymorphism was noted in alcohol dependence by Lutz et al., [12]. They found that the T - allele frequency increased from 0.28 in healthy control subjects to 0.33 in alcohol dependent patients suffering from mild withdrawal symptoms up to 0.40 in alcohol dependent men with a history of withdrawal seizure [12]. De Bree et al. [33] found that the frequency of substance

carrier genotypes of the MTHFR C677T gene could be explained by the following mechanism; MTHFR synthesizes 5-methyltetrahydrofolate, the major carbon donor in the remethylation of homocysteine to methionine. C to T mutation causes the substitution of valine for alanine at amino acid 223 and renders the enzyme less efficient (raising levels of homocysteine especially in TT genotype) and thermolabile [23,25].

Explanation for how hyperhomocysteinemia (associated with T carrier genotype of the MTHFR C677T gene in the current study) could harm brain and increase Heroin dependence can be extracted from findings of previous studies. MTHFR C677T gene is linked with low folate levels and is associated with increased incidence of neurotube defects [26]. Moreover, in patients with severe MTHFR deficiency cerebral demyelination was detected [25]. In addition, MTHFR is involved in remethylation of homocysteine to methionine which is central for dopamine methylation and the synthesis of neurotransmitters [11]. In addition, hyperhomocysteinemia is associated with global DNA hypomethylation [27]. Major life events occurring before the onset of Heroin dependence and/or chronic Heroin use were found to modify DNA methylation [28]. The polymorphisms within the MTHFR gene of homocysteine related neurotransmitters like taurine (play an important role in oxidative process). Homocysteine increase superoxide

Limitation of the study

Although we tried hard to exclude ethnicity in this study using through history taking but this may not be true. Moreover, the present study population was accessed at two private hospitals this may excluded a large portion of the population who could not be treated in private hospital. In our locality no public hospital treating addiction is present.

Conclusion

MTHFR polymorphism may contribute to the incidence and severity of heroin dependence, but that numerous other factors may play an equal or more significant role. In addition, plasma homocysteine may have a role in pathogenesis of Heroin related disorders. Multicenter studies and larger samples are recommended in further research to confirm the result of this study.

References

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