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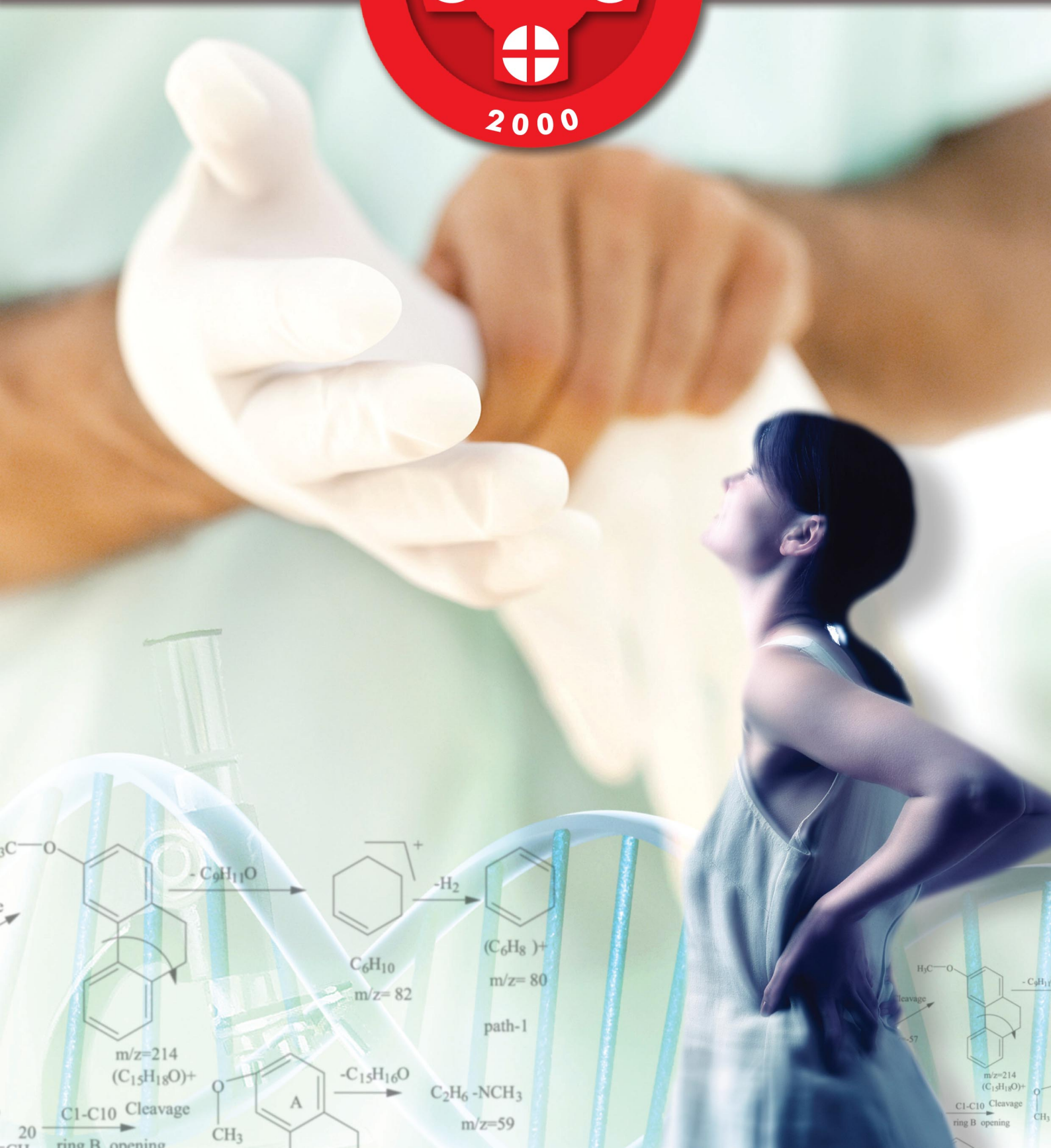
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# HOMOCYSTEINE: NEUROTOXICITY AND MECHANISMS OF INDUCED HYPEREXCITABILITY

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## HOMOCISTEIN: NEUROTOKSIČNOST I MEHANIZMI INDUKOVANE HIPEREKSCITABILNOSTI

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### ABSTRACT

Within the past four decades, investigators worldwide have established that the amino acid homocysteine (Hcy) is a potent, independent, novel and emerging risk factor for arteriosclerosis. In addition, Hcy is considered a vasotoxic and neurotoxic agent that interferes with fundamental biological processes common to all living cells. The aim of this article is to present data addressing central nervous system hyperexcitability in experimental models of homocysteine-induced seizures. These results demonstrate that acute administration of homocysteine and homocysteine-related compounds significantly affects neuronal activity, electroencephalographic (EEG) recordings and behavioural responses in adult Wistar male rats. It is suggested that hyperhomocysteinemia may generate similar effects on human brain activity. Homocysteine and D,L-homocysteine thiolactone are excitotoxic and elicit seizures through stimulation of NMDA receptors. In addition, they inhibit  $\text{Na}^+/\text{K}^+$ -ATPase activity in the cortex, hippocampus and brain stem. Nitric oxide (NO) acts as an anticonvulsant in D,L-homocysteine thiolactone-induced seizures and prevents D,L-homocysteine thiolactone-induced inhibition of  $\text{Na}^+/\text{K}^+$ -ATPase activity. EEG monitoring for the effects of ethanol on homocysteine-induced epilepsy support the idea that ethanol intake could represent one of the exogenous factors influencing brain excitability. Finally, it was found that homocysteine thiolactone significantly inhibits acetylcholinesterase (AChE) activity in rat brain tissue.

**Keywords:** homocysteine, CNS, hyperexcitability, rat

#### Abbreviations used:

AChE - acetylcholinesterase  
BHMT - betaine-Hcy S-methyltransferase  
CNS - central nervous system  
EEG - electroencephalography  
Hcy - homocysteine  
NO - nitric oxide  
SWDs - spike-wave discharges  
tHcy - total plasma homocysteine

### SAŽETAK

Četiri decenije istraživačkog rada širom sveta dovele su do ustanovljavanja semi-esencijalne sulfhidrilne aminokiseline homocisteina (Hcy) kao novog potentnog i nezavisnog faktora rizika za arteriosklerozu, kao i vazotoksične i neurotoksične supstance koja učestvuje u fundamentalnim biološkim procesima svojstvenim svim živim ćelijama. Cilj ovog rada je da prikaže dostupna saznanja na temu hiperekscitabilnosti centralnog nervnog sistema u eksperimentalnom modelu homocisteinom indukovane epilepsije. Rezultati pokazuju da akutna administracija homocisteina i homocisteinu-srodnih supstanci značajno utiče na nervnu aktivnost, elektroencefalografske (EEG) zapise i ponašanje odraslih mužjaka soja Wistar pacova, a smatra se da hiperhomocisteinemija može imati slične efekte i na aktivnost ljudskog mozga. Homocistein i D,L-homocistein tiolakton su ekscitotoksični i izazivaju epileptične napade stimulacijom NMDA receptora; oni indukuju značajnu inhibiciju aktivnosti  $\text{Na}^+/\text{K}^+$ -ATPaze u korteksu, hipokampusu i moždanom stablu. Azot monoksid (NO) deluje kao antikonvulzant u D,L-homocistein tiolakton indukovanim epileptičnim napadima i vrši prevenciju D,L-homocistein tiolakton indukovane inhibicije aktivnosti  $\text{Na}^+/\text{K}^+$ -ATPaze. EEG monitoring efekata etanola na homocisteinom indukovanu epilepsiju podržava ideju da akutni unos etanola može predstavljati jedan od faktora egzogenog uticaja na moždanu ekscitabilnost. Takođe je utvrđeno da D, L-homocistein tiolakton u značajnom procentu inhibira aktivnost acetilholinesteraze u mozgu pacova.

**Ključne reči:** homocistein, CNS, hiperekscitabilnost, pacov

#### Korišćene skraćenice:

AChE – acetilholin esteraza  
BHMT - betain-homocistein S-metiltransferaza  
CNS – centralni nervni sistem  
EEG - elektroencefalografija  
Hcy - homocistein  
NO – azot monoksid  
SWDs – "šiljak pražnjenja"  
tHcy – ukupni plazma homocistein

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## BACKGROUND

Within the past four decades, investigators worldwide have established that the amino acid homocysteine (Hcy) is a novel and independent risk factor for arteriosclerosis. In addition, Hcy is a vasotoxic and neurotoxic agent that interferes with the fundamental biological processes common to all living cells. Therefore, it has also been termed the “cholesterol of the 21<sup>st</sup> century.”

Metabolism of homocysteine is coordinately regulated to maintain a balance between remethylation and transsulfuration pathways, which are critical to maintaining low levels of this potentially cytotoxic sulphur containing amino acid (1). Homocysteine belongs to a group of molecules known as cellular thiols and is considered a “bad” thiol. Glutathione and cysteine, the most abundant cellular thiols, are considered to be “good” thiols (2). In the methylation pathway that occurs in all tissues, homocysteine acquires a methyl group to form methionine in a vitamin B12 dependent reaction, which is catalysed by methionine synthase. The second substrate required by methionine synthase is 5-methyltetrahydrofolate, which is formed by the reduction of 5,10-methylenetetrahydrofolate. This reaction is catalysed by the enzyme methylenetetrahydrofolate reductase (MTHFR). The TT genotype of this enzyme causes thermolability and reduces enzyme activity, which impairs the formation of 5-methyltetrahydrofolate. This reduced enzyme activity explains why this genotype is associated with increased Hcy levels when the folate status is relatively low. However, liver, kidney and the lens of the eye have the ability to convert Hcy to L-methionine through a vitamin B12-independent reaction catalysed by betaine-Hcy S-methyltransferase (BHMT). The CNS lacks BHMT and it is completely dependent on the folate and vitamin B12 pathway for the conversion of Hcy to L-methionine.

Homocysteine condenses with serine to form cystathionine in an irreversible reaction catalysed by the B6 containing enzyme cystathionine beta-synthase, which is known as the transsulfuration pathway. Hcy catabolism requires vitamin B6 and as a consequence, alterations in folic acid and B vitamin status impairs Hcy biotransformation. These alterations result in the synthesis of cysteine, taurine and inorganic sulphates that are excreted in urine.

Elevation of homocysteine levels is known to lead to the metabolic conversion and inadvertent elevation of homocysteine thiolactone, which is a reactive thioester representing less than 1% of total plasma homocysteine. In all cell types, from bacteria to human, homocysteine is metabolised to homocysteine thiolactone by methionyl-tRNA synthetase (3). Homocysteine thiolactone causes lethality, growth retardation, blisters and abnormalities in somite development by oxidative stress, which is one of important mechanisms for its toxicity to neuronal cells (4). The highly reactive and toxic homocysteine metabolite, homocysteine thiolactone, can be produced in two steps by enzymatic and/or non-enzymatic reactions in blood serum. The ability to detoxify or eliminate homocysteine thiolactone is essential for biological integrity (3, 4).

Total plasma Hcy (tHcy) consists of a pool of free homocysteine, homocystine, Hcy-S-S-Cys disulphide, protein bound N- and S-linked Hcy as well as their oxidised forms and Hcy-thiolactone (5-7). Under physiological conditions, less than 1% of the total Hcy is present in a free reduced form. Approximately 10–20% of total Hcy is present in different oxidised forms, such as Hcy-Cys and homocystine, which is a Hcy dimer. Plasma tHcy levels are affected by genetic, physiologic (age and sex) and lifestyle factors as well as various pathologic conditions (5, 6, 8). Hyperhomocysteinemia is defined as an elevated plasma total homocysteine (tHcy) concentration (>15  $\mu$ M).

Elevated tHcy is a recognised risk factor for cardiovascular disease (9) and has been linked to neurological diseases during aging, such as cognitive declines, cerebrovascular disease and stroke, vascular dementia and Alzheimer’s disease. In addition, elevated tHcy levels are linked with pathological brain functioning, such as mental retardation, depression, schizophrenia and memory impairment. Hcy is also a pro-thrombotic and pro-inflammatory factor, vasodilatation impairing substance and an endoplasmatic reticulum stress inducer (10, 11).

Mitrovic et al. (12) evaluated total plasma homocysteine in patients with angiographically confirmed coronary artery disease. In addition, they investigated the effects of homocysteine lowering therapy on endothelial function, carotid wall thickness and myocardial perfusion. Their study found that homocysteine levels decreased significantly (34%) with folic acid therapy and that endothelium function improved by 27% with this treatment. However, carotid structure and myocardial perfusion did not show any significant improvement in patients with confirmed coronary artery disease (13). Djuric et al. (14, 15) also investigated the link between homocysteine and folic acid, demonstrated folic acid induced coronary vasodilation and decreased oxidative stress in the isolated rat heart.

## HOMOCYSTEINE AND NEUROTOXICITY

Despite several theories, a complete understanding of Hcy toxicity remains unclear. In particular, the hypothesis that relates homocysteine to CNS dysfunction by way of overt neurotoxicity is based on the neuroactive properties of homocysteine. Elucidating the link between homocysteine and CNS dysfunction is vital for improving the treatment of homocysteine related CNS disorders. Hcy adversely affects brain functioning (i.e., mental retardation, dementia and memory impairment) and high tissue concentrations cause oxidative stress and excitotoxicity in neurons (16). When these symptoms are combined with homocystinuria, patients often present with convulsions (17). These studies provide evidence for a complex and multifaceted relationship between homocysteinemia and CNS disorders. Hcy is an endogenous compound that is neurotoxic at supraphysiological concentrations and induces neuronal damage and cell loss through excitotoxicity and apoptosis. These CNS



pathologies could be a direct consequence of the inability of cerebral tissue to metabolise Hcy through the betaine and transsulfuration pathways, thereby favouring Hcy accumulation in the nervous system (18). High brain concentrations of either Hcy or its oxidised derivatives have been shown to alter neurotransmission (19). The accumulation of homocysteine (at synapses or in the extracellular space) increases intracellular S-adenosylhomocysteine (SAH), which is a potent inhibitor of many methylation reactions that are vital for neurological function including the O-methylation of biogenic amines. Methylation of myelin basic protein and reducing the synthesis of phosphatidyl choline, which can lead to blood-brain barrier (BBB) disruption, are a few pathological processes that are possible in the absence of normal methylation patterns (20).

It has been recently suggested that Hcy toxicity is a consequence of its covalent binding to proteins, which interferes with protein biosynthesis and decreases the normal physiological activity of proteins. This modification of protein function is a process called homocysteinylation. Transthyretin is a plasma protein that is modified through homocysteinylation (3, 21). Other reports suggest that Hcy exerts its toxicity through the induction of endoplasmic reticulum (ER) stress. Increased intracellular Hcy concentrations are associated with both the alteration of redox balance and post-translational protein modifications through N- and S-homocysteinylation (22). Other studies suggest that Hcy induces the expression of superoxide dismutase in endothelial cells, which leads to the consumption of NO and impairs endothelial vasorelaxation (23).

## HOMOCYSTEINE AND SEIZURES

Experimental models of generalised clonic-tonic seizures induced by metaphit (1-[1(3-isothiocyanatophenyl)cyclohexyl] piperidine) (24, 25) and lindane ( $\gamma$ -1,2,3,4,5,6-hexachlorocyclohexane) (26, 27) are widely used for the study of epilepsy and the preclinical evaluation of potential antiepileptic treatments. In contrast to audiogenic epilepsy induced by metaphit with its typical and characteristic three degrees of intensity, the lindane model is similar to Hcy-induced epilepsy with a wide spectra of behavioural manifestations (e.g., hot plate reaction and Kangaroo position) incorporated into the rating scale from grades 0 - 4. Almost four decades ago, Sprince et al. (28) described that elevated levels of homocysteine arising from excess dietary methionine may induce epilepsy and lethality. The fact that increased homocysteine concentrations damage endothelial structures and the blood coagulation system during aging and in patients on antiepileptic drugs (29) justifies the attention directed toward the examination of homocysteine's neural effects.

Furthermore, experimental models of epilepsy may be induced by manipulations of  $\gamma$ -aminobutyric acid (e.g., bicuculline, corasol, picrotoxin and benzylpenicillin sodium) (30). In contrast, epilepsy can be induced by an opposite

mechanism such as by increasing cerebral excitatory neurotransmission. There are several proposed mechanisms by which exposure to excess D,L-homocysteine or D,L-homocysteine thiolactone induce seizures. The most reactive natural amino acid is homocysteic monocarboxylic, which is sulphur-containing and its homocysteic sulphinic acid derivatives are excitatory amino acids that have been proposed as candidates for brain excitatory neurotransmitters (31). However, answering the question as to whether D,L-homocysteine thiolactone should be included in the list of epileptogenic factors according to behavioural and electrophysiological responses will require additional studies and future human therapeutic trials. Increased levels of Hcy and its oxidised forms can provoke seizures by increasing activation of certain neuronal receptors such as N-methyl d-aspartate (NMDA),  $\alpha$ -amino-3-hydroxy-5-methyl-4-isoxazolepropionate (AMPA)/kainate ionotropic glutamate receptors or group I and group III metabotropic glutamate receptors (5). All of these glutamate receptors are expressed in hippocampal pyramidal cells and may directly induce or drive these cells over the threshold for excitotoxic cell death. Overstimulation of these receptors triggers cytoplasmic calcium pulses,  $\text{Ca}^{2+}$  influx and intraneuronal calcium mobilisation in the presence of glycine (32). Increased cytosolic  $\text{Ca}^{2+}$  concentrations affect enzyme activities and the synthesis of nitric oxide, which is a retrograde messenger that overstimulates excitatory amino acid receptors including glutamate, thereby lowering the convulsive threshold (33). However, expression of the NMDA receptor is not confined to neurons. Other cell types, including endothelial cells from cerebral tissue, can express this receptor. Free radicals induce the up-regulation of the NR1 subunit of the NMDA receptor, thereby increasing the susceptibility of cerebral endothelial cells to excitatory amino acids, which can lead to blood brain barrier disruption (34). Microglia are also subject to the toxic effects of Hcy (35). Hcy can induce convulsions in adults and in immature experimental animals through modulating the activity of metabotropic glutamate receptors (mGluRs) (36). Kubova et al. (37) stated that the main epileptic phenomenon in homocysteine thiolactone treated rats was emprosthotonic, flexion seizures, which are observed in young animals but never seen in rats older than 25 days. Homocysteine was shown to elicit minor (predominantly clonic) and major (generalised tonic-clonic) seizures during ontogenesis in immature Wistar rats (36, 37). It seems that Hcy exerts a direct excitatory effect comparable to the action of glutamate (17). In addition, Hcy has been shown to enhance either the release or uptake of other endogenous excitatory amino acids (36).

Classical anticonvulsants such as phenytoin, carbamazepine and valproic acid lower plasma folate levels and significantly increase homocysteine levels, which induces epileptogenesis and reduces the control of seizures in patients with epilepsy (38). On the other hand, increasing excitability with this native amino acid further contributes to epileptogenesis and disturbs the balance between excitation and inhibition in





the brain. Following the work of Djuric's study, Stanojlovic et al. administered homocysteine (i.p.) to adult rats, and for the first time, convulsive and non-convulsive seizures were recorded (39). This form of epilepsy was followed by two-wave patterns of seizures with one replacing the other. It has been suggested that D,L-homocysteine thiolactone may be considered as an excitatory metabolite, which is capable of becoming a natural convulsant if accumulated to a great extent in the brain (39). Hyperhomocysteinemia in awake adult male Wistar rats induced a generalised seizure disorder characterised by recurrent unprovoked clonic-tonic convulsions and absence-like seizures as well as epileptic electrical discharges. In addition to a prolonged median latency to the first seizure, the seizure incidence, median seizure episode severity and median number of seizure episodes per rat were significantly higher in all Hcy treated groups (39). Non-convulsive status epilepticus can occur from a variety of causes, including primarily generalised absence epilepsy, genetic origins (Wakayama or tremor epileptic rats) or pharmacologically (penicillin, pentylentetrazole or  $\gamma$ -hydroxy-butyric acid) induced models (40). Behavioural immobility during motor cortex activation and the occurrence of generalised spike-wave activity are among the most puzzling phenomena in absence epilepsy. According to one of numerous hypotheses addressing the development of spike-wave discharges (SWDs), these electrographic discharges may belong to the same class of phenomena such as sleep spindles. Sleep spindles are normally generated sleep rhythms that transformation one, two or more spindle waves into the spike component of the SWD (41). Cortico-thalamo-cortical oscillatory network connections may play a pacemaker role in the electrogenesis of oscillations and primarily represent the neurophysiological basis underlying the initiation and propagation of SWD (40).

In keeping with the well-known fact that rhythmic bursts of spikes represent an electrophysiological marker of a hyperexcitability, Folbergrova et al. (42) found a very poor electroclinical correlation between electroencephalographic (EEG) patterns and motor phenomena in immature rats. The epileptogenic process is closely associated with the changes in neuronal synchronisation, and non-convulsive, generalised epilepsy is characterised by brief episodes of unpredictable and unresponsive behaviours with a sudden arrest accompanied by SWDs. However, the spike-wave complexes of SWDs have a unique shape. Bilateral, high-voltage, synchronous, spindle-like electrical oscillations, which are common phenomenon during paroxysmal EEG attacks, are termed SWDs. These are associated with sudden motor immobility and minor clinical signs, such as the loss of responsiveness with rhythmic twitches of vibrissae or cervicofacial musculature and are seen after i.p. administration of D,L-homocysteine thiolactone in adult rats. Interestingly, absence-like animals showed no reaction to these different stimuli (e.g., audiogenic, tactile).

Stanojlović et al. (39) found poor electro-clinical correlations in Hcy treated rats. It is worth mentioning that electrographic seizure discharges were absent even during motor convulsions of grade 3 or 4. On the contrary, EEG seizures

without motor symptoms were regularly observed. These aforementioned EEG signatures were distinguishable from sleep spindles (10-16 Hz) in regard to their frequency, duration, morphology (sleep spindles are more stereotyped than SWD waves) and moment of occurrence. For example, SWDs occur during passive wakefulness, while sleep spindle-like oscillations occur during high amplitude delta activity (43). The poor prognosis for the Hcy group of animals appears to be a result of the combined systemic damage caused by continuous seizure activity as well as the excitotoxic and neurotoxic effects of homocysteine accompanied by the action of free radicals, including oxidative brain damage (42, 44). In most of the related studies, it is not quite clear whether the observed effects are due to homocysteine itself or to homocysteine metabolites. Notably, all of the above mentioned mechanisms induce cell death (45) and result in a high mortality rate (100%).

The basic function of the  $\text{Na}^+/\text{K}^+$ -ATPase is to maintain the high  $\text{Na}^+$  and  $\text{K}^+$  gradients across the plasma membrane of animal cells and it is essential for the generation of the membrane potential and maintenance of neuronal excitability (2). In addition, it has been demonstrated that there is a moderate inhibition of rat hippocampal  $\text{Na}^+/\text{K}^+$ -ATPase activity by D,L-homocysteine, which is not observed in the cortex and brain stem (46). In contrast, D,L-homocysteine thiolactone strongly inhibits  $\text{Na}^+/\text{K}^+$ -ATPase activity in cortex, hippocampus and the brain stem of rats, thereby affecting the membrane potential with deleterious effects for neurons. Considering that hyperhomocysteinemia increases formation of superoxide and hydrogen peroxide and that excitotoxicity indirectly provokes increased intracellular free radicals production, it is feasible that oxidative stress may also be associated with the CNS injury caused by homocysteine and D,L-homocysteine thiolactone (47).

Otherwise, nitric oxide (NO) is a highly reactive second messenger molecule synthesised in a number of tissues. It serves a key role in interneuronal communications via modulating release of classical neurotransmitters and influencing the excitability status of neurons (33). It is produced from L-arginine by the action of a family of enzymes known as NO synthases (NOS). Neural (nNOS) and endothelial NOS (eNOS) are  $\text{Ca}^{2+}$ /calmodulin-dependent enzymes, while inducible NOS (iNOS) shows  $\text{Ca}^{2+}$ -independent properties. N-nitro-L-arginine methyl ester (L-NAME) is a non-selective NOS inhibitor commonly used to decrease NO levels. Recently, it has been determined that NO serves a role in the mechanisms underlying D,L homocysteine thiolactone induced seizures. This was demonstrated by testing the action of L-arginine (NO precursor) and L-NAME (NOS inhibitor) on behavioural and EEG manifestations of D,L homocysteine thiolactone induced seizures (48). Following Djuric's study, Hrcnic et al. (48) showed functional involvement of NO in the convulsive activity of D,L-homocysteine thiolactone induced seizures in adult rats. Pretreatment with L-NAME in a dose-dependent manner increased seizure incidence and severity and shortened latency time to the first seizure following injection with a subthreshold dose of D,L-homo-



cysteine thiolactone (5.5 mmol/kg, i.p.). Nitro indazole is an inhibitor of neuronal nitric oxide synthase that attenuates pilocarpine-induced seizures (49). NO has been shown to modulate NMDA receptor activity by interacting with the -SH group of its redox modulatory site via S-nitrosylation. This modification results in the downregulation of this receptor complex (19) and prevents the neurotoxic effects of an excessive  $\text{Ca}^{2+}$  influx during homocysteine induced overstimulation of NMDA and mGluRs I receptors. In addition, Kim et al. (50) demonstrated that NO ameliorates homocysteine's adverse effects of S-nitrosylation in cultured rat cortical neurons. Moreover, NO induces a reduction of glutamate by activation of glial cells. Many experimental studies have demonstrated co-localisation of NOS and GABA and have suggested that basal NO levels induce a depression of inhibition, while high concentrations of NO increase GABA release. In contrast to homocysteine, which increases oxidative stress by the production of reactive oxygen species, NO can act as a neural protector. This neuroprotective property is due to the formation of S-nitroso-L-glutathione, which is an antioxidant and NO scavenging molecule (51). However, the role of NO in epileptogenesis has been studied in different experimental models and reported results are highly contradictory. Namely, the proconvulsive role of NO has been demonstrated in the lindane model of convulsions and several others (52). Thus, it seems that the activity of NO depends on the animal strain, the seizure model employed and the type and dose of drugs used in order to modify cerebral NO levels.

Furthermore, it has been demonstrated that when L-arginine is applied alone, it significantly increases the activity of  $\text{Na}^+/\text{K}^+$ -ATPase activity in the hippocampus, the cortex and the brain stem. However, when L-arginine is applied prior to D,L homocysteine thiolactone (8 mmol/kg), this completely reverses its inhibitory effect (48). The same holds true for its effects on  $\text{Mg}^{2+}$ -ATPase activity in the rat cortex and the brain stem. L-NAME increases  $\text{Na}^+/\text{K}^+$ -ATPase activity in the cortex and the brain stem but not in the hippocampus. When L-NAME was administered prior to homocysteine thiolactone (5.5 mmol/kg), it increased the activity of both  $\text{Na}^+/\text{K}^+$ -ATPase and  $\text{Mg}^{2+}$ -ATPase in the rat cortex and the hippocampus.

Changes in total spectral power density after ethanol alone and together with D,L-homocysteine thiolactone in adult rats were examined (53). Recording electrical activity from the brain represents a measure of both brain function and dysfunction. Ethanol is used as a social substance and is the second most widely used psychoactive substance in the world after caffeine. The influence of ethanol on the central nervous system depends on the dose, drinking pattern, tolerance and other factors. While chronic ethanol consumption is followed by a series of seizures during the withdrawal period, acute ethanol intake exerts mainly inhibitory effects on the CNS and is usually associated with an increase of seizure threshold (54). Rasic-Markovic et al. (53) found that ethanol's action on electrographic pattern is biphasic, which is char-

acterised by potentiation of epileptiform activity in one dose range and depression in another one. Low ethanol doses causing euphoria and behavioural arousal are associated with desynchronisation of the EEG, decreases in the mean amplitude and increases in theta and alpha activity. In addition, ethanol increases mean total spectral power density 15 min and 30 min after administration. Ethanol affects voltage-gated ion channels, second messenger systems and a variety of different neurotransmitter systems such as glycine, acetylcholine as well as monoamines and neuropeptides systems (55). Two major amino acid neurotransmitter systems, GABA and glutamate, as well as aspartate are affected by ethanol. Acute administration of ethanol inhibits NMDA-induced  $\text{Ca}^{2+}$  influx, cyclic GMP production, neurotransmitter release and reduces NMDA-evoked neurotoxicity (56, 57).

Pre-treatment with MK-801, which is a NMDA receptor antagonist, showed a tendency to reduced the incidence of convulsions, latency to the first seizure onset and the severity of seizure episodes; however, there was no statistical significance when compared to the D,L-homocysteine thiolactone treated group. Nevertheless, the median number of seizure episodes was significantly decreased by MK-801 when compared to the D,L-homocysteine thiolactone treated group. On the other hand, ifenprodil, which is another type of NMDA receptor antagonist, decreased the latency to the first seizure onset and increased the median number of seizure episodes. The majority of seizure episodes in the ifenprodil (72.1%) and MK-801 (73.1%) groups were significantly different compared to the D,L-homocysteine thiolactone treated group (36.0%). Our findings suggest that D,L-homocysteine thiolactone induces seizures through the stimulation of NMDA receptors in the central nervous system but other mechanisms (i.e., NO signalling) may also be involved (58).

Finally, limited data exist in the literature regarding the effects of homocysteine and D,L-homocysteine thiolactone on the activity of the acetylcholinesterase (AChE) enzyme in the blood, but practically no data exist regarding the influence of these compounds on this enzyme in the brain and heart. Recent results showed a significant reduction in AChE activity in all tissues obtained from rats treated with D,L-homocysteine thiolactone compared to the enzyme activity of the control group. In addition, these results also showed that the blood enzyme activity was the lowest (12%) after treatment, while the enzyme activity was slightly higher in the brain (27.8%) and heart samples (86.3%). Therefore, it was concluded that D,L-homocysteine thiolactone significantly inhibited AChE activity in the heart and brain tissue but not in the blood of the rat (59).

Overall, these studies clearly demonstrate that acute administration of homocysteine and especially D,L-homocysteine thiolactone elicit seizures in adult rats, affect neuronal activity, EEG recordings and behavioural responses. These effects have been linked to the stimulation of NMDA receptors, inhibition of the  $\text{Na}^+/\text{K}^+$ -ATPase, inhibition of AChE activity and the functional involvement of NO during D,L-homocysteine thiolactone induced seizures in adult rats.





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While dysfunction of the prefrontal cortex has been repeatedly implicated in the pathophysiology of schizophrenia, the role of serotonin in this brain region in schizophrenia is unclear. We therefore examined the effects of local serotonin depletion in the medial prefrontal cortex on psychotomimetic drug-induced locomotor hyperactivity and prepulse inhibition, two animal models of aspects of schizophrenia. Pentobarbital-anaesthetised (60 mg/kg, i.p.) male Sprague-Dawley rats were stereotaxically micro-injected with 0.5 µl of a 5 µg/µl solution of the serotonin neurotoxin 5,7-dihydroxytryptamine into the medial prefrontal cortex. Two weeks after the surgery, rats underwent behavioural testing. When compared to sham-operated controls, rats with medial prefrontal cortical lesions did not show changes in either psychotomimetic drug-induced locomotor hyperactivity or prepulse inhibition. However, following the administration of the serotonin neurotoxin into the medial prefrontal cortex, the concentration of serotonin was reduced by 60%. These results suggest that serotonin depletion in the medial prefrontal cortex does not lead to dysregulation of psychotomimetic drug-induced locomotor hyperactivity and does not

**Saline**

Time	Sham	mPFC
Basal	~1500	~1500
0-30	~1000	~1000
30-60	~1000	~1000
60-90	~500	~500

**Amphetamine**

Time	Sham	mPFC
Basal	~2000	~2000
0-30	~5000	~5000
30-60	~8000	~8000
60-90	~4000	~4000

**Amphetamine**

Time	Sham	mPFC
Basal	~2000	~2000
0-30	~5000	~5000
30-60	~8000	~8000
60-90	~4000	~4000



# THE EFFECT OF SEROTONERGIC LESIONS IN THE MEDIAL PREFRONTAL CORTEX ON PSYCHOTOMIMETIC DRUG-INDUCED LOCOMOTOR HYPERACTIVITY AND PREPULSE INHIBITION IN RATS

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## ABSTRACT:

While dysfunction of the prefrontal cortex has been repeatedly implicated in the pathophysiology of schizophrenia, the role of serotonin in this brain region in schizophrenia is unclear. We therefore examined the effects of local serotonin depletion in the medial prefrontal cortex on psychotomimetic drug-induced locomotor hyperactivity and prepulse inhibition, two animal models of aspects of schizophrenia. Pentobarbital-anaesthetised (60 mg/kg, i.p.) male Sprague-Dawley rats were stereotaxically micro-injected with 0.5  $\mu$ l of a 5  $\mu$ g/ $\mu$ l solution of the serotonin neurotoxin 5,7-dihydroxytryptamine into the medial prefrontal cortex. Two weeks after the surgery, rats underwent behavioural testing. When compared to sham-operated controls, rats with me-

dial prefrontal cortical lesions did not show changes in either psychotomimetic drug-induced locomotor hyperactivity or prepulse inhibition. However, following the administration of the serotonin neurotoxin into the medial prefrontal cortex, the concentration of serotonin was reduced by 60%. These results suggest that serotonin depletion in the medial prefrontal cortex does not lead to dysregulation of subcortical dopaminergic activity and does not cause aberrant responses to environmental stimuli.

**Keywords:** schizophrenia, serotonin, medial prefrontal cortex, prepulse inhibition

**Running title:** Serotonin, the medial prefrontal cortex and behaviour

## Abbreviations used:

5,7-DHT- 5,7-dihydroxytryptamine

5-HT- 5-hydroxytryptamine, serotonin

5-HT<sub>1-7, A-F</sub>- serotonin receptor subtypes

ANOVA- analysis of variance

D<sub>1-5</sub>- dopamine receptors 1-5

DRN- dorsal raphe nucleus

GABA-  $\gamma$ -aminobutyric acid

HPLC- high pressure liquid chromatography

i.p.- intraperitoneal

MRN: median raphe nucleus

mPFC- medial prefrontal cortex

NMDA- N-methyl-D-aspartate

PPI- prepulse inhibition

PP- prepulse intensity

PP8- prepulse of 8 dB

s.c.- subcutaneous

SEM- standard error of the mean

VTA- ventral tegmental area

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## INTRODUCTION

Schizophrenia is a chronic and severe psychiatric disorder that generally occurs in late adolescence or early adulthood. Approximately 1% of the population worldwide is affected by schizophrenia, placing a heavy burden on society, both in terms of emotional suffering and economic loss [1]. Schizophrenia is the classic example of a disorder that always has psychosis as one of its features [2]. As psychotic episodes are extremely debilitating, management and treatment aim to reduce and eliminate these dramatic personality changes, consisting of irritability, confusion and paranoia associated with hallucinations and delusions. Antipsychotic drugs are used to treat nearly all forms of psychosis, including schizophrenia. It has been generally accepted that the mechanism by which antipsychotic drugs decrease hallucinations and delusions is mediated at least in part by dopamine  $D_2$  receptor blockade [3-5]. Moreover, atypical antipsychotic drugs, such as olanzapine, display a unique neuropharmacological profile; they minimise psychoses by interacting with a number of neurotransmitter and receptor systems through binding at multiple receptor sites. Olanzapine has a high affinity for serotonin 5-HT<sub>2A</sub>, dopamine, cholinergic, histamine and  $\alpha$ 1-adrenergic receptors [6]. However, the most important mechanisms underlying the clinical properties of atypical antipsychotic drugs appear to be mediated by interactions with the serotonin 5-HT<sub>2A</sub> receptor subtype [7, 8].

Serotonin is one of the major neurotransmitters in the human brain and plays a central role in the regulation of a wide range of behaviours, such as mood, eating and the stress response [9-11]. The serotonergic projections arising from the brainstem raphe nuclei form the largest and most complex efferent system in the human brain [12, 13]. The axons of dorsal raphe nucleus (DRN) neurons contribute to the majority of the serotonergic innervation in the frontal cortex, ventral hippocampus and striatal regions [14, 15], while the axons of median raphe nucleus (MRN) serotonergic neurons are more abundant in the dorsal hippocampus and the cingulate cortex [16, 17]. The hypothalamus, the substantia nigra and the nucleus accumbens receive serotonergic innervation from both nuclei [18, 19]. This organisation of the serotonergic neuronal population suggests that serotonin is involved in the regulation of different functional systems, such as the motor, limbic and somatosensory systems [13]. Thus, it is not surprising that atypical antipsychotic medications target multiple brain serotonin receptor subtypes. As it is very difficult to assess alterations of serotonergic transmission in the pathophysiology of psychiatric disorders in the living human brain, animal models are needed.

Animal models of psychiatric disorders, including schizophrenia, rely on mimicking specific aspects or symptoms associated with the disease [20-22]. The two most widely used models are locomotor hyperactivity induced by psychotomimetic drugs and prepulse inhibition.

Psychotomimetic drugs, such as amphetamine and phencyclidine, can induce abnormal behaviours in animals and mimic certain aspects of psychotic disorders in humans [20, 23-25]. Amphetamine, an indirectly acting sympathomimetic, causes increased dopamine release from presynaptic terminals [26], and hyperlocomotion induced by amphetamine is dependent upon intact subcortical dopamine activity in the nucleus accumbens [27]. In contrast, phencyclidine interferes with multiple neurotransmitter systems [28]. Phencyclidine acts as a non-competitive antagonist at the ion channel associated with the N-methyl-D-aspartate (NMDA) glutamate receptor and also indirectly facilitates dopaminergic and serotonergic transmission [29]. Similar mechanisms are also activated in humans by phencyclidine [30]. Prepulse inhibition of the acoustic startle response is an operational measure of sensorimotor gating that is disrupted in patients with schizophrenia [31, 32] and in rats treated with drugs that facilitate dopaminergic activity [33-35]. Furthermore, prepulse inhibition is reduced in rats treated systemically with serotonin releasers, such as fenfluramine, direct 5-HT<sub>1A</sub> receptor agonists [36-38] and glutamate receptor antagonists, such as phencyclidine [39]. The prepulse inhibition-acoustic startle reflex model in rats offers a unique opportunity to assess attentional and information processing deficits in schizophrenia, as modulation of the startle responses is similar among mammalian species [40]. In animals, usually the whole body startle response is measured after exposure to acoustic or tactile stimuli, while in humans the eyeblink component of the startle response is measured [40].

There is a growing body of evidence that suggests that the hippocampus, amygdala and prefrontal cortex play an important role in the pathogenesis of schizophrenia. The activity of these brain regions may cause changes in subcortical dopaminergic activity and therefore lead to the inappropriate initiation of behavioural responses to external stimuli. We have previously reported that serotonergic projections into the hippocampus and amygdala are differentially involved in the regulation of psychotomimetic drug-induced locomotor hyperactivity and prepulse inhibition [41, 42]. As serotonergic projections from both raphe nuclei innervate the prefrontal cortex, in addition to the hippocampus and amygdala, the aim of the present study was to determine whether serotonergic lesions of the prefrontal cortex caused behavioural changes similar to those produced by lesions of the hippocampus and/or amygdala.

In humans, dysfunction of the prefrontal cortical areas, with which the medial prefrontal cortex of the rat is comparable, is related to psychopathology of schizophrenia and other psychiatric disorders (for a review, see [43]). A wealth of evidence from studies in animals and humans indicates that the medial prefrontal cortex (mPFC) is a key component of the cortico-limbic-striatal circuits that generate pathological emotional behaviour [44, 45]. The various subdivisions of the mPFC appear to serve separate and distinct functions. For example, ven-



tral regions of the mPFC (the prelimbic and the infralimbic cortices) have been associated with diverse emotional and cognitive processes [43, 46, 47]. The ventral mPFC is also of interest, as it has been strongly implicated in the expression of behavioural and autonomic responses to emotionally relevant stimuli [48]. As dysfunction of the prefrontal cortex has been repeatedly implicated in the pathophysiology of schizophrenia [49-52], studies in the rat have focused on elucidating the role of this region in paradigms such as locomotor activity and prepulse inhibition. Dopaminergic lesions and intra-mPFC infusion of selective dopamine receptor antagonists have been reported to disrupt prepulse inhibition [53, 54], whereas the intra-mPFC infusion of amphetamine has been shown to decrease systemic amphetamine-induced increases in locomotor activity in the open field test [55]. However, none of these studies have addressed the importance of serotonergic innervation of the mPFC in neural circuitry involved in the regulation of motor behaviour and prepulse inhibition. Therefore, the present study investigated the effects of local lesions of serotonergic projections into the mPFC on psychotomimetic drug-induced locomotor hyperactivity and prepulse inhibition.

## MATERIALS AND METHODS

### Experimental animals

A total of 25 male Sprague-Dawley rats (Department of Pathology, University of Melbourne), weighing 250-300 g at the time of surgery, were used in this study. The animals were housed under standard conditions in groups of two or three with free access to food and water. They were maintained on a 12 h light/dark cycle (lights on at 0700 h) at a constant temperature of 21°C. One week prior to the surgical procedure, the animals were handled each day over a five-day period. The experimental protocol and surgical procedures were approved by the Animal Experimentation Ethics Committee of the University of Melbourne, Australia.

### Drugs and solutions

D-amphetamine sulphate (Sigma Chemical Co., St. Louis, MO, USA) and phencyclidine HCl (PCP, Sigma) were dissolved in a 0.9% saline solution and injected subcutaneously (s.c.) into the nape of the neck. Desipramine HCl (Sigma) was dissolved in distilled water and injected intraperitoneally (i.p.) 30 min prior to the neurotoxin microinjection. All doses are expressed as the weight of the salt and were administered in an injection volume of 1 ml/kg body weight. The serotonergic neurotoxin, 5,7-dihydroxytryptamine (5,7-DHT) (Sigma), was dissolved in 0.1% ascorbic acid (BDH Chemicals, Kilsyth, VIC, Australia) in saline to prevent oxidation of the neurotoxin. Carprofen (50 mg/ml, Heriot AgVet, Rowville, VIC, Australia) was diluted in 0.9% saline to a dose of 5 mg/kg and injected s.c. immediately after the surgical procedure.

### Surgical procedure

The rats were pretreated with 20 mg/kg desipramine, 30 min prior to surgery, to prevent the destruction of noradrenergic neurons by 5,7-DHT [56]. The rats were subsequently anaesthetised with sodium pentobarbitone (60 mg/kg i.p., Rhone Merieux, QLD, Australia). The rats were mounted in a Kopf stereotaxic frame (David Kopf Instruments, Tujunga, CA, USA) with the incisor bar set at -3.3 mm [57]. The skull surface was exposed, and a small hole was drilled. A 25 gauge stainless-steel cannula, which was attached to a 10 µl glass syringe and connected via polyethylene tubing mounted in an infusion pump (UltraMicroPump, World Precision Instruments, Sarasota, FL, USA), was lowered into the mPFC. With bregma set to zero and the stereotaxic arm at 0°, the coordinates were as follows: mPFC lesions (n=13 for behavioural experiments, n=2 for histology): 3.2 mm anterior, 0.7 mm lateral and 4.5 mm ventral to bregma. A volume of 0.5 µl of 5,7-DHT (5 µg/µl) was infused over a period of 2 min on each side. Sham-operated controls (n=10) underwent the same surgical procedure and received an equal volume of vehicle solution. The injection volumes and rate of infusion were selected to minimise non-specific damage at the site of injection. Movement of the meniscus in the cannula was monitored to ensure successful infusion. After infusion, the cannula was left in place for a further 2 min to avoid backflow of the solution up the injection path. After lesioning, the skin was closed with silk-2 sutures (Cynamid, Baulkham Hills, NSW, Australia), and the animals were administered 5 mg/kg of carprofen, a non-steroidal, anti-inflammatory analgesic, to reduce post-operative inflammation and discomfort. The rats were placed on a heated pad until they recovered from the anaesthesia. After the surgery, the rats were allowed to recover for two weeks, during which they were handled regularly and health checks were made two to three times a week.

### Experimental design and apparatus

Behavioural tests were performed starting two weeks after the surgery, and each session included random numbers of 5,7-DHT-lesioned rats and sham-operated rats. Locomotor activity was monitored using eight automated photocell cages (31 x 43 x 43 cm, h x w x l, ENV-520, MED Associates, St. Albans, VT, USA). The position of the rat at any time was detected with sixteen evenly spaced infrared sources and sensors on each of the four sides of the monitor. The addition of a photobeam array above the subject added a second plane of detection to the system to detect rearing and vertical counts. This infrared beam array thus defined an X, Y and Z coordinate map for the system. The sensors detected the presence or absence of the infrared beam at these coordinates. Every 50 msec, the software checked for the presence or absence of the infrared beam at each sensor, allowing for the very precise tracking of the movement of a subject. Several types of behavioural responses were recorded, including distance moved, ambulation, stereotypy and rearing. Ambulatory counts con-





sisted of consecutive interruptions of at least four beams within a period of 500 ms. Small, repetitive beam breaks within a virtual box of 4 x 4 beams around the rat were recorded as stereotypic counts. Recordings of photocell beam interruptions by the rat were taken every 5 min and stored by the computer software. Three locomotor activity tests were done after treatment with saline, 0.5 mg/kg of amphetamine or 2.5 mg/kg of phencyclidine administered in a random order. These locomotor activity tests were done with three to four day intervals to prevent habituation due to repeated testing and to allow for clearance of the drugs. Prior to any drug manipulation, the rats were placed in the locomotor photocell cages for 30 min to establish baseline locomotor activity and allow for habituation to the test environment. After 30 min of spontaneous baseline activity, the rats were injected and locomotor activity was recorded over a further 90 min, generating a total session time of two hours. For the purpose of this paper, locomotor activity data were expressed as cumulative data from 30 min periods and presented as a time course of distance moved during the 30 min before injection and 90 min after injection.

After locomotor activity experiments, rats were tested for prepulse inhibition. This was done using six automated startle chambers (SR-LAB, San Diego Instruments, San Diego, CA, USA) consisting of clear Plexiglas cylinders, 9 cm in diameter, resting on a platform inside a ventilated, sound-attenuated and illuminated chamber. A speaker mounted 24 cm above the cylinder produced both continuous background noise at 70 dB and the various acoustic stimuli. Whole-body startle responses of the animal in response to acoustic stimuli caused vibrations of the Plexiglas cylinder, which were then converted into quantitative responses by a piezoelectric accelerometer unit attached beneath the platform. The percent prepulse inhibition was calculated as  $100 \times \{[\text{pulse-alone trials} - (\text{prepulse} + \text{pulse trials})] / (\text{pulse-alone trials})\}$  [31]. At least one day before the prepulse inhibition testing, rats underwent a pretest session where they were exposed to the testing cylinders and the testing protocol for the first time. This session was conducted to allow rats to habituate to the testing environment. Each rat was placed into the chamber for a 5 min acclimation period with a 70 dB background noise level that continued throughout the session. A single prepulse inhibition session lasted for about 45 min and consisted of high- and low-intensity stimulus combinations with a continuous background noise of 70 dB. The session started and ended with a block of ten pulse-alone trials of 115 dB. These blocks, together with twenty pseudo-randomly presented pulse-alone trials during the prepulse inhibition protocol, were used to calculate the basal startle reactivity and startle habituation. Prepulses were presented for 20 ms and differed in intensity. Prepulse inhibition was assessed by the random presentation of 115 dB pulses, ten each of prepulse-2, -4, -8, 12 and -16 and ten 'no-stimulus' trials. For example, prepulse-8 (PP8) is a 20-ms prepulse of 8 dB above the background noise, i.e., 78 dB, followed 100 ms

later by a 40-ms 115 dB pulse [58]. The interval between trials varied (10 - 37 s) to prevent conditioning of the responses. A microcomputer and an interface assembly that controlled the delivery of acoustic stimuli digitised and recorded the readings.

### **Tissue preparation for histology and high pressure liquid chromatography (HPLC)**

At the end of the experiments, rats were killed by decapitation and the brains were removed from the skull. The brains were placed on a cold plate. First, the frontal cortex was dissected bilaterally [59], and then the mPFC was dissected out. For the histological assessment of the location of the injection sites, 20- $\mu\text{m}$ -thick sections of the medial prefrontal cortex of two mPFC-lesioned rats were cut on a cryostat and mounted onto gelatin-coated glass slides. The sections were then stained with cresyl violet (ProSciTech, Thuringowa, QLD, Australia) and examined microscopically to verify the location of the tips of the infusion cannulas.

The HPLC measurements of the tissue serotonin (5-HT) concentration were carried out in 23 animals. The dissected structures were weighed and stored in Eppendorf tubes at  $-80^{\circ}\text{C}$  until the biochemical assays were performed. The tissue samples were homogenised in 500  $\mu\text{l}$  of 0.1 M perchloric acid by ultrasonication and centrifuged at 15,500 g for 5 min. A 50  $\mu\text{l}$  aliquot of the supernatant was injected into a high pressure liquid chromatography (HPLC) system to determine the content of 5-HT (ng/mg tissue of wet weight). The HPLC system consisted of a Waters Model 510 Solvent delivery system, a Waters U6K injector, an Alphasorb C18 125A 10 U 150\* 3.9 mm column and a Column & Spectra-Physics 970 D-A1 fluorescence spectrometer. The output signal from the fluorescence detector was analysed with the chromatography software package, 810 Baseline, version 3.31. The mobile phase used consisted of 9.8 g/l  $\text{KH}_2\text{PO}_4$ , 1.0 g/l  $\text{Na}_2\text{EDTA}$ , 5% acetonitrile and 1 ml/l triethylamine. The pH of the mobile phase solution was adjusted to 3.0 with 1 M HCl. Subsequently, the solution was filtered and degassed and delivered to the HPLC at a flow rate of 1 ml/min. Prior to sample testing, the following standards for 5-HT were run through the system: 12.5 ng/ml, 25 ng/ml, 50 ng/ml, 100 ng/ml and 200 ng/ml. Calibration curves were constructed, and the level of 5-HT in tissue samples was calculated relative to these standards. Each run lasted 8 min and the retention time for serotonin was 2.6 min.

### **Statistical analysis**

Data were expressed as the mean  $\pm$  the standard error of the mean (SEM). All of the statistical analyses were performed using the statistical software package SYSTAT 9.0 (SPSS Inc., Chicago, IL, USA). All of the data were analysed using an analysis of variance (ANOVA) with repeated measures where appropriate. In the locomotor activity experiments, data were summed in 30-min blocks, and these blocks were used to assess the main effects of the lesion type (group), the treatment with amphetamine or phency-



clidine (time) and the interactions between these factors. In this analysis, the time effect was a repeated-measures factor. The baseline and the drug-induced locomotor activity were part of the data set analysed. In the prepulse inhibition experiments, the factors were group and habituation (four blocks of ten startle responses) or group and prepulse (five different prepulse intensities), where habituation and prepulse were repeated-measures factors. After calculating ANOVAs for all of the surgery groups, subsequent pairwise ANOVAs were performed where needed. A 'p-value' of  $p < 0.05$  was considered to be statistically significant. For HPLC measurements, a one-way ANOVA was used, followed by the Bonferroni-corrected t-test comparison.

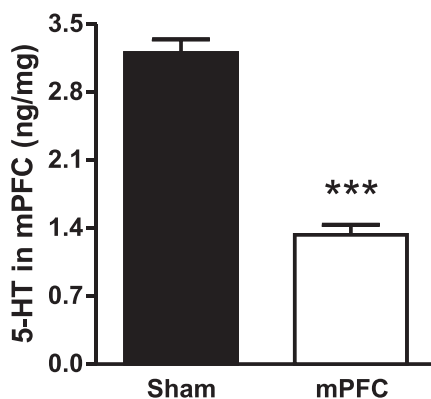
## RESULTS

### Histology: injection sites

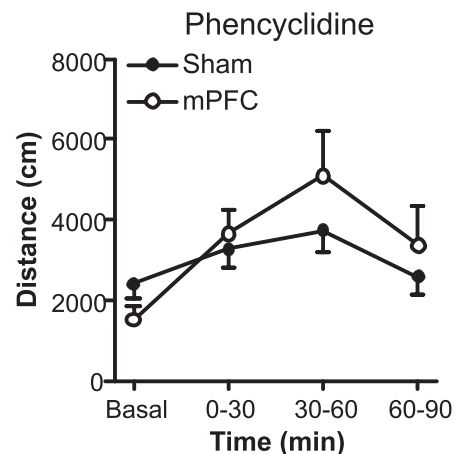
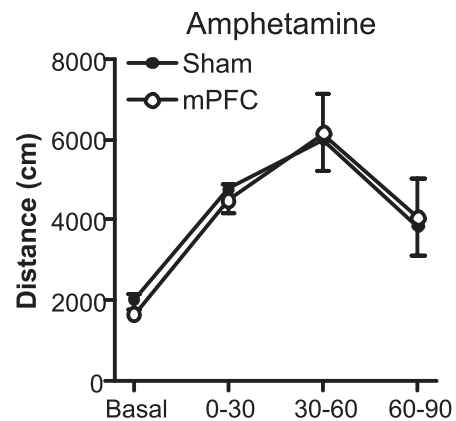
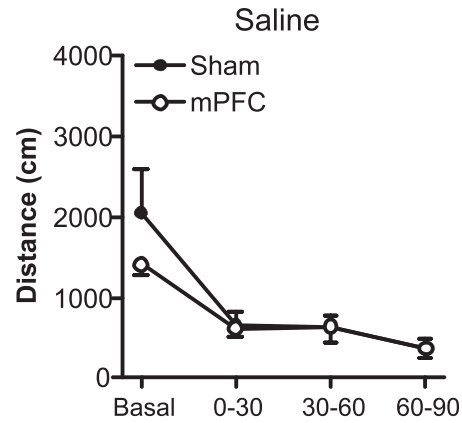
Inspection of cresyl violet-stained brain sections from two animals revealed that the tip of the infusion cannula was situated within the boundaries of the mPFC, as delineated by the Paxinos and Watson rat brain atlas [57].

### HPLC: 5-HT depletion

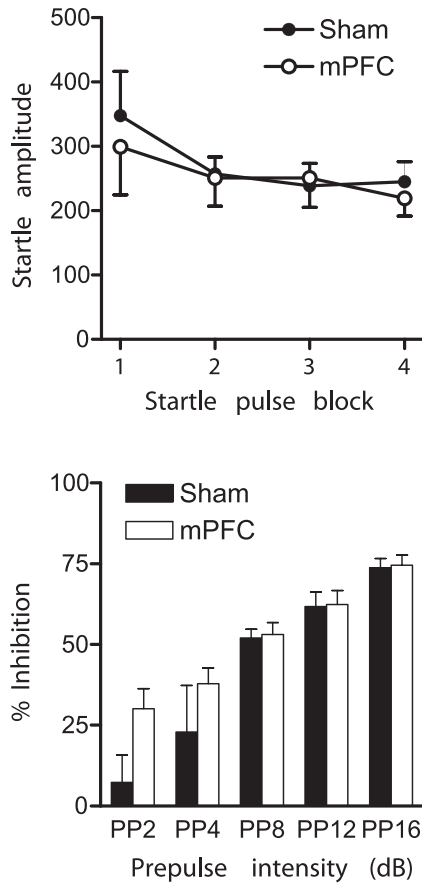
The behavioural data of four mPFC-lesioned rats with partial depletions of 5-HT (<50%), as measured by HPLC, were excluded from the study. The final group size for the analysis was  $n=10$  for the sham-operated group and  $n=9$  for the mPFC-lesioned group. In the mPFC-lesioned rats, the local injection of 5,7 DHT caused a marked reduction of the 5-HT concentration in the mPFC. After the microinjection of 5,7-DHT into the mPFC, the concentration of 5-HT was reduced by 60% (Figure 1). An ANOVA revealed a significant reduction of the 5-HT concentration in the mPFC-lesioned rats compared to the controls ( $F_{1,17}=92.3, p < 0.001$ ).



**Figure 1** 5-HT content in the mPFC after sham surgery or 5,7 DHT microinjection into the mPFC. The data are expressed as the average 5-HT concentration (ng/mg of tissue wet weight)  $\pm$  SEM. \*\*\* $p < 0.001$  for the difference in the 5-HT concentration between the mPFC-lesioned rats and the sham-operated control rats as indicated by ANOVA.



**Figure 2** Time course of the effects of a subcutaneous injection of saline, 0.5 mg/kg of amphetamine or 2.5 mg/kg of phencyclidine on locomotor activity in sham-operated rats and rats with 5,7 DHT-induced mPFC lesions. Locomotor hyperactivity is expressed as the distance moved (cm)  $\pm$  SEM for sham-operated ( $n=10, \bullet$ ) and mPFC-lesioned rats ( $n=9, \circ$ ). There were no significant differences between the groups.



**Figure 3**  
Effect of sham surgery (n=10) or 5,7-DHT lesions of the mPFC (n=9) on startle amplitude, startle habituation and prepulse inhibition. The top panel illustrates the basal startle reactivity and the startle habituation. The data are expressed as mean startle amplitudes  $\pm$  SEM for each of the four blocks of ten 115 dB pulses. The bottom panel illustrates the prepulse inhibition of the acoustic startle response. The prepulse inhibition is expressed as the % inhibition  $\pm$  SEM at different prepulse intensities. There were no significant differences between the groups as indicated by ANOVA.

### The effects of the microinjection of 5,7-DHT on amphetamine- and phencyclidine-induced locomotor hyperactivity

The locomotor hyperactivity caused by treatment with amphetamine or phencyclidine was not significantly different between sham-operated rats and mPFC-lesioned rats (Figure 2). After treatment with either amphetamine or phencyclidine, there was an expected main effect of time ( $F_{2,34}=10.4$ ,  $p<0.001$  and  $F_{2,34}=5.4$ ,  $p=0.01$ , respectively), reflecting the increases in activity caused by these treatments. The lack of a main group effect or a time  $\times$  group interaction suggested that the time course of the effects of either amphetamine or phencyclidine was not altered after mPFC-lesions. Lesioned animals tended to show increased phencyclidine responses (Figure 2, bottom panel); however, this was seen in only two out of nine rats. The analysis of the 5-HT concentrations in these two rats did not reveal any differences in the level of 5-HT depletion in comparison

with the other rats. After a saline injection, the locomotor activity levels were very low, and there was no significant difference between the lesioned group and the sham-operated group (Figure 2).

### The effects of the microinjection of 5,7-DHT on the startle response, habituation and prepulse inhibition

The startle amplitude in the pulse-alone trials and the habituation of the rats were not different between the mPFC-lesioned rats compared to the sham-operated controls (Figure 3). An ANOVA revealed that there was no significant main effect of group or of the habituation  $\times$  group interaction. Figure 4 appeared to suggest that the mPFC-lesioned rats showed an increase in prepulse inhibition at PP2 and PP4 compared to the sham-operated controls; however, an ANOVA indicated that there was no significant main effect of group or of the prepulse  $\times$  group interaction.

## DISCUSSION

The behavioural assessment of rats with serotonergic lesions of the mPFC, using the locomotor hyperactivity and prepulse inhibition paradigms, revealed that there were no differences between the lesioned and control groups. The present findings suggest that normal regulation of locomotor activity and prepulse inhibition is independent of serotonin release from terminals in the mPFC.

### Serotonin in the mPFC and psychotomimetic drug action

As previously reported [41], it is likely that serotonin release from terminals in the dorsal hippocampus has an inhibitory effect on glutamatergic projections to the core of the nucleus accumbens. Disruption of this inhibition may lead to increased glutamatergic transmission in the nucleus accumbens and enhancement of phencyclidine-induced hyperlocomotion. In addition, enhancement of glutamatergic transmission has been shown to have an opposite effect on amphetamine-induced hyperlocomotion [41].

It has been shown that phencyclidine enhances glutamate release in the prefrontal cortex to compensate for the blockade of NMDA receptors, which leads to the overstimulation of postsynaptic non-NMDA glutamate receptors [60, 61]. Additionally, the acute administration of phencyclidine promotes dopamine release in the prefrontal cortex [62]. It has been reported that group II metabotropic glutamate receptor agonists and non-NMDA receptor antagonists diminish phencyclidine-induced hyperlocomotion by preventing the responses of pyramidal cells of the rat mPFC [60, 61]. We suggest that serotonin depletion in the mPFC does not influence phencyclidine-induced dopamine or glutamate release in this brain region. Furthermore, our results suggest that glutamatergic transmission in the nucleus accumbens is not altered after the manipulation of serotonin release in the mPFC. We postulate that the effect of en-





hanced glutamatergic transmission on the amphetamine response depends on the balance of NMDA and non-NMDA receptor activation interacting with dopamine D<sub>1</sub> and D<sub>2</sub> receptors. Both NMDA and non-NMDA receptors in the nucleus accumbens are located at the presynaptic level on glutamatergic and dopaminergic terminals arising from the cortex, the hippocampus and the midbrain, respectively (for a review see [63]). Our results suggest that serotonin depletion in the mPFC does not influence either dopaminergic or glutamatergic transmission within the nucleus accumbens.

### Serotonin in the mPFC and prepulse inhibition

Prepulse inhibition is thought to be mediated by a prefrontocortico-limbic-striato-pallidal circuit in which the mPFC plays an important role [64]. Manipulations that decrease dopamine levels in the mPFC disrupt prepulse inhibition [53, 54], presumably by disinhibition of the descending glutamatergic projections [64]. The mPFC sends direct glutamatergic projections to the nucleus accumbens and the ventral tegmental area (VTA) [65], from which dopaminergic projections ascend to the nucleus accumbens. Stimulation of the mPFC increases dopamine release in the nucleus accumbens, probably via the VTA [66, 67]. It is therefore possible that disinhibition of the mPFC glutamatergic output to the VTA increases dopamine release in the nucleus accumbens and thereby reduces prepulse inhibition [68]. Our results suggest that serotonin depletion in the mPFC does not alter mPFC output neurons to either the VTA or the nucleus accumbens, thus having no effect on sensorimotor gating.

In conclusion, serotonin depletion in the mPFC does not lead to the dysregulation of subcortical dopaminergic activity and does not cause aberrant responses to environmental stimuli. It is therefore clear that the behavioural effects of raphe lesions described in our initial studies [69, 70] are not mediated by serotonin depletion in the mPFC.

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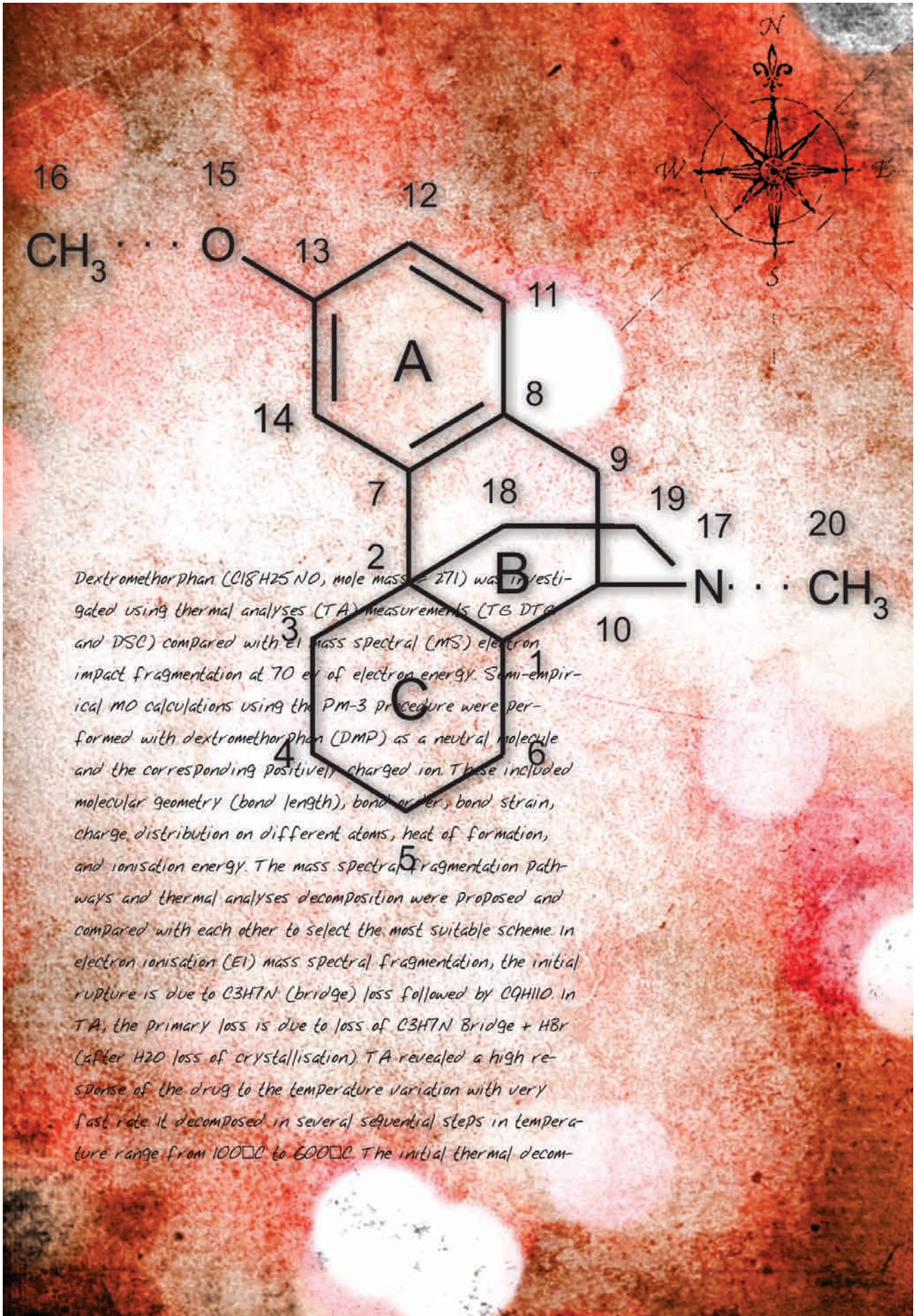


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# STRUCTURAL INVESTIGATION OF DEXTROMETHORPHAN USING MASS SPECTROMETRY AND THERMAL ANALYSES COMBINED WITH MO CALCULATIONS

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## ABSTRACT

( $C_{18}H_{25}NO$ , mole mass = 271) was investigated using thermal analyses (TA) measurements (TG/DTG and DSC) compared with EI mass spectral (MS) electron impact fragmentation at 70 eV of electron energy. Semi-empirical MO calculations using the PM-3 procedure were performed with dextromethorphan (DMP) as a neutral molecule and the corresponding positively charged ion. These included molecular geometry (bond length), bond order, bond strain, charge distribution on different atoms, heat of formation, and ionisation energy. The mass spectral fragmentation pathways and thermal analyses decomposition were proposed and compared with each other to select the most suitable scheme. In electron ionisation (EI) mass spectral fragmentation, the initial rup-

ture is due to  $C_3H_7N$  (bridge) loss followed by  $C_9H_{11}O$ . In TA, the primary loss is due to loss of  $C_3H_7N$  Bridge + HBr (after  $H_2O$  loss of crystallisation). TA revealed a high response of the drug to the temperature variation with very fast rate. It decomposed in several sequential steps in temperature range from 100°C to 600°C. The initial thermal decomposition was similar to that obtained using MS. So, it is possible to select the proper pathway using both techniques. This comparison was successfully confirmed by MO calculation. The structural-reactivity effect on metabolism and the biological effect compared with codeine are discussed.

**Keywords:** Dextromethorphan structure, Mass Spectrometry, Thermal analyses, MO calculation

## INTRODUCTION

Dextromethorphan (DMP, methoxy-methyl morphinan) is an antitussive agent used in many nonprescription cough and cold medications [1]. DM has an IUPAC name (+)-3-methoxy-17-methyl-(9 $\alpha$ , 13 $\alpha$ , 14 $\alpha$ )- morphinan ( $C_{18}H_{25}NO$ , m/z=271) and is a synthetic product that is chemically related to codeine [2]. The drug is a safe, oral antitussive that is widely available [3 – 6]. The structural formula and numbering system of the drug is given in Fig. 1.

A number of methods have been reported to measure DM in biological fluids, including high-performance liquid chromatography (HPLC) [7, 8], gas chromatography (GC) [9, 10] and gas chromatography–mass spectrometry (GC–MS) [11–14, 1]. The latter technique is highly sensitive [5].

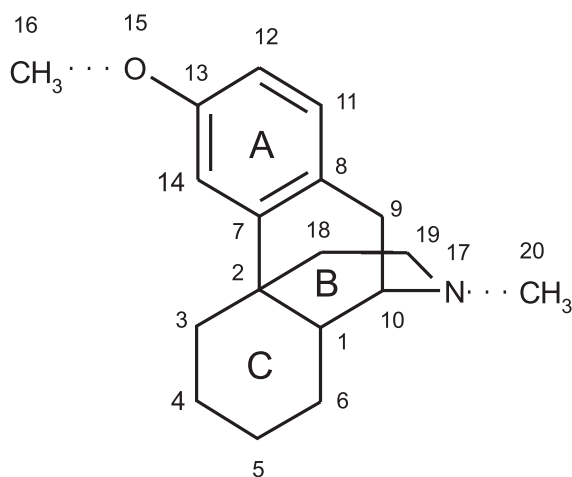
Rapid advances in biological sciences have led to an increased demand for chemical and structural information from biological systems. Mass spectrometry plays a pivotal role in the structural characterisation of biological molecules [15]. The technique is important because it provides a substantial amount of structural information using

a small amount of sample. Moreover, the techniques offer comparative advantages in speed and productivity for pharmaceutical analysis [16]. In contrast, thermal analysis delivers extremely sensitive measurements of heat change, which can be applied on a broad scale with pharmaceutical development. These methods provide unique information relating to thermodynamic data of the system studied [17]. The increasing use of the combined techniques in TA could provide more specific information and thus facilities a more rapid interpretation of the curves obtained [17].

In the electron ionisation (EI) mass spectra, the fragmentation consists of competitive and consecutive unimolecular fragmentation [18]. The fragmentation of the ionised molecule depends mainly on their internal energy [19]. The thermo-gravimetric TG/DTG analysis provides quantitative information on weight losses due to decomposition and/or evaporation of low molecular weight fragments as a function of time and temperature. In conjunction with mass spectrometric analysis [20–22], the nature of the released fragments

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**Dextromethorphan drug (DMP)**

Figure 1. Structure and numbering system of DMP for, C, N and O skelton.

may be deduced, thus greatly facilitating the interpretation of thermal degradation processes. In contrast, the computational quantum chemistry could provide additional information about the atoms and bonds, which could be used successfully in an interpretation of experimental results [23]. The application of the computational quantum chemistry in addition to experimental results (MS and TA) gives valuable information about the atoms and bonds, which helps in the description and prediction of the primary fragmentation site of cleavage and subsequent ones [24–28].

Although the literature is wealthy in information related to the biological activities of DMP and its metabolites *in vivo* and *in vitro* [3–6], there seems to be a lack of any correlation between chemical behaviour and its electronic structure.

The main aim of this work was to perform an experimental and theoretical investigation of DM using thermal analyses (TA) and EI mass spectral (MS) fragmentation at 70 eV. In addition, MO calculations were performed using the PM3 procedure on the neutral molecule and charged molecule to investigate geometrical parameters (bond length, bond order, bond strain, heats of formation ionisation energy) and charge distribution.

The calculations correlated the experimental results [TA and MS] to obtained information regarding the stability of the drug and the prediction of the site of primary fragmentation and subsequent ones. This study could be helpful in establishing a quantitative and qualitative structure–activity relationship for the drug, which is the main objective of this work. It is worth mentioning that there is scant literature concerning such a comparative study.

Moreover, the authors tried to correlate and discuss the structure reactivity and relationship between DM and Codeine [26], which are chemically related to each other [2] and which affect the O and OH on MS and TA and MO parameters. In addition, they attempted to find the metabolite structures in the mass spectrum and correlated the biological behaviour of the two drugs.

## EXPERIMENTAL

### Mass spectrometry (MS)

The electron ionisation (EI) mass spectrum of DMP was obtained using a Shimadzu GC-MS-Qp 1000 PX quadruple mass spectrometer with an electron multiplier detector equipped with the GC-MS data system. The direct probe for solid material was used in this study. The sample was put into a glass sample micro vial by a needle ( $\approx 1 \mu\text{g}$  max). The vial was installed on the tip of the DP containing the heating cable and was inserted into the evacuated ion source. The sample was ionised by an electron beam emitted from the filament with the generated ions being effectively introduced into the analyser by the focusing and extractor lens systems. The MS was continuously scanned, and the spectra obtained were stored. The electron ionisation mass spectra were obtained at an ionising energy value of 70 eV, ionisation current of 60  $\mu\text{A}$  and a vacuum greater than  $10^{-6}$  torr.

### Thermal analysis (TA)

The thermal analyses of DMP were made using a conventional thermal analyser (Shimadzu system of DTA-50 and 30 series TG-50). The mass loss for 5 mg of sample and the heat response to changes in the sample were measured from room temperature up to 600°C. The heating rate in an inert argon atmosphere was 10°C min<sup>-1</sup>. These instruments were calibrated using indium metal as a thermal stable material. The reproducibility of the instrument reading was determined by repeating each experiment more than twice.

### Quantum chemical calculations

The MO calculations were performed using a semi-empirical molecular orbital calculation. The method used in these computations was the parametric PM-3 method described by Stewart [29]. The geometry of all stable species studied was completely optimised with respect to all geometrical variables using the Eigen vector following (EF) routine [30]. The program was run under the molecular orbital calculation package MOPAC2000 by Stewart [31] for microcomputers.

## RESULTS AND DISCUSSION

The chemistry and reactivity of pharmaceutical drugs are of great interest because of their importance in treating various diseases. Knowledge of the thermal decomposition mechanism of the drug is very important to understand the chemical processes in biological systems. It is difficult to establish the exact major fragmentation pathway in EI using conventional MS. However, the combination of experimental techniques (TA and MS) and MO calculation is very important to understand the following topics:





- 1 -The primary site fragmentation process and its major fragmentation pathways in both techniques.
- 2 -The stability of the drug as a neutral molecule in the solid-state phase and molecular ion in the gas phase.
- 3 -Selection of the most probable decomposition pathway in both TA and MS.

### Thermal analysis (TA)

The TG/DTA curves of DMP (Fig. 2) were displayed between 25°C and 600°C. The drug has the general formula  $C_{18}H_{25}NO.HBr.H_2O$  in solid-state phase with a molecular weight (MW) of 370.3 and melting point (m.p.)  $\cong 125^\circ\text{C}$ .

It is clear from the thermal survey of the drug that there are four mass losses in the TGA (Table 1). The first one appeared at 100-150°C with a mass loss % of 4.1, which may be attributed to the loss of water present in the drug (cal. mass loss % = 4.8). The second mass loss was approximately 32.5% and occurred at 150-350°C. The third one was 54.57% and occurred at 350-450°C. The final one was 5.68% and occurred at 450-600°C, which may be attributed to the loss of the remainder. These mass losses were observed at the three main peaks at values: 120.8°C, 287.2°C and 378.6°C in DTG, respectively. These mass losses also appeared as two endothermic peaks in the DTA. The first one appeared at 110.77°C, requiring an energy  $E_1 = +242.6$  J/g, and the second one appeared at 362°C, requiring an energy  $E_2 = +356.75$  J/g. The fragmentation of this compound and its chemical changes occurring after this thermal degradation appeared at 555.56°C as a very strong exothermic peak requiring an energy  $E_3 = 513.5$  J/g.

### Mass spectral behaviour of dextromethorphan (MS)

The electron impact (EI) mass spectrum for DMP drug at 70 eV was recorded and investigated. A typical mass spectrum (barograph) of the drug is shown in Fig. 3.

Scheme 1 shows the proposed principal fragmentation pathways of DMP following electron impact at 70 eV. In addition, prominent ions and their relative intensities with some common ions formed from codeine [26] are listed

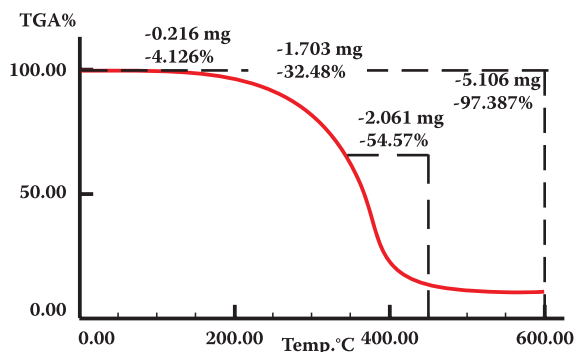


Fig.2a : TGA curve of Dextromethorphan Hydrobromide

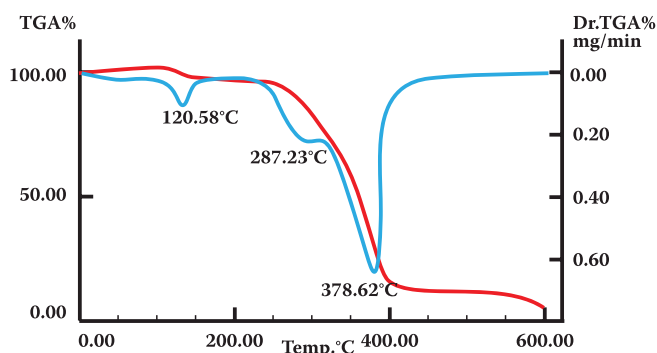


Fig.2b : Dr.TGA of Dextromethorphan Hydrobromide

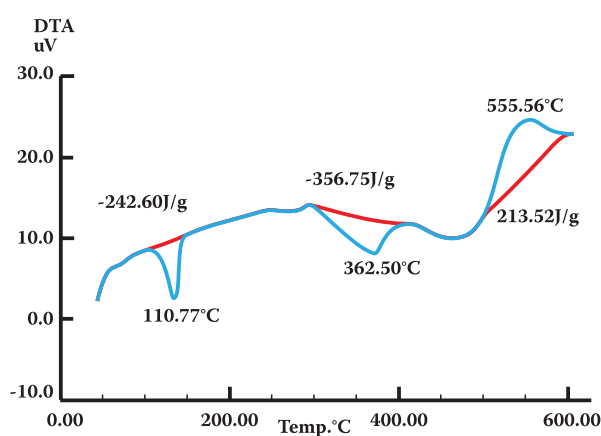
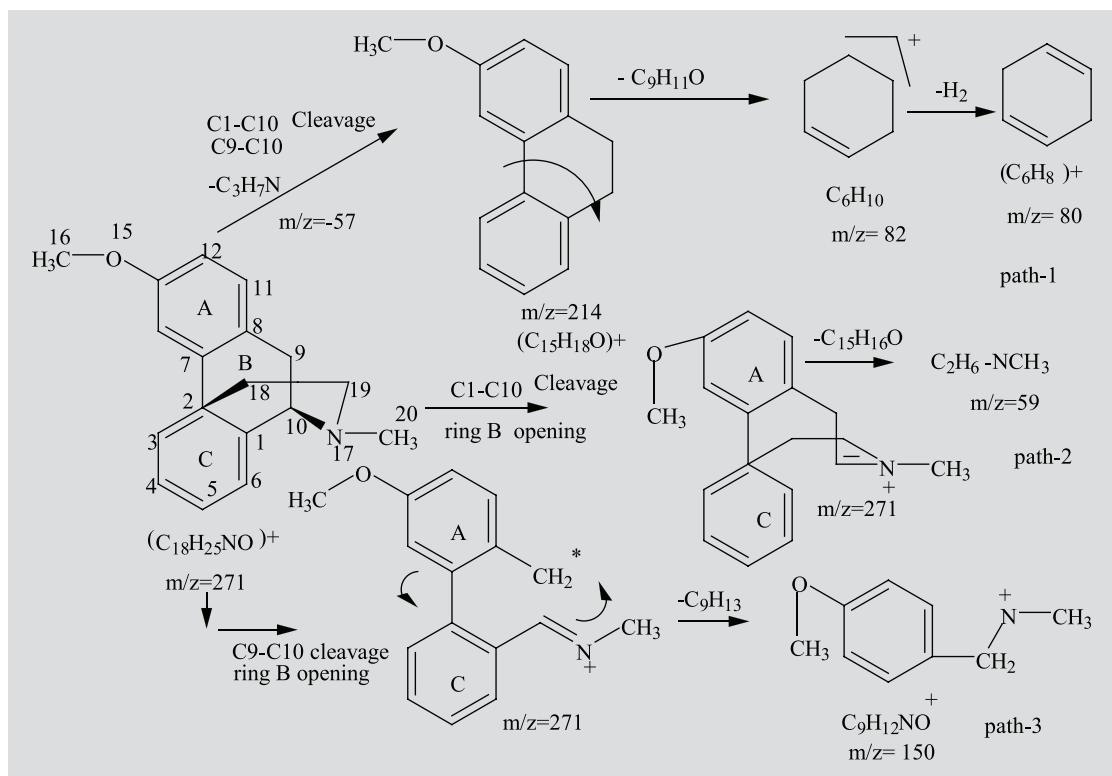


Fig 2 C: DTA of Dextromethorphan Hydrobromide

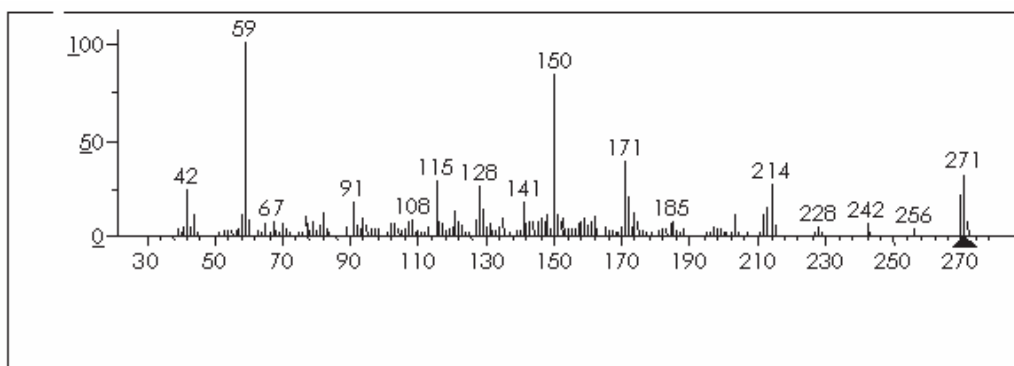
Table 1: Structure identification of Dextromethorphan Hydrobromide

TGA		DTA		
Wt. loss%	Temp.range °C	Peak temp. °C	Description	E J/g
4.10	100-150	118.77	Endothermic	+242.6
32.50	150-350	367.6	endothermic	+365.75
54.57	350-450	555.56	St.exothermic peak	-613.5
5.68	450-600			
$\Sigma$ Wt.loss %=96.85				

\*Dr. TGA gives three main peaks temperatures at 120.58, 287.23 and 378.62 oC.



**Scheme (1):** Mass spectral fragmentation pathways of Dextromethorphan



**Fig. (3):** Mass spectrum of DMP at 70 eV.

Dextromethorphan				Codeine		
m/z	R.I.	Structure	Process	m/z	Process	R.I.
271	31.6	$C_{18}H_{25}NO^+$	$M^+$	299	$M^+$	100%
270	20.3	$C_{18}H_{24}NO^+$	$[M - 1]^+$	298	$[M - 1]^+$	14.3
214	26.6	$C_{15}H_{18}O^+$	$[M - C_3H_7N]^+$	242	$[M - C_3H_7N]^+$	9.3
150	83.0	$C_9H_{12}NO^+$	$[M - C_9H_{13}]^+$			
82	12.5	$C_6H_{10}^+$	$[M - C_3H_7N - C_9H_8O]^+$			
59	100%	$C_3H_9N^+$	$[M - C_{15}H_{16}O]^+$			

**Table 2:** The prominent ions in the mass spectrum of DMP drug in comparison with codeine drug [26].



in Table 2. The mass spectrum of DMP showed that the molecular ion ( $C_{18}H_{25}NO^+$ ,  $m/z = 271$ ) signal represented approximately 30% of the base peak (= 100%) at  $m/z = 59$  [ $C_3H_9N^+$ ], indicating a moderate stability of the molecular ion in gas phase compared with codeine [26].

The main fragmentation pathway for DMP after ionisation at 8.83 eV consists of three principal pathways (paths 1-3) as rationalised in scheme 1. The formation of the fragment ion at signal  $m/z = 214$  (path 1, scheme 1) with R.I. = 26.6% is mainly due to the formation of  $[M-C_3H_7N]^+$  as a loss of the  $C_3H_7N$  bridge by disruption of the  $C_2-C_{18}$  and  $C_{10}-C_{17}$  bonds. It is worth mentioning that this fragment ion,  $C_{15}H_{18}O^+$  ( $M-C_3H_7$ ), was formed directly from the molecular ion. This formation was confirmed by the full scan positive ion mass spectrum using tandem mass spectrometry (MS/MS) [6] as a prominent peak. Subsequent cleavage of the  $C_7-C_2$  and  $C_1-C_{10}$  bonds (ring B) occurred forming the  $C_6H_{10}^+$  fragment ion ( $m/z = 82$ , R.I. = 12.5%) (path 1 scheme 1).

Two important possible modes of fragmentation (paths 2 and 3 scheme 1) are due to the cleavage of the two bonds  $\beta$  to the nitrogen atom (i.e., cleavage of  $C_1-C_{10}$  and  $C_9-C_{10}$  competitively forming two important fragment ions in the mass spectrum.) First, cleavage of the  $C_1-C_{10}$  bond and formation of the most prominent fragment (the base peak) at  $m/z = 59$  (path 2) of formula  $C_3H_9N$ . Also, cleavage of the  $C_9-C_{10}$  bond formed the second prominent peak at  $m/z = 150$  (R.I. = 83.0%) with the structure  $C_9H_{12}NO$  (path 3).

Structural determinations for the compounds of unknown structure are generally based upon information obtained through the use of a variety of modern instruments. The instrumental methods usually include mass spectrometric studies, and the techniques are well known to organic chemists and scientists [32].

It is known that DMP is chemically related to codeine [2] but that it differs in biological effect. DMP is an antitussive agent used in many nonprescription cough and cold medications [1]. Codeine has an analgesic effect [26]. Biotransformation (metabolites) *in vivo* and *in vitro* of the two drugs has been investigated by many investigators [3-6]. Metabolism studies often lead to compounds with unknown structures that are related to the material under study. The chemical structure and metabolic pathway of DMP were rationalised [5] in Fig. 3.

The metabolism of DMP is primarily by *o*-demethylation to dextrophan. DMP is also metabolised to 3-methoxymorphinan and 3-hydroxymorphinan.

It was noted that from mass spectrum of DMP, the signal due to  $C_2H_4$  loss was observed with minor abundance  $m/z = 243$ , R.I. = 1.21, which may be due to the formation of hydroxymorphinan.

Comparing the fragmentation pathway of DMP to codeine [26], one can notice that the two drugs are similar in the initial processes (i.e., loss of the bridge  $C_3H_7N$  and the ring opening of the two bonds  $\beta$  to the nitrogen atom). The subsequent fragmentation differed because of the presence of different substituent groups (O and OH).

## Computation

Molecular orbital calculations gave valuable information regarding the structure and reactivity of the molecules and molecular ions. The computational data supports the experimental data. The parameters calculated using the MO calculation include geometry, bond order, bond strain, charge distribution, heat of formation and ionisation energy.

The investigation of the molecular structure of DMP with the common formula  $C_{18}H_{25}NO$  was of interest in this work, and this investigation aimed to predict the weakest bond cleavage and the stability of the neutral molecule by TA and the charged molecular by MS.

Table 3 shows the comparison of computed bond length, bond order and bond strain using the PM3 method for neutral and molecular cationic forms. Table 4 shows a comparison of computed partial charges of neutral and charged species. The computations reveal some important results.

- 1 The charge density localised on the nitrogen atom increased from -0.076 to 0.489 from a neutral to charged atom, which reveals that the electron disruption upon ionisation at 8.83 eV occurs in the nitrogen atom. No appreciable change in the charge on the other atom was due to ionisation of the molecule.
- 2  $C_2-C_{18}$  is the lowest bond order for neutral molecule (0.964) and charged cationic form (0.967).

## Correlation of TA, MS and MO-calculations.

DMP is an essential cough and cold medication [1]. The drug is chemically related to codeine [2]. As indicated previously [32], a determination of initial bond cleavage is an important first step.

The scope of this investigation was restricted to a search or prediction and discerned the features of initial bond disruption during the course of fragmentation of DMP. Empirical observations indicate that the course of subsequent fragmentation is determined to a large extent by the initial bond disruption of the molecular ion in MS [34]. It is quite acceptable to say that the computational quantum chemistry can provide additional data, which can be used successfully to interpret both TA and MS experimental results. These theoretical data are particularly valuable for mass spectral scientists. These scientists study gas-phase species, which can be handled much more easily by quantum chemistry [33].

The mass spectrum of DMP in gas-phase ion revealed three competitive processes (1 - 3, scheme 1).

PM3 calculation (Table 3) revealed that  $C_{10}-N_{17}$  was the weakest bond in the charged system (weakest bond order at 0.961, large bond length = 1.484 Å and bond strain). Moreover, the weakness of this bond ( $C_{10}-N_{17}$ ) was due to the presence of a lone pair on the nitrogen. Table 3 shows that the electron disruption upon ionisation occurred from the nitrogen, which increased the weak-





Bond	Bond length (Å)		Bond order		Bond strain (k Cal/mol)	
	Neutral	Ionic	Neutral	Ionic	Neutral	Ionic
C1-C2	1.541	1.542	0.965	0.970	0.206	0.189
C1-C6	1.525	1.531	0.984	0.983	0.049	0.208
C1-C10	1.533	1.542	0.973	0.962	0.049	0.101
C2-C3	1.548	1.547	0.969	0.971	0.457	0.639
C2-C7	1.518	1.520	0.969	0.974	0.252	0.290
C3-C4	1.519	1.519	0.993	0.996	0.034	0.071
C4-C5	1.518	1.515	0.992	0.997	0.107	0.004
C5-C6	1.526	1.522	0.987	0.992	0.174	0.095
C7-C8	1.403	1.405	1.376	1.365	0.082	0.140
C7-C14	1.395	1.393	1.424	1.442	0.084	0.097
C8-C9	1.493	1.491	0.990	0.997	0.035	0.058
C8-C11	1.396	1.397	1.409	1.401	0.044	0.047
C9-C10	1.533	1.535	0.971	0.959	0.073	0.115
C11-C12	1.388	1.387	1.429	1.444	0.007	0.003
C12-C13	1.389	1.401	1.373	1.360	0.015	0.010
C13-C14	1.400	1.405	1.371	1.351	0.012	0.015
C13-C15	1.382	1.372	1.029	1.067	0.038	0.038
C15-C16	1.406	1.409	0.987	0.980	0.008	0.009
C10-N17	1.503	1.484	0.972	0.961	0.226	0.192
C2-C18	1.545	1.547	0.964	0.967	0.118	0.178
C18-C19	1.524	1.525	0.990	0.980	0.075	0.031
N17-C20	1.480	1.450	0.997	1.013	0.057	0.063
C19-N17	1.492	1.460	0.980	0.994	0.092	0.059

**Table 3:** Comparison of computed bond length, bond order and bond strain (k Cal mol<sup>-1</sup>) using PM3 method for neutral and molecular cation.

ness of the bond. After an easy disruption of this bond, the subsequent bond is C<sub>2</sub>-C<sub>18</sub> (bond order = 0.967, bond length = 1.547 Å and bond strain = 0.178 K cal mol<sup>-1</sup>). Disruption of these two bonds (C10-N<sub>17</sub> and C<sub>2</sub>-C<sub>18</sub>) formed a fragment ion at m/z = 214 by loss C<sub>3</sub>H<sub>7</sub>N (bridge) from the molecular ion. The subsequent loss of C<sub>9</sub>H<sub>11</sub>O (rupture of the B ring) formed the fragment ion C<sub>6</sub>H<sub>10</sub> at m/z = 82, which can lead to a loss of H<sub>2</sub> to form the fragment ion C<sub>6</sub>H<sub>8</sub> at m/z = 80 (process 1, scheme 1). One can conclude that process 1 is the principal one for DMP, which can help to interpret TA decomposition.

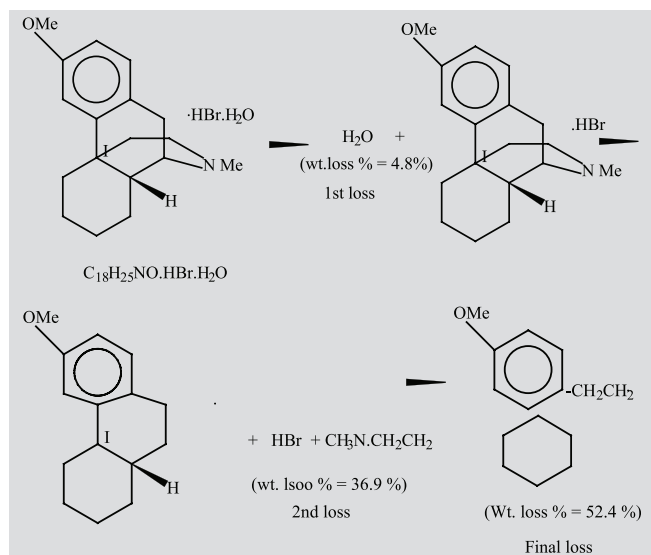
The primary TA decomposition of DMP (wt loss = 4.1%) between 100 °C and 150°C was mainly due to loss of water. It was followed by a mass loss of 32.5% between 150 °C and 350°C. On the basis of the MO calculation (Table 2), C<sub>2</sub>-C<sub>18</sub> was the first bond cleavage of neutral system having the lowest bond order at 0.964 with a long bond length (1.545 Å) and large bond strain (0.118 k cal mol<sup>-1</sup>) followed by C<sub>10</sub>-N<sub>17</sub> (bond order = 0.972, bond length = 1.503 Å and bond strain = 0.226 k cal mol<sup>-1</sup>). As a result of the disruption of these two bonds, C<sub>2</sub>-C18 and C<sub>10</sub>-N<sub>17</sub> lost the bridge C<sub>3</sub>H<sub>7</sub>N. The wt loss of 32.5%

resulted from the bridge and HBr molecule (C<sub>3</sub>H<sub>7</sub>N + HBr). The subsequent third mass loss of 54.57% occurred at 350-450°C and may be attributed to the loss of OCH<sub>3</sub>C<sub>6</sub>H<sub>4</sub>CH<sub>2</sub>CH<sub>2</sub> (i.e., C<sub>9</sub>H<sub>11</sub>O) via the cleavage of the B-ring (C<sub>2</sub>-C<sub>7</sub> and C<sub>1</sub>-C<sub>10</sub>).

The final mass loss of 5.68%, which occurred at temperature range 450-600°C, may be attributed to the loss of the remainder of the drug molecule. It is possible to rationalise the thermal degradation scheme 2 as follows.

The dextromethorphan (DMP-HBr) neutral compound is stable up to 150°C before being fragmented by the loss of the bridge C<sub>3</sub>H<sub>7</sub>N + HBr, whereas codeine fragments at 160°C [26]. In contrast, the molecular ion of DMP is RI = 30%, while the molecular ion of codeine is considered the base peak (RI = 100%). This high stability of codeine may be due to the presence of OH and O atoms in its skeleton.

This work provides further insights into utility of experimental TA and MS techniques and a theoretical investigation with MO calculation using the semi-empirical PM3 procedure on DMP. From the practical and theoretical techniques, we concluded that the primary fragmentation of DMP is initiated by the loss of C<sub>3</sub>H<sub>7</sub>N + HBr in



**Scheme (2) :** Thermal Decomposition of Dextromethorphan Hydrobromide Drug

Atom	Partial Charge	
	Neutral	cation
C1	- 0.073	- 0.072
C2	0.013	0.014
C3	- 0.107	- 0.106
C4	- 0.093	- 0.097
C5	- 0.091	- 0.096
C6	- 0.098	- 0.094
C7	- 0.019	- 0.047
C8	- 0.102	- 0.174
C9	- 0.080	- 0.050
C10	- 0.048	- 0.144
C11	- 0.073	- 0.042
C12	- 0.181	- 0.164
C13	0.087	0.124
C14	- 0.150	- 0.142
O15	- 0.189	- 0.177
C16	0.051	0.047
N17	- 0.076	0.489
C18	- 0.095	- 0.116
C19	- 0.077	- 0.169
C20	- 0.094	- 0.198

**Table 4:** Comparison of computed partial charge using PM3 procedure for neutral and molecular ion for DMP drug.

TA and  $C_3H_7N$  in MS. MO-calculation confirmed these processes. Subsequent cleavage in both techniques is due to the cleavage of the B ring and loss of the neutral fragment  $C_9H_{11}O$ . DMP can be completely dissociated between 150 and 600°C in TA, while in MS, the molecule

can be fragmented after a few electron volts above the ionisation energy of 8.83 eV. Theoretical calculation can help to select the most probable fragmentation pathway in both techniques.

Signal at  $m/z = 243$  and R.I. = 1.2% may have the structure  $C_{16}H_{21}NO$  (hydroxymorphinan structure), which is the metabolite of DMP by the loss of  $CH_2 + CH_2$ .

Compared with codeine, we noticed the following occurrences due to the presence of O and OH atoms.

- 1 -DMP is less stable than codeine in TA (decomposes  $\cong 10^\circ C$  before codeine).
- 2 -The stability of the molecular ion of DMP in gas phase (MS) is about 30% compared with codeine (100%).
- 3 -The biological behaviour of DMP is significantly different from codeine due to the presence of two excess O-atoms in the skeleton of the latter, which may help its transport to the brain and its differential effect.

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## EXPERIMENTAL MODELS OF DIABETES MELLITUS

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## EKSPERIMENTALNI MODELI DIJABETES MELITUSA

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### APSTRAKT

Early studies on pancreatectomised dogs confirmed the central role of the pancreas in the homeostasis of glycemia and resulted in the discovery of insulin. Today, hundreds of different animal models are used in experimental studies of diabetes. The aim of this review is to present experimental models of type 1 and type 2 diabetes. In preparing for this review, we searched the electronic databases **Medline, Highwire, and Hinari**.

The majority of the experiments are conducted on rodent models (mice and rats). Selective inbreeding resulted in the development of numerous models with pathogenic characteristics and the manifestation of type 1 and 2 diabetes and the related phenotypes of obesity and insulin resistance. In addition to analyzing the pathogenic mechanisms of the disease and its complications, these models are used to evaluate new treatment solutions as well as the transplantation of beta cells and disease prevention. New animal models have been created using techniques based in molecular biology and genetic engineering: transgenic, knockout and tissue-specific knockout models. These are very powerful methods, which may lead to exciting results in the future.

**Key words:** animal models, experimental diabetes, mouse, rat

### SAŽETAK

Prve studije rađene na psima kojima je prethodno uklonjen pankreas potvrdile su centralnu ulogu pankreasa u homeostazi glikemije i doprineli otkriću insulina. Danas, veliki broj različitih eksperimentalnih modela se koristi u istraživanjima u oblasti dijabetesa. Cilj ovog revijskog rada je da predstavi neke od ovih modela, koristeći kao literaturu originalne i revijske radove elektronskih baza **Medline, Highwire i Hinari**.

Većina laboratorija koristi selektivnim inbridingom dobijene miševe i pacove, kao i transgene i nokaut životinje dobijene genetskim inženjeringom, koje zbog svojih karakteristika (npr. gojaznost, rezistencija na insulin) predstavljaju idealne eksperimentalne modele za dijabetes tip 1 ili 2. Ovi eksperimentalni modeli koriste se kako za ispitivanje patogeneze dijabetesa tako i za ispitivanja novih terapijskih sredstava u lečenju šećerne bolesti.

**Ključne reči:** animalni modeli, eksperimentalni dijabetes, miš, pacov



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## INTRODUCTION

Since 1880, when von Mering and Minkowski removed the pancreas from a dog displaying symptoms of diabetes mellitus (1), many experiments have been conducted on mice, rats, rabbits, and dogs, which were all used as experimental models for this disease.

The rodents (mice and rats) are most frequently used in laboratory research of type 1 and type 2 diabetes because they are biologically and genetically similar to people, their reproductive potential is high, their life span is short, and they are cheap and easy to handle (1). The main drawback of these models is that sometimes they are unable to successfully simulate pathological processes occurring in humans; in those situations, the use of cats, dogs, pigs or primates is justifiable.

Models of experimental diabetes are:

1. induced (experimental)
2. spontaneous (genetically induced) or
3. transgenic and knock-out models (resulting from genetic engineering).

Hyperglycemia is induced by:

1. surgical method (partial or total pancreatectomy)
2. non-surgical method (using a diabetogenic toxin in experimental animals) (Table 1) (2).

## MATERIALS AND METHODS

We searched the electronic databases *Medline*, *Highwire*, and *Hinari*. Selected papers, both original and review, have been summarized and cited with regard to their scientific relevance for this research field.

### EXPERIMENTAL MODELS OF TYPE 1 DIABETES MELLITUS

#### A) Induced models

##### Multiple low-dose streptozotocin-induced diabetes

Streptozotocin (STZ) is a derivative of nitrosourea that has been isolated from *Streptomyces achromogenes*. It is a strong alkylating agent that affects glucose trans-

port (the role of glucokinase) and induces numerous lesions in the beta-cell DNA chain (3-5). It can selectively bind to glucose-transport protein 2 (GLUT2), which is predominantly expressed on the membranes of pancreatic beta-cells (3).

Toxicity of the streptozotocin on beta-cells is complex and includes genetic and non-genetic mechanisms (6). Nitric oxide (NO) and other free radicals influenced by streptozotocin result in spontaneous lesions of the DNA chain (7). Damaged DNA chains activate poly-(ADP-ribose) polymerase (PARP) as a reparatory mechanism, which in case of over-activation may have proapoptotic effects (8). PARP knockout mice were shown to be resistant for streptozotocin-induced diabetes (8).

In addition, the toxicity of streptozotocin is associated with a drastic decrease of insulin reserves within beta cells, resulting from the fast catabolism of nicotinamide-adenine-dinucleotide+ (NAD+), a substrate for PARP activation (9). Decrease of NAD+ and ATP+ provokes significant depletion of insulin secretion as well as other basic processes inside the cell. While excessive activation of PARP represents a proapoptotic stimulus, the reduction of necessary cell energy results in cellular necrotic death.

A single high dose of streptozotocin provokes diabetes in rodents, possibly as a result of the direct toxic effect. Multiple low-dose streptozotocin-induced diabetes (when streptozotocin is administered in lower doses (40 mg/kg) during five consecutive days) is an experimental model of type 1 diabetes and immune-mediated insulinitis and is widely used for studying the immunological background of this disease (10).

#### Partial pancreatectomy

The partially pancreatectomised rat was one of the first experimental models of diabetes (11). The surgical removal of 90% of the pancreatic tissue resulted in a slow development of hyperglycemia over the course of 12 weeks to 20 months (12). The period of post-surgical euglycemia is followed by the development of chronic hyperglycemia as a consequence of a further failure of the remaining beta cells.

The first stage of disease refers to glucose intolerance followed by a compensatory hyper-function of beta-cells, the exhaustion of the remaining beta cells and their further breakdown 14 weeks after surgery (12). Altered beta cell gene expression is associated with a decreased production of islet amylin polypeptide (IAPP), transcription factors, ionic canals and pumps, including GLUT2, glucokinase, mitochondrial glycerophosphate dehydrogenase (mGPD) and pyruvate carboxylase (13). A reduction of insulin secretion follows even in conditions of high concentrations of glucose and even after application of insulin secretagogues (14).

Histopathological findings demonstrate hypertrophy and fibrosis of pancreatic islets related to the level of hyperglycemia (13). Insulinitis, demonstrated as the infiltration of inflammatory cells, may be depressed, thus slowing down the process of cell degeneration (15).

Toxin	Diabetogenic effect
Streptozotocin	Hyperglycemia, glucose intolerance
Alloxan	Hyperglycemia, glucose intolerance
Chlorozotocin	Hyperglycemia, glucose intolerance
Dithizone	Hyperglycemia, glucose intolerance
Vacor	Hyperglycemia
Oxine	Hyperglycemia
Methylnitrosourea	Glucose intolerance
Cyproheptadine	Glucose intolerance

Table 1. Toxins with diabetogenic effect



## B) Spontaneous models

The most frequently used spontaneous models of type 1 diabetes are the non-obese diabetic (NOD) mouse and the bio-breeding (BB) rat. These two strains are the result of laboratory inbreeding (keene cross-breeding) of a large number of generations; they develop the disease spontaneously, similar to type 1 human diabetes.

### NOD mouse

The NOD mouse is an inbred, homozygous strain discovered in 1974 in Shionogi Research Laboratories in Osaka, Japan. It was a result of keene cross-breeding from normoglycemic strains used for cataract research (Jcl-ICR mouse) and bred in sterile conditions, free from species-specific pathogens (16). The NOD mouse developed insulinitis detectable by electronic microscopy after 2 weeks, and within 4-5 weeks it becomes visible under standard microscopy (17). The infiltration is mainly characterized by the presence of a large number of CD 4+ T lymphocytes, while CD 8+ T and B lymphocytes, as well as NK cells are not so numerous (18). Insulinitis is progressive and accompanied by the destruction of beta cells and a decrease of insulin levels in serum. In contrast to the disease in the human population, these strains have mild ketoacidosis and can survive for weeks without insulin substitution (1). Incidence of this disease differs between genders (90% among females and 60% among males) (1).

Similar to the human population, the main autoantigens in pathogenesis of this disease, recognized by secreted antibodies, are insulin, glutamic acid decarboxylase (GAD), and tyrosine phosphatase (ICA512, known as IA-2) (19).

The genetic background of diabetes in this model highlights the role of a major histocompatible complex (MHC) class II gene that codes for molecule I-A<sup>g7</sup> (20-21). The expression of MHC molecule class I refers to proteins K<sup>d</sup> and D<sup>b</sup> (22). An alteration of INS gene expression, controlling the expression of insulin in the thymus and the deletion of T reactive lymphocytes in the case of central tolerance, is related to an initiation of the autoimmune process (22). A similar situation is observed with the cytotoxic T lymphocyte antigen 4 (CTLA-4) gene, which codes for a negative signal molecule for CD8+ T cells and controls the activation and expansion of T lymphocytes (22). The protein tyrosine phosphatase non-receptor type 22 (PTPN22) gene, associated with autoimmune diabetes, codes for tyrosine phosphatase (LYP or PEP), a negative regulator of T cells (22). Variations of genes coding for IL2 and CD25 FoxP3<sup>+</sup> that a key role in maintaining immune homeostasis are common for both human and NOD mice (22).

### BB rat

The BB rat was discovered in 1974 within the commercial campaign of the Company Bio Breeding Laboratories, Ottawa. It develops diabetes spontaneously (decrease of body weight, polyuria, polydipsia, hyperglycemia and insulopenia) during puberty—approximately the 12<sup>th</sup> week of life (1). Similar to the human population, the BB rat develops serious/fatal ketoacidosis unless insulin substitution is administered. (1)

The disorder is autoimmune, and insulinitis is characterized by the infiltration of T lymphocytes, B lymphocytes, macrophages and NK cells (23). Typical for this species is extreme lymphopenia associated with a decreased number of T lymphocytes with an elevated expression of ADP-ribosyltransferase 2 (ART2); this has not been registered in the human population (24). A transfusion of histocompatible T ART2+ lymphocytes prevents spontaneous hyperglycemia in this model (24).

Recent studies show that BB rat diabetic syndrome is a complex, polygenic disease that may share additional susceptibility genes aside from MHC class II with human type 1 diabetes (25). The expression of two main susceptibility genes is responsible for the development of diabetes in the BB rat: MHC (RT1) class II u haplotype (Iddm1) and the Gimap5 (GTPase immunity associated protein family member 5) gene, a key genetic factor for lymphopenia in spontaneous BB rat diabetes (Iddm2) (25).

## EXPERIMENTAL MODELS OF TYPE 2 DIABETES

### A) Induced models

#### C57BL/6J mouse fed a high-fat diet

This model was originally introduced by Surwit et al. in 1988 as a robust model of impaired glucose tolerance (IGT) and early type 2 diabetes (26). C57BL/6J mice fed a high-fat diet (58% energy by fat, a “Western diet”) developed hyperglycemia, hyperinsulinemia, hyperlipidemia and increased adiposity (27). High-fat diet results in insulin resistance with compensatory hyperinsulinemia. After one week on the diet, baseline plasma glucose and insulin are significantly elevated and intravenous glucose tolerance test (IVGTT) shows reduced glucose elimination and impaired insulin secretion (28). The model thus shows two important pathophysiological characteristics for impaired glucose tolerance (IGT) and type 2 diabetes: insulin resistance and islet dysfunction.

In this model, the weight gain is due primarily to an increase in mesenteric adiposity. In high-fat diet-fed mice, energy intake is higher and metabolic efficiency index is lower compared to normal diet-fed mice (28). Despite obesity, plasma leptin levels are significantly lower than in the control group in the absence of hyperphagia (29). Increased serum lipid levels develop concomitantly with the development of hyperglycemia, thus deteriorating insulin resistance and producing lipid deposits in non-adipose cells, including beta cells and skeletal muscle cells (30). Microscopically, a high-fat diet induces hypertrophy of pancreatic islets (31).

This model is suitable for examining novel therapeutic interventions. The dipeptidyl peptidase-IV (DPP-IV) inhibitor is efficient in improving glucose tolerance and insulin secretion in the high-fat diet-fed mouse model (28).

**Partial pancreatectomy** as a surgical method of inducing diabetes was described above.





## B) Spontaneous models

### Goto Kakizaki Rat

The Goto Kakizaki (GK) rat is skinny, diabetic rat developed in Japan in 1975 from Wistar species (6). Both insulin resistance and impaired insulin secretion are present in this model (32). GK rats have a decreased numbers of beta cells at birth, which is probably the consequence of their apoptosis during embryonic development (16-18 day). However, GK rats are still normoglycemic and without marked changes in islet morphology at birth (33). The characteristics of diabetes type 2 in this model are mild hyperglycemia (9 mmol/l) in the fourth week and the increase of basal insulin secretion and insulin resistance followed by a decrease in glucose tolerance (34). GK rats are resistant to food intake restrictions (35).

Starting from the 8<sup>th</sup> week, insulin islets show signs of fibrosis with a marked elevation of hyperglycemia (36). In addition, the expression of CD38 and GLUT2 proteins is reduced in GK islets (37-38). Microscopically, numerous macrophages (major histocompatibility complex class II<sup>+</sup> and CD68<sup>+</sup>) and granulocytes were found in and around pancreatic islets (39). Elevated islet IL-1 $\beta$  activity in GK rat promotes cytokine and chemokine expression, leading to the recruitment of innate immune cells (40). Treatment of GK rats with IL-1 receptor antagonists decreases hyperglycemia, reduces the proinsulin/insulin ratio and improves insulin sensitivity (40).

The most important factor for the disorder of insulin secretion is the dysfunction of glucose metabolism connected with a deficiency of enzymes that control oxidative glycolysis (6). The GK rat develops some features that can be compared with the complications of diabetes seen in humans (1).

### Zucker Fatty and Zucker Fatty Diabetic Rat

The Zucker Fatty (ZF) rat has a defect of a gene that codes for leptin receptors (*fa/fa*), resulting in the development of obesity and hypertension with associated renal and cardiovascular disease (6). The Zucker Fatty Diabetic Rat (ZDF) has the identical genetic mutation and, additionally, a mutation which leads to the spontaneous development of hyperglycemia in males during the 7<sup>th</sup>-10<sup>th</sup> week (41). The females become hyperglycemic only after high-fat diet, and it is supposed that this additional mutation is expressed in beta cells (42).

Leptin suppresses insulin secretion and controls food intake. The *fa/fa* mutation of the leptin receptor results in insulin resistance with severe glucose intolerance (6). ZF and ZDF rats are hyperphagic due to the reduction in leptin signalling that results in obesity (6). These animals, in adulthood, become extremely dislipemic with increased concentration of free fatty acids, cholesterol, and triglycerides in plasma with a consequent development of diabetes mellitus complications (cardiovascular, renal, and peripheral neuropathy) (6, 43-44).

Pancreatic islets in adults show increased beta cell activity, with marked characteristics of a prediabetic state (6). After the onset of diabetes, pancreatic islets become irreg-

ular as a result of hyperplasia, hypertrophy and infiltration of inflammatory cells (45). An almost complete absence of beta cells was detected by the 14<sup>th</sup> week of age (45).

Reduced expression of mGDP and pyruvate carboxylase activity was noticed in this model (46).

### Otsuka Long Evans Tokushima Fatty Rat

The Otsuka Long Evans Tokushima Fatty Rat (OLETF) is derived by inbreeding from the glucose-intolerant Long-Evans colony of rats (47). These animals develop hyperglycemia slowly, by 40 weeks of age, with greater incidence in males than in females (1). Adult animals are mildly obese and hyperglycemic, with increased appetite and renal glomerulosclerosis. These changes are the result of a deletion of the gene for cholecystokinin 1 receptors. Diet and exercise prevent type 2 diabetes in this model (48).

Histopathologic changes inside the islets during the 8<sup>th</sup>-10<sup>th</sup> week show an infiltration of inflammatory cells followed by tissue damage. Beta cell hyperplasia and proliferation occur between the 20<sup>th</sup> and 40<sup>th</sup> weeks, while atrophy and fibrosis of the islets start after the 40<sup>th</sup> week (47). From the 72<sup>nd</sup> week, inflammatory cytokines (tumour necrosis factor- (TNF- ), interleukin-1 $\beta$  (IL-1 $\beta$ ) and interleukin-6 (IL-6)) are detectable by immunohistochemical method (49).

### Psammomys obesus

The *Psammomys obesus* is rat from North America and the Middle East. It usually feeds on a low-calorie diet and demonstrates the strong effect of a high-calorie diet by developing hyperglycemia within 4-7 days (6).

Pathogenesis of the disorders goes through four typical stages: normoglycemia and normoinsulinemia, normoglycemia with hyperinsulinemia (indicating insulin resistance), hyperglycemia with hyperinsulinemia, and hyperglycemia with hypoinsulinemia (50). The last stage is characterized by marked apoptosis and a reduction of beta cell mass (51). Returning to a low-calorie diet results in the recovery of beta cells, indicating that the irreversible loss of beta cell mass is a late event (52).

## TRANSGENIC AND KNOCKOUT MODELS OF DIABETES MELLITUS

In experimental diabetes, numerous transgenic and knockout models are being used. These models have been created by genetic manipulations using either transgene insertion and/or targeted gene-deletion approaches.

Still, these techniques have some limitations. Some homozygotes may die in the womb, which makes studies in adults impossible. Several genes could function differently during embryogenesis and in adulthood. Finally, some models' gene expression is altered in soft tissues, and therefore it is impossible to analyze their function in the desired tissue. Tissue-specific knockout models for diabetes, in which the gene being studied is only knocked out in specific tissues, were created to address this issue.



Mouse/rat model	Phenotype	Reference
IR +/-	Mild insulin resistance	53, 54
IR -/-	Severe hyperglycemia, neonatal death from ketoacidosis	54
IRS-1 -/-	Mild insulin resistance, normoglycemia, elevated serum triglycerides, hypertension, growth retardation	55
IRS-2 -/-	Insulin resistance, hyperglycemia, reduced beta-cell mass	56
IRS-1 -/- IRS-3 -/-	Hyperglycemia, lipatrophy, decreased plasma levels of leptin, hyperinsulinemia	57
IRS-1 +/- IR +/-	Insulin resistance, beta-cell hyperplasia	58
GK -/-	Hyperglycemia, perinatal death	59
GK +/-	Glucose intolerance	59
GLUT4 -/-	Mild changes in glucose hemostasis, growth retardation, reduction of fat, cardiac hypertrophy	60
PPAR $\gamma$ +/-	Mild insulin resistance	61
PPAR $\gamma$ 2 -/-	Insulin resistance, reduction of white adipose tissue	62
PPAR $\alpha$ -/-	No phenotypical defect without pharmacological intervention	63
hIAPP +/+ rat	Decreased proliferation and increased apoptosis of beta cell, hyaline replacement of islet cells	64

homozygous (-/-), heterozygous (-/+)

**Table 2.** Transgenic and knockout models of Diabetes mellitus

The first knockout models of insulin resistance aimed at the disruption of major molecules in insulin receptor (IR) signaling. Heterozygous IR knockout mice show 50% of IR expression, which is sufficient for the maintenance of physiological blood glucose concentrations (53). In contrast, homozygous IR-deficient mice rapidly develop diabetic ketoacidosis and die within 3-7 days after birth, showing the central role of IR in glucose metabolism (54). Aside from IR knockouts, various molecules of the signal transduction pathway initiated by IR have been disrupted, such as insulin receptor substrates (IRS-1, IRS-2, IRS-3) (55-58). Insulin resistance and insulin secretory defects have also been presented by a generation of mice deficient in beta cell glucokinase (GK). The heterozygous beta cell GK knockout mouse is characterised by decreased insulin secretion in response to glucose (59). The homozygous knockout mouse model with a disrupted gene for the glucose transporter 4 (GLUT4), which mediates glucose transport in muscle and adipose tissue in response to insulin, is characterised by mild changes in glucose hemostasis, growth retardation, reduction of fat and cardiac hypertrophy (60). The important physiological role of peroxisome proliferator-activator receptors (PPAR $\alpha$  and PPAR $\gamma$ ) was deduced from findings identifying the PPARs as primary targets of thiazolidinedione insulin sensitizers and fibrates that have been used in the successful treatment of diabetes and dyslipidemia. Diverse knockout models have been created to study the function of single PPARs isoforms (61-63). Similar to other mouse models, a transgenic human islet amyloid polypeptide-expressing rat has been recently developed (64). This model shows pancreatic amyloidosis, insulin resistance, and reduced  $\beta$ -cell mass with resulting hyperglycemia (64). Some of the transgenic/knockout and tissue-specific knockout models are presented in Table 2.

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## DIAGNOSTICS AND THERAPY OF LEFT VENTRICULAR HYPERTROPHY IN HAEMODIALYSIS PATIENTS

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## DIJAGNOSTIKA I LEČENJE HIPERTROFIJE LEVE KOMORE KOD BOLESNIKA NA HEMODIJALIZI

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### ABSTRACT

*Cardiovascular diseases present a leading cause of death in patients treated with haemodialysis. The rate of cardiovascular mortality in this population is approximately 9% on an annual basis, with left ventricular hypertrophy, ischemic heart diseases and heart failure having the highest rates of mortality. Left ventricular hypertrophy is present in 75-80% of haemodialysis-treated patients. The most important risk factors for the progression of left ventricular hypertrophy are: hypertension, arteriosclerosis, secondary aortic stenosis, anaemia, increased volume of extracellular fluid and increased blood flow through the vascular access for haemodialysis. Left ventricular hypertrophy is present when the left ventricular mass index on echocardiography exceeds 131 g/m<sup>2</sup> in males and 100 g/m<sup>2</sup> in females. Left ventricular hypertrophy is a risk factor of unfavourable outcome in patients treated with haemodialysis. The identification of patients with increased risk of progression of left ventricular hypertrophy, the timely implementation of adequate treatment, and the realisation and maintenance of targeted values of risk factors decelerates the progression of the hypertrophy and leads to the regression of existing left ventricular hypertrophy, the reduction of cardiovascular morbidity and mortality rates and the improvement of the quality of life of patients treated with haemodialysis.*

**Key words:** risk factors, left ventricular hypertrophy, echocardiography, haemodialysis

### SAŽETAK

*Kardiovaskularne bolesti su vodeći uzrok smrti bolesnika koji se leče hemodijalizom. Stopa kardiovaskularnog mortaliteta kod ovih bolesnika iznosi približno 9% godišnje, a među kardiovaskularnim bolestima najveća je prevalencija hipertrofije leve komore, ishemijske bolesti srca i srčane slabosti. Hipertrofiju leve komore ima 75-80% bolesnika koji se leče hemodijalizom. Najznačajniji faktori rizika za razvoj hipertrofije leve komore su: hipertenzija, arterioskleroza, sekundarna aortna stenoza, anemija, povećan volumen ekstracelularne tečnosti i povećan protok krvi kroz vaskularni pristup za hemodijalizu. Hipertrofija leve komore je prisutna ako je na ehokardiografskom pregledu indeks mase leve komore veći od 131 g/m<sup>2</sup> kod muškaraca i veći od 100 g/m<sup>2</sup> kod žena. Hipertrofija leve komore je faktor rizika za nepovoljan ishod bolesnika koji se leče hemodijalizom. Izdvajanje bolesnika koji imaju povećan rizik za razvoj hipertrofije leve komore, pravovremena primena odgovarajućeg lečenja, ostvarivanje i održavanje ciljnih vrednosti faktora rizika usporavaju razvoj i dovode do regresije postojeće hipertrofije leve komore, do smanjenja stope kardiovaskularnog morbiditeta i mortaliteta, i poboljšanja kvaliteta života bolesnika koji se leče hemodijalizom.*

**Ključne reči:** faktori rizika, hipertrofija leve komore, ehokardiografija, hemodijaliza

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## INTRODUCTION

Cardiovascular diseases present a leading cause of death in patients treated with haemodialysis [1, 2]. The rate of cardiovascular mortality in this population is approximately 9% on an annual basis, with left ventricular hypertrophy, ischemic heart diseases and heart failure having the highest rates of mortality [1, 2].

## METHODS

For this study, the Medline database was used. The following key words were used for searching: risk factors, left ventricular hypertrophy, echocardiography, and haemodialysis. About 40.000 references were found dealing with these types of problems. Systematic review articles and well-controlled clinical studies were singled out. Editor's letters and uncontrolled clinical studies were not used. Systematic review articles were used and analysed by the National Kidney Foundation Kidney Disease Outcome Quality Initiative (NKF KDOQI), which confirmed the validity and quality of the selected references.

## RESULTS

### Aetiopathogenesis of left ventricle hypertrophy

The most important risk factors of left ventricle hypertrophy in haemodialysis patients are hypertension, arteriosclerosis, adult aortic stenosis, anaemia, increased blood flow through the vascular access for haemodialysis and increased volume of extracellular fluid [3, 4]. Pressure overloads of the left ventricle due to hypertension, aortic stenosis and arteriosclerosis of blood vessels can cause thickening of the left ventricle wall without the enlargement of the ventricle diameter (concentric left ventricle hypertrophy) [5, 6]. Volume overloads of the left ventricle due to increased sodium and water intake, anaemia and increased blood flow through the vascular access for haemodialysis ( $Q_{AV} \geq 1000$  ml/min) cause the ventricle wall to become thickened and cause an enlargement of ventricle diameter (eccentric left ventricle hypertrophy) [5, 6]. The development of left ventricle hypertrophy and fibrosis of the myocardial interstitium can also be affected by factors that do not depend on the *afterload* and *preload*, such as microinflammation, oxidative stress, the activation of the intracardial renin-angiotensin system-RAS, the vitamin D deficiency and uncontrolled secondary hyperparathyroidism, scheme 1 [7-10]. Left ventricle hypertrophy goes through two phases. Adaptive hypertrophy is actually the response to increased tension stress to the left ventricle wall and has a protective function. When volume and pressure overload the left ventricle to the point of cardiac muscle failure, adaptive hypertrophy becomes maladaptive left ventricle hypertrophy, with myocyte loss, heart failure and, eventually, death [11].

## Diagnostics of left ventricular hypertrophy

Morphological and functional disorders of the left ventricle in haemodialysis patients include systolic disorder, concentric hypertrophy, eccentric hypertrophy, left ventricle dilation and diastolic dysfunction. The systolic function of the left ventricle is disturbed when echocardiography reveals that the fractional shortening of the left ventricle (FSLV) is  $\leq 25\%$  and that the ejection fraction of the left ventricle (EFLV) is  $\leq 50\%$  [12-14]. Concentric hypertrophy of the left ventricle is defined by a left ventricle mass index- (LVMi)  $> 131$  g/m<sup>2</sup> in males and  $>100$  g/m<sup>2</sup> in females and a relative wall thickness of the left ventricle (RWT)  $> 45\%$  [12-14]. Eccentric hypertrophy of the left ventricle is characterised by an increased left ventricle mass index (LVMi  $> 131$  g/m<sup>2</sup> in males and  $>100$  g/m<sup>2</sup> in females) and an RWT  $\leq 45\%$  [12-14]. An echocardiographic estimation of the left ventricle function in diastole is based upon the measurement of the velocity of early (E) and late (A) components of blood flow through the mitral valve and their relative ratio E/A. There are three types of left ventricle diastolic dysfunction: relaxation disorder [ $V_{maxE}/V_{maxA} < 1,0$ ; E wave deceleration time ( $DT_e$ )  $> 250$  ms; and isovolumetric relaxation of the left ventricle (IVRT)  $> 100$  ms]; pseudonormalisation and restriction disturbance [ $V_{maxE}/V_{maxA} > 1,6$ ;  $DT_e < 150$  ms; and IVRT  $< 60$  ms] [15]. Cardiac Magnetic Resonance Imaging (CMRI) is the golden standard for remodelling of the left ventricle and for the estimation of myocardial interstitium fibrosis in haemodialysis patients [16].

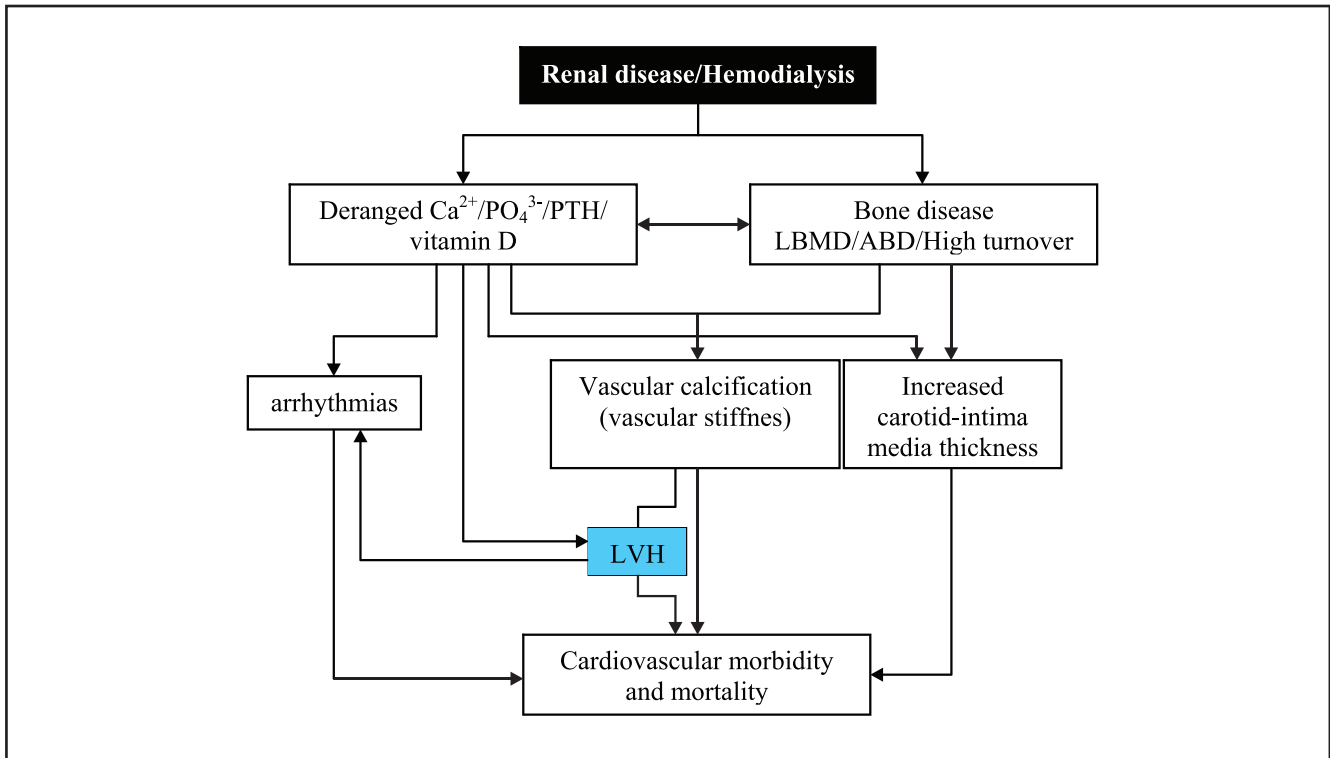
### Clinical importance of left ventricular hypertrophy

Left ventricle hypertrophy is accompanied by disordered diastolic function (50-60% of haemodialysis patients), by ventricle arrhythmias (dispersion interval (QTd)  $> 50$  ms), by *de novo* ischemic heart disease (in patients with LVMi  $> 160$  g/m<sup>2</sup>) and by unfavourable outcomes in haemodialysis patients [17-19]. A high left ventricle mass index (LVMi  $> 120$  g/m<sup>2</sup>) and an LV mass/volume ratio (LVMi/iEDV)  $> 2,2$  g/ml are predictors of a poor outcome in dialysis patients with a normal LV volume (iEDV  $\leq 90$  ml/m<sup>2</sup>) and a normal systolic function (FS  $> 25\%$ , EF  $> 50\%$ ) [20, 21].

### Therapy of left ventricular hypertrophy

Well-timed risk factor detection and adequate therapy help regression of left ventricle hypertrophy in haemodialysis patients [22-25]. Results of large, well-controlled clinical trials stress that the management of anaemia and blood pressure control, the correction of metabolic disorders of calcium, phosphate, vitamin D and parathormone (parathyroid hormone), as well as the individualisation of treatment of haemodialysis as the most important factors in the successful management of left ventricle hypertrophy [22-25].

Anaemia is an important cause of left ventricle hypertrophy, and a decrease in haemoglobin of 10 g/l is



**Shema 1.** Impact of calcium and phosphate metabolism disorder on progression of cardiovascular morbidity and mortality in hemodialysis patients

associated with an increase in the LVMi of 10 g/m<sup>2</sup> [25, 26]. Erythropoietin treatment and iron supplementation should maintain the haemoglobin levels at 110 - 120 g/l (Hct = 33 -36%) and ferritin levels at 200-500 ng/ml [25, 26]. Hypertension is an important risk factor for the development of left ventricle hypertrophy, and an increase in the mean arterial blood pressure of 10 mmHg increases the left ventricle mass index by 7,2 g/m<sup>2</sup> [27]. In haemodialysis patients, before haemodialysis, the arterial blood pressure should be ≤ 140/90 mmHg and should be ≤ 130/80 mmHg afterwards. Angiotensin I convertase blockers, angiotensin II receptor blockers and β-blockers should be used in the treatment of hypertension [27].

Arteriosclerosis increases peripheral vascular resistance, overloads the left ventricle because of increased blood pressure and acts as a stimulus for left ventricle remodelling [28]. The most important risk factors for the development of arteriosclerosis (media calcification) and for heart valve calcification in haemodialysis patients are hyperphosphatemia, increased solubility products and secondary hyperparathyroidism [29-31]. Calcification of the arterial medial wall, increased artery stiffness, heart valve calcification and decreased opening of the aortic valve leaflets leads to the development of left ventricle concentric hypertrophy, and an increased serum parathormone level also leads to a significant myocardial interstitial fibrosis (fibroblast proliferation, increased creation and deposition of extracellular ma-

trix proteins in the myocardial interstitium) in haemodialysis patients [29-31]. Hygienic and diet regimens, phosphate binders, active vitamin D3 metabolites, and calcimimetics should maintain the phosphate concentration at < 1,6 mmol/l, the solubility product ([Ca<sup>2+</sup>] x [PO<sub>4</sub><sup>3-</sup>]) at ≤ 4,4 mmol<sup>2</sup>/l<sup>2</sup>, the concentration of 25(OH) D<sub>3</sub> at > 30 ng/ml and the concentration of iPTH od at 100-300 pg/ml (iPTH < 500 pg/ml) [29-34].

Haemodialysis treatment significantly reduces left ventricle hypertrophy. Every-day short-term haemodialysis (6 times weekly, 3h) for 12 months decreases left ventricle hypertrophy by 30%, as compared to conventional haemodialysis (3 times weekly, 4h) [24]. Clinical trial results show that night-time haemodialysis (3 times weekly, 6-8h) during the six-month period significantly reduces left ventricle hypertrophy compared to conventional haemodialysis (3 times weekly, 4h) [25].

Recognising patients with an increased risk for the development of left ventricular hypertrophy and implementing adequate treatment to achieve target values of risk factors decreases cardiovascular morbidity and mortality and improves the quality of life in patients receiving regular haemodialysis treatments.

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# THE RELATIONSHIP BETWEEN SPORTS ENGAGEMENT, BODY MASS INDEX AND PHYSICAL ABILITIES IN CHILDREN

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## ABSTRACT

*Hypokinesia, which is a consequence of modern technological development, is now one of the three biggest health risk factors. Whereas physical inactivity and obesity in adults is subject of concern, these factors in children presage an even greater public health problem because many health-related problems and life-threatening diseases begin in childhood and adolescence.*

*It has been noted that regular physical activity modulates vascular endothelial function via an effect on basal nitric oxide (NO) production, thus leading to a significant reduction in the incidence of cardiovascular disease (1). Exercise can also increase energy expenditure and can create a negative energy balance. Conversely, scientific data regarding the effect of school-based physical activity interventions on body*

*mass index (BMI) in children are conflicting (2, 3). Moreover, there is insufficient clinically useful scientific data on BMI in children and its possible association with motor skills. With both sexes, motor abilities improve from physical development in adolescents, and regular sports training influences fitness and motor abilities.*

*The aim of our study was to evaluate the morphological status and physical abilities of 13-year-old children in order to investigate the relationship between sports engagement, BMI and physical abilities. The results of our study should give coaches important parameters in the selection of young athletes, as well as provide them with information that may be useful in training and programming assignments.*

## MATERIALS AND METHODS

The research was carried out with a group of 67 boys from an elementary school. The boys were 12-13 years old and had differing sports training experiences.

After an assessment of morphological characteristics (height, weight, BMI), the children were subjected to a battery of motor tests designed to test speed and coordination (Jumping and going through Swedish boxes, slalom with 3 balls, hand tapping, 50-m sprint, 30-m flying start sprint and 20-m slalom sprint (4)).

## RESULTS AND DISCUSSION

The average BMI of the children who participated in the study was  $19.2 \pm 3.4$ . After transforming BMI data to appropriate percentiles according to the formulas proposed by the National Center for Health Statistics of United States (5), we classified each subject into one of four groups: underweight (athletes vs. non-athletes: 7.7%

vs. 17.8%), healthy weight (69.2% vs. 46.4%), overweight (12.8% vs. 21.4%) and obese (10.2% vs. 14.3%). Although BMI is thought to measure excess weight rather than excess body fat, if appropriate cut-off points are used, a high BMI level is a moderately sensitive and very specific indicator of excess adiposity among children (6).

Since hypokinesia in children and adolescents is thought to be a growing problem, especially in urban places, it is encouraging that 58.2% of the boys in the study stated that they participated in a sport activity. To determine if sport activity influences morphological status of young 13-year-old boys, we applied a contingency analysis ( $\chi^2$ -test) and found no significant difference between children who train for a sport and those that do not. These results are in agreement with many other studies that have found that, in contrast to biological growth, the absence of involvement in sports activities in children is not the main factor that influences their morphological status and physical fitness. Moreover, an analysis of morpho-functional skills of young





teens and premature soccer players (14 and 15 vs. 16 and 17 year olds) revealed that the aerobic capacity ( $VO_2$  max) and BMI significantly changed with age, but not with the training experience of the examined athletes (7).

Regarding the influence of morphological status of 13-year-old boys on motor performance, the results of our study suggest that morphological status, estimated by BMI, is an important predictor of physical abilities in children. This finding is in agreement with the results of other studies (8). Discriminative analysis (Kruskal-Wallis test, Table 1) showed that subjects with different morphological status had significantly different performance in 5 out of the 6 tests performed, of which two were designed to test speed and three tested coordination. However, canonical discrimination, used to determine the influence of morphological status on latent motor function, showed significant differences only in coordination tests.

Regarding the influence of sports engagement on motor performance in our participants, a Man-Whitney test (Table 2) showed that, compared with those that did not train, boys who train for a sport had significantly better results only in 2 motor tests, which were confirmed by factor analysis to be valid tests of coordination. These results suggest that the morphological status of 13-year-old children better predicts coordination performance than speed performance.

The results of our study suggest that involvement of adolescents in sport activities has much less influence on motor abilities than that of natural biological growth and maturation. The age at which the boys in this study were tested falls within the first phase of adolescence, which is characterised by sudden and intermittent skeletal maturation without being followed by adequate skeletal muscle development. That is one of the reasons why boys of this age experience a fall in coordination performance. In other words, motor development in 13-year-old boys does not depend on training experience but instead relies on genetics and acceleration of growth.

It should be emphasised that the small number of participants in this study limited our ability to make any significant anthropological generalisations. However, the sample size did not affect the ability to address the main aim of this study, which was to assess the predictive value of morphological status and involvement in sports as parameters that affect motor performance.

Variable	Athletes	Non-athletes	Sig.
Jumping and going through Swedish boxes (sec)	15.85	16.58	.097
Hand tapping (taps)	69.28	64.07	.018*
Slalom with 3 balls (sec)	22.04	25.86	.004*
50-m sprint (sec)	9.24	9.49	.088
30-m flying start sprint (sec)	5.38	5.50	.065
20-m slalom sprint (sec)	4.76	4.88	.132

**Table 2.** Mean values of results that boys who train or do not train some sport achieved on motor tests.

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**Table 1.** Mean values of results that subjects with different morphological status achieved on motor tests.

Variable	Underweight	Healthy weight	Overweight	Obese	Sig.
Jumping and going through Swedish boxes (sec)	15.60	15.37	17.59	20.23	.000*
Hand tapping (taps)	60.00	69.00	73.09	73.75	.040*
Slalom with 3 balls(sec)	28.65	22.26	21.61	28.29	.004*
50-m sprint (sec)	9.90	9.15	9.47	10.28	.002*
30-m flying start sprint (sec)	5.82	5.37	5.53	6.01	.008*
20-m slalom sprint (sec)	5.03	4.77	4.97	5.05	.204



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