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The effect of NAPQI on neurotransmitter transporters

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## Table of contents

ACKNOWLEDGEMENTS .....	3
ABSTRACT .....	4
ZUSAMMENFASSUNG .....	5
1. INTRODUCTION .....	6
1.1. Neurotransmitters and Neurotransmitter Transporters .....	6
1.2. Structure and function of neurotransmitter transporters.....	6
1.3. Monoamine transporters and psychostimulant action .....	7
1.3.1. Serotonin transporter .....	9
1.3.2. Dopamine transporter .....	11
1.3.3. Norepinephrine transporter .....	12
1.4. $\gamma$ -Aminobutyric acid (GABA) transporter .....	13
1.5. Psychostimulant action of cocaine on monoamine transporters.....	15
1.6. Paracetamol .....	15
1.6.1. Pharmacokinetics or metabolism of acetaminophen .....	16
1.6.2. Acetaminophen mechanism of toxicity .....	17
AIMS OF THE THESIS .....	19
2. MATERIAL AND METHODS.....	20
2.1. Cloning .....	20
2.1.1. Generating competent E. coli bacteria .....	20
2.1.2. QuikChange site-directed mutagenesis .....	20
2.1.3. Transformation .....	21
2.1.4. DNA/Plasmid isolation .....	22
2.2. Cell culture .....	22
2.2.1. Cultivation .....	22
2.2.2. Cell counting/seeding .....	22
2.2.3. Transfection .....	23
2.3. Pharmacological Assays .....	23
2.3.1. Uptake assay .....	23
2.3.2. Inhibition assay .....	24
2.4. Materials .....	25
2.4.1. Standard solutions/buffers .....	25

3.	RESULTS .....	26
3.1.	Evaluation of the inhibitory capacity of NAPQI on different mutated serotonin transporters 26	
3.2.	Time dependent uptake inhibition assay.....	27
3.3.	NAPQI is a competitive inhibitor of uptake of SERT .....	27
3.4.	Increase of serotonin shifts the inhibition curve for NAPQI .....	28
3.5.	Effects of NAPQI on other transporters: GABA.....	29
4.	DISCUSSION.....	30
4.1.	Evaluation of inhibition capacity of NAPQI on different mutated serotonin transporters ..	30
4.2.	Time dependent uptake inhibition assay.....	31
4.3.	NAPQI is a competitive inhibitor of SERT.....	31
4.4.	Increasing of serotonin shifts the inhibition curve for NAPQI .....	31
4.5.	Effects of NAPQI on the GABA transporter.....	32
5.	LIST OF FIGURES.....	33
6.	REFERENCES .....	34

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## ABSTRACT

Paracetamol (Acetaminophen) is in use as an analgesic drug for decades, although its mechanism of action is not understood yet. Among various theories, one states that the toxic metabolite of Paracetamol NAPQI (N-acetyl-p-benzoquinone imine) can modify TRP-channels, leading to analgesia via this pathway. NAPQI is a cysteine-modifying reagent that, under therapeutic concentrations of paracetamol, binds to the sulphur group of glutathione and is excreted. When the glutathione stores are depleted, NAPQI can irreversibly bind cysteine residues of other proteins, rendering them non-functional. This can possibly result in life-threatening liver failure.

The analgesic action of paracetamol also seems to include the serotonergic system. Serotonin and norepinephrine can blunt pain sensation via descending neurons. The serotonin transporter (SERT), which is essential for the reuptake of serotonin, has been reported to be susceptible to cysteine modifying reagents.

Therefore, we investigated a possible effect of NAPQI on SERT by performing radioligand uptake assays in cells stably expressing wildtype SERT. Furthermore, we generated alanine mutants of cysteines in SERT to investigate the mechanism of NAPQI.

Our results show that the metabolite NAPQI is capable of inhibiting the uptake of serotonin. Interestingly, the double mutant C109A/C357A was inhibited by NAPQI similar to wildtype. Furthermore, the inhibition of SERT was clearly reversible and competing with serotonin.

The transporter for  $\gamma$ -aminobutyric acid, GAT, was tested for a similar effect by NAPQI. Results show that, unlike SERT, GAT was not inhibited.

We conclude that NAPQI inhibits SERT in a specific manner, and, unlike its action on TRP channels, it does so in a reversible and competitive manner. Blockade of SERT could increase the analgesic effect of descending antinociceptive neurons, providing an additional explanation for the analgesic mechanism of paracetamol.

## ZUSAMMENFASSUNG

Paracetamol (Acetaminophen) ist ein seit Jahrzehnten gebräuchliches Schmerzmittel, obwohl dessen Wirkmechanismus nicht aufgeklärt ist. Laut einer vieler Theorien kann NAPQI (N-Acetyl-p-Benzoquinonimin), der toxische Metabolit des Paracetamol, TRP-Kanäle modifizieren und so zu einer analgetischen Wirkung führen. NAPQI ist eine Cystein-modifizierende Substanz, die bei therapeutischen Konzentrationen von Paracetamol nach Bindung an die Schwefelgruppe von Glutathion ausgeschieden wird. Bei Erschöpfung des Glutathion-Speichers kann NAPQI irreversibel an Cystein-Seitenketten von anderen Proteinen binden, die daraufhin ihre Funktion verlieren. Dieses Ereignis kann möglicherweise zu einem Leberversagen führen.

Die analgetische Wirkung des Paracetamol könnte das serotonerge System der Schmerzempfindung auch mit einschließen. Serotonin und Norepinephrin können die Schmerzempfindung mit Hilfe efferenter Neuronen abtumpfen. Der Serotonintransporter SERT ist für die Wiederaufnahme von Serotonin zuständig und empfindlich auf Cystein-modifizierende Substanzen.

Aufgrund dieser Datenlage wurde mit Hilfe von Radioliganden-Aufnahme Assays eine mögliche Wirkung von NAPQI auf SERT in Zellen mit Wildtyp-SERT untersucht. Zusätzlich wurden Alanin-Mutanten einiger Cysteine des SERT erstellt, um die Wirkung des NAPQI durch die Cysteine festzustellen.

Die Ergebnisse dieser Arbeit zeigen, dass der Metabolit NAPQI die Aufnahme von Serotonin hemmen kann. Die Doppelmutante C109A/C357A wurde von NAPQI wie bei Wildtyp inhibiert. Wir fanden zusätzlich heraus, dass die Hemmung von SERT deutlich reversibel und kompetitiv mit Serotonin war. Als Kontrolle wurde der Transporter für  $\gamma$ -Amino-Buttersäure (GABA) auch nach einer möglichen Wirkung des NAPQI überprüft, wobei hier eine irreversible Hemmung festgestellt wurde.

Mit Hilfe dieser Arbeit haben wir eine spezifische Hemmung von SERT durch NAPQI gezeigt, die im Gegensatz zu TRP Kanälen reversibel und kompetitiv ist. Diese Hemmung könnte die analgetische Wirkung auf antinozizeptive Neuronen erhöhen und so auf einen zusätzlichen Wirkmechanismus von Paracetamol hindeuten.

# 1. INTRODUCTION

## 1.1. Neurotransmitters and Neurotransmitter Transporters

Before the mid-nineteenth century, scientists assumed that neurons communicate by electrical signalling. The work of Alan Hodgkin and Andrew Huxley in 1952 (Hodgkin & Huxley 1952) confirmed that action potentials are transferred through the nerve cells by ion permeation. Otto Loewi identified the transmission via Vagustoff and Henry Dale identified that the Vagustoff to be Acetylcholine (Dale and Dudley, 1929). Loewi and Dale won the 1936 Medicine and Physiology Nobel Award for this important discovery.

Neurotransmitters are small chemical compounds also known as chemical messengers, which enable neurotransmission. They transmit signals from one nerve cell (neuron) to another “target” neuron.

Major neurotransmitters include:

- Amino acids such as glutamate, D-serine,  $\gamma$ -aminobutyric acid (GABA), glycine
- Peptides such as somatostatin, opioid peptides
- Monoamines such as serotonin (SER, 5-HT), dopamine (DA), norepinephrine (noradrenaline; NE, NA), epinephrine (adrenaline), histamine.

Neurotransmitters are classified into “inhibitory” or “excitatory” depending on their effect on the target cell: if they hyperpolarize the target cell, they are inhibitory (they increase the threshold for the cell to fire an action potential), whereas if they depolarize it, they are excitatory. The ion selectivity of the corresponding receptors at the postsynaptic membrane determines whether the cell is hyperpolarized or depolarized. Neurotransmission, also called synaptic transmission, is the process by which neurotransmitters are released by a neuron (the presynaptic neuron), and bind to and activate the receptors of another neuron (the postsynaptic neuron). Neurotransmission is essential for the process of communication between two neurons.

The typical course of a neurotransmission is as follows: when an action potential of an excited neuron reaches the synapse, it leads to depolarization of the presynaptic membrane, followed by influx of  $\text{Ca}^{2+}$  into the intracellular compartment. The rise of  $\text{Ca}^{2+}$  induces neurotransmitter-filled vesicles to fuse with the presynaptic membrane, releasing their content into the synaptic cleft (Südhof 2012). The released neurotransmitters bind to neurotransmitter receptors at the postsynaptic membrane (membrane of the target cell), leading to the depolarisation or hyperpolarisation of the cell. (Vanhatalo & Soinila 1998).

Receptor-mediated responses may result from the influx of ions via ionotropic receptors or lead to activation of intracellular signalling cascades through G protein-coupled receptors (metabotropic receptors).

## 1.2. Structure and function of neurotransmitter transporters

After release of neurotransmitters and receptor activation, neurotransmitters in excess have to be actively removed from the synaptic cleft by either enzymatic processing or reuptake by transporters which are high-affinity integral membrane proteins located on the presynaptic membrane (Masson et al. 1999).

Transporters for amino acids (e.g. SLC1, SLC7, SLC17), nucleotides (e.g. SLC28, SLC35), sugars (e.g. SLC2, SLC50, SLC37), and drugs (e.g. SLC47), play an important role in maintaining crucial processes within and between cells via controlling uptake and efflux (Colas et al., 2016) . The solute carrier

family 6 (SLC6) includes transporters for the inhibitory neurotransmitters such as GABA and glycine, proteinogenic amino acids, the metabolic compound creatine, the osmolytes taurine and betaine, as well as transporters for serotonin, dopamine and norepinephrine (Beuming et al. 2006). In the human genome, the solute carrier family 6 (SLC6) consists of 20 members (Chen et al. 2004; Bröer 2006). SLC6 transporters are not only located on neurons and glia, but also in many non-neural tissues. Furthermore, plenty of family members are expressed excessively in the kidney.

In the SLC6 family, approximately half of the transporters use the electrochemical potential of extracellular  $\text{Na}^+$  ions as an energy source in order to transport their substrate across membrane (Table 1). Extracellular chloride ions are also co-transported, but the exact role of chloride in the transport process has not yet been fully determined. The bacterial orthologue LeuT has been first crystallized and is known to be chloride-independent (Zomot et al. 2007).

One interesting phenomenon exists in the literature regarding sodium/chloride-coupled neurotransmitter transporters: they can operate in two directions. As compared with other substances, amphetamines are well known to induce reverse transport, which has been recorded for many transporters (Roux & Supplisson 2000). By virtue of the low concentration of sodium ions in the cytoplasm compared to the extracellular space, reverse transport is per se a rare event. Nevertheless, in the event of large amounts of substrates transported, the resulting ion conductivity becomes much larger and reverse transport is quite likely to occur (Sitte & Freissmuth, 2010). For example, presence of releasing substrate, such as amphetamines, trigger reverse transport. For this effect, it seems that both the N-terminus and the oligomerization are necessary. Truncation experiments of the N terminus showed that the amphetamine-induced reverse transport in SERT was abrogated when a certain N-terminal stretch was missing (Sucic et al., 2010). This was also found in in DAT (Seidel et al. 2005). On N-terminal serines, PKC- and amphetamine-stimulated phosphorylation of DAT might to contribute to this mechanism by changing the conformation of DAT affirmative for reverse transport (Cervinski et al., 2005).

### **1.3. Monoamine transporters and psychostimulant action**

Monoamine transporters belong to the SLC6 gene family and are responsible for the uptake of biogenic amines. This subfamily consists of the serotonin transporter (SERT, *SLC6A4*), the dopamine transporter (DAT, *SLC6A3*) and the norepinephrine transporter (NET, *SLC6A2*).



Human gene name	Protein name	Predominant substrates	Transport type/coupling ions	Predominant tissue distribution	Link to disease
SLC6A1	GAT1	GABA	C/Na <sup>+</sup> , Cl <sup>-</sup>	GABAergic neurons in central and peripheral nervous system, some non-neural tissues	Epilepsy, schizophrenia
SLC6A2	NET	norepinephrine, dopamine	C/Na <sup>+</sup> , Cl <sup>-</sup>	Central and peripheral nervous system, adrenal gland, placenta	Depression, orthostatic intolerance, anorexia nervosa, cardiovascular diseases
SLC6A3	DAT	dopamine	C/Na <sup>+</sup> , Cl <sup>-</sup>	Brain (dopaminergic neurons)	Parkinsonism, substance abuse, ADHD
SLC6A4	SERT	serotonin	C/Na <sup>+</sup> , Cl <sup>-</sup> , K <sup>+</sup>	Central and peripheral nervous system, epithelial cells, platelets	Anxiety, depression, autism, substance abuse

**Table 1:** Features of the members of the solute carrier 6 (SLC6) gene family. Figure adapted by (Chen et al., 2004).

Monoamine transporters share a predicted structure (Kristensen et al. 2011) which includes more than 600 amino acids and twelve hydrophobic membrane-spanning segments (Blakely et al. 1991). The transport of substrate generally requires 1 sodium (Na<sup>+</sup>) and 1 chloride (Cl<sup>-</sup>) ion (symport) against 1 potassium (K<sup>+</sup>) ion (antiport) (Pacholczyk et al. 1991; Guastella et al. 1990; Ramamoorthy et al. 1993).

Monoamine signalling is regulated by a different set of macromolecules, which includes biosynthetic enzymes, secretory proteins, pre- and postsynaptic receptors, transporters and ion channels.

They play a modulatory role in neurotransmission and are associated with various physiological functions such as regulation of movement, mood, attention and sleep. Disturbances in the regulation and function of the monoamines result in pathological conditions like sleep disturbances, Parkinson's disease and childhood parkinsonism, depression and other mood-related disorders (Kurian et al., 2011; Ng et al., 2015)

Monoamine transporters are pharmacological targets of antidepressants and psychostimulant drugs (Jayanthi & Ramamoorthy 2005). There is a large number of published studies describing that selective monoamine transporter inhibitors (e.g., paroxetine, mazindol, imipramine, desipramine, nisoxetine) are beneficial in the treatment of numerous brain-related disorders such as anxiety disorders, mood disorders, schizophrenia, personality disorders, eating disorders, Parkinson's disease, Alzheimer's disease, depression, addiction and abuse, ADHD (attention deficit hyperactivity disorder), cocaine dependence, stroke, obesity, chronic pain, migraine, epilepsy, narcolepsy and multiple sclerosis (Sasaki-Adams & Kelley 2001; Zhou, Zhang, et al. 2003; Zhou, Kläß, et al. 2003; Smith et al. 1999; Zhang et al. 2002; Fleishaker 2000; Wong et al. 2000; Van Moffaert & Dierick 1999; Plewnia et al. 2002; Versiani et al. 2002).

These psychostimulant drugs not only inhibit the monoamine uptake via their transporters, but are also taken up by the transporters, leading to a reverse transport of monoamine transporters via a non-exocytotic mechanism (Sulzer et al. 2005; Sitte & Freissmuth 2010). Some of those substances (e.g. D-amphetamine,) are still largely used in the treatment of attention deficit hyperactivity disorder (ADHD) and prescribed for weight control and narcolepsy (Olfson et al., 2013) . In the central and peripheral nervous system, each monoamine controls different physiological and behavioural functions. Serotonin regulates aggression, mood, motivation, appetite, sleep and sexual activity. In addition, changes in the serotonin signalling have been linked to mental illnesses related to these biological processes (Coccaro 1989; Owens & Nemeroff 1994). Besides this, serotonin also has important peripheral actions, which include mediation of vasoconstriction, placental and gastrointestinal functions. Dysregulation of dopamine transmission is associated with attention/hyperactivity disorder (ADHD), addiction, schizophrenia, Tourette's syndrome and Parkinson's disease (Bannon et al. 1995). Mood, attention, arousal, stress-responsiveness and affective disorders are controlled by norepinephrine (Klimek et al., 1997; Leonard, 1997; Ressler & Nemeroff, 1999; Schildkraut, 1965). Selective serotonin reuptake inhibitors (SSRIs) such as escitalopram and paroxetine (Wong et al. 1995) and selective norepinephrine reuptake inhibitors (SNRI) such as reboxetine (Andersen et al. 2009) are successfully used in treatment of depressive disorder; methylphenidate is used in the treatment of ADHD; DARIs (dopamine reuptake inhibitors) are highlighted in the treatment of obesity.

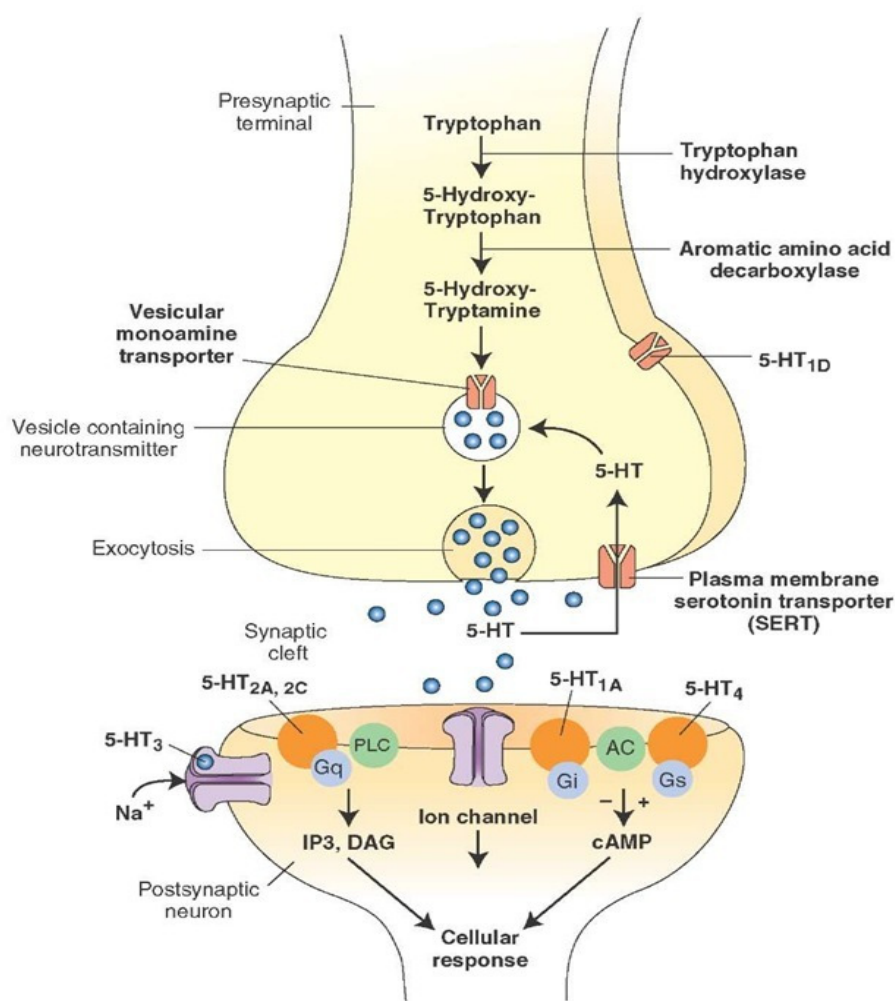
Regulation of transporters can take place via phosphorylation-dependent and -independent post-translational modifications. Post-translational modifications can

- a. alter intrinsic transport activity,
- b. modify transporter turnover,
- c. modulate exocytic fusion of transporter
- d. modulate sequestration of transporter from the plasma membrane via regulating endocytic machinery pathways.

In addition, transporter regulation can also happen through their association with other interacting proteins via phosphorylation-dependent or -independent pathways (Eriksen et al. 2010).

### **1.3.1. Serotonin transporter**

The Na<sup>+</sup>/Cl<sup>-</sup>-dependent serotonin transporter (SERT) plays a role in the uptake of synaptic 5-HT. This is the main process of terminating serotonergic neurotransmission. A single gene, SLC6A4, encoding SERT (Ramamoorthy et al. 1993) is responsible for extracellular 5-HT clearance (Blakely et al. 1991; Hoffman et al. 1991; Lesch et al. 1993). Human SERT (hSERT) consists of 630 amino acids.



**Figure 1:** Synthesis and release of serotonin. cAMP = cyclic adenosine monophosphate; AC = adenylyl cyclase; DAG = diacylglycerol; G<sub>i</sub>, G<sub>q</sub>, G<sub>s</sub> = different G-proteins; IP<sub>3</sub> = inositol triphosphate; 5-HT<sub>1A</sub>, 5-HT<sub>1D</sub>, 5-HT<sub>2A</sub>, 5-HT<sub>2C</sub>, 5-HT<sub>3</sub>, 5-HT<sub>4</sub> = different 5-HT receptors (figure from What-When-How.com).

Besides serotonergic neurons, SERT is expressed in peripheral tissue (Lesch et al. 1993; Rudnick 1977) including specialized cells of the gut (Gordon & Barnes 2003), placenta (Balkovetz et al. 1989), lung (Paczkowski et al. 1996), blood lymphocytes (Faraj et al. 1994; Gordon & Barnes 2003) and also platelets (Jayanthi et al. 2005; Carneiro & Blakely 2006; Carneiro et al. 2008).

Additionally, findings of modified SERT expression in various types of psychopathology point out the importance of SERT in maintaining normal brain function (Murphy et al. 2004). SERT also interacts with psychostimulants such as cocaine, which are inhibitors, while others like amphetamine derivatives including fenfluramine and 3,4-methylenedioxymethamphetamine (MDMA or "Ecstasy") are alternative substrates for SERT (Rudnick & Wall 1992).

Serotonin is one of the key components in the relay of pain, which is possible by descending serotonergic pathways. Especially in the periphery, it enhances the inflammatory signals and contributes hyperalgesia, the excessive sensation (Sommer, 2010).

### **1.3.2. Dopamine transporter**

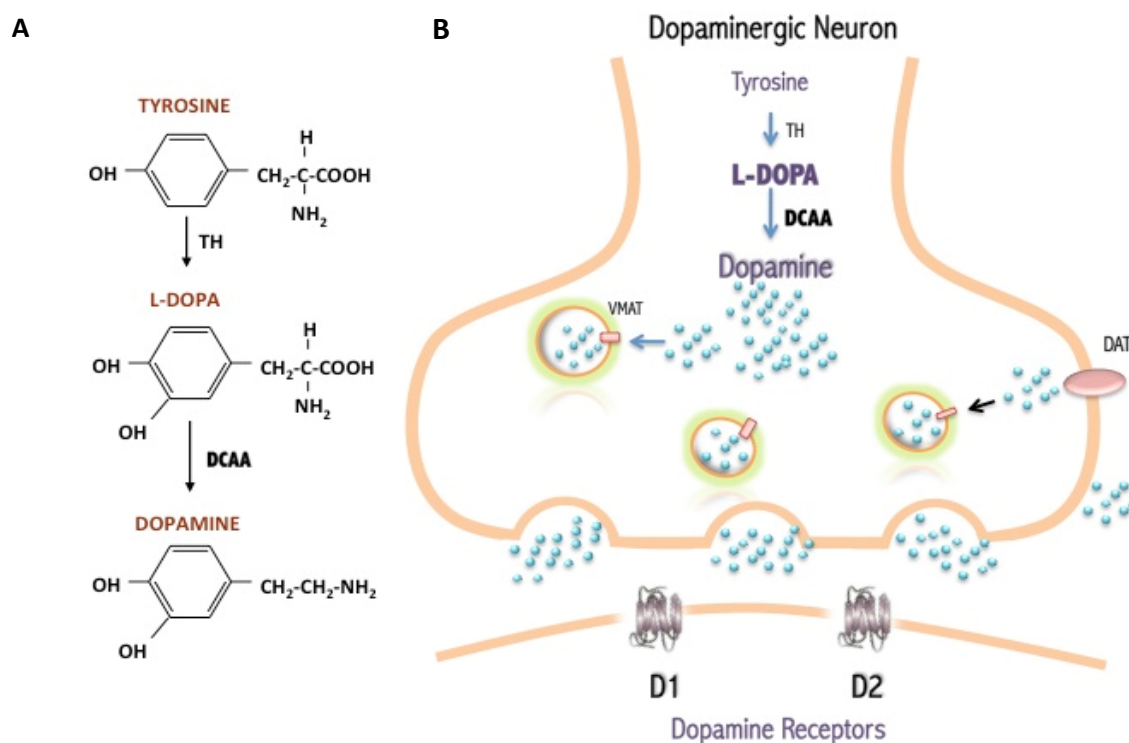
In and around the synapse, the concentration of dopamine (DA) is reduced by reuptake. The dopamine transporter (DAT) is a main regulator of dopamine neurotransmission and is located in chromosome 5p15.3 (Giros et al. 1992), encoding 620 amino acids.

DAT is mainly expressed in dopaminergic neurons (as shown in figure 2) and predominantly present in the brain. DAT is also found in the periphery, including lymphocytes (Amenta et al. 2001). DAT knockout mice exhibit hyperactivity, dwarfism, cognitive deficits, sleep dysregulation, alterations in gut motility, skeletal abnormalities and stereotypical behaviors (Gainetdinov & Caron 2003; Giros et al. 1996).

Furthermore, the dopamine transporter is the main molecular target for several psychostimulants such as cocaine, amphetamine and methamphetamine (Ritz et al. 1987; Kuhar et al. 1991).

The basis of DAT for normal DA clearance and signalling states that DAT dysfunction may apply to various brain disorders associated with dysregulation of DA transmission such as schizophrenia, affective disorders and addiction. Human neuroimaging and genetic studies demonstrate modified DAT availability or function in attention deficit hyperactivity disorder (ADHD). Two ADHD medications, methylphenidate and amphetamine target DAT (Logan et al. 2007; Volkow et al. 2001). Abnormal DAT regulation and function correlated with ADHD in which DAT coding variant A559V was identified (Mazei-Robison & Blakely 2005; Mazei-Robison et al. 2008).

Decreased mesocorticolimbic DA transmission is one of the identified symptoms of depression (Nestler & Carlezon 2006). Some medications with proven antidepressant effects in humans (e.g., nomifensine, amineptine) are DA uptake inhibitors.



**Figure 2:** A, Pathway of biosynthesis of dopamine.; B, The fate of dopamine in a typical dopaminergic neuron after biosynthesis. Dopamine is stored in intracellular vesicles after uptake through the vesicular monoamine transporters (VMAT). After release, dopamine binds to dopamine receptors D1 and D2. Reuptake of excess dopamine is ensured with the dopamine transporter (DAT). Figure adapted from intechopen website.

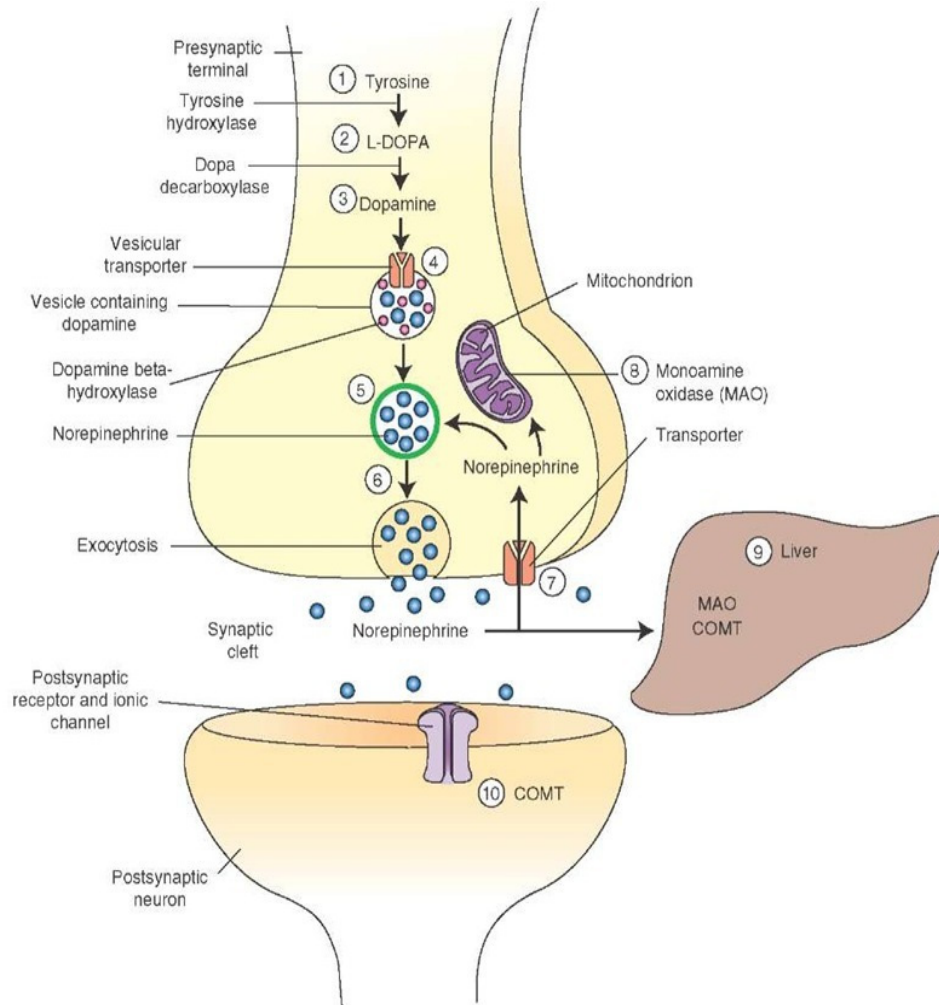
### 1.3.3. Norepinephrine transporter

Norepinephrine (NE), also known as noradrenaline, is the main neurotransmitter of the sympathetic nervous system (Zhou 2004; Schroeter et al. 2000).

It is taken up by the norepinephrine transporter (NET) which is encoded by the SLC6A2 gene in humans (Pacholczyk et al. 1991) and is located in chromosome 16q13-q21 (16q12.2) (Brüss et al. 1993).

It has been demonstrated that NET is selectively expressed on NE nerve terminals which enable temporal and spatial control of the activity of NE (Foote et al. 1983; Moore & Bloom 1979). Several studies have demonstrated that NET is also expressed in peripheral tissue (e.g., adrenal glands, placenta, vas deferens (Jayanthi et al. 2002; Schroeter et al. 2000; Sung et al. 2003). NE is also a target for psychostimulants such as cocaine and amphetamines (Jayanthi et al. 2002; Pacholczyk et al. 1991). Previous studies have found out that the modified NET function is associated with mood, cardiovascular disorders and attention (Esler et al. 2006; Haenisch et al. 2009; Hahn et al. 2008; Kim et al. 2006; Rumantir et al. 2000; Shannon et al. 2000). In addition to that, NE plays an essential role in human physiology and pathology such as mood and sleep regulation (Young & Landsberg 1998). Therefore, NET is used in the treatment of mood and cognitive disorders (Blier 2001; Bönisch & Brüss 2006).

Moreover, norepinephrine signalling is controlled by NET, thereby, a crucial homeostatic mechanism and its dysregulation via psychostimulants is thought to provide the neurochemical and behavioral effects of psychostimulants (Dipace et al. 2007) and chronic stress (Miner et al. 2006).



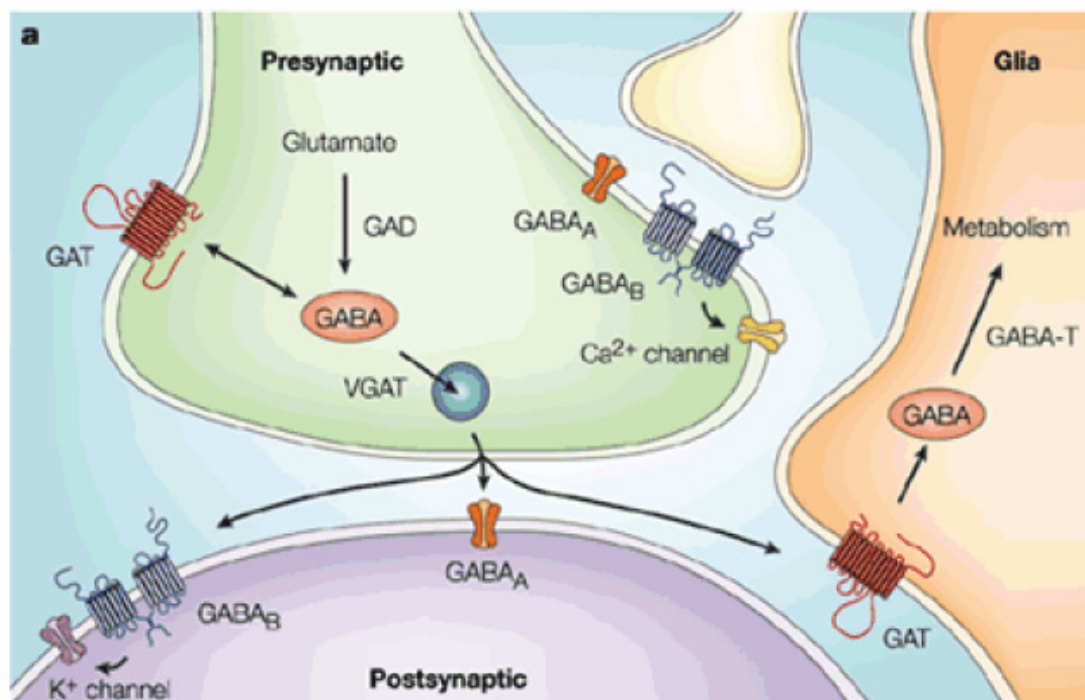
**Figure 3:** Synthesis and release of norepinephrine. COMT = catechol-O-methyltransferase (figure from what-when-how website)

#### 1.4. $\gamma$ -Aminobutyric acid (GABA) transporter

Gamma-aminobutyric acid (GABA) transporters are also members of the SLC6 family (Liu et al. 1993). GABA is stored in small vesicles and they are transported through the vesicular GABA transporter (VGAT). By way of fusion of the small vesicles to the presynaptic membrane, GABA is released into the synaptic cleft, where it can bind to GABA<sub>A</sub> or GABA<sub>B</sub> receptors (fig. 4). Afterwards, in surrounding glia cells or in the pre-synapse, where it is stored in vesicles or degraded via Succinic semialdehyde dehydrogenase (SSADH) and GABA-transaminase (GABA-T), it is taken up again quickly by GABA transporters, (Owens & Kriegstein 2002).

GABA is converted from glutamate by glutamic acid decarboxylase. It is the major inhibitory neurotransmitter in the mammalian central nervous system (Krnjević, 2004; Danbolt, 2001), whereas glutamate is the major excitatory neurotransmitter. Balance of GABAergic and

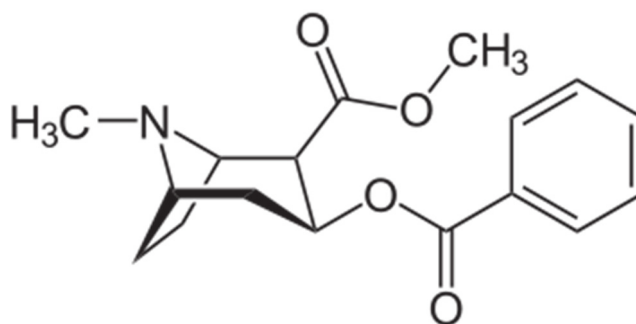
glutamatergic signals is crucial in the functioning of the central nerve system. Therefore, an inequality of these two neurotransmitter systems is presumably involved in many if not all central nerve system disorders to some scope, making the GABAergic system an applicable target for CNS drugs (Olsen 2002). Manipulation of the GABAergic neurotransmission is generally related to some disorders such as insomnia (Ebert et al. 2006), epilepsy (Galanopoulou 2010; Macdonald et al. 2010), and anxiety disorders (Kalueff & Nutt 2007). In addition to this, there is proof of involvement of this receptor in schizophrenia and some other neuropsychiatric disorders (Lewis et al. 2008; Möhler 2009; Vinkers et al. 2010). It is involved, directly or indirectly, in the most aspects of normal brain function such as memory, cognition, and learning, as well as in the development in of tolerance and addiction (Enoch 2008; Tan et al. 2010). Moreover, it may also prove beneficial to target GABA receptors in the treatment of neuropathic pain (Munro et al. 2009; Zeilhofer et al. 2009).



**Figure 4:** The structure of GABAergic synapse. GABA is synthesized from glutamate by the glutamic acid decarboxylase GAD, and stored in vesicles via transport through the vesicular GABA transporter (VGAT). Upon the depolarisation of the neuron, GABA is released into the synaptic cleft where it activates the GABA receptors GABA<sub>A</sub> and GABA<sub>B</sub>. Subsequently, it is taken up via the GABA transporter (GAT) into either glia cells or the presynaptic neuron. They are either restored in vesicles or degraded via the transaminase GABA-T to be further metabolised. Figure from Owens and Kriegstein, 2002.



### 1.5. Psychostimulant action of cocaine on monoamine transporters



**Figure 5:** Structure of Cocaine

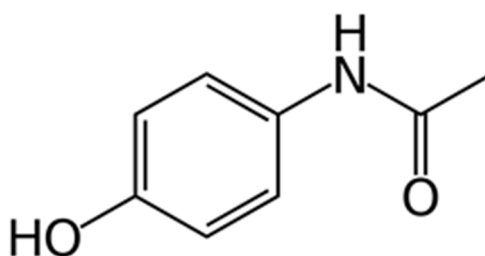
Cocaine (benzoylmethylecgonine, fig.5) is a tropane alkaloid obtained from the Andean plant *Erythroxylum coca*. It was used by indigenous people in South America for a long time and became popular all around the world by way of Sigmund Freud's reports in 1884 and introduction of Coca Cola in 1886 (Sulzer et al. 2005). Its usage as a local anaesthetic by Freud's suggestions and Koller's demonstration widely led to the discovery of its toxic effects (Y.A. et al., 2001).

In 1910, Alfred Fröhlich and Otto Loewi from the Institute of Pharmacology of Vienna were among the first who tried to shed light on the pharmacological role of cocaine (Sulzer et al. 2005).

Moreover, drugs of abuse such as cocaine and amphetamines target monoamine transporters. For DAT, NET and SERT, cocaine acts as a non-selective inhibitor with a relatively high affinity (Eshleman et al. 1999). It increases the synaptic concentration of all monoamines by stimulating the release and inhibiting the reuptake, triggering the cocaine reward especially through the increase of dopamine (Rothman et al., 2001)

For various monoamine transporters several cocaine analogues have been developed with both higher affinity and improved selectivity to treat cocaine addiction, of which some are still debated (Newman & Kulkarni 2002; Dutta et al. 2003; Loland et al. 2008).

### 1.6. Paracetamol



**Figure 6.** Chemical structure of paracetamol

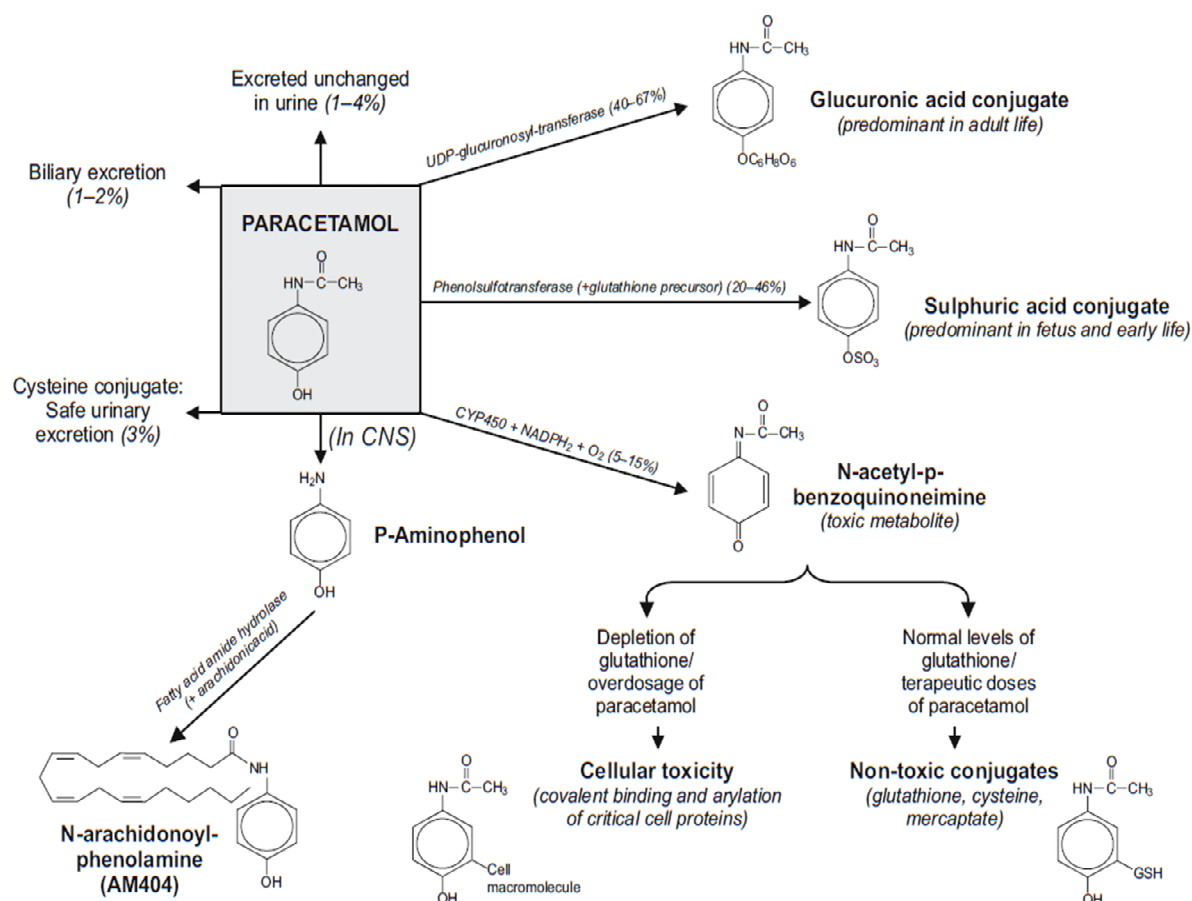
Paracetamol (Fig.6), also known as Acetaminophen in North America, derives from its chemical name para-acetylamino phenol (APAP) and acts antipyretic and analgesic. APAP intoxication is the most common cause of death in the United States (Bronstein et al. 2009).

This substance was first synthesised in Germany in 1878 by Morse and clinically used by Von Mering as an antipyretic in 1887 after the search for a less toxic compound than acetanilide, the only antipyretic and analgesic medication in that time (Bertolini et al. 2006).



In 1948 Brodie and Axelrod found paracetamol to be the major metabolite for the analgesic action of acetanilide (Brodie & Axelrod 1948). Since then, it has become one of the most commonly used analgesic and antipyretic drug (Anker & Smilkstein 1994; Meredith & Goulding 1980).

### 1.6.1. Pharmacokinetics or metabolism of acetaminophen

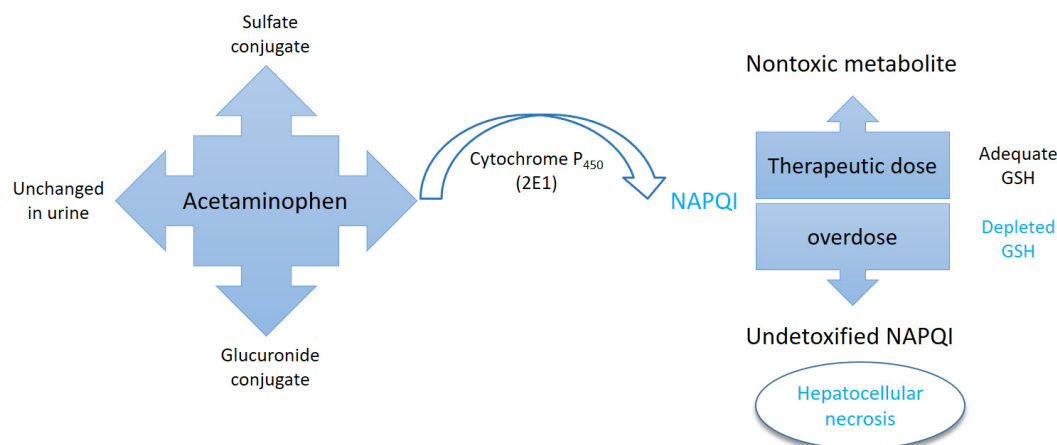


**Figure 7:** Metabolism of Paracetamol (figure from CNS Drug Reviews)

The main organ for the metabolism of paracetamol is the liver which eliminates approximately 25% of the therapeutic dose of paracetamol by first pass metabolism (Clements et al. 1978). In adults, APAP is safe at the therapeutic doses. Approximately 90% of paracetamol is conjugated with glucuronide (40-67%), sulphate (20-46%) and cysteine (3%) to inactive and harmless metabolites (fig. 7) (Bertolini et al. 2006). Around 5% is eliminated unaltered. The remaining 5% of APAP is oxidized via the cytochrome P450 (CYP450) (including CYP1A2, CYP3A4, and mainly CYP2E1) enzymes to the electrophilic and toxic metabolite N-acetyl-p-benzoquinoneimine, NAPQI (Nelson 1990). NAPQI is inactivated enzymatically via glutathione-S-transferases or non-enzymatically by binding glutathione, resulting in non-toxic cysteine or mercaptate conjugates, which are then eliminated (Miller et al. 1976).

The typical dose of paracetamol for the desired analgesic and antipyretic effects is 650-1000 mg for adults, which can be taken every 4 hours, up to a suggested maximum daily dose of 4 g. At therapeutic doses of acetaminophen, NAPQI is detoxified by glutathione and is excreted after

glucuronidation as the mercapturic acid into urine (Miller et al. 1976; Potter et al. 1985; James et al. 2003). After an overdose, glutathione is depleted and the increasing amount of NAPQI covalently binds to cysteine residues on proteins as 3-(cysteine-S-yl)-acetaminophen (APAP-CYS) (Streeter et al. 1984). When glutathione stores are depleted below a critical rate (almost 30% of normal stores) free NAPQI can attack nucleophilic sites on essential cellular macromolecules, inducing a series of events that may lead to cell death (Mitchell et al. 1973b) (fig. 7).



**Figure 8:** Toxicity mechanism of Acetaminophen

Hepatotoxicity involves factors such as increased frequency of paracetamol dosing, elongated or excessive dosing, increased capacity for P450 activation to NAPQI, decreased glutathione existence, or decreased sulfation and glucuronidation capacity. Another important factor of increased risk is chronic ethanol abuse (3 or more alcoholic drinks per day). Although, the combination of hepatic enzyme induction and glutathione depletion appears to increase paracetamol toxicity in chronic alcoholism, acute alcohol ingestion reduces toxic metabolic activation based on competitive inhibition and depletion of cytosolic NADPH (nicotinamide adenine dinucleotide phosphate) and therefore plays a preventive role in hepatotoxicity (Draganov et al. 2000; Teschke et al. 1979; Zimmerman & Maddrey 1995; Prescott 2000; Thummel et al. 1988; Tredger et al. 1985).

It was observed that simultaneous use of paracetamol and alcohol may increase the CYP2E1-mediated metabolism of paracetamol to NAPQI. In non-alcoholics, NAPQI is detoxified by conjugation with glutathione. In contrast, in alcoholics, the combination of CYP2E1 induction and glutathione depletion produce NAPQI accumulation (Draganov et al. 2000).

Some drugs such as carbamazepine, phenobarbital, phenytoin, primidone, and rifampicin in particular also induce cytochrome P450 enzymes (Miners et al. 1984) and may thus increase the formation of NAPQI.

### 1.6.2. Acetaminophen mechanism of toxicity

Although paracetamol is one of the safest analgesics when taken at the suggested therapeutic dose (Jackson et al, 1984), approximately 200 humans die every year of fulminant hepatic failure from APAP overdose (Rowden et al. 2006)

Nevertheless, paracetamol has been in clinical use for over 30 years and it is still unclear how the exact mechanism of acetaminophen toxicity happens (Shannon et al. 2009). Present literature implies N-acetyl-para-benzoquinone-imine (NAPQI) as the primary metabolite responsible for hepatotoxicity. The liver is the target organ for APAP toxicity because this is where it is detoxified.

Although NAPQI is toxic, it can be neutralized by conjugation with glutathione (GSH) (Jaeschke & Bajt 2006). However, in case of an overdose of acetaminophen, the generation of NAPQI exceeds the detoxification capacity of GSH, which leads to covalent binding of the sulfhydryl group to cysteines of cellular proteins (Cohen et al. 1997; Bond 2009).

Furthermore, N-Acetyl cysteine (NAC) is an acetylated cysteine residue that provides L-cysteine, the amino acid required for the biosynthesis of glutathione after being deacetylated in the liver (Viña et al. 1978). NAC is an important antidote used to treat overdoses of acetaminophen by leading to an increase in glutathione (GSH) levels, which protect against endogenously generated reactive oxygen species and reactive species such as NAPQI (Bessemers & Vermeulen 2001).

Animal experiments have shown that the administration of high dose of acetaminophen to mice causes centrilobular necrosis leading to death. Post administration of NAC effectively protects mice against this acetaminophen-induced hepatotoxicity (Williamson et al. 1982; Nagasawa et al. 1984; Roberts et al. 1987)

Recently, NAPQI has been shown to act on transient receptor potential (TRP) channels which are a family of non-selective cation permeable channels, permeable to calcium ions (Hofmann et al. 2000; Inoue 2005; Nilius & Voets 2005; Venkatachalam & Montell 2007).

TRPA1 is a unique sensor of noxious stimuli and a probable drug target for analgesics (Story et al. 2003; Bautista et al. 2006; Kwan et al. 2006; Jordt et al. 2004; Bandell et al. 2004).

Chemical activation of TRPA1 produces pain, and causes hyperreactivity and irritation in skin and visceral organs (Baraldi et al. 2010; Holzer 2011; Viana 2011). However, there are symptoms that indicate that TRPA1 is present in the spinal cord, and its role in spinal processing of nociceptive input is not certain (Kim et al. 2010).

The findings that TRPA1 is activated by electrophilic compounds (Jordt et al. 2004; Bandell et al. 2004) gave rise to speculate that compounds such as NAPQI and p-benzoquinone (p-BQ), p-AP and APAP indirectly activate TRPA1 on primary sensory neurons. The configuration of these electrophilic compounds in vivo is catalysed by a couple of enzymes, containing COX, cytochrome P450 (CYP450) monooxygenases, peroxidases, many of which are existing in the central nervous system (Chen et al. 2008; Dahlin et al. 1984; Prescott et al. 1981; Pascoe et al. 1988; Shinoda & Aoyama 2007).

TRPV1 activators are capable of producing spinal analgesia (Eimerl & Papir-Kricheli 1987) and TRPV1 is coexpressed with TRPA1 on a subpopulation of nociceptive sensory neurons (Fernandes et al., 2012). This led David Anderson and coworkers to investigate whether the analgesic effect of APAP could include the activation of TRPA1 expressed on the central terminals of primary afferent neurons. For this reason, they investigated the role of TRPA1 in the antinociceptive effect of systemic and intrathecal administration of APAP and identified whether its metabolites such as p-BQ and NAPQI, produce antinociception via interacting with TRPA1 in the spinal cord. Indeed, results showed that NAPQI and p-BQ are potent activators of TRPA1. This indicates that there is a new role of TRPA1 in synaptic signalling and that spinal TRPA1 activation through a non-antinociceptive mechanism promotes the analgesic activity of acetaminophen (Andersson et al. 2011).

## AIMS OF THE THESIS

NAPQI binds to the sulfhydryl group of glutathione at therapeutical doses of paracetamol. Once the glutathione stores of the body are emptied, it covalently binds to cysteine residues of proteins, leading to 3-(Cysteine-S-yl) adducts (Mitchell et al. 1973b; Jollow et al. 1973). The cysteine-modifying agent 2-(aminoethyl)-methanethiosulfonate hydrobromide (MTSEA) was reported to inhibit the function of SERT (Androutsellis-Theotokis et al., 2001), and cysteine mutants protected the transporter from MTSEA inhibition. Accordingly, we wondered whether NAPQI can similarly affect the function of the transporter and whether the effects can be prevented by mutations in cysteine residues of SERT.

Hence, the aims of this thesis hence were following:

1. Does NAPQI affect wildtype SERT in its uptake? If so, how does this change with cysteine mutants thereof?
2. If yes, is the effect
  - time- or concentration-dependent?
  - competitive or non-competitive?
3. Does NAPQI similarly affect other transporters such as the GABA transporter GAT?
4. Can we recapitulate the effects of MTSEA on SERT?

## 2. MATERIAL AND METHODS

### 2.1. Cloning

#### 2.1.1. Generating competent *E. coli* bacteria

Chemical competent *E. coli* bacteria were prepared as following: 10 ml LB medium (containing 10µg/ml tetracycline) was inoculated with a bacterial *E. coli* colony (XL10) from a tetracycline added agar plate. That preculture was incubated at 37°C overnight under continuous agitation, and 2ml of that was transferred into an autoclaved Erlenmeyer-flask including 200 ml prewarmed LB. The flask was incubated at 37°C until OD550 of 0.5 was reached, which indicates exponential growth of bacteria. Once the cells had reached this phase, they were placed them on ice and kept them there for 10 minutes. The bacteria were harvested by centrifugation in sterile centrifugation for 15 minutes (4°C 1500g). The supernatant was poured off and then supernatant and resuspended in 20 ml of ice-cold TSS buffer (Trypton 1%, Yeast Extract 0.5%, NaCl 100mM, Polyethylenglycol(PEG) 10%, DMSO 5%, MgCl<sub>2</sub> 50 mM, pH 6.5). Afterwards, 5 ml glycerol was added and 50µl of competent bacteria were distributed in sterile Eppendorf-tubes, immediately frozen in liquid nitrogen and stored at -80°C.

#### 2.1.2. QuikChange site-directed mutagenesis

Mutant transporters were generated by the QuikChange Site-Directed Mutagenesis kit from Agilent Technologies (Santa Clara, CA). The QuikChange Site-Directed Mutagenesis kit was used to make point mutations by replacing amino acids. The mutagenesis was performed by using two synthetic oligonucleotide complementary primers. Table 2 shows the amount of each component used for the PCR reaction.

Reagents	Volume
10× reaction buffer [(100mM KCl, 100mM(NH <sub>4</sub> ) <sub>2</sub> SO <sub>4</sub> , 200mM Tris-HCl (pH 8.8), 20mM MgSO <sub>4</sub> , 1%Triton® X-100, 1mg/ml nuclease-free BSA)]	2,5 µl
dsDNA template	5-50 ng
Quick Solution	0,75 µl
dNTP mix	0,5 µl
Primer	10 µl
PfuTurbo DNA polymerase (2.5 U/µl)	1 µl
ddH <sub>2</sub> O	Add 25 µl

**Table 2:** Quantities used for site-directed mutagenesis for each reaction. This procedure utilizes a double-stranded DNA (dsDNA) vector within localized gene of interest and two complementary primers containing the desired mutation.

A plasmid containing the gene of interest was used as a template. Both strands of the plasmid were amplified introducing the desired mutation by PCR (see table 3 for the PCR reaction).

Temperature (°C)	Time	Cycles
95	2 min	1x
95	20 sec	20x
58	20 sec	
68	4 min 30 sec	
72	5 min	1x
4	Endless	1x

**Table 3:** Quickchange PCR protocol

After the PCR reaction, the samples were treated with the endonuclease DpnI for 30 minutes at 37°C. The DpnI endonuclease which uses 5'-Gm6ATC-3' as a target sequence is specific for methylated and hemimethylated DNA. This enzyme was used to digest only the parental methylated and hemimethylated DNA template. As DNA from PCR reactions is not methylated, it is resistant to this enzyme. Therefore, only mutated DNA should stay unharmed.

Next, these mutated plasmids were transformed into competent bacteria and then they were grown over night on LB agar dishes which contained selected antibiotics appropriately to the antibiotic resistance gene on the plasmid.

### 2.1.3. Transformation

Transformation is a useful method to bring the desired plasmid into the bacterial cells in order to express the gene of interest. Plasmids were transformed into competent bacteria XL10. 100µl of competent bacteria cells (as described in section 8.1) were thawed on ice. After that 5µl of the DNA was added and gently mixed by flicking the bottom of the tube a few times. DNA/competent cell mixture was kept on ice for 20 minutes for incubation then a heat shock each transformation tube was given a heat shock of 42°C for 45 seconds. The time can be altered depending on the cell type (45 second is usually ideal, but this differs depending on the competent cells you are using). Following this, the tubes were put back on ice for 2 minutes. That procedure allowed plasmid DNA to enter the competent bacteria. 1 ml of LB (Lysogeny Broth) medium was added. Transformed bacteria were incubated for one hour at 37°C. During the incubation period bacteria expressed the plasmid-encoded gene for antibiotic resistance. Next, the bacteria were centrifuged for 5 minutes at 10,000 rpm (MiniSpin by Eppendorf), the supernatant was poured off and the pellet was resuspended gently. Then all of the transformation was spread on an LB agar plate containing the appropriate antibiotic. Lastly, bacteria were grown overnight in an incubator.

To obtain the best chance of getting single colonies, 50µL of transformation was added on one plate and the rest on a second plate (a method used in place of centrifuge).

#### **2.1.4. DNA/Plasmid isolation**

DNA was purified from liquid bacteria cultures (2ml), using NucleoSpin® Plasmid QuickPure Kit (Marcheray&Nagel) according to the manufacturer's protocol. After bacteria were lysed, both chromosomal DNA and protein were denaturated and precipitated, while the small bacterial DNA plasmids stayed in the solution and bound to a column in order to elute.

For qualitative analysis DNA was isolated from 2ml overnight LB cultures (Mini prep), using the NucleoSpin® Plasmid QuickPure Kit (Marcheray&Nagel) following the manufacturer's protocol. After this, lysis of bacteria, proteins and genomic DNA is precipitated and the plasmid of interest was bound to a column followed by elution later on.

DNAs obtained via "mini preps" were used for sequencing and diagnostic restrictions digest. In order to perform Midi Prep, proper plasmids were transformed and larger amounts of plasmid were isolated using 250 ml LB cultures.

### **2.2. Cell culture**

#### **2.2.1. Cultivation**

Products are required for cell culture (Media, Trypsin, serum and antibiotics) were purchased from Sigma-Aldrich, (St. Louis, MO), cell culture dishes were from Sarstedt (Germany).

HEK293 cells (The human embryonic kidney 293 cell line was used) were cultured in Dulbecco's modified Eagle's medium (DMEM; Sigma Aldrich) supplemented with High Glucose (4.5 g/L) containing 10% (w/v) fetal calf serum (Serum is an extremely complex solution of albumins, globulins, growth inhibitors, and growth promoters) and 1% Penicillin/Streptomycin (10,000 units penicillin/10mg/streptomycin/ml) on polystyrene culture dishes. Maintenance of stable cell lines: 50 µg/ml G418 was added to DMEM. Cells were kept in an incubator (CO2 Incubator CB, Binder) at 37°C in 5% CO2 conditions. CO2 was needed to control pH because cell physiology is highly sensitive to pH variations.

For this cultivation, the cells were incubated in an incubator overnight, then the medium was aspirated by use of a pump. In order to get rid of excess medium and dead unattached cells and cell debris, the cells were washed with 1x phosphate buffered saline (PBS; 1 mM MgCl2; 0, 1 mM CaCl2). In order to detach the cells from the culture dish, adherent cells had to be trypsinized, which is why 1ml of Trypsin/EDTA (0.5g trypsin/0.2g EDTA; Sigma-Aldrich) was added. The cells were observed under the microscope (generally for after about 3 to 5 minutes). When the cells pulled away from each other and rounded up, trypsin was neutralized by adding 8 ml of prewarmed cultivation medium. The dissociated cells were transferred into a sterile 50 ml centrifuge tube (Greiner) and harvested by centrifugation for 5 minutes at 1000 rpm. The neutralized dissociation solutions were aspirated from the cell pellet and the cells were resuspended in 8 mL fresh, pre-warmed growth medium.

#### **2.2.2. Cell counting/seeding**

As mentioned in section 3.2.1, cells were detached by adding Trypsin/EDTA solution (1ml per 10 ml cultivation medium), harvested by centrifugation and resuspended in fresh cultivation medium.

It was ensured that, the cell suspension to be counted was well mixed by either gentle agitation of the falcon tube or vortex. Before the cells had a chance to settle at all, cells were taken out of

approximately 10 to 15µl of the cell suspension and added to the object slide which consisted of 9 large squares which are subdivided into 16 smaller squares each.

Cells were counted by moving the hemocytometer to the next set of 16 corner squares and in this way counting was continued until all 16 corners were counted by using a hand tally counter. The average cell count was determined and multiplied by 10,000 (10<sup>4</sup>). The final value being the number of cells per ml in the original cell suspension.

### **2.2.3. Transfection**

#### **JetPrime® DNA transfection (Polyplus transfection)**

In this study, we used jetPRIME® as a transfection reagent which is well known for use in common experiments. HEK293 cells were seeded at 2.5x10<sup>6</sup> cells per 10cm dish 24h before transfection. The number of seeded cells depends on the cell type. Furthermore, cell confluence should be around 50% to 70% on the day of transfection to have high transfection efficiency. Appropriated amount of jetPrime® transfection reagent and DNA were added into the cells respectively and incubated for 4h. The medium of cells was changed 4h post-transfection and kept in an incubator overnight. The next day, the transfection efficiency was verified under the microscope and the cells were cultured for the two days following.

#### **Generation of stable cell lines**

The antibiotic G418 was used to select the cells that had integrated the plasmid coding for a selection marker and the gene of interest into their genome. The first step in this process was that cells were selected by 250µg/ml G418 for 10 days and the selection medium was changed several times to remove the dead cells. When resistant clones appeared, clones were picked and reseeded to facilitate growth. New cell lines were tested for functional expression of SERT by fluorescent microscopy as well as functional assays. Once generated and cultivated, to avoid the exclusion of the plasmid after a long cultivation period, cells that stably expressed the constructs were propagated in the medium containing 50µg/ml G418 per 10 ml.

## **2.3. Pharmacological Assays**

### **2.3.1. Uptake assay**

In order to test the functional activity of the serotonin transporter, a pharmacologically relevant uptake assay was performed using tritiated substances. Until recently, radioactively labeled compounds were used to measure serotonin, norepinephrine and dopamine transporter uptake. Following the standard protocol, transfected cells or stable cell lines (hSERT expression was induced by addition of tetracycline (TET/antibiotic) the day before) were seeded on Poly-D-Lysine (PDL purchased by Sigma-Aldrich) coated 48well cultivation dishes. 1x10<sup>5</sup> cells per well were seeded in a final volume of 500µl standard cultivation medium and incubated overnight at 37°C and 5% CO<sub>2</sub>.

The following day the cultivation medium was removed by aspiration and cells were covered with 500µl pre-warmed to Krebs-HEPES Buffer pH 7.3 at room temperature. Then cells were incubated at room temperature with 100µl of the uptake solution which contained a fixed concentration of the tritiated substance and different amounts of the unlabeled substance. In order to determine non-



specific uptake, cells were pre-incubated with 10 $\mu$ M paroxetine (for hSERT) for 5 min following incubation with [ $^3$ H] 5HT for 1 min at room temperature. And also in order to get specific uptake,  $K_m$  and  $V_{max}$  values were determined via unlabeled substrate which was used at increasing concentrations to dilute the specific activity of [ $^3$ H] 5HT. Thus, 0,2-100 $\mu$ M [ $^3$ H] 5HT (containing 0.2 $\mu$ M [ $^3$ H]) was added and incubated for 1min at RT.

Incubation time was dependent on the predicted transport rate of each transporter. After the incubation period, cells were washed with 500 $\mu$ l ice cold KHB (in order to stopped uptake) and immediately cells were lysed by adding 500 $\mu$ l 1% (w/v) SDS and the cell suspension was transferred to scintillation-vials and 2 ml of the scintillation cocktail (Rotisint<sup>®</sup> eco plus LSC-Universalcocktail; Roth) was added and counted in a beta counter. The scintillation cocktail absorbs emitted energy produced by the disintegration of the radioisotopes and re-emits it as flashlight. These signals can be detected and measured in a  $\beta$ -counter (TRI-CARD 2300 RT (Packard)).

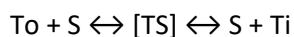
Data were analyzed by GraphPadPrism5.0 software.  $V_{max}$  and  $K_m$  values are calculated by non-linear regression curve fitting.

The  $K_m$  value or Michaelis-Menten constant is described as the substrate concentration at half-maximum enzyme velocity ( $V_{max}/2$ ). To characterize enzyme reactions, the  $K_m$  value defines half of the substrate binding sites to be occupied, which is in regard to the concentration of the substrate needed for performing a catalytic reaction (Berg et al. 2007). The reaction mechanism can be simply described as



where the concentrations of the enzyme E and of the substrate S are considered to transition to an intermediate complex of enzyme and substrate ES, which, through the activity of the enzyme, gives rise to the product P and the free enzyme (REFERENCE).

These kinetic parameters also describe the affinity of a substrate to a transporter. Therefore, a similar reaction can be applied, consisting of the outward-facing ( $T_o$ ) and the inward conformation ( $T_i$ ) of the transporter, resulting in:



Here, TS represents all intermediate conformations of the transporter from the bound state with substrate until the transition into the inside of the cell (REFERENCE).

### 2.3.2. Inhibition assay

The main purpose of this method is to analyze the potency of a putative antagonist to inhibit the transport of a substrate. It is characterized by the  $IC_{50}$  value that is defined as the concentration of the antagonist needed for 50% of inhibition of substrate transport.

Substrate concentration was kept constant and blockers were used at increasing concentrations according to expected  $IC_{50}$  values. After incubation reaction was stopped by washing the cells with ice cold uptake buffer, lysed by 1%SDS and cell suspension was transferred to scintillation vials and 2 ml of liquid scintillation cocktail was added and counted in a beta counter. In order to evaluate the data, GraphPadPrism<sup>®</sup> software was used. “% Uptake relative to control” values were plotted against logarithmic concentrations of NAPQI and analyzed by “One site competition” to calculate  $IC_{50}$  and  $K_i$  values.

## **2.4. Materials**

### **2.4.1. Standard solutions/buffers**

#### **Phosphata buffered saline (PBS)**

2.7 mM KCl, 1.5 mM KH<sub>2</sub>PO<sub>4</sub>, 137 mM NaCl, 4.3 mM Na<sub>2</sub>HPO<sub>4</sub> × 2H<sub>2</sub>O, (pH 7.3)

#### **TSS buffer**

Trypton 1%, Yeast Extract 0.5%, NaCl 100 mM, Polyethylenglycol (PEG) 10%, DMSO 5%, MgCl<sub>2</sub> 50 mM, pH 6.5

#### **Dulbecco's modified eagle's medium (DMEM; Sigma Aldrich)**

Supplemented with High Glucose (4.5 g/L) containing 10% (w/v) fetal calf serum (Serum is an extremely complex solution of albumins, globulins, growth inhibitors, and growth promoters) and 1% Pen Strep (10,000 units penicillin/10 mg/streptomycin/ml)

#### **Krebs-HEPES buffer (KHB)**

25 mM HEPES; 120 mM NaCl; 5 mM KCl (Merck); 1,2 mM CaCl<sub>2</sub> (Merck); 1,2 mM MgSO<sub>4</sub> (Merck); 5 mM D-glucose (Roth) (pH 7,3)

#### **Lysogeny Broth medium (LB)**

NaCl 10%, Peptone 10%, Yeast extract 5%, ddH<sub>2</sub>O appr. Vol.

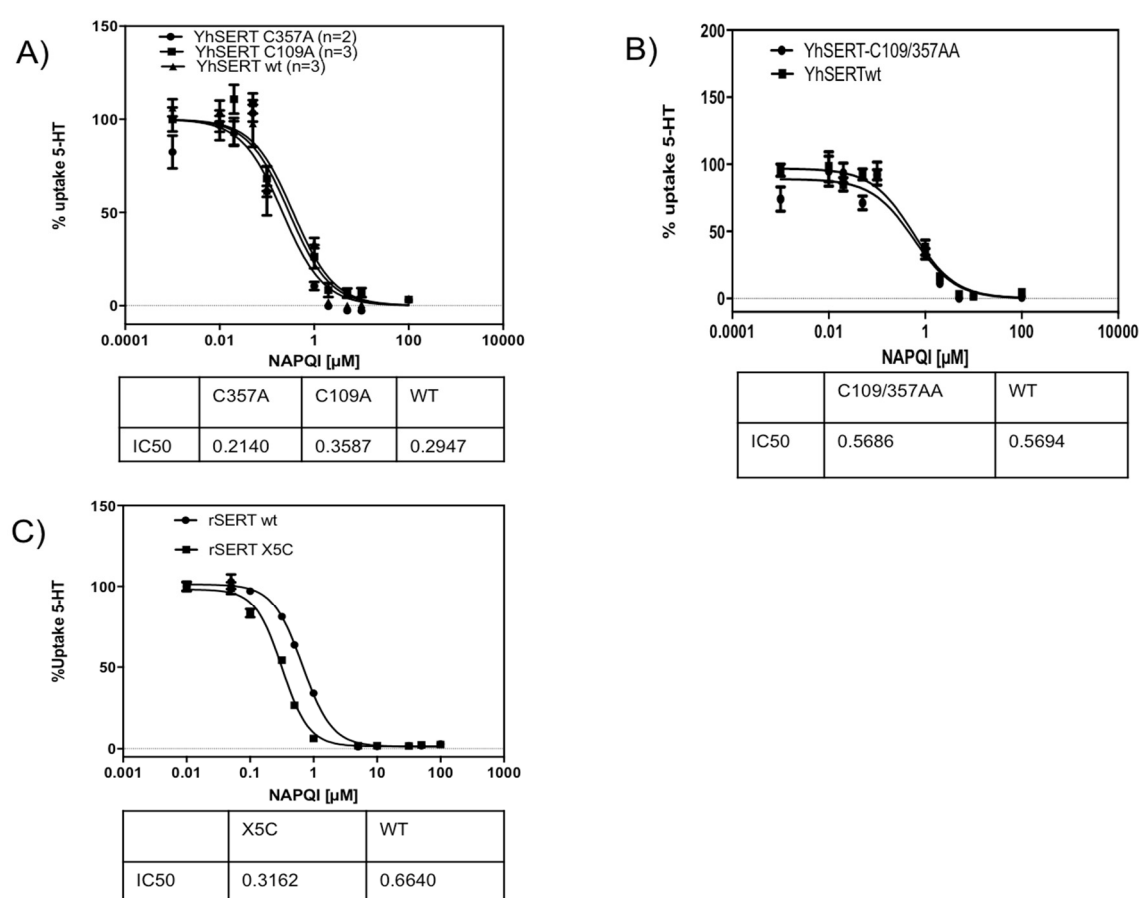
#### **5×Poly-D-lysine**

25 mg Poly-D-lysine (sigma), ddH<sub>2</sub>O ad. 100 ml

### 3. RESULTS

#### 3.1. Evaluation of the inhibitory capacity of NAPQI on different mutated serotonin transporters

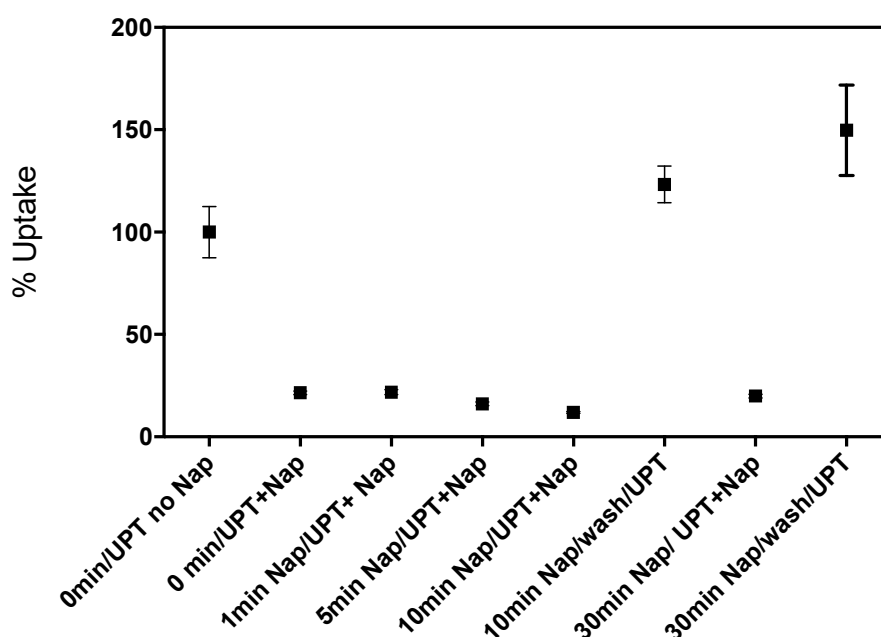
Serotonin transporter (SERT) comprises a single reactive external cysteine residue at position 109 (Chen et al. 1997) and seven predicted cytoplasmic cysteine residues. Therefore, I performed three inhibition uptake assay with alanine mutants of the cysteine residues C109 and C357 and a double alanine mutant thereof. In addition, I used the quintuple cysteine mutant of the rat serotonin transporter C15A/C21A/C109A/C357I/C622A that was formerly described by the lab of Gary Rudnick for the sensitivity to another cysteine-modifying agent (Androutsellis-Theotokis et al. 2001). Figure 8 shows that neither mutation of single cysteine residues nor the multiple mutants had any effect on inhibitory potency of NAPQI.



**Figure 8:** Evaluation of inhibition capacity of NAPQI on different mutated serotonin transporters. Inhibition of [ $^3$ H] 5-HT uptake by NAPQI was determined in HEK293 cells transiently transfected with wild type and mutant cDNAs respectively using jetPrime (as described under Materials and Methods). A) Single mutant C357A, C109A B) double mutant C109/C357AA and C) mutant rat SERT. Cells were incubated with 0.1 $\mu$ M [ $^3$ H] 5-HT and increasing concentration of NAPQI and background was measured at 10 $\mu$ M paroxetine. Data, plotted according to the sigmoidal model. Mean and SEM from a representative experiment run in triplicate are shown.

### 3.2. Time dependent uptake inhibition assay

SERT-expressing cells were pre-incubated with 1 $\mu$ M NAPQI for various times. Uptake inhibition was then performed in the presence or absence of NAPQI. Even without preincubation NAPQI reduced the uptake of 5-HT by 70% (Fig.10 0min/UPT+Nap). Preincubation for longer periods up to 30 min did only result in a slightly higher reduction of uptake (max 80%). Covalent modification of a protein would result in an effect, which could be observed even if the modifying drug is removed. When we exchanged the preincubation buffer with buffer containing no NAPQI, uptake was immediately back to almost 100%. This clearly indicates that NAPQI does not covalently modify the transporter but rather inhibits uptake in a reversible manner.



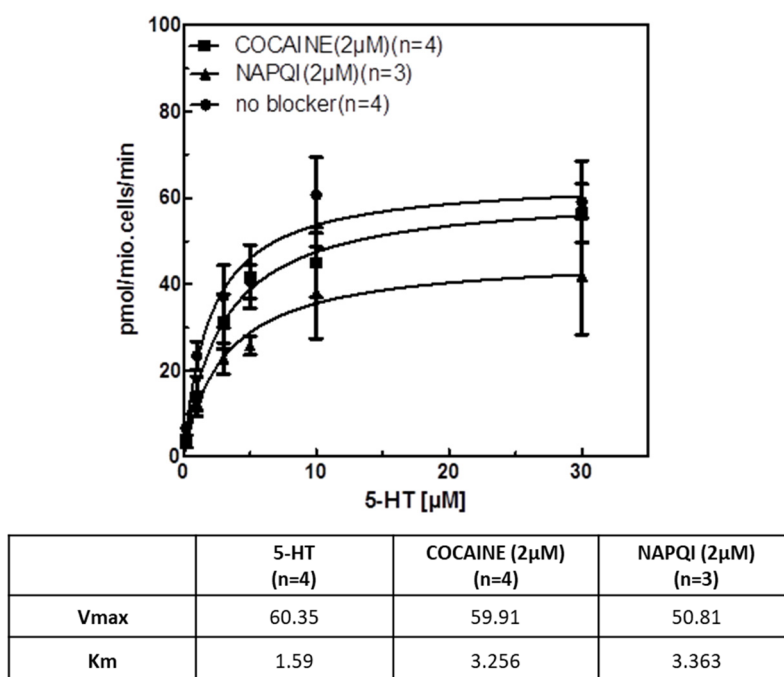
**Figure 10:** Time dependent uptake inhibition assay. Assay was performed in transiently transfected HEK293 cells and cells were exposed to 1 $\mu$ M NAPQI for different time points (0/1/5/10/30 minutes) and various washing steps. Uptake incubation was performed in the presence or absence of NAPQI. Samples were plotted % versus time [min]

### 3.3. NAPQI is a competitive inhibitor of uptake of SERT

Our hypothesis was that NAPQI binds to transporters irreversibly and modifies them, inhibiting the function of the transporter. However, the previous results showed that NAPQI binds to the transporters reversibly. Thereby, we wondered whether NAPQI inhibits the transporters in a competitive or non-competitive manner. In order to answer this question, we performed an uptake assay using increasing concentrations of substrate in the absence or presence of cocaine or NAPQI at indicated concentrations. To investigate whether NAPQI is a competitive inhibitor, we used cocaine because binding studies of cocaine, imipramine, and the SSRIs paroxetine and citalopram imply the presence of more than one binding site on the SERT (REF). Cocaine, imipramine and citalopram act as competitive inhibitors of 5-HT reuptake (Sur et al. 1998).

If the uptake reaches the same  $V_{max}$  (maximum velocity) in the presence and absence of the inhibitor, while the value of  $K_m$  (Michaelis-Menten constant) increases, this is a sign for competitive

enzyme inhibition. If  $V_{max}$  is decreased while  $K_m$  is not altered this would point to a non-competitive inhibition. In order to check if NAPQI is competitive inhibitor, saturation uptake was performed. As expected, cocaine inhibits the uptake of 5-HT in a competitive way. NAPQI interestingly shifted the  $K_m$  value to the right but also reduced  $V_{max}$  (Fig. 11).



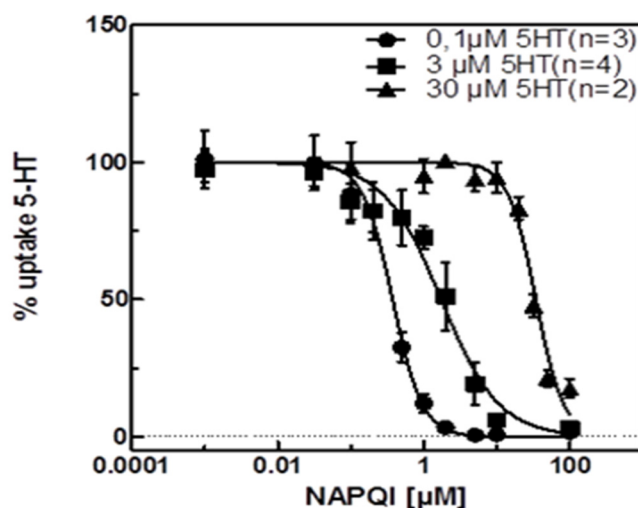
**Figure 11:** Saturation uptake of serotonin in the presence or absence of cocaine. Uptake experiments were performed using increasing concentrations of substrate (0.2, 1, 3, 5, 10, 30, 100μM of [ $^3$ H] 5-HT) in the absence or presence of cocaine or NAPQI, each 2 μM. Cells were transfected with wild-type SERT. Cells were incubated with 0.2 μM [ $^3$ H] 5-HT, in the presence or absence of 10μM paroxetine to determine nonspecific uptake. Table below shows calculated  $V_{max}$  and  $K_m$  for each fit.

### 3.4. Increase of serotonin shifts the inhibition curve for NAPQI

The results in figure 11 point towards a mixed competitive inhibition. In order to investigate this in detail, we performed the experiments in a different way where we increased substrate concentrations and tested for a potential shift in the  $IC_{50}$  values of NAPQI.

In the Cheng-Prusoff Equation " $K_i = IC_{50} \div (1 + [S] / K_m)$ "  $K_i$  is the inhibition constant for a drug where 50% of receptors are occupied, while the  $IC_{50}$  value defines the experimentally found value where 50 % of the enzymatic reaction (uptake) is inhibited.  $[S]$  defines the concentration of the substrate (5-HT) used while  $K_m$  is the affinity of the ligand. This means that  $IC_{50}$  is dependent on  $K_m$  and the ligand concentration.

As  $IC_{50} = K_i * (1 + [S]/K_m)$  and in most experiments the ligand is used below its  $K_m$  value, this infers that  $IC_{50} \sim K_i$ . If the concentration of the ligand is now increased,  $IC_{50}$  will increase as  $K_m$  and  $K_i$  are constant values. The  $K_m$  value for 5-HT was determined as 1.6 μM (Fig 12).

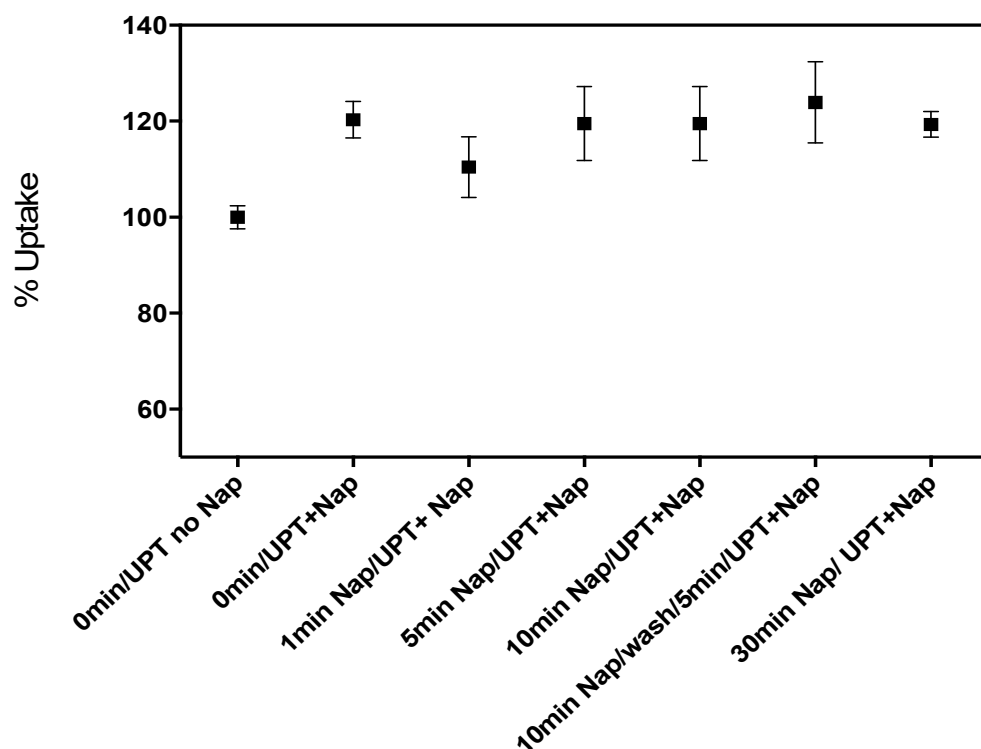


	0.1 $\mu$ M 5-HT (n=3)	3 $\mu$ M 5-HT (n=4)	30 $\mu$ M 5-HT (n=2)
IC <sub>50</sub>	0.3729	1.732	34.25

**Figure 12:** Uptake inhibition experiments in the presence of increasing serotonin concentrations. Experiments were performed using 0, 1 $\mu$ M, 1 $\mu$ M and 10 $\mu$ M serotonin on SERT with increasing concentrations of NAPQI. Background was measured at 10 $\mu$ M paroxetine. Data, plotted according to the hyperbolic model, are shown as means of a representative experiment carried out in triplicates. Table below indicates the calculated IC<sub>50</sub> values for each 5-HT concentration.

### 3.5. Effects of NAPQI on other transporters: GABA

Gamma-aminobutyric acid (GABA) and glutamate are the major inhibitory and excitatory neurotransmitters in the mammalian central nervous system (Krnjević 2004; Danbolt 2001). The reuptake transporter especially for GABA is target of various therapeutically used drugs. We wanted to test whether NAPQI inhibits uptake of not only monoamines but also of other neurotransmitters. As the exact mechanism of paracetamol is not identified yet, involvement of the GABAergic system could still be a possible explanation, as targeting of GABA receptors shows benefits in the treatment of neuropathic pain (Munro et al. 2009; Zeilhofer et al. 2009). However, even at a high concentration of NAPQI (30 $\mu$ M) for increasing time periods no reduction of GABA uptake could be observed (fig. 13).



**Figure 13:** Uptake inhibition of the GABA transporter. Time-dependent uptake inhibition of HEK293 cells stably expressing rat GAT at different time points (0/1/5/10/30 minutes) and a washing step (wash). Uptake inhibition was performed in the absence or presence of 30  $\mu$ M NAPQI (NAP). 10  $\mu$ M Tiagabine was used to block GAT. Samples were plotted as % uptake versus time in minutes.

## 4. DISCUSSION

### 4.1. Evaluation of inhibition capacity of NAPQI on different mutated serotonin transporters

As mentioned before, NAPQI binds to the Sulfhydryl group of glutathione under therapeutical doses of Paracetamol. In case of taking overdose of Paracetamol, glutathione stores are emptied and thus NAPQI binds sulfhydryl groups on cysteine residues of other proteins (Mitchell et al. 1973a; Jollow et al. 1973). Binding of NAPQI to TRPA1 showed functional alterations of the channel (Andersson et al. 2011). As we found out that NAPQI inhibits SERT the most logical explanation would have been a similar covalent modification. The serotonin transporter consists of a single reactive external cysteine residue at position 109 and 7 predicted cytoplasmic cysteines (Chen et al. 1997). Furthermore, some cysteine residues have been indicated to be important for expression of SERT (Sur et al. 1997). Previous experiments by Jae-Won Yang (institute of pharmacology) have shown using mass spectrometry that at least a rather C-terminally located cysteine (C622) can be modified by NAPQI (data not shown). As it was shown before, deletion of the last 16 amino acids of SERT does not affect its function (El Kasaby et al 2010). Therefore, the modification of cysteine 622 will most likely not be responsible for the inhibition of SERT by NAPQI. Based on this, we performed uptake inhibition experiments to investigate inhibition effect of NAPQI on serotonin transporter mutants. However, as shown in Fig.8, inhibition effect of NAPQI still continued although the cysteine at position 109 were not present. According to Andreas Androutsellis's publication, his outcomes

indicated that when 2 cysteine residues were mutated at position 109 and 357, in contrast to wild type SERT, the mutants are resistant against the inhibition effect of the cocaine analogue MTSEA (2-(aminoethyl)methanethiosulfonate hydrobromide) (Androutsellis-Theotokis et al. 2001). Therefore, we mutated two cysteine residues at position C109 and C357 and after that 5 cysteines from rat SERT which were shown to react with other cysteine-reactive compounds. On the contrary, as shown in Fig.8, our mutated cysteine residues did not affect the susceptibility of SERT to NAPQI. Therefore, we can conclude that modification of these residues is not the mechanism underlying the inhibition of SERT by NAPQI.

#### **4.2. Time dependent uptake inhibition assay**

To find out whether or not NAPQI covalently binds to SERT, we performed uptake inhibition experiment in the presence or absence of NAPQI. When we incubated our cells with NAPQI and then washed it away, the transport capacity came back immediately (Fig 10). Taken this result, together with the fact that removing accessible cysteine residues by mutagenesis did not change the inhibitory profile of NAPQI, one can clearly state that NAPQI inhibits the 5-HT uptake in a reversible way. A similar effect of NAPQI was observed for the uptake of dopamine via DAT while the uptake of MPP<sup>+</sup> via the norepinephrine transporter NET was not altered at all (data not shown). These experiments were performed parallel to my thesis by other colleagues at the institute of pharmacology.

#### **4.3. NAPQI is a competitive inhibitor of SERT**

However, we explored that NAPQI reversibly inhibits SERT, we performed the saturation uptake of serotonin in the presence or absence of an inhibitor. In a competitive inhibition, the competitive inhibitor shows similarity to the substrate and it binds to the active site of the protein and substrate and inhibitor cannot bind to the active site at the same time. Therefore, the substrate is prevented from binding to the same active site of enzyme. Competitive inhibition is characterised by competitive inhibitors that can be overcome by an increasing the amount of substrate. Although competitive inhibitor could change the  $K_m$  value, an enzyme will have the same  $V_{max}$  as in the presence or absence of an inhibitor (Berg et al. 2002).

In order to examine whether or not NAPQI is a competitive inhibitor we used cocaine. Because cocaine, imipramine and citalopram behave like competitive inhibitors of 5-HT reuptake (Sur et al. 1998).

As shown in figure 11,  $V_{max}$  of serotonin in the presence and absence of cocaine and NAPQI were similar: in the presence of NAPQI,  $K_m$  of serotonin was 2-fold increased and shifted to the right like cocaine. This could point towards a mixed competitive inhibition.

#### **4.4. Increasing of serotonin shifts the inhibition curve for NAPQI**

In accordance with competitive inhibition, and because an inhibitor can act like a substrate and bind to the active site of the enzyme, it prevents binding of the substrate, and more substrate is used to compete with inhibitor. Thus, the concentration of a drug which displaces 50% of the specific binding of the substrate ( $IC_{50}$ ) may vary between experiments by experimental conditions like substrate or inhibitor concentration. Further, via using Cheng-Prusoff equation (Cheng & Prusoff 1973), the  $IC_{50}$  value is converted to the inhibition constant ( $K_i$ ) and this quantitative measure shows how much of drug or inhibitor is needed to inhibit the transporter. Due to this reason, we



performed uptake inhibition experiment to see whether or not serotonin and NAPQI compete with each other. The experimental data extracted from figure 12 show that the expected values are roughly correct for 0.1 and 3 $\mu$ M. The experimental IC<sub>50</sub> determined at 30 $\mu$ M 5-HT is too high. This could have been because of the concentrations of NAPQI not being higher than 100 $\mu$ M. This was because of its limitation of solubility and the data points being inaccurate in this region. Slight reduction of 5-HT might help to circumvent this problem and would be a possibility to check, but was not performed during my thesis.

When we calculate our IC<sub>50</sub> values using Cheng-Prusoff equation which was shown in Result section, it indicates that, in case of increasing concentrations of serotonin, IC<sub>50</sub> value of serotonin becomes larger, although the concentration of NAPQI was in a micromolar range. Thus, these results again demonstrate that NAPQI is a competitive inhibitor.

We conclude that NAPQI, the toxic metabolite of Paracetamol, is capable of blocking SERT in a reversible competitive way. Blockage of SERT would increase the analgesic effect of descending antinociceptive neurons, providing a new perspective for the analgesic mechanism of Paracetamol.

Therefore, we propose the inhibition of SERT as a putative mechanism for the analgesic action of paracetamol. Although paracetamol is in use as an analgesic drug for decades, the precise mechanism of its action remains unclear. One explanation points to the serotonergic system to be involved in this mechanism (Karandikar et al., 2016; Tjølsen et al., 1991; Pini et al., 1996). Descending serotonergic neurons are known to play an important role in the inhibition of pain sensation. Given the possibility that paracetamol at therapeutical doses can be transformed to NAPQI in the nervous system as proposed by Andersson et al (Andersson et al. 2011) it could reach local concentrations high enough to block 5-HT reuptake into synapses. This would result in an elevated and prolonged 5-HT concentration in the synaptic cleft, thereby increasing the antinociceptive action of 5-HT. A similar effect is the reason why antidepressants show antinociceptive properties (Mochizuki 2004; Richeimer et al. 1997; Micó et al. 2006). Although it has to be proven that paracetamol really increases 5-HT concentrations in the nervous system and therefore NAPQI might be the active component responsible for the antinociceptive action, the experiments performed within this thesis might provide the first experimental data to explain the mode of action of paracetamol.

#### **4.5. Effects of NAPQI on the GABA transporter**

GABA is an inhibitory neurotransmitter and localized both presynaptically in primary afferents and postsynaptically in dorsal horn interneurons in the mammalian spinal cord (Malcangio & Bowery 1996).

The GABAergic system plays an outstanding role in presynaptic inhibition of primary afferents, therefore influencing sensory transmission, nociception, and motor activity on both pre- and postsynaptic levels (Barker & Nicoll 1972; Polc 1982; Cattaert et al. 1992; Stuart & Redman 1992).

Due to these features of GABA, we wanted to investigate whether NAPQI has an effect on this transporter. Our uptake inhibition experiments indicate that NAPQI has no effect on the GABA transporter (GAT). Human GAT has 38% homology with human SERT (uniprot alignment) and, compared to SERT, has less cysteine and histidine residues in the extracellular and intracellular loops.

## 5. LIST OF FIGURES

Figure number	Source <sup>1</sup>
Figure 1	<a href="http://what-when-how.com/neuroscience/neurotransmitters-the-neuron-part-4/">http://what-when-how.com/neuroscience/neurotransmitters-the-neuron-part-4/</a>
Figure 2	<a href="http://www.intechopen.com/books/a-synopsis-of-parkinson-s-disease/pathophysiology-of-l-dopa-induced-dyskinesia-changes-in-d1-d3-receptors-and-their-signaling-pathway">http://www.intechopen.com/books/a-synopsis-of-parkinson-s-disease/pathophysiology-of-l-dopa-induced-dyskinesia-changes-in-d1-d3-receptors-and-their-signaling-pathway</a>
Figure 3	<a href="http://what-when-how.com/neuroscience/neurotransmitters-the-neuron-part-4/">http://what-when-how.com/neuroscience/neurotransmitters-the-neuron-part-4/</a>
Figure 4	Owens and Kriegstein, 2002
Figure 5	<a href="https://en.wikipedia.org/wiki/Cocaine">https://en.wikipedia.org/wiki/Cocaine</a>
Figure 6	Self-designed
Figure 7	CNS Drug Reviews, Vol.12, No.3-4, 2006
Figure 8	<a href="https://en.wikipedia.org/wiki/Paracetamol">https://en.wikipedia.org/wiki/Paracetamol</a>

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<sup>1</sup> as of July 2019

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