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# The subthalamic nucleus in motor and affective functions

An optogenetic in vivo-investigation

**ADRIANE GUILLAUMIN** 





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#### Abstract

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The basal ganglia form a group of subcortical interconnected nuclei involved in motor, limbic and cognitive functions. According to the classical model of the basal ganglia, two main pathways exert opposing control over movement, one facilitating movement and the other suppressing movement. The subthalamic nucleus (STN) plays a critical role in this function, and has also been implicated in reward processing. Despite ample knowledge of the role of the STN in motor dysfunctions in relation to Parkinson's disease, less is known about STN's natural role in healthy subjects.

The studies described in this thesis aimed to address the functional role of the STN in its natural neurocircuitry by using a transgenic mouse line which expresses Cre recombinase under the *Pitx2* promoter. The *Pitx2* gene is restricted to the STN and the use of Pitx2-Cre mice thereby allows selective manipulation of STN neurons by using optogenetics. By expressing Channelrhodopsin (ChR2) or Archaerhodopsin (Arch) in Pitx2-Cre neurons, we could optogenetically excite or inhibit STN Pitx2-Cre neurons and investigate the role of the STN in motor and affective functions. We showed that optogenetic inhibition and excitation of the STN induce opposite effects on motor activity. STN excitation reduced locomotion while STN inhibition enhanced locomotion, thereby providing experimental evidence to classical motor models postulating this role. We also showed that optogenetic excitation of the STN induces potent place avoidance, a behaviour relevant to aversion. Projections from the STN to the ventral pallidum (VP) exist that when excited induced the same behaviour. The VP projects to the lateral habenula (LHb), a structure known for its role in aversion. A glutamatergic multisynaptic connection between the STN and the LHb was confirmed.

Aversive behaviour is also mediated by the hypothalamic-mesencephalic area. The *Trpv1* gene is expressed within the posterior hypothalamus. By applying optogenetics in a Trpv1-Cre mouse line, projection patterns to limbic brain areas were identified, and optogenetic excitation of Trpv1-Cre neurons was found to induce place avoidance.

The STN and posterior hypothalamus are thereby demonstrated as new players in the aversion neurocircuitry, while the long-assumed role of the STN in motor behaviour is confirmed. To enable future analyses of how STN manipulation might rescue motor and affective deficiency relevant to human disorders, a neuronal degeneration mouse model was generated.

To conclude, the results presented in this thesis contribute to enhanced neurobiological understanding of the role played by the STN in motor and affective functions.

Keywords: Subthalamic nucleus, optogenetics, basal ganglia, locomotion, Parkinson's disease, aversion

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"There is a single lit everywhere."  — Isaac Asimov	light of science,	and to bright	en it anywhere	e is to brighten
				$\grave{A}$ ma famille,

## List of Papers

This thesis is based on the following papers, which are referred to in the text by their Roman numerals.

- Guillaumin, A., Serra, G.P., Georges, F., Wallén-Mackenzie, Å. (2020): Optogenetic investigation into the role of the subthalamic nucleus in motor control.
   BioRxiv. doi: 10.1101/2020.07.08.193359.
   Submitted to journal.
- II Serra, G.P., Guillaumin, A., Baufreton, J., Georges F., Wallén-Mackenzie Å. (2020): Aversion encoded in the subthalamic nucleus.

**BioRxiv**. doi: 10.1101/2020.07.09.195610. *Submitted to journal*.

- III Guillaumin A., Vlcek B., Dumas S., Serra G.P., Wallén-Mackenzie Å. (2020): Anatomical-functional analysis of the spatially restricted Transient receptor vanilloid-1 (Trpv1)-positive domain within the medial hypothalamic-mesencephalic area.

  Manuscript.
- IV Guillaumin A. & Wallén-Mackenzie Å. (2020): Optimization protocol for the 6-OHDA model of Parkinson's disease in wild-type mice Manuscript.

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## **Abbreviations**

**6-OHDA** 6-hydroxydopamine

μg microgram microliter

**AAV** adeno-associated virus

**Aldh1a1** Aldehyde dehydrogenase 1 family, member A1

AP antero-posterior Arch Archaerhodopsin

**BNST** bed nucleus of the stria terminalis

Calb1 Calbindin 1

ChR2 Channelrhodopsin 2
Cre Cre recombinase
EP entopeduncular nucleus

**eYFP** enhanced yellow fluorescent protein

**FA** formaldehyde

f/f flanked by floxed sites
D1R dopamine receptor subtype 1
D2R dopamine receptor subtype 2

**DA** dopamine

**DBS** deep brain stimulation

DIG digoxigenin DV dorso-ventral

**EF1a** human elongation factor 1alpha promoter

EP entopeduncular nucleus GABA gamma-aminobutyric acid

**GP** globus pallidus

GPi globus pallidus interna
Grp Gastrin-releasing peptide
HFS high-frequency stimulation
IF interfascicular nucleus

**IMAO-B** inhibitor of the monoamine oxidase B

IPF interpeduncular fossa
LFS low-frequency stimulation
LHA lateral hypothalamic area

LHb lateral habenula MAB maleate buffer

**mAcbSh** medial part of the nucleus accumbens shell

MFB median forebrain bundle

mg milligram

mlmedial lemniscusMLmedio-lateralmmmillimeter

**MM** mammillary bodies

MnM medial mammillary nucleus, median part

**mPFC** medial prefrontal cortex

**MPTP** 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine

MSN medium spiny neurons NAc nucleus accumbens

nl nanoliter nm nanometer

PAG periaqueductal gray matter PBP parabrachial pigmented nucleus

PBS phosphate buffer saline
PCR polymerase chain reaction
PD Parkinson's disease
PFA paraformaldehyde

PIF parainterfascicular nucleus

Pitx2 Paired-like homeodomain 2 transcription factor

PH posterior hypothalamus PHA posterior hypothalamic area

PN paranigral nucleus
pSTN parasubthalamic nucleus
RLi rostral linear nucleus
RM retromammillary nucleus

**RMM** retromammillary nucleus, medial part **RML** retromammillary nucleus, lateral part

RMTg rostromedial tegmental nucleus
SNr substantia nigra pars reticulata
SNc substantia nigra pars compacta
SSC saline-sodium citrate buffer

STN subthalamic nucleus SuM supramammillary nucleus

tg transgenic allele

**Trpv1** Transient receptor potential cation channel subfamily V

member 1 or vanilloid receptor 1

Vglut2 Vesicular glutamate transporter 2 Vglut3 Vesicular glutamate transporter 3

VP ventral pallidum VTA ventral tegmental area

wt wild-type allele ZI zona incerta

#### Introduction

All along our existence, we as human beings, but also any living animal, make decisions that will shape and orientate the course of our life. Those decisions are the final results of complex processes related to positive and negative emotions, motivation and memories, and will be implemented by following an adequate selection of actions to reach a goal. For example, our behaviour will change and adapt depending on sensory stimuli we perceive in different contexts: It is -20°C outside, drinking a warm tea by a fireplace would sound very pleasant and a series of actions will be implemented to reach this goal. However, if it was 35°C outside, our behaviour will adapt in a very different manner and we would probably end up in a swimming pool instead. Those behaviours, which include affective, cognitive and motor (executive) functions, are rendered possible by an assembly of subcortical nuclei called the basal ganglia that are connected to the cerebral cortex to form top-down control loops. Among those interconnected nuclei, the subthalamic nucleus (STN) plays an important and central role as an excitatory input to the GABA nuclei of the basal ganglia. In this thesis, studies have been performed in order to identify and analyse the role of the STN in two of the main functions of the basal ganglia: Motor and affective functions.

#### The basal ganglia

The basal ganglia are a group of interconnected subcortical nuclei: The striatum (caudate and putamen), pallidum, substantia nigra (SN) and the STN. The major input to the basal ganglia is the cerebral cortex which projects to the striatum and STN, while the internal part of the pallidum and the substantia nigra *pars reticulata* (SNr) are considered the basal ganglia output structures and project to the thalamus. The basal ganglia consists of three parallel loops corresponding to their role in affective, associative and motor functions (Alexander, Crutcher and DeLong, 1990). The motor loop is the best known and most well studied loop of the three while fewer studies have investigated the affective (or limbic) and associative loops. Accordingly, the

basal ganglia are mostly known for their role in movement-related behaviours, and their disorders.

#### The motor loop

The motor loop is necessary for executing movements. The cortex sends glutamatergic projections to the medium spiny neurons (MSNs) of the dorsal striatum. The dorsal striatum is primarily a GABAergic structure which, in addition to excitatory cortical input, is modulated by dopamine (DA) projections from the substantia nigra *pars compacta* (SNc). Via the striatum, two pathways function in parallel with opposite effects on movement: 1) *The direct pathway* which starts from MSNs in the striatum expressing the post-synaptic DA receptor D1 (D1R) that project directly to the output structures of the basal ganglia, the SNr and the globus pallidus *interna* (GPi, or EP for entopeduncular nucleus in rodents). This GABAergic pathway inhibits the SNr/EP which leads to disinhibition of the ventro-lateral thalamus, and consequently, to promotion of movement; and, 2) *The indirect pathway* which is initiated from MSNs expressing the post-synaptic DA D2 receptor (D2R) and projecting to the globus pallidus *externa* (GPe, or GP in rodents). The GP is a GABAergic nucleus which sends strong inhibitory projections to the

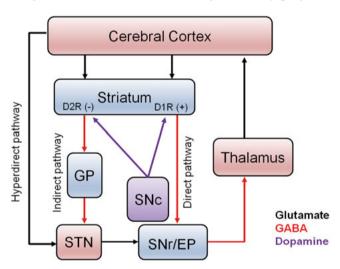


Figure 1: Simplified representation of the basal ganglia circuitry with the direct, indirect and hyperdirect pathways. Excitatory structures are in red, inhibitory structures in blue and modulatory structures in purple. GP: globus pallidus; STN: subthalamic nucleus; SNc: substantia nigra pars compacta; SNr: substantia nigra pars reticulata; EP: entopeduncular nucleus; D1R: dopamine receptor D1; D2R: dopamine receptor D2.

STN. MSNs of the indirect pathway inhibit the GP which leads to the disinhibition of the STN, an excitatory nucleus. The STN excites the output structures of the basal ganglia (SNr/EP) which in turn inhibit the ventro-lateral thalamus and consequently suppresses unwanted movements. In addition to the direct and indirect pathways, a third pathway, *The hyperdirect pathway*, by-passes the striatum and serves as a fast stopping pathway. In this pathway, the STN receives direct projections from the cortex.

The direct, indirect and hyperdirect pathways regulate executive functions according to the classical basal ganglia model (Figure 1). The direct pathway facilitates movement while the indirect and hyperdirect pathways act as a "brake" and suppress movement. The normal functioning of the basal ganglia is therefore necessary to perform adequate movements, like walking or reaching for a glass of water.

#### The limbic loop

The limbic loop engages the same brain structures as the motor loop, but different areas within these. Instead of the dorsal striatum, the limbic loop engages the ventral striatum, known as the nucleus accumbens (NAc). The NAc receives DA projections from the ventral tegmental area (VTA) instead of the SNc, and GABAergic projections from the ventral pallidum (VP) instead of the GP. The limbic loop conveys affective functions such as reward-related behaviours. The cortex sends excitatory input to the NAc, which contains MSNs, just as the dorsal striatum of the motor loop. Direct, indirect and hyperdirect pathways are also similar to as in the motor loop but here target the medial and ventral aspects of the pallidal and subthalamic structures: The ventral pallidum (VP) and the medial tip of the STN, also known as the limbic tip. In the limbic loop, the SNr/EP send projections to the medio-dorsal nucleus of the thalamus.

## The cognitive loop

The cognitive loop, also called the associative loop, is the third parallel loop of the basal ganglia which slightly differs anatomically. Indeed, this loop involves the dorso-lateral part of the prefrontal cortex which sends projections to the anterior caudate (anterior striatum in rodents). GABAergic neurons from the anterior caudate innervate the GPi and SNr which in turn inhibit the medio-dorsal and ventral-anterior nuclei of the thalamus. The involvement of the basal ganglia in cognitive and associative functions is well

known (Brown, Schneider and Lidsky, 1997). Several studies have shown its role in goal-directed behaviours, decision-making, action selection and attention (Middleton and Strick, 2000; Rogers *et al.*, 2001; Stocco, 2018; Rusu and Pennartz, 2020). Impairment of the cognitive loop leads to cognitive symptoms observed in many brain disorders implicating the basal ganglia, like PD and obsessive compulsive disorder (OCD), further discussed below (Benzina *et al.*, 2016; O'Callaghan and Lewis, 2017).

## Basal ganglia-related disorders

Given the importance of the basal ganglia in motor, cognitive and affective functions, dysregulation of basal ganglia pathways is strongly associated with disorders and diseases, including Parkinson's disease, hemiballismus, chorea, Huntington's disease and Obsessive compulsive disorder. Below follows a short description of Parkinson's disease and Obsessive compulsive disorder, both of which are considered in the studies of this thesis.

#### Parkinson's disease

Parkinson's disease (PD) is the second most common progressive neuro-degenerative disease after Alzheimer disease with a prevalence of 1 to 2 per 1000 (Tysnes and Storstein, 2017). There are two main forms of PD: Familial PD and idiopathic PD. The onset of familial PD starts before the age of 50 and is often the consequence of a mutation. The idiopathic form of PD appears later in life, above 60 years, and is the consequence of neurodegeneration of the dopaminergic neurons in the SNc. The death of SNc DA neurons leads to decreased DA release in the dorsal striatum, which in turn affects basal ganglia function (Obeso et al., 2017; Tysnes and Storstein, 2017; Khan et al., 2019).

The cause of DA cell degeneration is unclear but has been proposed to depend on a mixture of genetic background and environmental factors (Marras, Canning and Goldman, 2019). DA cell degeneration causes dysregulation of the basal ganglia with over-activation of the indirect pathway over the direct pathway. Changes in firing activity of basal ganglia nuclei has been observed, in particular in the STN where the firing pattern becomes irregular (Bergman *et al.*, 1994; Benazzouz *et al.*, 2002). This dysfunction leads to progressive motor symptoms such as bradykinesia (slowness of movements), akinesia (failure to make a movement, freezing), tremor, difficulty to initiate

movement, and rigidity. Non-motor symptoms are also present in PD, sometimes before the appearance of motor symptoms: Olfactory dysfunction, apathy, mood disorders, sleep disorders, cognitive changes and autonomous-related functions like constipation and urination (Khoo *et al.*, 2013; Obeso *et al.*, 2017).

The main histological feature of PD is the presence of protein aggregates of abnormally folded alpha-synuclein. Alpha-synuclein is a naturally occurring protein in the brain, however, in PD, misfolded alpha-synuclein proteins aggregate into large complexes, forming so called Lewy bodies. These accumulate progressively in cerebral structures, including the DA neurons of the SNc (Braak *et al.*, 2003). Already in the early 2000's, Braak and colleagues hypothesized that alpha-synuclein could travel via the vagus nerve from the gut to the brain, and several studies have recently provided evidence supporting this hypothesis (Liddle, 2018; Kim *et al.*, 2019; Elfil *et al.*, 2020).

Current treatments for PD focus mostly on replacing the loss of DA by dopaminergic agonists or levodopa, a precursor to DA. These treatments are often combined with inhibitors of the monoamine oxidase B (IMAO-B) to prevent the degradation of DA. Another treatment used for patients suffering an advanced-stage PD is deep brain stimulation (DBS) of the STN which aims to stabilize aberrant STN activity by applying high-frequency electrical stimulation (will be discussed more below). Finally, another type of strategy to treat PD consists of transplanting mesencephalic DA neurons derived from human pluripotent stem cells, also called hPSC-derived mesDA neurons. This method aims to replace the loss of mesencephalic neurons in the SNc of PD patients by transplanting hPSC-derived mesDA neurons directly into the putamen (striatum). Pre-clinical studies have shown interesting results in animal models of PD. DA release from the transplanted hPSCderived mesDA neurons was observed as well as improved motor symptoms (Grealish et al., 2014; Chen et al., 2016). The translation of this method to PD patients seems promising with one clinical study recently initiated (Cyranoski, 2018; Parmar, Grealish and Henchcliffe, 2020).

Intensive research is still on-going to find better treatments for PD. To do so, researchers use various methods to generate animal parkinsonian models. Among them, neurotoxin-based models are the most common. For example, application of either of the toxins 1-methyl-4-phenyl-1,2,3,6-

tetrahydropyridine (MPTP) or 6-hydroxydopamine (6-OHDA) is commonly used to generate experimental PD models in non-human primates and rodents, primarily mice and rats. 6-OHDA is a neurotoxin commonly injected into the median forebrain bundle (MFB), the dorsal striatum, or the SNc with the objective to induce degeneration of the nigrostriatal pathway and mimic the loss of DA neurons in PD. 6-OHDA injections lead to degeneration of SNc DA neurons while VTA DA neurons are substantially less affected, for reasons not entirely known. Unilateral injection of 6-OHDA leads to unilateral degeneration of SNc DA neurons, and, consequently, the experimental animals display strong motor impairments with ipsilateral rotations (Boix, Padel and Paul, 2015; Park *et al.*, 2015).

#### Obsessive compulsive disorder

Obsessive compulsive disorder (OCD) is a common and chronic disorder characterized by excessive and uncontrollable thoughts (obsessions) and/or behaviours (compulsions). Since these obsessions and compulsions can go on for several hours a day, OCD has a large negative impact on everyday life.

The symptoms of OCD often appear in early adulthood with an earlier onset in boys than girls. The cause of OCD is unknown but seems multifactorial: Genetic background, environmental factors like childhood trauma, and even infections (Williams and Swedo, 2015; Robbins, Vaghi and Banca, 2019; Stein et al., 2019). Imaging studies have shown that affected brain areas include the orbital frontal cortex and the basal ganglia, in particular the caudate nucleus (Baxter, 1987; Pauls et al., 2014; Haber, 2016). Two current hypotheses aim to explain the neurobiological basis of OCD: "The cognitive hypothesis" which suggests a dysfunction in the valence attribution to a goal-directed behaviour and its outcome; and "the habit hypothesis" which postulates that OCD symptoms are due to a shift from goal-directed behaviours to excessive habit formation, with the compulsive action preceding the obsessive thoughts (Gillan and Robbins, 2014). Besides impairment in cognitive control and goal-directed and habit imbalance, emotional vulnerability including anxiety has long been considered an important factor in the etiology of OCD (Robbins, Vaghi and Banca, 2019). In addition to OCD, compulsive behaviour is also included in the symptom domain of substance abuse disorder and pathological gambling (Figee et al., 2016).

## Deep brain stimulation

Early in the 1960's, a study from Albe Fessard and colleagues reported that high-frequency stimulation (HFS), also called Deep brain stimulation (DBS) in humans, of the ventro-intermediate thalamic nucleus was efficient for treating tremor in PD patients (Benabid et al., 1991). The results were similar to those obtained by surgical lesioning of the area, but more efficient than a thalamotomy and with the important advantage of being reversible. The DBS method used consisted of implanting electrodes in the desired cerebral structure and sending short pulses (60-100 µs) at a frequency ranging from 100 to 185 Hz. DBS was later on improved by Benabid and colleagues in the late 1980's, and applied in the 1990's in non-human primate models and subsequently in PD patients for the treatment of motor symptoms (Benabid et al., 1994). Among different brain areas used as implantation sites for the stimulating electrodes, including thalamic nuclei, the GP and the STN, the STN was found to be the most efficient area for improving motor symptoms in PD, in particular when associated with levodopa treatment (Vizcarra et al., 2019). While motor improvement in PD is more efficient when applying DBS in the STN (STN-DBS) compared to GPi (GPi-STN) (Odekerken et al., 2016), GPi-STN has the advantage that patients need not adjust their medication to avoid treatment-induced dyskinesia (Vitek, 2002). Today, DBS treatment is a recommended clinical approach to alleviate motor symptoms in PD at the advanced-stage when DA neurons have substantially degenerated, and DA-based treatments no longer are efficient.

#### Theories of DBS mechanisms

The mechanisms underlying the beneficial effects of DBS remain to fully resolve, and are therefore still debated. Because STN-DBS induces similar clinical outcome as a lesion or a STN blockade (Bergman, Wichmann and DeLong, 1990; Luo, 2002), the main theory is that DBS inhibits neuronal activity. Neuronal inhibition induced by STN-DBS was first observed in rats and subsequently in non-human primates and humans in the 1990's. Several studies have provided evidence supporting the hypothesis that STN-DBS induces an inhibition of STN neurons and STN output structures. Studies in humans, monkeys and rodents have shown a decrease in the firing rate of STN neurons and STN output structures upon HFS (Tai *et al.*, 2003; Filali *et al.*, 2004; Meissner *et al.*, 2005). The mechanisms are unclear but several hypotheses exist: 1) Activation of the presynaptic inhibitory fibres innervating the STN (Boraud *et al.*, 1996; Deniau *et al.*, 2010; Chiken and Nambu,

2013); 2) block depolarization (Bikson *et al.*, 2001); and 3) inactivation of voltage-gated currents (Beurrier *et al.*, 2001; Shin *et al.*, 2007).

Despite strong evidence supporting the "inhibition theory", other studies have shown the opposite, that STN-DBS is, in fact, excitatory. First, studies have shown that HFS of the STN causes release of glutamate in some of the output structures of the STN (Lee *et al.*, 2004) as well as an increase in firing rate and c-fos levels in some of the target structures of the STN, namely the GP, EP and SNr (Hashimoto *et al.*, 2003; Galati *et al.*, 2006; Reese *et al.*, 2011; Shehab *et al.*, 2014). Second, STN-DBS triggers antidromic activation of cortical areas and induces changes in oscillatory activities and synchronicities between the basal ganglia nuclei (Li *et al.*, 2007; Moran *et al.*, 2011; Degos *et al.*, 2013). Finally, it has been proposed that the beneficial effects on movement observed in PD patients could come from the disruption of the abnormal activity in the indirect and hyperdirect pathways during PD, which would allow regulation of the aberrant hyperactivity displayed by the STN (Chiken and Nambu, 2016).

#### Side-effects upon DBS treatment

Despite giving rise to great improvements in motor control, observations have shown that STN-DBS can cause adverse side-effects, primarily in limbic and cognitive functions (Kim, Jeon and Paek, 2015; Serranová *et al.*, 2019). DBS is a non-selective electrical stimulation method through which all neural structures that come in contact with the HFS will be affected. Because of this lack of specificity, not only STN neurons, but also cerebral structures surrounding the STN as well as passing fibres will be reached by the electrical stimulation. In addition, as discussed further below, the STN is likely composed of several internal domains, or territories, that are differentially involved in motor, cognitive and affective functions. Thus, depending on the precise position of the DBS electrodes in the subthalamic area, the treatment can cause both alleviation of motor symptoms and unwanted side-effects (Petry-Schmelzer *et al.*, 2019).

The reasons for the variability in success rate and the appearance of sideeffects are not entirely known. It has been suggested that the clinical outcome of STN-DBS in PD patients depends on the stimulation of different domains within the STN, the electrical parameters of the stimulation, damages along the trajectory of the electrodes, changes in the medication and/or the progressive nature of the disease. For example, it has been shown that electrodes placed in different positions within the STN motor domain induce different clinical outcomes depending on the antero-posterior location (Serranová *et al.*, 2013). Several studies have shown mixed results in post-surgery depression and apathy with either improvements or deteriorations (Czernecki, 2005; Kalteis *et al.*, 2006; Le Jeune *et al.*, 2009; Pariwatcharakul *et al.*, 2013; Pinsker *et al.*, 2013; Robert *et al.*, 2014; Accolla and Pollo, 2019). Non-motor side-effects that are regularly observed are depression, apathy, weight gain, mood changes, worsened verbal fluency, hyper/hypomania and impulse control disorder (Bronstein *et al.*, 2011; Witt, Daniels and Volkmann, 2012; Nassery *et al.*, 2016).

## The subthalamic nucleus, STN

The STN is a bilateral, small and dense nucleus located between the zona incerta dorsally and the cerebral peduncle ventrally, posteriorly to the EP and anteriorly to the midbrain. In humans, it is considered as a "closed" nucleus, meaning that STN dendrites are mainly restricted within the nucleus itself, except on its medial aspect where it opens on the lateral hypothalamic area (LHA) (Figure 2). The STN primarily contains glutamatergic projection neurons expressing the *Vglut2/Slc17a6* gene encoding the Vesicular glutamate transporter 2 (VGLUT2). VGLUT2, together with VGLUT1 and VGLUT3, form a family of proteins in which members enable packaging of glutamate into presynaptic vesicles for neurotransmitter release. The three VGLUTs show different distribution patterns in the brain. The *Vglut2* gene is predominantly expressed in neurons of the STN, thalamus, hypothalamus,

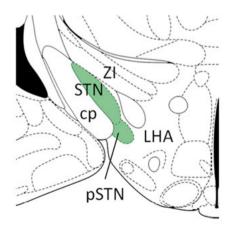


Figure 2: Schematic representation of the subthalamic nucleus (STN) and para-STN nucleus (pSTN) in green on a coronal plan at -2.06 mm from the bregma. ZI: zona incerta; cp: cerebral peduncle; LHA: lateral hypothalamic area. Adapted from the Franklin & Paxinos atlas.

brainstem and some forebrain structures; the VGLUT2 protein localizes to synaptic vesicles in the presynaptic terminals of these neurons (Herzog *et al.*, 2001; Kaneko and Fujiyama, 2002; Oliveira *et al.*, 2003; El Mestikawy *et al.*, 2011).

While VGLUTs show broad distribution patterns and cover the vast extent of glutamatergic neurons in the brain, the *Pitx2* gene encoding the Paired-like homeodomain 2 (PITX2) transcription factor shows high selectivity for the STN. The expression of the *Pitx2* gene in the STN was identified already in the early 2000's in studies also showing that, in the mouse, *Pitx2* gene expression is required for the primary neuronal migration from the hypothalamus to form the STN, and consequently necessary for its development (Martin *et al.*, 2004a; Skidmore *et al.*, 2008a). In the adult rodent, Pitx2 mRNA can also be found in the adjacently located para-STN (pSTN) and in some hypothalamic and mammillary neurons, but at substantially lower levels than in the STN (Wallén-Mackenzie 2020). Pitx2 mRNA overlaps to near-100% with Vglut2 mRNA, further identifying the glutamatergic nature of the Pitx2-positive STN neurons (Schweizer *et al.*, 2016).

The primate STN is often divided in three internal anatomical-functional domains, or territories, corresponding to the three loops of the basal ganglia: A dorso-lateral domain involved in motor functions, a ventral domain involved in associative/cognitive functions and the medial tip, the limbic tip, for limbic functions (Figure 3). This tripartite macro-architecture, the so called *tripartite model*, was first characterized by anatomical tracing studies followed by clinical results obtained in STN-DBS treatment of PD, and the improvement of imaging technologies (Lambert *et al.*, 2012; Alkemade, Schnitzler and Forstmann, 2015). Indeed, the location of DBS electrodes in the dorso-lateral domain of the STN is crucial to alleviate motor symptoms in PD patients while limiting side-effects. However, a strict tripartite organization of the STN in humans and non-human primates is debated (Alkemade, Schnitzler and Forstmann, 2015).

Several studies have suggested that the macro-architecture of the STN and many other cerebral structures is more intermingled in rodents than in humans and non-human primates with the overlap of different neuronal subpopulations (Figure 3) (Mallet et al., 2007; Alkemade, Schnitzler and Forstmann, 2015; Janssen et al., 2017). Here, the identification of intermingled cellular organization, rather than domain-type organization, in other basal ganglia structures such as the GP (Hegeman et al., 2016) and in midbrain structures like the VTA (Viereckel et al., 2016a; Poulin et al., 2018), does not lend support to the tripartite model. Furthermore, studies in humans and monkeys investigating the distribution of various proteins and mRNAs in the STN have found contradicting results with both clear expression in one of the STN domain for some mRNA/proteins like parvalbumin and calretinin (Parent et al., 1996; Augood et al., 1999) and homogeneous expression in the whole STN for others, like tyrosine hydroxylase (TH), prepro-Enkephalin B, GABA<sub>B</sub> and GABA<sub>A</sub> receptors (Kultas-Ilinsky, Leontiev and Whiting, 1998; Charara et al., 1999; Hedreen, 1999; Aubert et al., 2007).

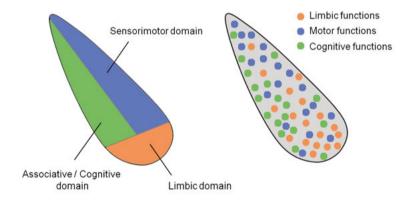


Figure 3: Schematic representation of the internal organization of the STN according to the tripartite model (left) and the intermingled hypothesis (right). Left: Blue, the sensorimotor domain (dorso-lateral); green, the associative/cognitive domain (ventro-medial); orange, the limbic domain (medial, also known as the limbic tip). Right: An overlap of intermingled subpopulations of neurons across the entire STN but the preservation of some topographic organization.

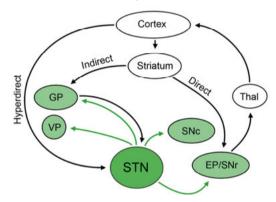
#### Afferent projections to the STN

The inputs and outputs of the STN were mostly investigated in the 1980's and 1990's by using neuronal tracers like the retrograde cholera toxin subunit B, CTB, and the anterograde protein from *Phaseolus vulgaris*, PHA-L.

Studies using neuronal tracing methods showed that the STN receives excitatory input from various cortical areas and the centro-median parafascicular complex of the thalamus, and inhibitory input from the GP and VP (Figure 4). Monoaminergic projections from the dorsal raphe nucleus (serotonin) and SNc (DA) were also identified in several publications (Parent and Hazrati, 1995).

#### Efferent projections from the subthalamic nucleus

The STN sends glutamatergic projections to both basal ganglia and non-basal ganglia structures. Substantial projections reach the SNr and EP, and the STN also communicates reciprocally with the GP. To a lesser extent, STN neurons innervate the SNc, VP and the pedunculopontine nucleus (PPN) (Figure 4) (Schweizer *et al.*, 2016; Fife *et al.*, 2017). In addition to these well-known target structures, tracing studies have shown projections from the STN to the striatum, the thalamus, and the VTA (Nauta and Cole, 1978; Kita and Kitai, 1987; Groenewegen and Berendse, 1990).



*Figure 4:* Illustration of the STN circuitry within the basal ganglia. The STN projects to the globus pallidus (GP), substantia nigra *pars compacta* (SNc) and *pars reticulata* (SNr), the entopeduncular nucleus (EP) and the ventral pallidum (VP).

#### Motor functions of the STN

"Despite the current interest, still very little is known about the STN's normal function in relation to movement"

(Alkemade, Schnitzler, and Forstmann 2015).

The role of the STN in motor functions has long been investigated in relation to PD. However, as stated in the quote above, knowledge of its natural role in movement is surprisingly limited (Alkemade, Schnitzler and Forstmann, 2015). Instead, most knowledge is derived from various disease models, and through clinical observations. For example, Whittier and Mettler showed already in 1949 that a lesion of the STN induces hemichorea/ballism in monkeys (Whittier and Mettler, 1949; Carpenter, Whittier and Mettler, 1950). Later on, studies confirmed these findings with more selective lesions. Crossman and colleagues showed that injection of bicuculline, a GABA<sub>A</sub> receptor antagonist, into the STN, induced uncontrollable and irregular limb movements, also called hemiballism (Crossman, Sambrook and Jackson, 1984). This finding was confirmed by Beurrier and colleagues by lesioning the STN in healthy monkeys (Beurrier et al., 1997). Several studies in rodents and monkeys have used neurotoxin-based models to induce Parkinsonism. It has been shown that subthalamotomy in MPTP-lesioned or 6-OHDA-lesioned rodents and monkeys abolishes the toxicity-induced motor symptoms (Bergman, Wichmann and DeLong, 1990; Aziz et al., 1991; Beurrier et al., 1997; Chang et al., 2003; Darbaky et al., 2003; Marin et al., 2013). Similar results were observed in PD patients after subthalamotomy or pallidotomy (Laitinen, Bergenheim and Hariz, 1992; Lozano et al., 1995; Heywood and Gill, 1997). According to the classical basal ganglia model, this alleviation of motor symptoms in PD is derived from the removal of STN over-activation which lowers the neuronal activity of GPi/EP neurons, and consequently disinhibits the thalamus and promotes movement.

Beyond STN-DBS in PD and STN-HFS studies in animal PD models, only few studies have investigated the normal function of the STN. Instead, most studies have used STN-HFS in non-human primates or rodents to better understand the mechanisms of STN-DBS in PD. However, some STN-HFS studies have included healthy rodents as control groups and showed that unilateral STN-HFS induces contralateral rotations and dyskinesia (Bergmann *et al.*, 2004; Boulet *et al.*, 2006).

In addition to electrical stimulation models, another method to experimentally study the function of a brain structure is conditional mouse genetics. Mice in which glutamate packaging and release has been reduced in STN neurons, by the combination of Pitx2-Cre mice (Martin *et al.*, 2004a) with the floxed allele of *Vglut2* (Wallen-Mackenzie, 2006), resulted in hyperlocomotion characterized by increased horizontal and vertical (rearing) activities without

affecting gait and fine motor coordination (Schweizer 2014, 2016, Pupe, Schweizer, Mackenzie, 2014). These studies thereby provided indirect support for the role of STN glutamatergic neurotransmission in locomotor regulation, in accordance with the basal ganglia model of movement regulation.

Over the past decade, optogenetics has been implemented to study the STN. However, in contrast to other cerebral structures, the behavioural effect of optogenetic excitation or inhibition of the STN has only rarely been investigated in normal, healthy animals. Instead, the vast majority of optogenetic studies of the STN have been performed in 6-OHDA lesioned rodents. For example, it has been shown that optogenetic inhibition of the STN in 6-OHDA lesioned rats improves forelimb akinesia and levodopa-induced dyskinesia (Yoon et al., 2014, 2016). Also in 6-OHDA lesioned rodents, activation of cortico-subthalamic afferent projections with HFS above 100 Hz reverses the induced motor symptoms while low frequency stimulation (LFS) worsen the motor symptoms, or has no effect (Gradinaru et al., 2009; Sanders and Jaeger, 2016). In healthy mice. Tian and colleagues observed hyperkinesia, decreased exploratory activity and stereotyped movements (grooming) upon activation of GABAergic GP neurons with 20 Hz optogenetic stimulation. The same study also showed that activation of subthalamic glutamatergic terminals in the GP induces hyperkinesia and stereotyped movements such as grooming and dystonia-like behaviours (Tian et al., 2018). A hypothesis aiming to explain the opposite behavioural outcome between STN-HFS and STN-LFS in PD animal models postulates that spontaneous activity is still present during STN-LFS while STN-HFS suppresses spontaneous activity to replace it with a different and regular pattern (Garcia et al., 2005).

Finally, only one study has investigated the effect of direct optogenetic excitation of the STN in healthy mice, and could demonstrate that brief activation of the STN was sufficient to interrupt an on-going licking behaviour while inhibition of the STN could suppress the interruptive effect of surprise (Fife *et al.*, 2017). This study was important as it directly pin-pointed the role of the STN in pausing/stopping movements that have already been initiated.

#### Affective and associative functions of the STN

Since the discovery of STN-DBS in the 90's, most studies have focused on motor functions of the STN. However, increasing research is now directed towards the role of the STN in affective functions, not least due to the concern for non-motor symptoms in PD. In addition, there is increasing interest in the potential role played by the STN in addiction and OCD (Bari and Robbins, 2013; Pelloux and Baunez, 2013; Creed, 2018; Pelloux et al., 2018; Rappel et al., 2018). The role of the STN in limbic functions is not as well understood as for motor functions. Several studies have shown that the nonmotor side effects induced by STN-DBS could come from the stimulation of the limbic and/or associative domains of the STN itself, and not from surrounding regions or fibres (Temel et al., 2005, 2006; Mallet et al., 2007; Haegelen et al., 2009; Baláž et al., 2011). This is an interesting observation which points towards the importance of increasing the understanding of precisely how the STN is involved in regulation of the various functions it has been associated with, i.e. motor, cognitive/associative and limbic/affective functions. Improved knowledge of the anatomical-functional organization of the STN is clearly of essence.

In the context of affective functions, the STN is intrinsically connected to limbic brain structures. The STN receives its cortical projections from the prefrontal cortex area, well known for its role in cognitive and limbic functions (Haynes and Haber, 2013), and from the VP, a central structure of the reward system which projects directly to the VTA and the lateral habenula, LHb (Root et al., 2015; Wulff et al., 2019). In turn, the STN sends direct projections to limbic structures such as the VP, or via relay structures such as the EP, to the LHb, a pathway involved in evaluating action outcomes (Stephenson-Jones et al., 2016). The LHb itself is known for its role in regulating negatively motivated behaviours and for its contribution to psychiatric diseases like major depression disorder and addiction (Lecca et al., 2017; Hu, Cui and Yang, 2020). Additional structures of the limbic system are affected by STN-HFS with an increase of DA release in the NAc shell (NAcbSh) and NAc core (NAcbC), and a decrease of GABA in the VTA (Winter, Lemke, et al., 2008). Further, mice lacking Vglut2 in Pitx2-Cre STN neurons also displayed an increase in dopamine transporter (DAT) capacity in the NAcbSh (Schweizer et al., 2016).

Similar to the majority of studies investigating the role of the STN in motor function, most studies interested in the role of the STN in limbic and cogni-

tive functions have used STN-HFS, primarily in PD models. Several studies in rodents have demonstrated a role of the STN in reward processing: Electrophysiological recordings in the STN showed that neurons responded differently to positive and/or negative stimuli as well as to reward omission. In addition, changes in the activity of such STN subpopulations predicted the availability of reward, pointing towards a role of STN neurons in reward prediction error (Breysse, Pelloux and Baunez, 2015). Further, it has been shown that STN-HFS decreases the motivation for cocaine (Baunez et al., 2005; Rouaud et al., 2010) and prevents the re-escalation of heroin intake in rats (Wade et al., 2017), confirming the importance of the STN in addictive behaviours. Another study showed that lesioning of the STN affects the processing for positive and negative reinforcers like saccharine and lithium chloride (Pelloux et al., 2014). These findings are comparable to results obtained by Schweizer and colleagues in which mice with reduced Vglut2 levels selectively in Pitx2-Cre neurons showed decreased sugar consumption compared to controls, but without reduced reward-related learning, memory, motivation or ability for task-switching (Schweizer et al., 2016).

Besides affecting reward-related behaviours, several lines of evidence suggest that the STN is a key structure in regulating compulsive behaviours. In rodents, compulsive behaviours are characterized by stereotyped movements defined as motor responses that are repetitive, invariant, and seemingly without purpose or goal, for example excessive self-grooming, licking or self-gnawing (Kelley, 1998; Kalueff et al., 2016). Studies using STN-HFS or STN lesioning have shown a decrease in compulsive behaviours in different OCD and autism animal models (Baup et al., 2008; Winter, Mundt, et al., 2008; Klavir et al., 2009; Chang et al., 2016). In contrast to STN inhibition or inactivation, disinhibition of the STN induces strong self-grooming (Tian et al., 2018). Several animal studies have demonstrated that compulsive behaviours can be induced by activating different pathways within, or connected to, the basal ganglia: Overactivation of the ventral thalamus via injections of bicuculline in monkeys triggers repetitive and time-consuming motor acts (Rotge et al., 2012); activation of the anterior part of the GPe via bicuculline injections also gave rise to compulsive behaviours in monkeys (Grabli, 2004); HFS, but not LFS, of the GP and the EP, reduced excessive selfgrooming in an OCD rat model (Klavir, Winter and Joel, 2011); and lesion of the VP and GP impaired grooming without affecting its sequential organization (Cromwell and Berridge, 1996). The conclusions of these studies demonstrate the role of basal ganglia-related structures in compulsive behaviours, often characterized in rodent models of human disorders by excessive self-grooming. However, the involvement of many cerebral structures suggests that compulsive behaviours rely on complex circuitries which might process different aspects of compulsivity.

## The ventral pallidum, VP

The VP, also referred as substantia innominata, is a structure located in the basal forebrain. The VP is about 2 mm long in mice, and extends along the antero-posterior axis from below the NAc to the anterior commissure, the bed nucleus of the stria terminalis (BNST) and the GP in its most posterior aspect. The VP was first described as an extension of the GPe but the presence of strong substance-P GABAergic fibres from the NAc distinguishes the VP from the dorsal aspect of the pallidum (Haber et al., 1985; Zahm, 1989). Another difference between the VP and GP is the cellular heterogeneity of the VP with a majority of GABA neurons, but also glutamatergic and cholinergic neurons. GABAergic and glutamatergic VP neurons project reciprocally to the NAc and STN, and to the VTA and the LHb, while cholinergic VP neurons target preferentially the prefrontal cortex and the basolateral amygdala (Jones, 2004; Unal, Pare and Zaborszky, 2015). GABAergic and glutamatergic VP neurons seem to project to the same target structures, with some exceptions, and regulate the value of a stimulus in an opposite manner (Faget et al., 2018). Indeed, as a part of the limbic loop of the basal ganglia, the VP plays an important role in reward- and aversive-related behaviours and motivated behaviours and also, to a lesser extent, in motor and cognitive behaviours (Root et al., 2015; Saga et al., 2017). The VP is well known for its role in addiction to drugs of abuse and has recently been proposed as a possible target for DBS as a new treatment strategy (McGovern and Root, 2019).

## The lateral habenula, LHb

The habenula is a pair of medially positioned nuclei located dorsally of the thalamus. It is divided into two parts: the medial habenula and the lateral habenula, LHb. Neurons of the LHb are mainly glutamatergic and express the *Vglut2* or *Vglut3* genes, but some studies have also shown the presence of inhibitory interneurons expressing the *Glutamic acid decarboxylase 2* (*GAD2*) gene and the *GABA transporter 1* (*GAT1*) gene that trigger inhibito-

ry responses when stimulated optogenetically (Zhang et al., 2018; Flanigan et al., 2020; Webster et al., 2020). However, the presence of functional GABA neurons in the LHb was recently questioned by Wallace and colleagues who confirmed a low expression of GAD2 and GAT1 but also showed the absence of expression of the SlcSlc32a1 and Slc18a2 genes, encoding mRNAs of the Vesicular inhibitory amino acid transporter, Viaat, and Vesicular monoamine transporter 2,Vmat2, respectively (Wallace et al., 2020).

The LHb receives input from many forebrain structures including the medial prefrontal cortex (mPFC), lateral hypothalamic area (LHA), VP and EP. LHb neurons project to midbrain and brainstem structures including the SNc, VTA, rostromedial tegmental nucleus, dorsal raphe nucleus and locus coeruleus (Hu, Cui and Yang, 2020). The LHb gained substantial attention when Hikosaka and colleagues discovered the role of the LHb in negative reward processing (Hikosaka, 2010). Indeed, with its connections to both the dopaminergic and serotoninergic system, the LHb regulates many essential functions including reward-seeking behaviours, avoidance-like behaviours, sleep, anxiety, stress and pain. Dysfunction of the LHb is consequently involved in several severe neuropsychiatric disorders, including both addiction and major depression disorder (Matsumoto and Hikosaka, 2009; Hikosaka, 2010; Baker *et al.*, 2016; Lecca *et al.*, 2017; Shabel *et al.*, 2019; Hu, Cui and Yang, 2020).

A couple of studies have shown that STN-HFS can either activate or inhibit LHb neurons and that it induces c-fos expression in the LHb (Tan et al., 2011; Hartung et al., 2016). Curiously, however, no direct connection has ever been demonstrated between the STN and the LHb. Some studies have shown that the VP and/or the EP could be relay structures between the STN and the LHb. Indeed, it was recently shown that a subpopulation of VP neurons expresses Vglut2 and innervates the LHb. Furthermore, while general optogenetic stimulation of the VP structure induced place preference, selective activation of the glutamatergic subpopulation within the VP instead triggered place avoidance (Faget et al., 2018; Tooley et al., 2018). Even though a STN → VP → LHb pathway thus seems likely, the only projections actually demonstrated are the glutamatergic projections from the STN to the VP, and from the VP to the LHb. However, no study has shown that glutamatergic VP neurons projecting to the LHb receive glutamatergic input from the STN. Another possible pathway explaining the activation of the

LHb via STN stimulation is through the EP. The EP is mostly a GABAergic structure but also contains a subset of neurons releasing only glutamate or both glutamate and GABA (Stephenson-Jones *et al.*, 2016). It was recently shown that the limbic tip of the STN projects to EP neurons that synthesize glutamate, and that in turn send projections to the LHb (Stephenson-Jones *et al.*, 2016; Wallace *et al.*, 2017, 2017; Li, Pullmann and Jhou, 2019). To summarize, STN stimulation may lead to responses in the LHb via indirect projections.

## The medial hypothalamic-mesencephalic area

#### The ventral tegmental area, VTA

The VTA is a midline structure located ventrally of the third ventricle in the midbrain, and it is flanked bilaterally by the SNc and SNr. The VTA is composed of several subnuclei: The interfascicular nucleus (IF) medially, the parabrachial pigmented nucleus (PBP) laterally, the paranigral nucleus (PN) and the parainterfascicular nucleus (PIF) ventral to the PBP, and more rostrally, the ventral area rostral nucleus (VTAr) (from Franklin & Paxinos atlas). Some studies group the IF subnucleus with the caudal aspect of the hypothalamus (Cavanaugh *et al.*, 2011), but most commonly, the IF is considered a VTA subnucleus. In addition, the rostral and caudal linear nuclei are close to the VTA, and sometimes considered VTA subnuclei, especially the rostral linear nucleus (Figure 5).

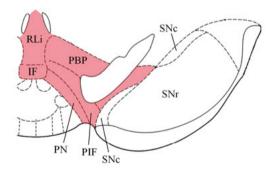


Figure 5: Schematic representation of the ventral tegmental area at -3.40 mm from Bregma. The VTA consists of an assembly of several nuclei in the medial aspect of the ventral midbrain (red area). RLi: rostral linear nucleus; IF: interfascicular nucleus; PBP: parabrachial pigmented nucleus; PN: paranigral nucleus; PIF: parainterfascicular nucleus; SNc: substantia nigra pars compacta; SNr: substantia nigra pars reticulata.

While long believed to be composed of DA neurons only, the VTA has recently gained attention as a strongly heterogeneous structure in which glutamatergic and GABAergic neurons are intermixed with the DA neurons, and also with neurons able to co-release these neurotransmitters (Trudeau *et al.*, 2014). The distribution of these different types of neurons varies across subnuclei. The density of DA neurons increases from the midline to the lateral part of the VTA, with the exception of the IF, while glutamatergic neurons are highly concentrated in the medial VTA and become gradually more sparse in the lateral VTA (Morales and Root, 2014; Morales and Margolis, 2017). DA neurons express the *Th* gene encoding Tyrosine hydroxylase (TH), the rate-limiting enzyme in DA synthesis, which is commonly used to identify these neurons, and VTA glutamatergic neurons are characterized by *Vglut2* gene expression. Th/Vglut2 mRNA co-labeling has often been used to distinguish DA-glutamate co-releasing neurons, most common in the medial VTA.

The VTA is well known for its role in motivated behaviour, reward and aversion through projections to various forebrain structures: NAc, olfactory tubercule, PFC, amygdala, hippocampus, BNST, VP, LHb, locus coeruleus and periaqueductal gray matter (Morales and Margolis, 2017). While these pathways have been carefully mapped for VTA DA neurons over several decades, the more recently discovered VTA glutamatergic neurons have been shown to have a similar projection pattern (Hnasko et al., 2012). VTA GABA neurons are mostly interneurons that regulate DA neuron activity (Tan et al., 2012; Creed, Ntamati and Tan, 2014). Through substantial projections to the NAc, VTA DA neurons are involved in reward processing. Indeed, the firing rate of VTA DA neurons increases upon delivery of an unexpected reward but also in expectation of a reward, and this induces DA release in the NAc. Moreover, when a reward is systematically paired with a neutral stimulus, the increased response of DA neurons shifts in time from the delivery of the reward to the reward-predicting stimulus. This mechanism forms the basis of reward-prediction and error evaluation (Schultz, Dayan and Montague, 1997).

In addition, VTA neurons co-releasing DA and glutamate in the mAcbSh modulate a behaviour switch depending on positive or negative reinforcers (Mingote *et al.*, 2019). Several studies over the past decade have shown that DA-glutamate co-releasing neurons are important for reinforcement processing of both natural rewards, such as sugar, and addictive drugs, such as cocaine (Birgner *et al.*, 2010; Alsio *et al.*, 2011; Fortin *et al.*, 2012; Papathanou *et al.*, 2018). VTA glutamatergic neurons projecting to the LHb

have been shown to mediate aversion (Root *et al.*, 2014; Lammel *et al.*, 2015) while VTA GABA neurons have been proposed to locally inhibit DA neurons (Tan *et al.*, 2012).

#### The hypothalamic-mesencephalic area

Glutamatergic neurons of the midbrain are not restricted to the VTA but form a rostro-caudal continuum reaching the posterior hypothalamic area (PHA). The hypothalamus develops from the ventral diencephalon and is composed of three main regions: the rostral thalamus with preoptic areas, the tuberal hypothalamus containing the lateral hypothalamic area (LHA) and the infundibulum, and a posterior region which consists of the mammillary bodies (MM), tuberomammillary, retromammillary nucleus (RMM), supramammillary (SuM), and posterior hypothalamic nucleus (PH) (Figure 6) (Saper and Lowell, 2014).

The hypothalamus contains both GABAergic and glutamatergic neurons. involved in autonomic brain functions such as sleep/wake cycle (Sapin et al., 2010; Luppi and Fort, 2019), thermoregulation (Contreras et al., 2016; Ishiwata and Greenwood, 2018), food intake (Schwartz et al., 2000), nociception (Akerman, Holland and Goadsby, 2011), aversion (Lazaridis et al., 2019) and more. These various functions arise from the many diverse subnuclei of the hypothalamus and recent studies have identified neuronal subpopulations expressing different molecular patterns (Chen et al., 2017). Furthermore, a study identified a neuronal population expressing the Trpv1 gene encoding the Transient receptor potential cation channel subfamily V member 1 (TRPV1) located in the ventral hypothalamic-mesencephalic area (Viereckel et al., 2016b). In mice, Trpv1 is highly expressed in the mes-di-encephalon between E14 and P3 but is only weakly expressed in the adult (Viereckel et al., 2016b; Dumas and Wallén-Mackenzie, 2019). In the adult mouse, Trpv1 expression is restricted to a rostro-caudal band from the PH and SuM to the medial nuclei of the VTA (IF, PN, PIF and RLi) and is mainly glutamatergic (Figure 6) (Cavanaugh et al., 2011; Viereckel et al., 2016b).

It is unclear if the neuronal subpopulation expressing the *Trpv1* gene plays a distinct role in any of the known functions of the hypothalamus and no study has looked at the projections of this subpopulation. Only one anatomical study using PHA-L showed that the SuM strongly projects to the hippocampus, the septum, the dorsal raphe nucleus, the dorsomedial hypothalamic

area, the thalamus, preoptic areas, BNST, VP, VDB, DEn and cortical areas (Vertes, 1992).

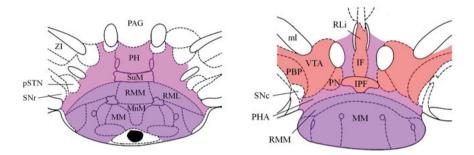


Figure 6: Schematic representation of the hypothalamic-mesencephalic area at two different bregma levels, -2.70 mm on the left and -3.08 mm on the right. The posterior hypothalamic area corresponds to the pink area, the mammillary bodies to the purple area and the VTA to the red area. ZI: zona incerta, pSTN: parasubthalamic nucleus; SNr: substantia nigra pars reticulata; PAG: periaqueductal gray matter; PH: posterior hypothalamus; SuM: supramammillary nucleus; RMM: retromammillary nucleus; MnM: median part of the mammillary nucleus; MM: medial mammillary nucleus; RML: lateral mammillary nucleus; RLi: rostral linear nucleus; VTA: ventral tegmental area; ml: lemniscus; PBP: parabrachial pigmented nucleus; SNc: substantia nigra pars compacta; PHA: posterior hypothalamic area; PN: paranigral nucleus of the VTA; IF: interfascicular nucleus; IPF: interpeduncular fossa.

#### Overall aim

It has long been established, primarily via clinical conditions and animal models of disease, that the STN is involved in motor, associative and limbic functions. But how does the STN, with its modest size, engage in such different functions? Further, which functions can really be pin-pointed as regulated by the STN during normal, non-pathological conditions? And, does the STN actually engage in so many different functions, or are some of these mediated by immediately surrounding brain structures or even passing fibres? For example, when stimulating DBS electrodes are placed in or even near the STN, which of the observed effects are due to stimulation of STN neurons? To understand this, the role of the STN under baseline conditions must be better understood than it is today. Further, each of these functions, motor, associative and limbic, contain many different aspects – which ones can actually be ascribed to the STN? For example, which particular aspects of motor control and movement involve the STN? The STN has also been implicated in reward processing but considering recent revelations that some components within the STN circuitry, as well as structures adjacent to the STN itself, engage in aversion, does the STN also play a similar role? Many questions remain to answer before the STN and its role in neurocircuitry and behavioural regulation can be fully decoded.

The overall aim of this thesis work has been to reveal neurobiological underpinnings of the STN and its network by experimentally addressing critical questions regarding the neurocircuitry of the STN, and its role in behaviour. Many aspects have been assumed as solved, such as the role of STN in motor control. However, experimental evidence has been largely lacking. In this thesis, some of the crucial STN queries have been addressed experimentally, with both expected and surprising findings as the result. While many issues remain to fully solve, the work presented brings new knowledge about the mouse STN that should help towards decoding this elusive but clinically important brain structure and its natural brain habitat.

## Material and Methods

	Study I	Study II	Study III	Study IV
Viral injections	X	X	X	NA
Cannula implantations	X	X	X	NA
In vivo electrophysiology	X	X	X	NA
Behavioural experiments	X	X	X	X
6-OHDA injections	NA	NA	NA	X
In situ hybridization	NA	NA	SD	X
Immunohistochemistry	X	X	X	X
Genotyping of mice	X	X	X	NA

**Table 1**: Overview of my contribution (X) to the different methods performed in the four studies described in this thesis. Behavioural experiments were also performed by additional co-authors. NA: not applicable. SD: performed by other co-author.

## Transgenic mice

The technology enabling gene mutations in mice, also called transgenic mouse technology, is now widely used in many fields of biology. It allows the manipulations of the genome in order to delete, insert, translocate or inverse a specific sequence of DNA. A commonly used method using transgenic mouse lines is the Cre-Lox technology developed in the 1990's by Tsien and colleagues (Tsien, 2016). By introducing the gene encoding the enzyme Cre recombinase downstream of a promoter of interest, specific manipulations of a cellular population in which the promoter is actively transcribed can be achieved. The Cre recombinase, commonly abbreviated as Cre, has the capacity to recombine short DNA sequences called Lox sites. Depending on the orientation of the Lox sequences, any DNA sequence which is flanked by Lox sites will be excised, inverted or translocated.

Transgenic mice expressing Cre recombinase are usually named after the promoter which controls expression of Cre expression, and they are referred to as strains or lines of mice. In the current studies, we have used two transgenic mouse lines: Pitx2-Cre (Study I and II) and Trpv1-Cre (Study III).

The Pitx2-Cre mouse line was created by Dr Martin and colleagues by crossing a pre-existing *Pitx2*<sup>creneo/+</sup> line with a FLPe recombinase-expressing transgenic mouse line to remove neomycin sequences that might interfere with the normal expression of *Pitx2* gene (Liu, 2003; Martin *et al.*, 2004b; Skidmore *et al.*, 2008b). Martin and colleagues maintained Pitx2-Cre mice on a C57NL/6J background. Upon import to our laboratory, the line has been bred in-house by mating Pitx2-Cre <sup>tg/wt</sup> males with C57BL/6NTac female mice.

The Trpv1-Cre mouse line (also called B6.129-Trpv1tm1(cre)Bbm/J) was developed by Dr Basbaum's laboratory (Cavanaugh *et al.*, 2011) and is available for purchase at The Jackson Laboratory. The mice contain a myctagged IRES-cre sequence inserted downstream of the *Trpv1* stop codon. This method ensures that the endogenous *Trpv1* coding sequence is not disrupted. Trpv1-Cre mice were bred in-house. The line was maintained by mating Trpv1-Cre<sup>tg/wt</sup> or Trpv1-Cre<sup>tg/tg</sup> males with C57BL/6NTac female mice.

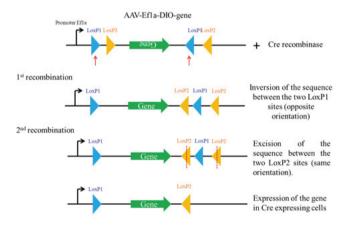
The genotype of Pitx2-Cre and Trpv1-Cre mice were confirmed by PCR analyses using DNA extracted from ear biopsies and Cre-directed primers (Forward primer: 5'-CACGACCAAGTGACAGCAAT-3' and reverse primer: 5'-AGAGACGGAAATCCATCGCT-3'). Both female and male mice were used in the experiments.

#### **Optogenetics**

Optogenetics is a technique combining genetic and optical methods which was developed to allow controlled manipulation of neuronal activity by applying light (Gradinaru *et al.*, 2007; Deisseroth, 2011). It became possible with the discovery of light sensitive proteins in microorganisms such as channelrhodopsins, bacteriorhodopsins and halorhodopsins. Light of a specific wavelength can activate each of these light-sensitive proteins and trigger an exchange of ions to, depending on the opsin, hyperpolarize or depolarize the neuron expressing the opsin. A commonly used channelrhodopsin, which has

been implemented in the current studies I-III, is the Channelrhodospin 2 (ChR2), a light-gated ion channel from the algae *Chlamydomonas* reinhardtii. ChR2 is a seven transmembrane protein containing a nonspecific cation channel which opens when receiving light at 473 nm. The flow of ions induces the depolarization of the neuron leading to the generation of action potentials. Another common opsin, used in the current study I, is the Archeorhodopsin 3.0 (Arch). This opsin is a proton pump from *Halorubrum* sodomense activated by a 532 nm wavelength light which induces strong photocurrents leading to hyperpolarization and inhibition of the neurons expressing the opsin (Gradinaru et al., 2010).

When applying optogenetics in rodents, selectivity is commonly achieved through the Cre-Lox system. Cre-transgenic mice are selected based on the promoter driving the expression of Cre. The DNA construct encoding the opsin contains Lox sites to allow recombination selectively in neurons expressing Cre (Figure 8). Through the Cre-Lox system, spatial selectivity can be achieved. This way, optogenetics allows a level of selectivity that is not possible by electrical stimulation. Even if light stimulation covers a broad area, only neurons expressing Cre and containing the floxed allele will be activated or inhibited when receiving the light (Figure 9).



*Figure 8*: Illustration of the Cre-Lox system used with a double-floxed with inverted orientation (DIO) strategy.

In the current studies, Cre-dependent Adeno-associated viruses (AAVs) (Studies I, II, III: AAV2-DIO-Ef1a-(hChR2(H134R))-eYFP; Study I: AAV2-EF1a-DIO-(Arch3.0)-eYFP) were injected to specifically target neurons expressing the Cre recombinase under the *Pitx2* or the *Trpv1* promoter, as explained above. The AAV2 serotype is known for its high transfection efficiency in neuronal cells and allowed us to visualize Pitx2-Cre and Trpv1-Cre positive cell bodies, their dendrites, axons and terminals. Here, a double-floxed strategy with inverted orientation (DIO) was used. AAV viruses carried a construct containing the gene for an opsin, either ChR2 or Arch, and the gene encoding the fluorescent reporter eYFP. Mice were injected in either the STN (Pitx2-Cre) or the PHA (Trpv1-Cre) allowing expression of the opsin and reporter genes in these brain structures (Figure 8).

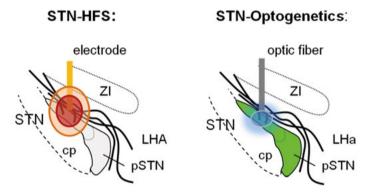


Figure 9: Illustration of two stimulating methods of the STN: Left, High-frequency stimulation (HFS) which non-selectively affects all structures that are in the vicinity of the electrode, including the STN but also possibly surrounding areas and passing fibers; Right, Optogenetic stimulation in Pitx2-Cre mice injected into the STN with optogenetic virus which allows selective manipulation upon light-stimulation. STN: subthalamic nucleus; pSTN: parasubthalamic nucleus; ZI: zona incerta; cp: cerebral

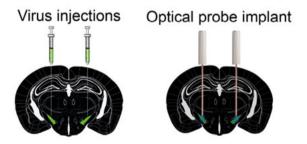
# Surgery and viral injections

All mice in Studies I, II and III received injections of a viral construct to express ChR2, Arch or only eYFP. Mice were anesthetized with a mixture of air/isoflurane (1.4-1.8 % isoflurane/air v/v) and placed in a stereotaxic apparatus. An anti-inflammatory solution (Carprofen, 5 mg/ml) was subcutaneously injected before starting the surgery. After applying a local anaesthetic (Marcain 1:3), the skin was incised to access the skull. Holes were then

drilled above the assigned stereotaxic coordinates. A virus, AAV2-EF1a-DIO-hChR2(H134R)-eYFP or AAV2-EF1a-DIO-Arch3.0-eYFP for experimental mice or AAV2-EF1a-DIO-eYFP for control mice, was injected at 100 nl.min-1 either in the STN (Studies I and II) or in the PHA (Study III). For behavioural experiments, optic cannulas were implanted to achieve intracranial light-stimulation, also referred to as photostimulation. Detailed protocols for each study are described below and in the manuscripts.

In Study I, AAV2-ChR2-eYFP, AAV2-Arch3-eYFP or AA2-eYFP was injected bilaterally in Pitx2-Cre mice in the STN. Mice are referred to as Pitx2/ChR2/eYFP mice, Pitx2/Arch/eYFP mice or Pitx2/eYFP-C (for ChR2 group) or Pitx2/eYFP-A (for Arch group) mice, respectively. Coordinates for the STN were: AP: -1.90 mm, ML: +/- 1.70 mm, DV: -4.65 and -4.25. A volume of two times 250 nl was injected at the two different depths. For behavioural experiments, optic cannulas were implanted above the STN and fixed with dental cement (Figure 10). Mice received another injection of Carprofen 20 to 24 hours after the surgery.

In Study II, AAV2-ChR2-eYFP or the control virus AAV2-eYFP was injected bilaterally in Pitx2-Cre mice in the STN. Coordinates for the STN were: AP: -1.90 mm, ML: +/- 1.70 mm, DV: -4.65 and -4.25 mm. A volume of two times 250 nl was injected at the two different depths. For behavioural experiments, optic cannulas were implanted above either the STN (Pitx2/ChR2/eYFP and Pitx2/eYFP mice) or the VP (Pitx2/ChR2/eYFP/VP and Pitx2/eYFP/VP mice) and fixed with dental cement (Figure 10). Coordinates for optic cannulas in the VP were AP: +0.45 mm, ML: +/- 1.55 mm, DV: -4.00 mm. Mice received another injection of Carprofen 20 to 24 hours after the surgery.



*Figure 10*: Schematic representation of bilateral virus injections and optic canula implantations in the STN of Pitx2-Cre mice.

In Study III, AAV2-ChR2-eYFP or AAV2-eYFP was injected unilaterally in Trpv1-Cre mice in the PHA. Mice are referred to as Trpv1/ChR2 and Trpv1/eYFP mice, respectively. Coordinates for the PHA were: AP: -2.80 mm, ML: -0.60 mm, DV: -4.40 mm, 8° angle in the frontal axis to avoid the sagittal vein and target the midline. A volume of 300 nl was injected. For behavioural experiments, an optic cannula was implanted unilaterally above the PHA and fixed with dental cement. Mice received another injection of Carprofen 20 to 24 hours after the surgery.

# Behavioural experiments

The optogenetic technology for control of neuronal activity was here combined with behavioural testing in order to characterize: 1) The effect on Pitx2-Cre STN neurons upon optogenetic excitation and inhibition on motor (Study 1) and limbic functions, primarily place preference (Study II); 2) the effect on Trpv1-Cre PHA neurons upon optogenetic excitation on behavioural output, primarily place preference (Study III). Four weeks after viral injections and optic cannula implantations, mice were progressively habituated to the experimenter, the testing environment, and the connection to optic fibres. Validated behavioural paradigms were implemented in the context of optogenetic stimulations in the relevant brain area to assess if these had measurable effects on the behavioural display of the mice. The Ethovision software, connected to a camera, was used to record the behavioural output of the mice.

Intra-cranial light-stimulation protocols: For all ChR2-based experiments except study III: 20 Hz, 5 ms pulse duration, 5 mW stimulation. Study III: 40 Hz, 10 ms pulse duration, 10 mW stimulation. For Arch-based experiments: continuous stimulation of 10 mW. Lasers were controlled by an Arduino Uno card, connected to the computer through a TTL box.

In Study I, mice were divided in two groups: 1) The Pitx2/ChR2 group, mice injected with ChR2-eYFP or control (eYFP) virus; 2) the Pitx2/Arch group, mice injected with Arch-eYFP or control (eYFP) virus. Pitx2/ChR2 mice and corresponding controls (Pitx2/eYFP-C) were connected to a blue laser (473 nm, CNI Lasers, Changchun, China) while Pitx2/Arch and corresponding controls (Pitx2/eYFP-A) were connected to a green laser (532 nm, CNI Lasers, Changchun, China).

In Study II, Pitx2/ChR2/eYFP, Pitx2/ChR2/eYFP/VP mice and corresponding controls were connected to a blue laser (473 nm) (CNI Lasers, Changchun, China).

In Study III, Trpv1/ChR2/eYFP mice and corresponding controls were connected to a blue laser (473 nm) (CNI Lasers, Changchun, China).

#### Motor-related tests

#### Open field test

The open field test was performed in Study I on Pitx2/ChR2 and Pitx2/Arch mice and corresponding controls as well as in Study III on Trpv1/ChR2 and respective controls. This test is commonly used test to assess general locomotion, exploratory activity and anxiety and was here used to assess if mice change these behaviours upon optogenetic stimulation. The test arena consists of a square plexiglas transparent box (55x55x22 cm) with a white floor. Centre and borders were virtually delimited (centre corresponds to 25 % of the total arena). The test was divided into four consecutive periods of 5 minutes: OFF-ON-OFF-ON where OFF periods correspond to the absence of intracranial light-stimulation and ON periods correspond to the presence of intracranial light-stimulation. Mice were placed at the centre of the arena and allowed to freely explore for 5 plus 20 min, corresponding to a 5 min habitation time plus the four periods. Distance moved, speed, time spent in centre and frequency in crossing the centre were recorded automatically, while rearing (the mouse standing on the hind legs and raise its forelimbs to explore/investigate a stimulus), self-grooming (the mouse cleans its nose, body and tail with its front paws in a specific sequence of movements) and jumping were recorded manually. All these behaviours involve the motor system and also recruits the limbic system: Rearing is related to exploration; selfgrooming is a stereotyped behaviour which can be a sign of impulsivity and compulsivity when too long and repetitive (Kalueff et al., 2016); a jumping behaviour can be the sign of an escaping behaviour because of aversion, anxiety or pain.

#### Rotarod

The rotarod test used in Study I consists of a rotating rod on which mice can walk. The test is used to assess gait and motor coordination, and was here used to assess if optogenetic stimulation of the STN affects these behaviours. Different speeds of the rotating rod as well as the onset of intracranial light-

stimulation were set by the experimenter. The rotarod test was performed in study I on Pitx2/ChR2 mice and corresponding controls. Mice were trained in four sessions (four days) to walk on the rod at different fixed-speed: 6, 8, 12 and 16 rpm. One trial consisted in a maximum of three attempts and was considered as a success when the mouse could walk on the rod for 120 s, and a failure when the mouse fell prior to full time. No intracranial light-stimulation was applied during the training days. After the training days, only mice that succeeded in the Pre-test could go for the test on the next day. The test day consisted in 4 trials at fixed 16 rpm speed with two OFF periods (no light-stimulation) alternating with two ON periods (light-stimulation) (OFF-ON-OFF-ON). Intracranial light-stimulation started 5 seconds after the mouse was placed on the rod and stopped either when the mouse fell or when the full trial of 120 s had elapsed.

#### Beam walk test

The beam walk test used in Study I measures the ability of mice to walk on a round wooden beam and is used to detect subtle motor skills deficits in motor coordination and balance. This test was performed in study I to assess these parameters upon optogenetic stimulation in Pitx2/ChR2 mice and corresponding controls. The test consisted of several training days where mice learnt to walk on a wooden horizontal 80 cm long beam divided into three successive segments called "starting area" (SA), recording area (RA), and "Goal zone" GZ. Trials were performed during which onset of intracranial light-stimulation differed. During the test day, the latency to fall and hind foot slips were recorded and summarized in a Neuroscore rating scale.

#### Limbic-related tests

#### Elevated plus maze

The elevated plus maze (EPM) used in Study II is a standard test to measure anxiety and exploratory behaviours and was implemented here to assess if these behaviours are affected upon optogenetic stimulation of the STN. corresponding Pitx2/ChR2/eYFP mice and controls well as Pitx2/ChR2/eYFP/VP mice and corresponding controls were assessed. The test arena consists of a 50 cm elevated plus shaped apparatus with two open arms (35 cm) and two closed arms (35 cm) that cross in the middle, forming a connecting centre area (Figure 11). In this test, mice choose between their natural preference for the protected closed arms and their curiosity to explore a new environment. Initially, each mouse was placed in the centre area, facing an open arm and was allowed to freely explore the apparatus for 10 minutes. Intracranial light-stimulation was applied during the whole duration of the test. The time spent and the number of entries in the closed arms, open arms and centre were recorded by the Ethovision software.



Figure 11: Overview of the elevated plus maze experiment with the plus shape of the apparatus on the top left corner and the delimitations of the closed arms (blue), open arms (red) and centre (grev).

#### Elevated plus maze avoidance

The EPM avoidance test is an in-house created variant of the standard EPM described above and was used in Study II on Pitx2/ChR2/eYFP mice and Pitx2/ChR2/eYFP/VP mice and corresponding controls. EPM avoidance was specifically designed in order to test if an optogenetic light-stimulation which induced aversion-like behaviour was more or less potent than the natural aversion for the open arms. Here, light-stimulation was only applied upon entry in any of the closed arms and stopped when the mouse left it. Time spent in closed arms, open arms and centre as well as the number of entries were recorded by the software.

#### Real-time place preference

The real-time place preference (RT-PP) test is an optogenetic variant of the classical conditioned place preference (CPP) paradigm. It is used to measure the preference, or the avoidance, to an optogenetic stimulus in real-time. The RT-PP apparatus used in the current Studies II and III consisted of three connected compartments: Two main compartments with visual and textural differences that serve as cues, and a small, transparent and neutral compartment connecting the two main compartments. One of the main compartments is paired with the optogenetic light-stimulation (light-paired-compartment). That is, intracranial light-stimulation is turned on upon entry into this chamber. Mice were initially placed in the neutral compartment before opening the accesses to the two main compartments whereupon they were allowed to freely explore the arena. If optogenetic stimulation of the neuronal population in question is perceived as rewarding, then mice are expected to spend more time and/or have more entries into the light-paired-compartment, whereas if it is perceived as aversive, less time and/or less entries is expected to take place. The time spent and the number of entries in all compartments were recorded and analysed to detect place preference or place avoidance.

In Study II, Pitx2/ChR2/eYFP, Pitx2/ChR2/eYFP/VP mice and respective controls were used to assess if intracranial light-stimulation of the STN or the VP gives rise to a place preference or place avoidance behaviour. The RT-PP test (Figure 12) consisted of two days for habituation and Pre-test (15 min) without any light-stimulation for reference followed by two days of conditioning (Cond1 and Cond2, 30 min) in which intracranial light-stimulation was applied upon entry into one of the main compartments (the light-paired compartment) and stopped when the mouse left it. This was followed by two test days during which no light was applied but the preference of the mice was recorded. Next followed two reverse conditioning days (Cond3 and Cond4, 30 min) during which the previously unpaired compartment was now paired with intracranial light-stimulation upon entry. This allows a measure of cognitive flexibility. Finally, a last test day, during which no light-stimulation was applied, was performed and place preference assessed

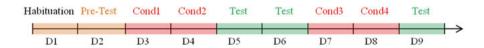


Figure 12: Experimental timeline of the RT-PP test used on Pitx2/Chr2 mice and controls (Study II).

In study III, Trpv1/ChR2 and control mice were analysed in the RT-PP test using a protocol consisting of four conditioning days and no reverse conditioning days (Figure 13). In addition, a stronger intracranial light-stimulation was used in this test: 40 Hz, 10 ms pulse duration and 10 mW light-stimulation protocol.



*Figure 13*: Experimental timeline of the RT-PP test used for Trpv1/ChR2 mice and controls (Study III).

# *In vivo* electrophysiology

Single-cell extracellular recordings were performed at least four weeks after Pitx2-Cre mice were injected with either AAV2-EF1a-DIO-hChR2(H134R)eYFP or AAV2-EF1a-DIO-Arch3.0-eYFP (Studies I and II). Mice were anesthetized with a mixture of air/isoflurane and placed in a stereotaxic frame. After subcutaneous injection of a local anaesthetic (lidocaine), the skin was incised to reveal the skull. Recording electrodes consisted of glass micropipettes (tip diameter 1-2 μm, resistance 10 to 15 MΩ) filled either with pontamine sky blue 0.5 M sodium acetate or neurobiotine 2% in 0.5 M acetate sodium solution. Optic fibres (100 µm diameter, Thorlabs) were either placed directly above the cerebral structure or glued onto the recording pipette to make an optrode. The distance between the tip of the recording pipette and the tip of the optic fibre ranged from 650 to 950 nm. Optic fibres were connected to either a blue (473 nm) or a green (532 nm) laser (CNI Lasers, Changchun, China) for optogenetic excitation or inhibition of Pitx2-Cre STN neurons. For all in vivo experiments, the power of the lightstimulation was calibrated with a power meter (PM100D, Thorlabs) to 5-8 mW for excitation and 9-10 mW for inhibition. Extracellular action potentials were recorded and amplified with an Axoclamp-2B and filtered (300 Hz/0.5 kHz). Single extracellular spikes were collected online (CED 1401, SPIKE2; Cambridge Electronic Design).

After baseline recording for 100 seconds, two different light-stimulation protocols were applied to optogenetically excite the STN in Pitx2/ChR2 mice: a 0.5 Hz, 5 ms light pulse duration protocol performed to create a peri-

stimulus time histogram (PSTH) and a 20 Hz, 5 ms light pulse duration protocol, referred as "behavioural protocol" similar to the light-stimulation protocol used in behavioural experiments. Similarly, different light protocols were applied to optogenetically inhibit STN Pitx2-Cre neurons in Pitx2/Arch mice: 0.5 Hz with 10 ms light pulse duration, 100 ms light pulse duration with 200 ms intervals, 100 ms light pulse duration with 1 s interval, 2 s light pulse duration with 8 s intervals and continuous stimulation for 20-50 s.

## STN optotagging

The objective of STN optotagging was to confirm the excitation and inhibition of STN neurons upon the same light-stimulation protocols that were used in the behavioural experiments. Custom-made optrodes were used to record the neuronal activity of STN Pitx2-Cre neurons in both Pitx2/ChR2 and Pitx2/Arch (Study I and II). Initial coordinates for the STN were: AP: -1.90 mm, ML:+/-1.70 mm and DV: from -4.10 to -4.70 mm. A deposit of sky blue pontamine was made at the last recording coordinates by iontophoresis (continuous current at -20  $\mu A$  for 20 to 30 min) for histological analysis.

## GP recordings

The GP is one of the main targets of the STN. It has been shown that optogenetic stimulation of STN neurons results in glutamate release in the GP (Viereckel, Konradsson-Geuken and Wallén-Mackenzie, 2018). The GP was therefore selected for electrophysiological confirmation of target area excitation upon STN optogenetic stimulation according to the protocols used in behavioural experiments (Study I). Coordinates for the optic fibre in the STN were: AP: -1.90 mm, ML: +/-1.70 mm and DV: -4.30 mm; Initial recording coordinates for the GP were: AP: -0.30 mm, ML: +/-1.50 mm and DV: from -3.00 to -4.30 mm. A deposit of sky blue pontamine was made by electrophoresis at the last recording coordinates for histological analysis.

## LHb recording

The objective was to stimulate STN neurons in Pitx2/ChR2/eYFP mice and record LHb neurons to validate a glutamatergic polysynaptic connection between the STN and the LHb (Study II). An optic fibre was placed above the STN. Upon light-stimulation of the STN, LHb neurons were recorded using a recording glass micropipette filled with either pontamine sky blue or

neurobiotine. Coordinates for the STN were the same as described above and initial recording coordinates for the LHb were: AP: -1.60 mm, ML: +/-0.60, DV: from -2.00 to -3.00 mm.

### 6-OHDA lesions

C57BL/6NTac wild-type mice were used to generate a 6-OHDA parkinsonian mouse model (Study IV). Mice were deeply anesthetized with a mixture of air/isoflurane (v/v), placed in a stereotaxic apparatus and injected subcutaneously with an anti-inflammatory drug (5 mg/ml, Carprofen). After injection of a local anaesthetic (0.17 % Marcain), the skin was incised to expose the skull. A hole was drilled above the coordinates of the median forebrain bundle, MFB: AP: -1.20 mm, ML: -1.10 mm and DV: -4.75 mm. Mice were unilaterally injected with either 1.85 mg of 6-OHDA diluted in a 0.9% NaCl 0.02% ascorbic acid solution or just the 0.9% NaCl 0.02% ascorbic acid solution for control mice. The 6-OHDA solution was kept in darkness during the whole procedure. For post-surgery care, mice received another injection of the anti-inflammatory drug (Carprofen) 20-24 hours after the surgery plus daily subcutaneous injections of saline (1 mL) until sacrifice. Mice were sacrificed at two different time points post-injection: 1) Two weeks postinjection (N=4 6-OHDA injected mice; N=4 control mice) and 2) Three weeks post-injection (N=4 6-OHDA injected mice; N=3 control mice).

In addition to these 15 mice, four additional mice were injected unilaterally with 6-OHDA. Two were used for histological analysis and two for behavioural analysis to estimate the severity of the lesion. One mouse received an intraperitoneal injection of amphetamine (2.5 mg/Kg) and one other mouse received saline. For behavioural assessment, each mouse was placed in a round arena (50 cm diameter) and allowed to habituate to the environment for 10 minutes. Thereafter, the number of ipsilateral turns was counted during 10 minutes. The results were expressed as number of ipsilateral turns per minute. A high ratio corresponds to a high percentage of DA neurons degeneration. Two or three weeks after injections, the cohort of 15 injected mice were sacrificed by cervical dislocation, brains extracted and snapped-frozen in -25/-30°C isopentane. Brains were cut with a cryostat and mounted on glass slides. Sections were stored at -80°C until usage.

# Histological analyses

## Immunohistochemistry

Upon completion of behavioural experiments, Pitx2-Cre mice (Studies I and II) and Trpv1-Cre mice (Study III) were transcardially perfused with 1x phosphate buffer saline (PBS) followed by 4% formaldehyde. Brains were extracted, post-fixed and 60 µm-thick sections were cut at the vibratome. Immunohistochemistry using standard protocols was performed to enhance the eYFP signal (Studies I, II and III) and to detect c-fos (Study I), TH (Study III) and neurobiotine (Study II). Pontamine sky blue naturally emits in the red part of the spectrum, no enhancement was needed (Study I and II). Slices were stained with DAPI for detection of cell nuclei, mounted and scanned using a NanoZoomer 2-0-HT.0 (Hamamatsu) scanner. Results were visualized using the NDPView2 software (Hamamatsu).

### *In situ* hybridization

*In situ* hybridization experiments were performed to detect mRNA using colorimetric (Study III, performed by other co-author) and fluorescent (Study III, IV) labelling.

In study IV, the selected mRNAs were standard markers for midbrain DA neurons: Th. Aldehyde dehydrogenase 1 (Aldh1a1), Calbindin1 (Calb1) and Gastrin-releasing peptide (Grp). In situ hybridization was performed on 16 um-thin frozen sections. Sections were thawed and post-fixed in 4% paraformaldehyde (4% PFA) for 10 min at room temperature (RT). After washes in PBS, sections were incubated in triethanolamine pH 8 (TEA) for 5 min followed by acetylation. Riboprobes (DIG labelled riboprobe, 50-75 ng/100 μl and fluorescein labelled riboprobe, 75-100 ng/100 μl) were denatured in a solution of 50% hybridization buffer / 50% formamide at 85°C for 10 min and thereafter applied onto the sections for a 16-18 hours in a humidified chamber at 65°C. After hybridization, sections were washed in 65°C saline sodium citrate buffer (SSC) baths (5X SSC followed by 0.2X SSC solutions) and a last wash with a 0.2X SSC at RT. Sections were then washed in 1x maleic acid buffer containing TritonX-100 (MABT) to decrease non-specific probe bindings. The fluorescein riboprobe was revealed by first incubating the sections in a blocking solution containing a blocking reagent (Roche), FBS heat inactivated and 5xMAB. Then, sections were incubated with an anti-fluorescein horseradish peroxidase-conjugated (POD) antibody (Roche) for one hour. An amplification step was performed by incubating the sections in TSA-biotin amplification buffer (Perkin Elmer) for 15 min. The fluorescein-TSA-biotin complex was revealed by adding Neutravidin-Oregon Green (Invitrogen) for 15 min. After rinsing the sections in TritonX-100 PBS (PBST), hydrogen peroxidase was inhibited by incubating the sections in 0.1M glycine pH 2.1 and in 3% hydrogen peroxide (H<sub>2</sub>O<sub>2</sub>). A similar procedure was performed to reveal the DIG riboprobe with an anti-DIG-POD antibody diluted in the blocking solution and TSA buffer plus Cy3 (Perkin Elmer). Sections were subsequently incubated for 10 min in 1/50000 DAPI solution and mounted with Fluoromount (Southern Biotech). Sections were scanned with NanoZoomer 2-0-HT.0 scanner using the NDP.scan 3.3 software (Hamamatsu) and visualized with the NDP.view2 software (Hamamatsu).

# Statistical analysis

### Behavioural experiments

For the open field test, EPM and RT-PP, statistical analysis was performed using a 2-way repeated measures (RM) ANOVA followed, when the p-value was below 0.05, by a Tukey or a Bonferroni Post-hoc analysis. In the open field test (Study I), a two-tailed Wilcoxon matched-pairs test was used to compare the number of jumps between Pitx2/ChR2 and control mice. Measured parameters were compared between days, compartments and the interaction between days and compartment for RT-PP experiments (Studies II and III). For EPM and EPM avoidance tests (Study II), the number of entries and the time spent in each arm were compared between groups (ChR2 *vs.* Controls), arms and the interaction between groups and arms. For the rotarod and the beam walk test (Study I), a Friedman test was used to analyse the data followed by Dunn's multiple comparison when needed.

All data is expressed on the plots as means  $\pm$  SEM; when appropriate, data was averaged for the two ON and OFF trials.

# In vivo electrophysiology experiments

For extracellular *in vivo* recordings in the GP (Study I) and the LHb (Study II), the non-parametric Friedman test was used to compare the firing rate

before, during and after optogenetic stimulation followed by a Dunn's multiple comparison.

# Study I

## Aim

The aim was to characterize the role of the STN in several aspects of movement by systematically applying optogenetic excitation and inhibition to the STN in Pitx2-Cre mice and analysing circuitry effects as well as impact on motor behaviour.

#### Results and discussion

The STN is a central excitatory nucleus within the indirect pathway of the basal ganglia. Despite knowledge of the role of the STN in movement in relation to PD, little is known on the natural role of the STN in the healthy subject. However, a few studies have shown that activation of the STN decreases locomotion, induces stereotyped movements and can interrupt an ongoing behaviour while its inactivation facilitates movement, induces contralateral rotations and suppresses the interruptive effect of surprise (Bergmann *et al.*, 2004; Schweizer *et al.*, 2014; Fife *et al.*, 2017; Tian *et al.*, 2018). None of the studies mentioned above investigated the effect of a direct optogenetic excitation or inhibition of the STN on general locomotion, coordination and fine motor movements

In this study, we used optogenetics in Pitx2-Cre mice to ensure selectivity for the STN over surrounding structures. By expressing excitatory (ChR2) and inhibitory (Arch) opsins in the STN structure, the STN could be directly, selectively and reversibly excited or inhibited, respectively. Control mice lacked the opsin in their optogenetic constructs but where otherwise identical to the experimental mice.

First, we confirmed STN projection target areas by visualization of eYFP-reporter positive fibres. Strong eYFP signal was visible in the GP, SNr and EP and weaker signal was observed in the VP, SNc and PPN.

To validate our optogenetic approach in Pitx2/ChR2 mice, we analysed c-fos expression in the STN after light-stimulation in the STN itself. We observed the presence of c-fos-positive neurons in the STN, below the tip of the optic fibre in Pitx2/ChR2 mice. As expected, no c-fos expression was found in control mice. Furthermore, we confirmed the excitation of Pitx2-Cre STN neurons in Pitx2/ChR2 mice by using in vivo extracellular electrophysiology recordings in the STN and in the GP. By simultaneously stimulating and recording STN neurons (optotagging), we could observe light-evoked action potentials when applying a 0.5 Hz 5 ms 5-8 mW stimulation protocol. In the GP, one of main targets of the STN, we observed both light-evoked action potentials with a 0.5 Hz 5 ms 5-8 mW light-stimulation protocol and an increase in the firing rate when using the same light-stimulation protocol as the one used for behavioural experiments (20 Hz, 5 ms, 5-8 mW). The increase in frequency lasted the whole duration of the light stimulation (100 s) with a return to baseline as soon as the light was off. Electrophysiological recordings in Pitx2/Arch mice were performed but could not confirm the inhibition of STN Pitx2-Cre neurons due to their low spontaneous activity.

Next, we assessed the effect of STN excitation and inhibition in different behavioural tests measuring various aspects of motor function. Results from the open field test showed that optogenetic excitation and inhibition of the STN has opposite effects on horizontal and vertical locomotion. Pitx2/Arch mice displayed an increased locomotor activity with increased distance moved, speed, time spent moving and rearing. In contrast, Pitx2/ChR2 mice showed a decrease in distance moved, speed, time spent moving and rearing. Surprisingly, a strong face-grooming was observed in Pitx2/ChR2 mice only seconds after the intracranial light-stimulation was ON. In contrast, Pitx2/Arch mice showed a decrease in the natural levels of self-grooming compared to controls. The results confirmed the long-assumed role of the STN in movement. That is, STN excitation has a reducing effect on locomotion while STN inhibition has an elevating effect on locomotion. The excessive face-grooming observed upon light-stimulation in Pitx2/ChR2 mice was surprising but is in accordance with results obtained by Tian and colleagues who showed that activation of glutamatergic STN terminals in the GP induces grooming (Tian et al., 2018).

To further investigate the reduced motor activity observed in Pitx2/ChR2 mice upon optogenetic excitation, we tested their general and fine motor coordination using the rotarod and beam walk tests. Upon light-stimulation,

Pitx2/ChR2 mice failed to walk on the rotating rod more than few seconds. Further, compared to control mice, Pitx2/ChR2 mice had difficulties in crossing the beam when light-stimulation was applied shortly prior to the test.

Finally, to further assess the impact of excitation *vs.* inhibition of the STN in basal ganglia function, we tested our hypothesis that unilateral light-stimulation of the STN in Pitx2/ChR2 and Pitx2/Arch mice would induce opposite rotations. According to previous studies and the classical basal ganglia model, activation of the STN would inhibit movement on the side of the stimulation and induce ipsilateral rotations, while inhibition of the STN would promote movement on the side of the stimulation and induce contralateral rotations. This was indeed what we observed: Unilateral optogenetic excitation or inhibition of the STN induced a strong rotational behaviour with ipsilateral rotations in Pitx2/ChR2 mice and contralateral rotations in Pitx2/Arch mice.

The results obtained in this study experimentally confirm the expected output upon STN excitation and inhibition as postulated by basal ganglia models. We show that optogenetic activation of the STN indeed reduces movement while its inhibition promotes movement. Beyond this finding, we also identify impaired motor coordination upon STN excitation, and, quite surprisingly, STN excitation directly and immediately induces face-grooming behaviour.

# Study II

### Aim

The primary aim of this study was to investigate the role of the STN in limbic functions related to reward and aversion. Upon identification of place avoidance induced by optogenetic STN excitation, the follow-up aim was to address putative neurocircuitry engaged in this unexpected behaviour.

#### Results and discussion

The STN has long been investigated in motor functions because of its role in PD. However, it has also been shown that the STN regulates reward-related behaviours and, more precisely, that it encodes positive or negative values (Baunez *et al.*, 2005; Rouaud *et al.*, 2010; Pelloux *et al.*, 2014; Breysse, Pelloux and Baunez, 2015; Schweizer *et al.*, 2016). Besides some evidence showing a role of the STN in encoding negative values (Breysse, Pelloux and Baunez, 2015), no study has ever shown a direct correlation between activation of the STN and aversive-like behaviours. Furthermore, based on the finding in Study I that STN excitation causes self-grooming behaviour, a putative correlate to compulsive behaviour in human OCD, it was of interest to assess if STN excitation caused aversion or anxiety or other types of affective or cognitive behaviours that could explain the stereotyped grooming.

We first applied optogenetics in the STN of Pitx2-Cre mice to excite this structure selectively. We validated our excitatory optogenetic approach with three complementary methods: We confirmed the presence of eYFP cell bodies in the STN and eYFP projections to the STN target areas; *in vivo* and *ex vivo* electrophysiological recordings in the STN (optotagging) were both implemented to confirm excitation of Pitx2/ChR2 STN neurons upon light-stimulation

Next, we tested the effects of STN optogenetic excitation in the RT-PP paradigm. Depending on the effect of the stimulation (pleasant or unpleasant),

mice would spend more or less time in the light-paired compartment. We observed a decrease in the time spent in the light-paired compartment and consequently an increased time spent in the unpaired-compartment. These results showed for the first time that excitation of the STN induces place avoidance

To determine if anxiety plays a role in this behaviour, we decided to test Pitx2/ChR2 mice in the elevated plus maze, EPM. STN-directed light-stimulation did not affect the time spent in either open or closed arms in Pitx2/ChR2 mice and controls. However, when applying the light-stimulation only in the closed arms (EPM avoidance), Pitx2/ChR2 mice avoided the open arms, naturally aversive for mice, and also the closed arms, where the light-stimulation was applied. The result was an increased time spent in the centre of the apparatus, in the light-unpaired cross between closed and open arms. These results confirmed that STN excitation induces strong avoidance behaviour.

Secondly, we hypothesized, based on previous studies, that this avoidance behaviour was mediated through a pathway reaching the LHb (Tan *et al.*, 2011; Hartung *et al.*, 2016; Stephenson-Jones *et al.*, 2016; Faget *et al.*, 2018; Tooley *et al.*, 2018). The STN projects directly to the VP and the EP which in turn project to the LHb. We decided to investigate the STN  $\rightarrow$  VP  $\rightarrow$  LHb pathway, first because it has never been shown that VP neurons receiving glutamatergic inputs from the STN project to the LHb, second because the VP is well known for its role in limbic functions, and third, because it was recently shown that activation of VP glutamatergic neurons projecting to the LHb induces place avoidance (Tooley *et al.*, 2018).

We first wanted to demonstrate if an indirect projection from the STN to the LHb exists at all. To do so, we used *in vivo* extracellular recordings with an optic fibre stimulating the STN and a recording pipette placed in the LHb. We observed evoked action potentials in the LHb in 50% of the recorded LHb neurons with 10.08 ms +/- 0.81 ms delay. Moreover, patch-clamp recordings showed excitation of VP neurons upon light stimulation, confirming STN glutamatergic projections on the VP.

Next, a new batch of Pitx2/ChR2 mice, in which ChR2 is expressed in the STN, were tested in the RT-PP and EPM paradigms, but now, the intracranial light-stimulation was directed to the VP instead of the STN. This would

allow us to selectively study the behavioural effect upon optogenetic excitation of the STN-VP terminals (Pitx2/ChR2/eYFP/VP mice). Also these new mice showed a decrease in the time spent in the light-paired compartment. Results in the EPM and EPM avoidance were also similar to the ones obtained in Pitx2/ChR2/eYFP mice, when the STN itself was stimulated. These results showed for the first time that the STN-VP pathway is involved in place avoidance.

Additional experiments analyzing this newly identified circuitry could be performed to confirm the role of this pathway in aversion and demonstrate the functional connection between the STN, the VP and the LHb. Also additional pathways between the STN and the LHb might contribute to the processing of aversive stimuli.

The main take-home message of this study is that excitation of the STN induces aversive behaviour, a finding which has potential clinical importance, not least in the STN-DBS context in which adverse side-effects include affective symptoms, including anxiety and depression.

# Study III

### Aim

The aim of this study was to characterize the neuronal population in the hypothalamic-mesencephalic area which expresses the *Trpv1* gene, its projection pattern and its functional role in limbic functions.

## Results and discussion

Several evidence support the hypothesis that different populations within the hypothalamic and mesencephalic areas mediate aversion and avoidance-type behaviours. Indeed glutamatergic, GABAergic and dopaminergic subpopulations of the VTA and hypothalamus have been shown to be coding for aversive or reward-related behaviours (Pupe and Wallén-Mackenzie, 2015; Chen et al., 2017; Morales and Margolis, 2017; Sharpe et al., 2017; Bimpisidis et al., 2019; de Jong et al., 2019; Lazaridis et al., 2019). The identification of such populations is of importance to improve our understanding of brain function in healthy and pathological conditions.

In this study, we describe the expression of a very restricted hypothalamic-mesencephalic neuronal population defined by its expression of the *Trpv1* gene from developmental stages. At E14.5, Trpv1 mRNA was found in Th-positive VTA neurons and in the PHA. Trpv1-positive neurons were mostly expressing *Vglut2* in the medial hypothalamic area and expressed *Th* in the VTA with some scattered neurons positive for Vglut2 mRNA. Trpv1 mRNA was not detected in any other brain regions. At P3, Trpv1 neurons were present along the midline from the PHA to the VTA. Trpv1 mRNA was mainly co-localized with Th mRNA in the medial nuclei of the VTA (IF, PN, PBP) and Trpv1/Vglut2 co-localization was also identified. In more rostral aspects of the VTA and the PHA, Trpv1 mRNA was mostly co-localized with Vglut2 mRNA. In the adult, Trpv1 mRNA was similarly distributed as in P3 but at a much lower level, making it nearly undetectable.

Next, we hypothesized that this Trpv1 subpopulation could play a role in aversion based on previous studies showing that glutamatergic neurons of both the midbrain and the hypothalamus (lateral hypothalamic area) are involved in aversion (Root *et al.*, 2014; Lammel *et al.*, 2015; Lazaridis *et al.*, 2019).

To selectively manipulate this neuronal population, we used Trpv1-Cre mice, originally imported from the Jackson Laboratory, and that we injected with optogenetic viruses, creating Trpv1/ChR2 mice. Firstly, by analysis of eYFP-positive projections throughout the brain, we identified the various brain areas innervated by Trpv1-Cre neurons. We observed strong eYFPpositive projections to the septal area, preoptic areas, CA1 and CA3 fields of the hippocampus, and the amygdalohippocampal nucleus (AHiAL). We also identified weaker but consistent projections to the endopiriform nucleus (DEn) and the fimbria of the hippocampus, as well as weaker and less consistent projections to the BNST, the dorsal tenia tecta (DTT), the mAcbSh and along the midline in the orbital and infralimbic cortices. Projections reaching forebrain structures were going through the median forebrain bundle, MFB, where transversally-sectioned eYFP fibres were visible in most injected Trpv1-Cre mice. Overall, the amount and density of projections was surprisingly high compared to the small amount of Trpv1-Cre cell bodies present in the hypothalamic-mesencephalic area, into which the opsin constructs were injected.

Secondly, we tested Trpv1/ChR2 mice in an RT-PP paradigm. According to the theory of this paradigm, if light-stimulation of Trpv1-Cre neurons of the hypothalamic-mesencephalic area is aversive, we should observe a decrease in the time spent in the light-paired compartment. Along the four consecutive conditioning days, Trpv1-Cre mice indeed decreased the time spent in the light-paired compartment, reaching significant difference on the fourth conditioning day. These results confirmed the involvement of Trpv1-Cre population in place avoidance. However, more studies will be required to fully understand this behaviour which was found progressive and which did not seem to result in significant learning.

Further experiments will also be needed to identify the pathways that support the observed place avoidance. In particular, it would be interesting to stimulate Trpv1-Cre terminals in septal or preoptic areas, two of the main target areas we observed for Trpv1-Cre neurons, and which are strongly involved in limbic functions with projections to the LHb (Yetnikoff *et al.*, 2015; Barker *et al.*, 2017).

# Study IV

## Aim

The aim of this study was to generate a 6-OHDA mouse parkinsonian model to enable future investigations of optogenetic excitation *vs.* inhibition of the STN in a parkinsonian paradigm.

## Results and discussion

The 6-OHDA parkinsonian mouse model is one of the most used methods to study PD. When injected intracerebrally into the MFB, the 6-OHDA neurotoxin induces a degeneration of DA neurons of the SNc and consequently of the nigrostriatal pathway. In accordance with the loss of SNc DA neurons, this parkinsonian model causes strong motor impairments. However, alphasynuclein aggregates are not observed in the 6-OHDA model.

In study I, we showed that optogenetic inhibition of Pitx2/Arch mice increased locomotor activity. In this study, we tested protocols to create a 6-OHDA mouse model to be able to, in future experiments, test if selective optogenetic inhibition of the STN can improve toxin-induced motor impairments. The model is also useful to compare the effect of STN inhibition with STN excitation, in order to evaluate STN stimulation paradigms. These types of experiments would allow further study of how selective STN manipulations might be used to rescue parkinsonian motor deficiency. Indeed, one previous study showed that optogenetic inhibition of the STN in 6-OHDA lesioned rats improves contralateral forelimb akinesia and levodopa-induced dyskinesia (Yoon *et al.*, 2016). In addition, we aimed to optimize the method to increase the survival rate and well-being of the mice by implementing two different time-duration protocols associated with post-operative cares. Also the amount of toxin was reduced compared to most studies, in order to improve the well-being of the mice.

First, we observed that unilateral injection of 6-OHDA in the MFB rapidly induced strong motor impairments characterized by spontaneous ipsilateral rotations, akinesia and bradykinesia. Despite these strong motor symptoms, the survival rate reached 88% (N=15/17 mice). This was likely a direct consequence of daily post-operative cares which lasted until sacrifice. Sacrifice took place either two or three weeks post-surgery.

Second, we compared the degeneration of DA neurons between mice sacrificed two or three weeks post-lesion. For both groups, we showed a strong decrease of TH immunoreactivity in the SNc and dorsal striatum (near 100%). However, we also noticed a partial decrease of TH in the VTA with the ventro-lateral part affected by the lesion while the medial nuclei of the VTA remained intact. In accordance with the partial degeneration of VTA DA neurons, the ventral striatum was also partially affected. The mAcbSh was nearly intact while only weak TH labelling was observed in the lateral nucleus accumbens shell (LAcbSh) and the nucleus accumbens core (AcbC).

In situ hybridization experiments showed similar results with a strong decrease of Th mRNA in the SNc and the ventro-lateral part of the VTA. No obvious differences were visible in Th mRNA between mice sacrificed two or three weeks after 6-OHDA injection. Besides Th, we used additional molecular markers for midbrain DA neurons (Grp, Calb1 and Aldh1a1) to identify potentially spared subpopulations. These markers are more restricted to the VTA than the SNc. As expected, Aldh1a1 mRNA was nearly absent in the SNc on the lesioned side of 6-OHDA mice. In the VTA, we observed the same pattern as for Th mRNA with a lower density of Calb1, Grp and Aldh1a1 in the lateral part.

To conclude, a protocol for generation of the parkinsonian 6-OHDA model in mice was formulated. We found that sacrifice only two weeks post-lesion is sufficient to induce a strong degeneration of SNc DA neurons and partial degeneration of VTA DA neurons, associated with the expected motor symptoms, while maintaining a high survival rate and well-being.

Further experiments will be necessary to allow implementation of this parkinsonian model in Pitx2/Arch mice. Indeed, mice would have to go through a surgery combining 6-OHDA injection, AAV-Arch-YFP injection and optic cannula implantation.

# Concluding remarks

This doctoral thesis aimed to investigate the role two neuronal populations in motor and limbic functions by using optogenetics, one expressing *Pitx2* and restricted to the STN and the other expressing *Trpv1* and restricted to the ventral hypothalamic-mesencephalic area.

First, we showed through in vivo and ex vivo electrophysiological experiments that optogenetic excitation induced excitation of STN neurons and its target structures in Pitx2-Cre mice. Next, we demonstrated that optogenetic excitation and inhibition of the STN induced opposite motor effects in a way that confirmed classical models of its role in the basal ganglia indirect and hyperdirect pathways. Surprisingly, selective STN excitation induced a robust self-grooming response and strong place avoidance. This is the first time that STN activation has been related with such an aversive-related behaviour. We hypothesized that this behaviour could be mediated through indirect connections to the LHb, a structure well-known for its role in aversion. Because the VP receives projections from the STN and, in turn, projects to the LHb, we hypothesized that an STN→VP→LHb pathway could be involved in the observed place-avoidance. In a follow-up study, we first confirmed, by using in vivo electrophysiology, an indirect excitatory connection between the STN and the LHb. Further, behavioural experiments indeed demonstrated place-avoidance behaviour when selectively exciting glutamatergic STN terminals in the VP. Together, these two studies are the first to experimentally provide direct evidence based on non-pathological conditions for the much-assumed role of the STN in motor regulation and to identify a direct role for the STN in inducing avoidance-related behaviour. These findings advance current knowledge of the STN and its neurocircuitry in behavioural regulation. In the context of PD, OCD and additional conditions in which the STN has been implicated, these revelations should help towards improving selectivity in current treatments based on electrical stimulations within the subthalamic area as well as provide increased knowledge as to the nature of aversive effects seen upon STN dysregulation and STN-based electrical treatment strategies.

Additional structures contribute to the aversion network in the brain, including the lateral aspect of the hypothalamus and the ventral midbrain, areas in which glutamatergic neurons defined by VGLUT2 reside. While the STN is anatomically close to the hypothalamus, a population of neurons expressing *Trpv1* is present in both the posterior aspect of hypothalamus and the ventral midbrain. This is a small population of heterogeneous neurotransmitter phenotype, primarily positive for VGLUT2. We could show that also this population, when stimulated with optogenetic excitation, induced place preference, albeit not as potent as when stimulation was aimed at the STN. The STN and the PHA are anatomically distinct but share their glutamatergic neurotransmitter phenotype. When comparing the projection target areas of the neurons expressing ChR2-eYFP upon stereotaxic injections in each respective area (STN and PHA), no joint target area was found, however, both structures projected to areas that in turn project to the LHb (Table 2).

Projection target areas/structures	Pitx2-Cre	Trpv1-Cre
Infralimbic and orbito-frontal cortices		X
mAcbSh		X
Septal areas		X
Preoptic areas		X
VP	X	
BNST		X
fimbria of the hippocampus (fi)		X
GP	X	
EP	X	
DEn		X
AHiAL		X
SNc	X	
SNr	X	
CA1 and CA3 fields of the hippocampus		X

Table 2: Summary of the structures/areas innervated by STN Pitx2-Cre neurons and Trpv1-Cre neurons of the hypothalamic-mesencephalic area as detected by histological analysis of eYFP reporter gene expression upon viral delivery of optogenetic constructs (Studies I-III). In dark purple, areas innervated by STN Pitx2-Cre neurons that project to the LHb. In light purple, areas innervated by Trpv1-Cre neurons that project to the LHb. mAcbSh: medial part of the nucleus accumbens shell; VP: ventral pallidum; BNST: bed nucleus of the stria terminalis; GP: globus pallidus; EP: entopeduncular nucleus; Den: endopirifom nucleus; SNr and SNc: substantia nigra pars reticulata and pars compacta; AHiAL: amydalohippocampal area.

Finally, to allow further study the effects of optogenetic inhibition of the STN in pathological conditions, a protocol for the 6-OHDA parkinsonian model was validated in mice. We hypothesise that inhibition of the STN would induce similar beneficial effects as STN-lesion by suppressing the overactivation of the indirect pathway and promoting movement.

To conclude, the results presented in this thesis add important insight to understand the role of the STN on motor control and limbic functions. In particular, we suggest that the STN might act as an important regulatory region upstream of the LHb in the neurocircuitry responding to aversive stimuli. A better comprehension of the basal ganglia and its role in executive and affective functions in healthy and pathological conditions is relevant for improving treatment prospects for PD and OCD.

# Future perspectives

The studies presented in this thesis increase the current understanding of the role of the STN in motor and affective functions. However, further experiments will be needed to fully explain the functional organization of the STN underlying these diverse functions. Follow-up studies could, for example, use molecular and viral tracing to further investigate the connection between the STN and the LHb. Also, the combination of optogenetics and Designer Receptor Exclusively Activated by Designer Drugs (DREADDs) and/or pharmacological substances could allow us to confirm the role of the STN → VP → LHb pathway in aversion.

Finally, the internal organization of the STN remains to solve. This will be critical to understand the functional connection between the STN and behavioural regulation. Does the STN consist of the three different domains hypothesised in the STN tripartite model or are neurons intermingled into subpopulations? While the STN can be identified histologically by expression of Pitx2, Vglut2 and additional genes, no molecularly distinct domains had for long been identified. Only this past summer was the first study published which could demonstrate the presence of distinct sets of gene expression patterns

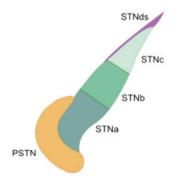


Figure 14: Schematic representation of the four internal domains recently identified within the mouse STN (Wallén-Mackenzie et al., 2020)

that anatomically identify domains within the STN (Wallén-Mackenzie *et al.*, 2020). Using Pitx2-Cre mice crossed with a floxed reporter and by implementing single-nuclei RNA sequencing, several clusters of genes were identified as differentially expressed. Upon histological analysis and anatomical mapping, it was clear that several genes show restricted expression within the STN structure, and by composite analysis, spatio-molecular maps

could be created revealing four distinct domains (STNa, STNb, STNc, and STNds, (Figure 14). In addition to providing new information supporting the presence of distinct domains in the mouse STN, these patterns should allow exciting new possibilities of selective optogenetic targeting *within* the STN using the newly identified promoters as selectivity-drivers in future Cre-Lox-based research applications centred around the STN.

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