

Decreased *in vivo* $\alpha 2$ adrenoceptor binding in the Flinders Sensitive Line rat model of depression

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ABSTRACT

Depression is a debilitating heterogeneous disorder and the underlying mechanisms remain elusive. Alterations in monoaminergic neurotransmission, including noradrenergic, have been implicated in the etiology of depression. Although depression is difficult to model in animals, the availability of animal models with face, predictive and construct validity permits more in-depth investigations resulting in a greater understanding of the disease. We investigated the role of noradrenaline (NA) and $\alpha 2$ adrenoceptors *in vivo* in a genetic model of depression, the Flinders Sensitive Line (FSL) rat. We determined baseline differences in NA receptor volume of distribution to $\alpha 2$ adrenoceptors in FSL, in comparison with two routinely used controls, Flinders Resistant Line (FRL) and Sprague–Dawley (SD) rats using positron emission tomography (PET) imaging and the carbon-11 labeled radioligand yohimbine. We demonstrate a 42–47% reduction in the binding of the tracer in the cortex, striatum, cerebellum, thalamus and pons of FSL rats compared to the two control groups. Our results suggest that the behavioral deficits expressed in the FSL depression model are associated with functional over-activity of the NA system.

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1. Introduction

According to the World Health Organization, clinical depression is estimated to debilitate 350 million people globally (Marcus et al., 2012). The core symptoms of depression are anhedonia, behavioral despair, changes in appetite, weight loss or weight gain, neuroendocrine disturbances, alterations in sleep architecture and anxiety-related behaviors (American Psychiatric Association, 2013). The last decades have seen many advances in therapeutic recourses and outcomes but the etiology remains elusive and is likely highly heterogeneous: while some presentations appear to have a more

endogenous nature, without apparent behavioral causes, others appear triggered by the cumulative effects of traumatic life events. It is unknown if the underlying neurochemistry varies with individual behavioural phenotypes, e.g. if an individual presenting with anxiety, irritability or high level of alarm and activity state has the same neurotransmitter imbalances as individuals with hopelessness and sadness. Similarly, there is no uniform pathology: the depressive disorders are heterogeneous, not associated with clear neuronal loss, and may stem more from a combination of genetic predisposition and environmental influences leading to neuronal dysfunction rather than obvious pathology (Kessler, 1997; Klengel and Binder, 2013). However, several decades of intense pharmacological research have demonstrated the contribution of monoaminergic transmission to the disease (Manji et al., 2001; Schildkraut, 1965).

Most effective pharmacological antidepressants affect to some degree one, and often more, of the main monoamines: serotonin (5HT), noradrenaline (NA) and dopamine (DA) (Marks et al., 2008). While the serotonin system in mood disorders has been subjected

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to scrutiny because of the empirical observation of the normalizing effects of most pharmacological antidepressants on serotonin levels, the relationship between α_2 adrenergic receptors and therapeutic effects has been recognized for decades (Cohen et al., 1980; Cottingham and Wang, 2012), and yet, the status of the α_2 adrenoceptor in depression remains unclear.

One of the limitations to the study of depression is the lack of appropriate animal models. There is no current adequate model of depression in animals that recapitulates the wide array of symptoms. Existing models use selected lesions, genetic manipulation, systemic or local administration of drugs to alter the function of specific neurotransmitters or brain regions, or application of pre-natal/early natal or long-term behavioral stressors to produce some of the behavioral observations, but without achieving the complete picture. In depression, it is however possible to use partial models to answer focused questions, for example to study neuroplastic changes or the involvement of monoamines in etiology, progression or therapeutic usefulness.

Among the various depression/anxiety rat models, the Flinders sensitive line (FSL) has many advantageous features. This rat model was developed by the selective breeding of Sprague Dawley (SD) rats for increased responses to an anticholinesterase agent (Overstreet, 1986; Overstreet and Russell, 1982). The Flinders resistant line (FRL) demonstrates a behavioral response similar to that of the original SD and is often used as a control line in study of FSL animals. Behaviorally, FSL rats are less active in novel environments, display sleep disturbances, reduced saccharin preference, and increased responsiveness to stress eg. anhedonia-like symptoms or increased immobility in the forced swim test, features reversible by chronic, but not acute, treatment with antidepressant drugs (Overstreet et al., 2005). FSL rats have reduced serotonin synthesis in the raphe nuclei and limbic areas implicated in depression (Hasegawa et al., 2006). Furthermore, the behavior of FSL rats improves with tricyclic antidepressants, serotonin reuptake inhibitors (SSRI) and ECT therapy (Chen et al., 2010; Jimenez-Vasquez et al., 2007). Interestingly, FSL rats also have lower vesicular monoamine transporter (VMAT2) in striatal and limbic areas (Schwartz et al., 2003). Taken together, the FSL rat is a genetic model of depression with good face and predictive validity (Overstreet et al., 2005; Overstreet and Wegener, 2013). However, while the role of NA in depression is well accepted, and desipramine is often used in this model as an antidepressant of choice (Roth-Deri et al., 2009; Zangen et al., 1999), no studies to date have directly investigated *in vivo* the role or status of NA and its receptors in FSL rats.

We have recently developed [^{11}C]yohimbine, an antagonist of the α_2 NA receptors, as a potential tracer for positron emission tomography (PET) brain imaging studies (Jakobsen et al., 2006). Yohimbine is a stimulant alkaloid naturally found in several plants. It has been used as a weight loss dietary extract and to treat sexual dysfunction as well as an adjunct to antidepressant therapy (Tam et al., 2001). Despite some antihypertensive properties, it generally increases blood pressure at rest, which is thought to be mediated via central antagonism of α_2 adrenergic receptors (Biaggioni et al., 1994). Our earlier studies have found that in tracer concentrations, yohimbine exhibits high selectivity for α_2 sites *in vivo* in pigs (Jakobsen et al., 2006) which is displaceable by amphetamine challenge (Landau et al., 2012a). Central α_2 NA receptors are widely expressed pre-synaptically primarily on NA cell bodies and dendrites in the locus coeruleus (LC). They are also widely expressed, mostly post-synaptically in every projection area of the NA neurons, throughout the entire cerebrum. As a consequence, high specific activity yohimbine is a valuable tool to assess relative alterations in the regional distribution and density of central α_2 receptors. In this study, we used [^{11}C]yohimbine to

investigate the differences in α_2 adrenoceptor binding in the FSL rat model of depression compared to control FRL and SD rats.

2. Materials and methods

2.1. Animals

Adult female FSL and FRL rats from breeding colonies at the Centre for Psychiatric Research, and Sprague Dawley (SD) rats from Taconic were used in this study (220–260 g). The animals were pair-housed and the individuals were issued from different dams. The FSL and FRL animals were bred in-house in same rooms and conditions and the SD were purchased as young animals and spent several weeks in the same environment as the FSL and FRL prior to scanning. The animals were kept on a normal 12-h light/12-h dark cycle and given free access to food and water. The study protocol was approved and regulated by the Danish Committee on Ethics in Animal Experimentation (authorization number: 2007/561-1378) and all efforts were made to reduce the number of animals used in this study and to minimize suffering.

2.2. MicroPET imaging

The rats ($N = 6$ per group) were initially anesthetized in an isoflurane (2%) chamber. An arterial catheter was placed in the femoral artery. The rats were then positioned prone in a plastic stereotaxic frame designed to fit in the gantry of the tomograph (microPET R4, CTI, Concorde Microsystems). Anesthesia was maintained with 1.8–2% isoflurane delivered through a mask fitted to the head holder throughout the procedure. As yohimbine is a substrate for the p-glycoprotein transport system of the blood brain barrier in rodents (Pearce et al., 1989), a non-specific p-glycoprotein inhibitor, cyclosporine-A (50 mg/kg IV) was administered 30 min prior to tracer injection to facilitate the penetration of the radioligand into the brain. After a 10 min transmission scan, the dynamic 90 min emission recording was initiated after bolus injection over 15–20 s of approximately 30–40 MBq (100–200 μL volume; injected mass 0.06–0.8 μg) of high specific activity [^{11}C]yohimbine. Arterial sampling was performed to determine the input function for data analysis about every 15 s during the first two minutes of the scan then at 3, 5, 10, 20, 30, 45, 60, 75 and 88 min post tracer injection. Three drops of blood (about 150 μL) were sampled at each time and was replaced with an equivalent volume of sterile saline. Body temperature during the scans was maintained around 36–36.5 °C using a heat lamp.

2.3. Analysis and statistics

A decay-corrected plasma time activity curve was produced from the arterial samples obtained during the study. Attempt at measuring metabolites yielded inconclusive data in all animals, suggesting that yohimbine was poorly if at all metabolized in rats. As a result, only the total plasma time activity curve was used as the input function for measurement of the volume of distribution V_T . The V_T was then divided by the free fraction in order to obtain the volume of distribution corrected for free plasma concentration (V_T/f_p), where f_p is the concentration of tracer in plasma that is not bound to plasma protein.

Determination of the plasma free fraction was done in a different set of SD, FRL and FSL rats. We elected to perform the measurements in a separate group of rats as, during the PET studies, plasma is already collected to construct the time activity curve and added sampling of the volume of plasma needed to adequately perform the free fraction measurements in triplicate could have significantly affected the animals' welfare as well as the acquisition of physiologically relevant data, due to potentially large decreases in blood volume and hematocrit. Thus, independent groups of isoflurane-anesthetized rats were processed in a manner similar as the animals used for the PET studies: they were injected with cyclosporine-A (50 mg/kg IV) and 30 min later were bled through intra-cardiac puncture to obtain sufficient amount of whole blood. To determine the plasma free fraction, f_p , measurement of plasma protein binding was performed using standard procedures as previously described (Gandelman et al., 1994). Briefly, the plasma and an equivalent amount of PBS were spiked with a small amount of [^{11}C]yohimbine and incubated for 10–15 min at room temperature. Aliquots of 50 μL of this solution were used to measure the total activity (unfiltered plasma) and the remaining volume was equally divided (~150 μL each) into three ultrafiltration devices (Centrifree® UF Device, Millipore, 30 kDa molecular weight cut-off). After centrifugation for 20 min at 10,000 g, aliquots of 50 μL of the ultrafiltrate were removed to measure the free activity (filtered protein-free plasma). The activity in the ultrafiltrates, plasma and PBS solutions was counted in a gamma counter (Packard Cobra Gamma Counter, Model D5003) and decay-corrected. The data from the triplicate measurements were averaged. A correction factor for device recovery was calculated from the PBS data as C, the ratio of the total activity (unfiltered) in the buffer to the activity in the filtered buffer. The free fraction f_p of [^{11}C]yohimbine was calculated as the ratio of the activity of the filtered ultrafiltrate to the activity of the unfiltered plasma multiplied by the correction factor C. The data from the individual animals were averaged to obtain a mean free fraction per group/strain.

MicroPET images were processed using Montreal Neurological Institute (MINC) software. Each scan was co-registered to an average rat brain atlas (Rubins et al., 2003). A set of regions of interest (ROI) (frontal cortex, striatum, thalamus, pons and cerebellum) was manually drawn on the atlas. This ROI template was applied to

the registered PET data to produce regional time activity curves. The total volume of distribution (V_T) for each region was obtained using the Logan graphical analysis (Logan et al., 1990) during the 30–90 min period of the scan, using each animal's plasma curve as the input function. The V_T and V_T/f_p were analyzed using a two way ANOVA with STRAIN and REGION as the factors, followed by a Bonferroni correction using Graphpad Prism version 5 for Mac OSX (GraphPad software Inc, La Jolla, CA).

3. Results

The tracer plasma free fraction in FRL (0.21 ± 0.02 mean \pm SD) and SD (0.21 ± 0.02) rats was not significantly different. It was significantly reduced ($P = 0.02$) in the FSL animals (0.17 ± 0.02). Averaged plasma activity curves were constructed for each group from the data of each individual animal corrected by the amount of activity injected per kg of body weight. There were no differences in the plasma total time activity curves between the 3 groups ($p = 0.85$, $F = 0.16$) (Fig 1).

Fig. 2 shows the group average time activity curves (normalized for injected activity per kg of body weight) for three cortical and subcortical alpha2 adrenoceptor binding regions (frontal cortex, striatum and thalamus) for all 3 groups. A three way ANOVA revealed no time \times strain interaction for any region ($F = 0.25$ to 0.44 , $Df = 32$, $p > 0.05$). The graph clearly shows the difference between the FSL animals and the 2 control groups in the different regions.

Two way ANOVA analysis of the V_T showed a significant effect of REGION ($F = 6.46$, $Df = 4$, $p = 0.0002$) and of STRAIN ($F = 54.03$, $Df = 2$, $p < 0.0001$) but no interaction ($F = 0.41$, $Df = 8$, $p = 0.91$). Post-Bonferroni correction of the group data revealed significantly lower V_T in all the regions in the FSL rats compared with SD rats. There was no significant difference between SD and FRL animals.

Two way ANOVA analysis of the V_T/f_p also demonstrated a significant effect of STRAIN ($F = 33.57$, $Df = 2$, $p < 0.0001$) and REGION ($F = 6.7$, $Df = 4$, $p = 0.0001$) and no interaction ($F = 0.29$, $Df = 8$, $p = 0.97$). Comparisons between STRAIN groups revealed statistically significant differences between the FSL and SD/FRL rats with FSL rats showing lowest V_T/f_p binding.

Table 1 compares the average values of V_T and V_T/f_p (SEM) for all 3 groups in all 5 regions. Fig. 3 shows the V_T/f_p for all 3 groups. Fig. 4 shows a co-registered parametric PET/MRI image of the V_T/f_p of one rat in each of the three groups.

4. Discussion

In this study, we demonstrated a reduced binding of yohimbine in FSL rats compared to FRL and SD control rats. This observation is consistent with a previous study demonstrating a two to threefold

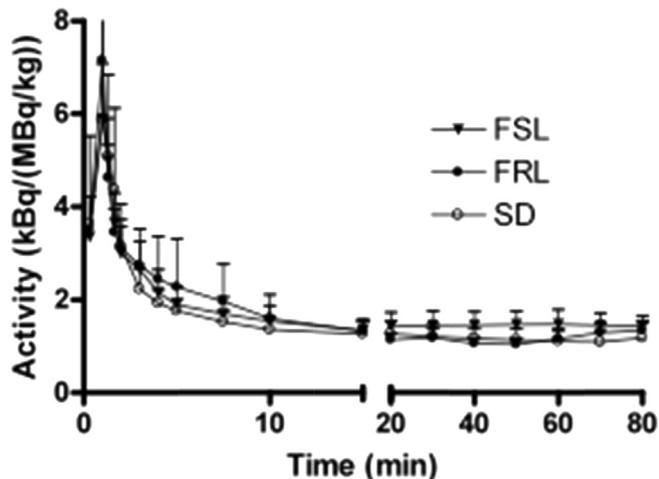


Fig. 1. Averaged total plasma activity curves in the 3 groups (\pm SEM). The early part of the curve was expanded to more clearly show the similarity between curves.

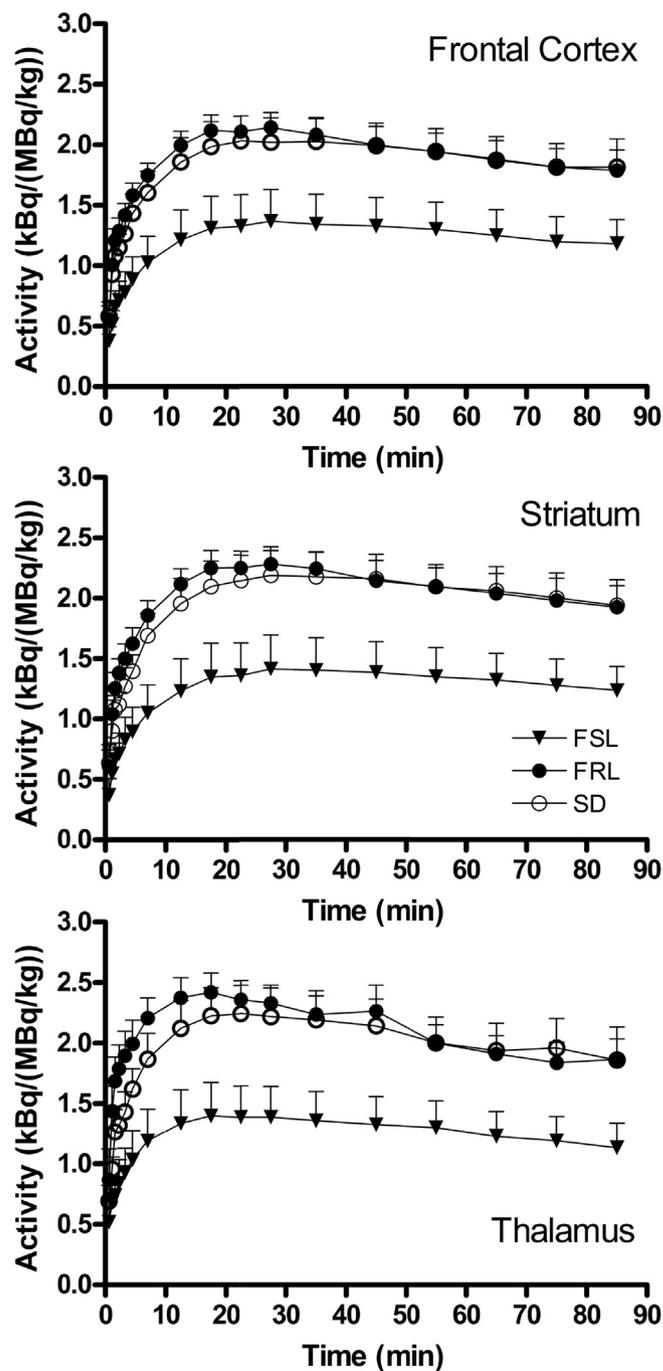


Fig. 2. Examples of averaged time activity curves in frontal cortex, striatum and thalamus in the 3 groups of animals. Activity is corrected for the amount of activity injected by kg of body weight (\pm SEM). The similar shape of the curves graphically demonstrates the lack of difference in perfusion kinetics.

increase in NA levels in the nucleus accumbens, prefrontal cortex and hippocampus in FSL rats compared to SD rats (Zangen et al., 1999). Indeed, we have previously demonstrated that yohimbine is highly sensitive to changes in endogenous NA (Landau et al.,

Table 1
Volume of distribution V_T and free-fraction corrected volume of distribution V_T/f_p averaged across the 5 regions. Data shown \pm SEM. The FSL column also shows the percent decrease in FSL binding compared to SD.

Units: (mL/cm ³)	SD	FRL	FSL
V_T	1.70 ± 0.09	1.71 ± 0.1	0.76 ± 0.04 (55%)
V_T/f_p	8.10 ± 0.42	8.13 ± 0.47	4.48 ± 0.24 (44%)

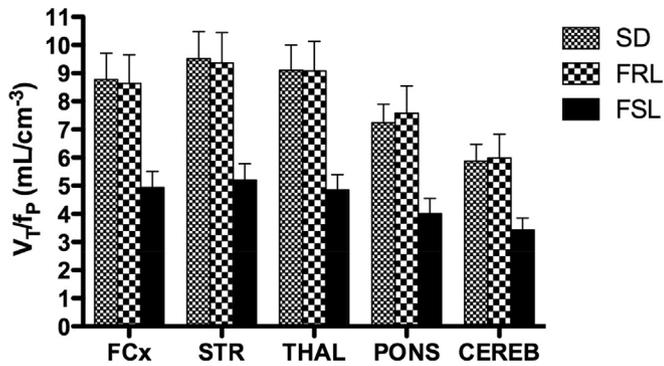


Fig. 3. Graphical representation of the volume of distribution corrected by the free plasma fraction values (V_T/f_p) in the control Sprague Dawley (SD), Flinders Resistant Line (FRL) and Flinders Sensitive Line (FSL) rats. Error bars indicate standard error.

2012b; Phan et al., in press), probably through a competition process similar to what is hypothesized for raclopride, a tracer of the DA D2 receptor routinely used as a surrogate marker of DA release (Laruelle, 2000). Decreased *in vivo* yohimbine binding would thus reflect this heightened endogenous synaptic NA concentrations and/or receptor down-regulation.

Initial studies demonstrated that, in rodents, yohimbine is a substrate for the ABC-cassette family of blood brain barrier transport systems that include P-glycoprotein, and control and reduce entry of drugs into brain parenchyma. Differences between the human and rodent isoforms (Zolnerciks et al., 2011) may account for species variation as yohimbine is not a P-glycoprotein substrate in human. Cyclosporine-A is a commonly used non-specific P-glycoprotein inhibitor that has often been used in rodents and primates to overcome low entrance of PET tracers into the brain. Administered prior to the tracer, it permitted the acquisition of images allowing reliable regional identification and data quantification of yohimbine binding.

We were unable to reliably detect plasma metabolites in any of the 3 groups, SD, FRL or FSL despite testing of several HPLC methods and columns. Nevertheless, the HPLC traces were remarkably similar between animals, leading us to conclude that there was little or no significant measurable metabolism of the tracer. Thus, we used the total plasma activity curve as the input to evaluate V_T . The plasma free fraction f_p was significantly lower in FSL rats compared to the FRL and SD controls. The reason for the difference in free fraction is unclear. A non-specific pharmacological drug like cyclosporine may have affected systemic pharmacokinetic factors and produced tracer–drug interactions independently of its effects on P-glycoprotein. This family of pharmacological agents is known to interact with metabolic functions, including liver enzymes. Kotsovolou et al. demonstrated significant differences in the expression of several major drug-metabolizing enzymes, notably in the CYP2 and CYP3 families of enzymes in FSL rats (Kotsovolou et al., 2010), which play a significant role in metabolizing numerous clinically used drugs, such as yohimbine and mirtazapine, another α_2 adrenergic antagonist. While all the rats received cyclosporine, interactions of an unknown nature may have altered the systemic pharmacokinetics of the tracer in FSL rats, a strain bred over generations to express specific behaviors and in which the genetic makeup is not fully understood. The lack of difference in total plasma activity between strains with a decreased free fraction in FSL animals suggests a decrease in the concentration of circulating free tracer in these rats, supported by the clear difference in the brain time activity curves in every brain region in the FSL compared to the control rats. The similarity of the shape of the curves between strains argues for an unaffected mechanism of

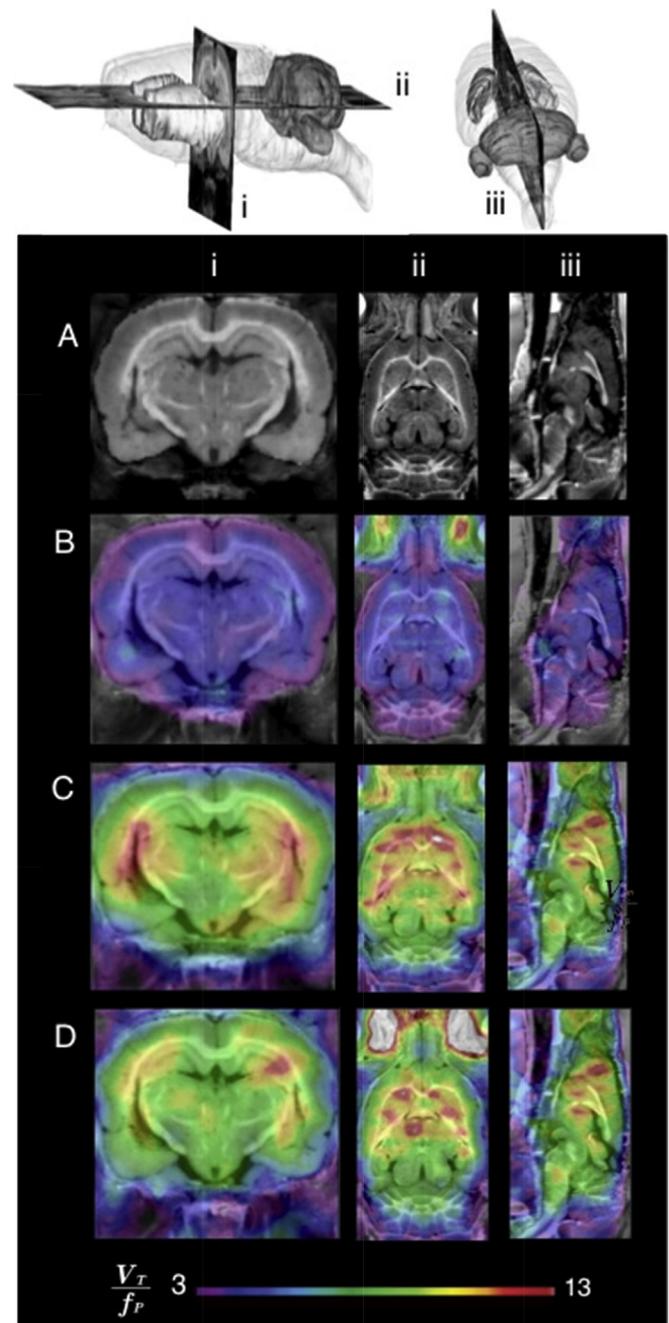


Fig. 4. Coronal (i), axial (ii) and (iii) sagittal parametric PET images superimposed on an MRI atlas (A) of the V_T/f_p of a representative FSL (B), FRL (C) and SD (D) rat. Note the reduced V_T/f_p in the FSL rat compared to the control groups.

brain entrance and little effect of blood flow on the differences between FSL and controls. The differences in binding between the control groups and FSL rats remained highly significant even after correcting the volume of distribution V_T by the tracer plasma free fraction f_p .

A large section of the clinical literature on mood disorders strongly supports α_2 receptor up-regulation as a result of dysfunction and/or loss of NA inputs from the LC and other brainstem and pontine nuclei (Ressler and Nemeroff, 1999). Increased density of α_2 adrenoceptors and decreased density of NA transporters are observed in the LC (Ordway et al., 2003), suggesting NA neuron loss. Receptor binding studies in the brains of suicide

victims demonstrate increased α_2 adrenoceptor binding and receptor mRNA levels (Escriva et al., 2004; Gonzalez-Maeso et al., 2002), and increased receptor agonist binding in the hippocampus and cerebral cortex (Gonzalez et al., 1994). Similarly, a preliminary in vitro autoradiography study in different groups of female SD, FRL and FSL animals using [3 H]-RX821002, another selective tracer of the α_2 receptors, performed in our lab as part of another study (Lillethorup et al., submitted) revealed either no significant change (thalamus, hippocampus, amygdala) or α_2 receptor up-regulation (cortical regions: 15–23%) compared to control SD.

While the post-mortem data are in apparent contradiction with our in vivo data, it is important to remember that in many cases, post-mortem autoradiography data reflect different processes than in vivo data: most significantly, the tissue is washed of any remaining endogenous ligand as part of the autoradiography method and the data represent the number of available receptors, not their functional state. The PET data are, on the contrary, acquired in a live subject in which all regulatory mechanisms are ongoing and the decrease in yohimbine binding may thus reflect a state of functional hyper-noradrenergic release. This hypothesis is supported by the increased levels of extracellular NA found in vivo in FSL rats compared to SD (Zangen et al., 1999). Furthermore, although in vivo PET data is often influenced by the choice of anesthetic, we have previously shown that yohimbine binding is fairly insensitive to changes in blood flow (Alstrup et al., 2013).

Increased markers of hyper-noradrenergia are also reported in some human depressive presentations (Brunello et al., 2003; Wong et al., 2000) and high levels of NA are often associated with heightened anxiety (Baldwin, 2006; Charney et al., 1984). Interestingly, in some contexts, anxiety and depression have been found to co-exist in FSL animals (Overstreet et al., 2004).

In conclusion, one may thus hypothesize that FSL animals represent a model of the anxious/depressive type of depression characterized by a functional state of high central levels of NA release, especially in stressful situations (pre-scan handling for example). This would suggest that the FSL rat model, like many animal models, while presenting a typical rodent depressive behavioral phenotype, reflects only a subset of the physiological underpinnings of depression in the human population and that more than one type of noradrenergic dysregulation may underlie the variable expression of mood disorders and behavioral dysfunctions.

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