Nandrolone supplementation promotes AMPK activation and divergent ¹⁸[FDG] PET brain connectivity in adult and aged mice

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Abstract

Decreased anabolic androgen levels are followed by impaired brain energy support and sensing with loss of neural connectivity during physiological aging, providing a neurobiological basis for hormone supplementation. Here, we investigated whether nandrolone decanoate (ND) administration mediates hypothalamic AMPK activation and glucose metabolism, thus affecting metabolic connectivity in brain areas of adult and aged mice. Metabolic interconnected brain areas of rodents can be detected by positron emission tomography using ^{18}F FDG-mPET. Albino CF1 mice at 3 and 18 months of age were separated into 4 groups that received daily subcutaneous injections of either ND (15 mg/kg) or vehicle for 15 days. At the *in vivo* baseline and on the 14th day, ^{18}F FDG-mPET analyses were performed. Hypothalamic pAMPK^{T172}/AMPK protein levels were assessed, and basal mitochondrial respiratory states were evaluated in synaptosomes. A metabolic connectivity network between brain areas was estimated based on ^{18}F FDG uptake. We found that ND increased the pAMPK^{T172}/AMPK ratio in both adult and aged mice but induced ^{18}F FDG uptake and mitochondrial basal respiration only in adult mice. Furthermore, ND triggered rearrangement in the metabolic connectivity of adult mice and aged mice compared to age-matched controls. Altogether, our findings suggest that ND promotes hypothalamic AMPK activation, and distinct glucose metabolism and metabolic connectivity rearrangements in the brains of adult and aged mice.

Key words: Aging; Mitochondria; ^{18}F FDG-PET; Anabolic-androgen steroids; Nandrolone decanoate; Neural metabolic connectivity

INTRODUCTION

The aging process is associated with a progressive decrease in physiological function, including abnormal nutrient sensing and cerebral hypometabolism. An exacerbation of these biochemical features may evolve to compromised synaptic integrity and connectivity, leading to neurofunctional deficits [1].

In particular, the hypothalamus plays a unique role in integrating brain and whole-body energy metabolism. Hypothalamic cells are sensitive to systemic metabolic signals, such as glucose, fatty acids, and hormones, including, insulin and leptin [2, 3]. Intracellularly, a molecular effector of these energy signals and endocrine regulation is AMP-dependent kinase (AMPK). Once activated, AMPK promotes increased glucose uptake, as well as induction of catabolic pathways leading to ATP production [4–6]. This allows AMPK to dynamically respond to changes in energy demand due to oscillations in synaptic activity, primarily orchestrating substrate flow toward mitochondrial oxidative phosphorylation.

As a metabolic hub connecting different cellular functions, mitochondrial dysfunction associated with aging is believed to contribute to key components leading to cellular senescence, such as exacerbated oxidative stress and promotion of apoptosis [7, 8]. Depending on the localization site at nerve cells, mitochondria can be more vulnerable to damage, exerting a major contribution to aging mechanisms [9]. Based on this, mitochondrial malfunction in synaptic terminals impairs the energy support for neurotransmitter release and uptake and control of ionic homeostasis, which are known substrates for the rupture of synaptic function and neural-cell connectivity [10, 11].

The maintenance of healthy synapses relies on adequate mitochondrial function as well as sufficient metabolic substrate inflow and utilization. Clinically,

hallmark neuroimaging findings in aging and age-related disorders include cerebral hypometabolism detected by fluorodeoxyglucose (^{18}F FDG). Associated with reduced topographical glucose uptake in the brain, diseases common in the elderly population, such as Alzheimer's and Parkinson's, display reduced cerebral connectivity assessed by PET- ^{18}F FDG, reflecting neuronal loss and synaptic dysfunction. Interestingly, abnormal energetic homeostasis during aging parallels altered hypothalamic activity [12, 13] and a reduction in hypophysis-hypothalamic-gonadal signaling, leading to reduced testosterone levels in males [14].

Remarkably, many physiological processes in the brain, from development to normal aging, are under the essential influence of the androgenic anabolic steroid (AAS) hormones produced in the gonads. For instance, androgen receptors (ARs) mediate synaptogenesis (synaptic density and dendritic spine number), which is a morphological correlate of neuroplasticity and likely cognitive function [15, 16]. The reported decrease in AAS during the course of aging has been the mechanistic basis for proposing exogenous hormone administration to hamper age-related physiological deficits [17–21]. Testosterone and its synthetic derivative nandrolone decanoate (ND) induce robust anabolic effects in various tissues, including the brain [22–24]. Based on these properties, administration of AAS has already been investigated to attenuate age-related brain pathologies, in which a decrease in bioenergetic function exerts a role in the progression of symptoms and the severity of disease [25–27]. On the other hand, testosterone and ND are often used in young adults to promote anabolic effects such as increased protein synthesis, skeletal muscle mass/strength and oxidative metabolism. However, such an induced metabolic boost is not limited to peripheral tissues and is frequently accompanied by behavioral effects, confirming that beyond morphological effects, AAS also mediates synaptic mechanisms. However, although

androgen hormones affect mitochondrial synaptic metabolism and function, the impacts of their use on AMPK-dependent hypothalamic metabolic control and brain connectivity during aging remain elusive.

In this study, we sought to investigate whether ND administration improves ^{18}F FDG uptake and metabolic connectivity in the brain in adult and aged mice.

METHODS

Animal procedures

Albino CF1 mice of 3 and 18 months of age were randomly selected to receive either subcutaneous administration of a synthetic analog of testosterone, nandrolone decanoate (18m/ND) (Decadurabolin[®], Schering-Plough, Brazil) at a dose of 15 mg/kg, or an equivalent volume of vehicle (Corn oil, Salada[™], Bunge, Brazil) once a day for 15 consecutive days [24, 28]. Animals were maintained in plexiglass cages, with an average of 4 littermates, and *ad libitum* access to water and standard chow under controlled temperature (22 ± 1 °C) and lighting (12 h light/12 h dark, lights on at 7 am) conditions. Following pharmacologic administration, animals were sacrificed, and the hypothalamus was dissected and homogenized for immunoblotting assays. The remaining left cerebral hemisphere was homogenized for synaptosome fractionation, which was used for mitochondrial respirometry. All procedures were performed with the approval of the UFRGS ethics committee for animal experimentation (CEUA UFRGS #33762)

Immunoblotting analysis

Dissected hypothalamic tissue was homogenized in NP-40 homogenizing buffer, submitted to 12% acrylamide/bis-acrylamide gel electrophoresis (SDS–PAGE) and transferred to and analyzed on nitrocellulose membranes. Membranes were blocked according to each antibody datasheet. Twenty-five micrograms of each sample were incubated overnight at 4°C with pAMPK α 2 (Cell Signaling®, USA, 1:1000) and AMPK α 2 (T172) (Cell Signaling®, USA, 1:1000) antibodies, respectively. After overnight incubation, membranes were washed with 1x TTBS and placed in a 1:25000 secondary HRP-conjugated IgG anti-rabbit or anti-mouse solution (GE Healthcare®, Sweden) according to the primary antibody datasheet specifications. Western blotting analysis is shown as the percent of control optic density, analyzed using ImageJ software, and normalized to the loading control β -actin HRP-linked primary antibody (1:10000, Sigma–Aldrich®, Saint Louis, MO, USA) according to the manufacturer's instructions.

Synaptosome preparation

Animals' left hemispheres were rapidly dissected and homogenized in 1000 μ L of isolation buffer (320 mM sucrose, 10 mM Tris, 0,1% BSA, pH 7.4) using a 10 mL glass potter. The brain tissue was gently homogenized with 10 strokes. The homogenate was immediately used for respirometry and protein quantification for subsequent normalization of oxygen consumption per mg of tissue.

For synaptosome isolation, homogenized samples were then centrifuged at 1330 $\times$ g for 3 min. The supernatant was carefully retained and then centrifuged at 21,200 $\times$ g for 10 min. The resulting pellet was resuspended, carefully layered on top of a discontinuous Percoll gradient, and centrifuged for 5 min at 30,700 $\times$ g [11, 12]. This results in a synaptosome-enriched band that was transferred and resuspended

in isolation buffer and centrifuged for an additional 16,900x g for 10 min. The resulting pellet was resuspended in 2 mL isolation buffer and centrifuged at 6,500x g for 10 min. The final pellet of the synaptosome fraction was resuspended in sucrose and Tris buffer (320 mM sucrose, 10 mM Tris, pH 7.4) and immediately used for further analysis as previously described [29, 30].

MicroPET-scan analysis

Two to six hours before the first subcutaneous injection of ND or VEH, mice were submitted to baseline [^{18}F]FDG uptake evaluation using positron emission tomography scans (microPET), performed at the Brain Institute (Porto Alegre, Rio Grande do Sul, Brazil). The same animals were resubmitted to [^{18}F]FDG uptake scans again 15 days later. Before both paired scans, animals were fasted overnight but had *ad libitum* access to water. Animals were placed in a controlled temperature environment, and an intraperitoneal injection of 200 μL of [^{18}F]FDG with a radioactive activity of approximately 240 μCi was administered, allowing conscious drug uptake for 40 min. Animals were then anesthetized (ketamine/xylazine 90 mg/kg and 7.5 mg/kg, respectively), and microPET scans were performed [31, 32]. Acquisition of ^{18}F FDG-PET images was mapped and quantified as SUV according to *mirrione* [33], and is expressed as the variation between paired baseline and 15th-day scans. Glucose uptake in the brain was topographically assessed in the following regions: (**RSTR**; **LSTR**: striatum, right and left, respectively.) (**CTX**: cortex) (**RHIP**; **LHIP**: hippocampus, right and left, respectively) (**THA**: thalamus) (**CB**: cerebellum) (**BFS**: basal forebrain & septum) (**HYP**: hypothalamus) (**RAMY**; **LAMY**: amygdala, right and left, respectively) (**BS**: brainstem) (**CG**: cingulate gyrus) (**SC**: superior colliculi) (**OLF**:

olfactory bulb) (**RMID**; **LMID**: midbrain, right and left, respectively) (**RIC**; **LIC**: inferior colliculi, right and left, respectively).

High-resolution respirometry analysis

High-resolution respirometry was performed using an O2k Oxygraph (Oroboros Instruments, Innsbruck, Austria) utilizing a SUIT (substrate, uncoupler inhibitor titration) protocol to assess specific mitochondrial respiratory states as described by Chance *et al* [34] and redefined by Gnaiger *et al* [35]. Synaptosomes obtained from the left cerebral hemisphere were submitted to a high-resolution respirometric analysis using an O2k oxygraph (Oroboros Instruments, Innsbruck, Austria). The analysis was performed in 300 μ L of synaptosomal fraction suspended in 2 mL of respiration buffer (KCl 100 mM, mannitol 75 mM, sucrose 25 mM, KH_2PO_4 5 mM, EDTA 0.05 mM, and Tris-HCl 10 mM, pH 7.4).

The ROUTINE respiration state was reached after the stabilization of oxygen consumption rates following sample addition and is sustained by endogenous energy substrates in which glucose represents a major contribution. After the ROUTINE state was achieved, KCN (1 mM) was added to inhibit complex IV, which limits oxygen consumption to nonoxidative reactions, allowing estimation of ROX. Basal respiration was then estimated as the difference between ROX and ROUTINE states.

Statistical analysis

Due to the nature of the presented data, an unpaired Student's t test was performed to assess differences, with significance considered when $p < 0.05$, and normality was investigated using the Kolgomorov-Smirnov normality statistical test. All results in the figures are shown as the mean $\pm$ SD. Pearson tests were performed to

investigate correlations between variables using the R package *psych* [36], with Bonferroni correction for multiple correlations. Correlation data are expressed as Pearson's r , and only correlations with $r > 0.9$ and $p < 0.01$ are shown.

Metabolic interconnected brain areas in rodents can be detected by positron emission tomography using ^{18}F FDG-mPET. Given this, we estimated metabolic networks through correlation of the $[^{18}\text{F}]$ FDG uptake values among different brain anatomical regions of interest (ROIs). The magnitude of the correlation implies the existence of a functional association. For this approach, the Pearson's r correlation of SUVs obtained in each anatomical region by microPET analysis was used.

Networks were built using the *RedeR* package [37]. Furthermore, we performed clustering with Euclidean distance and the complete linkage agglomeration method using the *pvclust* package [38]; 10000 bootstraps and 123 seed parameters were set to assess the uncertainty in hierarchical clustering. Additionally, heatmaps were obtained from the correlation plots.

RESULTS

Nandrolone upregulates hypothalamic AMPK in adult and aged mice.

Immunocontent analysis of hypothalamic pAMPK (Thr172) and AMPK in adult and aged mice revealed that ND significantly increased the pAMPK (Thr172)/AMPK ratio in adult mice (3m/ND vs. 3m/VEH, $p = 0.0274$; $t = 2.628$, $df = 9$), a proxy indicator of increased AMPK activation (Figure 1A, C). This regulatory effect of nandrolone was also observed in aged mice, with the hypothalamic pAMPK(Thr172)/AMPK ratio significantly increased in 18m/ND mice compared to 18m/VEH (18m/ND vs. 18m/VEH, $p = 0.0430$, $t = 2.403$, $df = 8$) (Figure 1B, C).

Nandrolone supplementation affects glucose metabolism and basal mitochondrial respiration

Brain ^{18}F FDG uptake after 15 days of nandrolone administration was significantly increased up to 1.5-fold in adult mice (3m/ND group) compared to age-matched controls injected with vehicle (3m/VEH) in all assessed areas (Table 1, and Figure 2A). In contrast, this increase in brain glucose metabolism promoted by ND was not observed in aged mice (18m/ND), compared to aged mice injected with vehicle (18m/VEH) in any of the assessed regions (Table 1 and Figure 2A).

Coupled with ^{18}F FDG analysis, synaptosomes were submitted to oxygraphic analysis to evaluate the oxidation of endogenous substrates, including glucose. Mitochondria from synaptosome preparations displayed increased oxygen consumption in adult mice supplemented with ND (3m/ND), compared to controls (3m/VEH), (Unpaired Student's t test, 3m/VEH vs. 3m/ND, $p=0.0046$, $t=3.748$, $df=9$) (Figure 2B). Aged mice supplemented with nandrolone (18m/ND) displayed no significant difference in routine respiration compared to age-matched controls (18 m/VEH) (unpaired Student's t test, 18m/VEH vs. 18m/ND, $p=0.4899$, $t=0.7237$, $df=8$) (Figure 2C).

A neural connectivity network using the between-region correlation of ^{18}F PET-FDG is shown in Figure 3. In this network, brain regions were clustered by correlation similarity into containers of closely related areas. The edge width between these containers is associated with connectivity among containers and their composing elements (regions). From this, we observed that ND administration in adult mice (3m/ND) promotes an overall loss of associations, which is visually evidenced by fewer interconnected regions (nodes), fewer intercorrelations inside the containers (links),

and finally, the loss of entire correlation clusters relative to age-matched controls (edge width) (3m/VEH) (Figure 3A-B). In contrast, ND supplementation in aged mice (18m/ND) did not elicit an expressive loss in the number of interconnected regions or in the degree of integration in the correlation network, indicated by the number of node and cluster links and edge width, compared to age-matched controls (18m/VEH).

Discussion

Healthy aging in humans is associated with a disruption in brain metabolism and connectivity and a reduction in androgen hormone levels. In this study, we found that age exerts influence over the ND-promoted metabolic rewiring in the brain but was not associated with AMPK activation.

Hormone replacement therapies are employed in aged populations for therapeutic purposes and target age-related declines in physiological function. Additionally, hormones such as ND are frequently used for the treatment of catabolic disorders and by young adults in drug abuse for aesthetic and performance purposes. The metabolic effects of the therapeutic and nontherapeutic use of these drugs on the brains of adult and aged populations are widely unknown.

Here, we demonstrated that ND supplementation induces increased AMPK phosphorylation, a key metabolic regulator in the hypothalamic brain area. This increase in AMPK activation (pAMPK^{Thr172}) following the acute use of steroid hormones has been described in other tissues, such as human cardiomyocytes [39], skeletal muscle[40], and the hypothalamic arcuate nucleus of guinea pigs [41]. Herein, we expanded this concept by demonstrating that AMPK activation is not restricted to the arcuate nucleus subregion but may be part of the entire hypothalamic signaling strategy to regulate energy homeostasis. Interestingly, such effects appear to be

sustained by ND over the course of at least 15 days in both adult and aged mice. This modulatory effect of steroid hormones may contribute to attenuating certain hallmark features of aging, such as abnormal nutrient sensing, which leads to imbalances in food intake and energy expenditure control in aged individuals [1]. In guinea pigs, activation of the arcuate nucleus AMPK by testosterone treatment led to not only adaptations in the orexigenic neuronal network and feeding patterns but also systemic metabolism, increasing body temperature up to 16 hours post injection through AMPK-dependent mechanisms [41].

The cellular effect of AMPK activation is the promotion of major catabolic pathways, generating endogenous substrates that culminate in increased mitochondrial respiration coupled with ATP synthesis [42, 43]. We showed that ND induces increased mitochondrial oxidation of endogenous substrates only in adult mouse synaptic terminals. Steroid hormones are known to modulate synaptic mitochondrial activity through the translocation of cytoplasmic androgen receptors to nuclei and by directly binding to mitochondrial androgen receptors [44, 45]. Studies have suggested that synaptic mitochondria in ND-treated mice display increased basal mitochondrial membrane potential and hydrogen peroxide production, a proxy indicator of increased endogenous substrate availability and utilization [22], a metabolic feature reinforced by our findings. However, an increase in basal mitochondrial respiration was not observed in ND-supplemented aged mice, indicating a certain level of resistance to this hormone, as evidenced by the limited metabolic responsiveness. A previous work published by Holloway *et al* [46] suggests that metabolic alterations observed in the skeletal muscle of aged males do not translate directly into mitochondrial respirometric alterations due to an increased insensitivity to ADP, leading to increased ROS production throughout the lifespan. We corroborate

this notion by showing that although ND supplementation promotes increased AMPK activation, this does not translate to equivalent elevation in basal mitochondrial respiration in aged mice.

Hypothalamic AMPK regulates glucose metabolism in both peripheral organs and the brain. In humans, testosterone has been described to increase ^{18}F FDG uptake, restoring regional hypometabolism in the brains of both adult men and women [18, 47]. These findings highlight the regulatory effect of androgen hormones in glucose metabolism through increased AMPK activation, which was previously shown in cardiomyocytes [48]. Here, we found that ND supplementation in adult mice increases brain glucose metabolism, further expanding the concept of androgen regulation over hypothalamic control of brain metabolism through AMPK activation.

Intriguingly, ND did not increase brain glucose metabolism in aged mice, which is not coincident with human studies previously reported in the literature. For instance, testosterone replacement therapy (TRT) effects on either cognitively impaired Alzheimer's disease patients or eugonadal patients have shown increases in cortical glucose metabolism [49, 50]. In our study, the lack of differences between ^{18}F FDG uptake and basal mitochondrial respiration after ND supplementation may shed light on differences in dose and time regimen influencing metabolic mechanisms in the brains of aged humans and mice. While we used 15 mg/kg ND for 15 days, those authors used doses ranging from 100mg (topical gel, daily) to 200 mg/kg (i.m., quarterly) for up to three months.

Although ND caused an approximately 1.5-fold increase in whole-brain glucose metabolism in adult mice, there was apparent disruption of between-brain area metabolic networks regardless of age, albeit at different intensities. Reductions in brain connectivity patterns in response to androgen steroid administration have been

shown using MRI and fMRI neuroimaging techniques. For instance, women injected with testosterone display reduced functional connectivity between the inferior frontal gyrus and the anterior cingulate cortex, which manifested by poorer performance in the 'Reading the Mind in the Eyes' test [51, 52]. This suggests that steroid hormone supplementation promotes a loss of connectivity through mechanisms involving alterations in blood flow, substrate uptake, and likely metabolic uncoupling, which may be triggers of synaptic abnormalities and likely neurofunctional deficits.

Conclusion

In summary, our findings suggest that the increase in AMPK activation in response to ND sustains increased glucose oxidation in the whole brain of adult mice. Overall, brain glucose metabolism was not improved by ND in aged mice despite AMPK activation. These findings suggest that ND promotes a distinct organization in the structure and hierarchy of metabolic connectivity among brain areas in both aged and adult mice.

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Competing Interests

The authors have no relevant financial or non-financial interests to disclose.

Author Contributions

All authors contributed to the study conception and design. Material preparation, data collection and analysis were performed by Nathan R. Strogulski, Afonso Kopczynski, Vitória Girelli De Oliveira, Randhall Bruce Carteri and Marcelo S. Rodolphi. Micro-PET FDG preparation, processing and analysis were performed by Giannina Venturin, Samuel Greggio and Jaderson C. da Costa. Marco A. de Bastiani performed the brain connectivity analysis. The first draft of the manuscript was written by Nathan R. Strogulski and all authors commented on previous versions of the manuscript. All authors read and approved the final manuscript.

Data availability

The datasets generated during and/or analysed during the current study are available in the Open Science Framework repository, DOI 10.17605/OSF.IO/K7CX3.

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Figures legends

Figure 1: AMPK activation in the hypothalamus. (A) pAMPK(T172)/AMPK

immunocontent ratio was increased in adult mice relative to age-matched controls (3m/VEH). **(B)** pAMPK(T172)/AMPK immunocontent ratio is increased in aged mice relative to age matched controls (18m/VEH) **(C)** Representative western blotting membrane of AMPK, pAMPK(T172) and loading control β -actin. (Unpaired Student's t test, * $p < 0.05$; 3m/VEH $n = 6$, 3m/ND $n = 5$, 18m/VEH $n = 5$, 18m/ND $n = 5$) Results are shown as the mean $\pm$ SD.

Figure 2: Mitochondrial respiration and ^{18}F FDG uptake. (A) ^{18}F FDG uptake values

of cerebral structures, showing increased uptake in young animals exposed to ND supplementation compared to all groups. Results are shown as the mean variation from baseline ^{18}F FDG uptake. (3m/VEH $n = 6$, 3m/ND $n = 5$, 18m/VEH $n = 5$, 18m/ND $n = 5$) **(B)** The routine oxygen consumption rate is increased in adult nandrolone injected mice (3m/ND) compared to aged-matched controls (3m/VEH) (Unpaired Student's t test, * $p = 0.0046$, $t = 3.748$, $df = 9$). **(C)** Nandrolone did not induce increased basal oxygen consumption rates in synaptic mitochondria of aged mice (18m/ND) compared to age-matched controls (18m/VEH) (Unpaired Student's t test, $p = 0.4899$, $t = 0.7237$, $df = 8$). Results are shown as the mean $\pm$ SD. (3m/VEH $n = 6$, 3m/ND $n = 5$, 18m/VEH $n = 5$, 18m/ND $n = 5$)

Figure 3: Nandrolone triggers metabolic connectivity rearrangements in the brains of adult and aged mice. (A, B) The correlation network of control adult mice (A) was drastically disrupted by 15 days of supplementation with nandrolone (B), exhibiting marked loss of connections between nodes and clusters, indicating compromised synchronous cerebral activity due to supplementation. **(C, D)** Nandrolone supplementation in aged mice alters connectivity, shifts connections, and possibly alters the hierarchy of cerebral interactions but does not lead to reduced cluster or node associations, indicating remodeling but not the overall loss of connections.

A**AMPK activation** **B****AMPK activation** **C**

A

Brain variation of ^{18}F FDG uptake

B

Routine Respiration

C

Routine Respiration

A

B

C

D

Region	Average Δ SUV		df	T-value	p-value
	3m/VEH	3m/ND			
RSTR	-0.2691 $\pm$ 0.66	1.4821 $\pm$ 0.52	9	-4.8070	0.0010
LSTR	-0.2498 $\pm$ 0.65	1.4906 $\pm$ 0.50	9	-4.8728	0.0009
CTX	-0.1957 $\pm$ 0.59	1.1387 $\pm$ 0.34	9	-4.4713	0.0016
RHIP	-0.1910 $\pm$ 0.65	1.4296 $\pm$ 0.43	9	-4.7440	0.0011
LHIP	-0.2768 $\pm$ 0.60	1.4302 $\pm$ 0.45	9	-5.2516	0.0005
THA	-0.2403 $\pm$ 0.69	1.5623 $\pm$ 0.44	9	-5.0065	0.0007
CB	-0.1233 $\pm$ 0.55	1.0318 $\pm$ 0.35	9	-4.0536	0.0029
BFS	-0.1619 $\pm$ 0.57	1.1068 $\pm$ 0.45	9	-4.0499	0.0029
HYP	-0.1415 $\pm$ 0.48	1.0046 $\pm$ 0.39	9	-4.2859	0.0020
RAMY	-0.0860 $\pm$ 0.48	0.9737 $\pm$ 0.34	9	-4.0947	0.0027
LAMY	-0.1494 $\pm$ 0.50	1.0082 $\pm$ 0.35	9	-4.3729	0.0018
BS	-0.1149 $\pm$ 0.52	1.1135 $\pm$ 0.32	9	-4.5572	0.0014
CG	-0.2406 $\pm$ 0.76	1.4737 $\pm$ 0.42	9	-4.4588	0.0016
SC	-0.2786 $\pm$ 0.72	1.4273 $\pm$ 0.47	9	-4.5196	0.0014
OLF	-0.1356 $\pm$ 0.62	1.2931 $\pm$ 0.35	9	-4.5362	0.0014
RMID	-0.1805 $\pm$ 0.70	1.4249 $\pm$ 0.40	9	-4.5122	0.0015
LMID	-0.2015 $\pm$ 0.63	1.4667 $\pm$ 0.47	9	-4.8982	0.0008
LIC	-0.2333 $\pm$ 0.74	1.5073 $\pm$ 0.36	9	-4.7574	0.0010
RIC	-0.2871 $\pm$ 0.67	1.2099 $\pm$ 0.54	9	-4.0392	0.0029

Region	Average Δ SUV		df	T-value	p-value
	18m/VEH	18m/ND			
RSTR	-0.0851 $\pm$ 0.66	-0.1602 $\pm$ 0.52	8	0.1458	0.8877
LSTR	-0.1071 $\pm$ 0.63	-0.1293 $\pm$ 0.99	8	0.0423	0.9673
CTX	-0.0315 $\pm$ 0.53	-0.1794 $\pm$ 0.79	8	0.3485	0.7365
RHIP	-0.0072 $\pm$ 0.65	0.0122 $\pm$ 0.99	8	-0.0366	0.9717
LHIP	0.0010 $\pm$ 0.63	-0.0062 $\pm$ 1.04	8	0.0133	0.9897
THA	-0.0253 $\pm$ 0.66	-0.0374 $\pm$ 1.05	8	0.0219	0.9831
CB	-0.0137 $\pm$ 0.61	-0.1460 $\pm$ 0.89	8	0.2748	0.7904
BFS	-0.0392 $\pm$ 0.53	-0.1178 $\pm$ 0.94	8	0.1637	0.8740
HYP	-0.0298 $\pm$ 0.50	0.0417 $\pm$ 0.83	8	-0.1656	0.8726
RAMY	0.0110 $\pm$ 0.46	-0.1009 $\pm$ 0.75	8	0.2838	0.7838
LAMY	-0.0969 $\pm$ 0.45	-0.0157 $\pm$ 0.79	8	-0.1994	0.8469
BS	-0.0097 $\pm$ 0.56	0.0063 $\pm$ 0.89	8	-0.0339	0.9738
CG	0.0007 $\pm$ 0.76	0.0691 $\pm$ 1.14	8	-0.1118	0.9137
SC	-0.0363 $\pm$ 0.73	0.0069 $\pm$ 1.06	8	-0.0752	0.9419
OLF	-0.0096 $\pm$ 0.62	-0.3017 $\pm$ 1.05	8	0.5339	0.6080
RMID	0.0447 $\pm$ 0.67	0.0323 $\pm$ 0.97	8	0.0235	0.9818
LMID	0.0007 $\pm$ 0.65	0.0122 $\pm$ 1.06	8	-0.0206	0.9841
LIC	-0.0182 $\pm$ 0.82	0.0281 $\pm$ 1.09	8	-0.0758	0.9414
RIC	-0.0313 $\pm$ 0.68	-0.1148 $\pm$ 0.94	8	0.1610	0.8761

Table 1: Descriptive statistics of the alterations promoted by nandrolone in 18 FDG uptake by brain regions in adult and aged mice. The variation of the baseline and 14 days of nandrolone (ND) or vehicle (VEH) treatment standardized uptake values (Δ SUV) is expressed as mean $\pm$ SD, respectively. * indicates

p-values <0.05, ** indicates p-values <0.01 and *** indicates p-values < 0.001 (Unpaired Student's T-test).

Abbreviations: df: degrees of freedom, (**RSTR; LSTR**: Striatum, right and left respectively.) (**CTX**: Cortex) (**RHIP; LHIP**: Hippocampus, right and left respectively) (**THA**: Thalamus) (**CB**: Cerebellum) (**BFS**: Basal Forebrain & Septum) (**HYP**: Hypothalamus) (**RAMY; LAMY**: Amygdala, right and left respectively) (**BS**: Brainstem) (**CG**: Cingulate Gyrus) (**SC**: Superior Colliculi) (**OLF**: Olfactory bulb) (**RMID; LMID**: Midbrain, right and left respectively) (**RIC; LIC**: Inferior Colliculi, right and left respectively).