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GRAPHICAL ABSTRACT

Design, synthesis and *in vivo* study of novel pyrrolidine-based 11β-HSD1 inhibitors for age-related cognitive dysfunction

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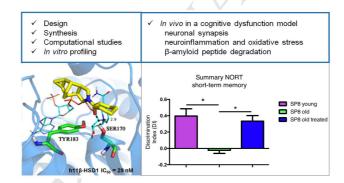
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Design, synthesis and in vivo study of novel pyrrolidine-based 11β-HSD1 inhibitors for age-related cognitive dysfunction

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Abstract:

Recent findings suggest that treatment with 11β-HSD1 inhibitors provides a novel approach to deal with age-related cognitive dysfunctions, including Alzheimer's disease. In this work we report potent 11β-HSD1 inhibitors featuring unexplored pyrrolidine-based polycyclic substituents. A selected candidate administered to 12month-old SAMP8 mice for four weeks prevented memory deficits and displayed a neuroprotective action. This is the first time that 11β -HSD1 inhibitors have been studied in this broadly-used mouse model of accelerated senescence and late-onset Alzheimer's disease.

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Keywords:

Glucocorticoids, 11β -HSD1 inhibitors, drug design, adamantane, polycyclic substituents, aged-related cognitive dysfunction, Alzheimer's disease, SAMP8 mouse.

Highlights:

- a) Several pyrrolidine-based polycyclic amides are prepared as 11β -HSD1 inhibitors.
- b) Amides are synthesized by coupling of pyrrolidines with carboxylic acids.
- c) The more potent compounds present low nanomolar IC_{50} .
- d) A candidate is administered to old SAMP8 mice for four weeks.
- e) The treatment prevents memory deficits and displays a neuroprotective action.

1. Introduction

Elevated glucocorticoids (GCs) exposure is widely accepted as a key factor in agerelated cognitive decline in rodents and humans [1-3]. High levels of GCs have been found in elderly individuals who exhibit learning and memory impairments. GC levels correlate with greater hippocampal atrophy, a region of the brain that is crucial for memory formation [3]. In contrast, low GC levels achieved through neonatal programming or adrenalectomy with exogenous steroid replacement in rats results in the prevention of memory impairments with aging [4].

Growing evidence also suggests that excessive glucocorticoid activity may contribute to Alzheimer's disease (AD), since elevated levels of circulating cortisol in AD patients are associated with more rapid disease progression [5-6]. In a rodent model of AD, systemic administration of GCs led to increases in β -amyloid and tau pathology, the two major histopathologic hallmarks of AD, suggesting a relationship between elevated GC levels and AD pathology [7]. Overall, these data suggest that reducing GC levels in the brain may relieve cognitive dysfunction in both aging and AD.

As in other tissues, the presence of GCs in the brain is not only dependent on adrenal secretion and diffusion from the circulation but also on intracellular metabolism [8]. 11β-hydroxysteroid dehydrogenase type 1 (11β-HSD1) catalyzes the regeneration of active GCs (cortisol in humans, corticosterone in rodents) from their inactive forms (cortisone and 11-dehydrocorticosterone, respectively), providing a local amplification of GC action [9-10]. 11β-HSD1 is highly expressed in fundamental brain areas for cognition, such as the hippocampus, cortex and amygdala [11-13]. By contrast, the isoenzyme 11β-hydroxysteroid dehydrogenase type 2 (11β-HSD2), which catalyzes the opposite reaction, plays an important role during development, as expression of 11β-HSD2 is relevant in fetal brain and placenta, but it has very limited expression in the adult brain [14-16].

Recent studies have demonstrated that aged mice with cognitive deficits show increased 11β-HSD1 expression in the hippocampus and forebrain, and that overexpression of 11β-HSD1 leads to a similar premature memory decline [17]. Conversely, 11β-HSD1 knock-out mice and even heterozygous null mice performed better in different behavioral tests, which suggests resistance to cognitive decline due to a neuroprotective effect of 11β-HSD1 inhibition [18]. Accordingly, this protection correlates with loss of the age-associated rise in intrahippocampal corticosterone, insinuating a role for 11β-

HSD1 in maintaining plasma corticosterone concentration [17]. Furthermore, acute and short-term treatments with 11β -HSD1 inhibitors have shown memory consolidation and improvements in cognitive function in aged mice and AD models [19-23]. Altogether, these findings suggest that 11β -HSD1 inhibitors provide a novel approach through a non-cholinergic mechanism to deal with these cognitive disorders.

In the present work, we report the results derived from a synthetic strategy, supported by molecular modeling studies, designed towards a novel family of potent 11β-HSD1 inhibitors, featuring unexplored pyrrolidine-based polycyclic substituents. The more potent compounds were characterized in terms of cellular potency, isoenzyme selectivity, human metabolic stability and predicted brain penetration to select a candidate for an *in vivo* study. For the first time in the context of 11β-HSD1 inhibitors, the Senescence Accelerated Mouse-Prone 8 (SAMP8) mice were used, as a naturally occurring mouse strain that displays a phenotype of accelerated aging as observed in AD and widely used as a robust rodent model of cognitive dysfunction [24-25].

2. Design, synthesis and in vitro evaluation of new inhibitors

Given that the 11β-HSD1 active site includes a hydrophobic pocket that can accommodate bulky lipophilic substituents, the introduction of a lipophilic group, such as adamantyl, has proven a successful strategy for the space filling of the cavity. Thus, several adamantyl-containing 11β-HSD1 inhibitors exhibit high affinity and potency and some of them (e.g. AZD8329 and ABT-384) have reached clinical trials (Figure 1) [26-34]. Although the evaluation of alternative polycyclic hydrocarbons may offer further opportunities for optimizing the space filling of the hydrophobic cavity, the use of other polycyclic substituents featuring different size or shape has only been briefly scrutinized (e.g. AMG-221 and MK-0736) [35-36].

Figure 1. Selected 11β-HSD1 inhibitors.

In the last few years, our research group has investigated new polycyclic substituents as surrogates of the adamantyl group, leading to inhibitors with promising results on multiple targets [37-45]. However, this strategy has not been successful yet in the case of human 11 β -HSD1 inhibitors [46]. Very recently, we have found that the *N*-(2-adamantyl)amide derivatives **1** and **2** (Figure 2), which are achiral analogues of PF-877423 (IC₅₀ = 4 nM) [47], are potent inhibitors of 11 β -HSD1 (IC₅₀ = 86 and 74 nM, respectively) [48]. Interestingly, the corresponding urea analog, **3**, was significantly less potent (IC₅₀ = 873 nM) [48]. Taking into account the simplicity of these three right-hand side (RHS) units, here we initially selected these fragments for finding alternative polycyclic substituents able to successfully replace the adamantyl group in 11 β -HSD1 inhibitors.

Figure 2. PF-877423 and related inhibitors.

The design of new inhibitors was initially based on a structure-activity relationship (SAR) investigation we previously adopted [46]. The structure of the putative inhibitor was partitioned into two parts: the polycyclic substituent, a surrogate of adamantyl, and the carbocyclic or heterocyclic ring, linked by an amido or urea unit, respectively. Since inspection of the available X-ray structures and preliminary docking studies (see below) indicated that the enzyme active site is large enough to accommodate a polycycle bigger than adamantane, we started our endeavour with previously synthesized amines 4, 5 and 10, three compounds that were successfully used to replace 1-adamantylamine in other targets (Scheme 1) [38]. These amines were then combined with a common RHS moiety to shed light on the effect of key structural features on the inhibitory action: i) primary vs secondary amine (e.g. 4 vs 10), ii) distance between the polycyclic ring and the nitrogen atom (e.g. 4 vs 5), iii) restraint of the conformational freedom (e.g. 5 vs 10), and iv) by incorporating two double bonds in the polycycle, which might form additional interactions in the binding site (e.g. derivatives of 10 vs its reduced analogues). Amides 6 and 7 were prepared in high yields by reaction of cyclohexane acyl chloride with amines 4 and 5, respectively. From amine 10, using either cyclohexanecarboxylic acid or 4-amino-3,5-dichlorobenzoic acid in combination with 1-hydroxybenzotriazole (HOBt) and N-(3-dimethylaminopropyl)-N'-ethylcarbodiimide (EDC) amides 11 and 15 were synthesized in moderate yields. Ureas 8, 9 and 13 were prepared from the required amine and N-chloroformylpiperidine in moderate to excellent yields. Finally, catalytic hydrogenation of 11, 13 and 15 furnished 12, 14 and **16**, respectively (Scheme 1).

Scheme 1. Amines 4, 5 and 10, and amides and ureas derived thereof. ^a

^aReagents and conditions: (i) cyclohexanecarbonyl chloride, anh. acetone, reflux, 3 h, 95% yield for **6**; 78% yield for **7**; (ii) 1-piperidinecarbonyl chloride, Et₃N, DCM, rt, overnight, 54% yield for **8**; quant yield for **9**; 71% for **13**; (iii) cyclohexanecarboxylic acid for **11** or 4-amino-3,5-dichlorobenzoic acid for **15**, HOBt, EDC, Et₃N, EtOAc, rt, overnight, 42% yield for **11**; 47% yield for **15**; (iv) H₂, Pd/C, abs. EtOH, rt, 5 h for **12** and **16**, 3 h for **14**, 78% yield for **12**; 72% yield for **14**; 89% yield for **16**.

A preliminary *in vitro* microsomal assay at 10 μ M compound concentration was performed to assess if the synthesized compounds were able to inhibit human 11 β -HSD1, and the IC₅₀ values were determined for those compounds presenting an inhibition higher than 50% (Table 1).

Compound	hHSD1 % inh at 10 μM	hHSD1 IC ₅₀ (μM)	
6	41	ND	
7	28	ND	
8	50	ND	
9	50	ND	
11	95	1.08	
12	100	0.29	
14	100	0.32	
15	98	2.77	
16	98	0.41	

Table 1. 11β-HSD1 inhibition by compounds 6-9, 11, 12 and 14-16. ^{a,b}

The analysis of the inhibitory potencies disclosed some SAR. First, while amides 6 and 7 displayed poor inhibitory activity, derivatives 11 and 12, featuring a pyrrolidine ring, showed low micromolar and submicromolar potency, respectively, reflecting a better fit within the hydrophobic pocket of the binding site. Second, replacement of the cyclohexyl ring of 12 by either a 1-piperidinyl substituent, as in 14, or a 4-amino-3,5-dichlorophenyl group, as in 16, retained the activity, further demonstrating that the adamantyl substituent may be replaced by other polycyclic groups. Third, no significant difference was found between the inhibitory activity of 12 and 14, and hence the replacement of the amide bridge by a urea within this particular polycycle does not seem to affect the potency. Finally, saturated hexacyclic pyrrolidines were more potent than their diene analogues (compare 12 vs 11 and 16 vs 15). Overall, aliphatic amides 11 and 12 were slightly more potent than aromatic amides 15 and 16 (compare 11 vs 15 and 12 vs 16).

^a11β-HSD1 inhibition was determined in mixed sex, human liver microsomes (Celsis In-vitro Technologies) by measuring the conversion of ³H-cortisone to ³H-cortisol in a cortisol-scintillation proximity assay. ^bPercentage inhibition was determined relative to a control system in the absence of inhibitor (see Experimental section for further details). ND, not determined.

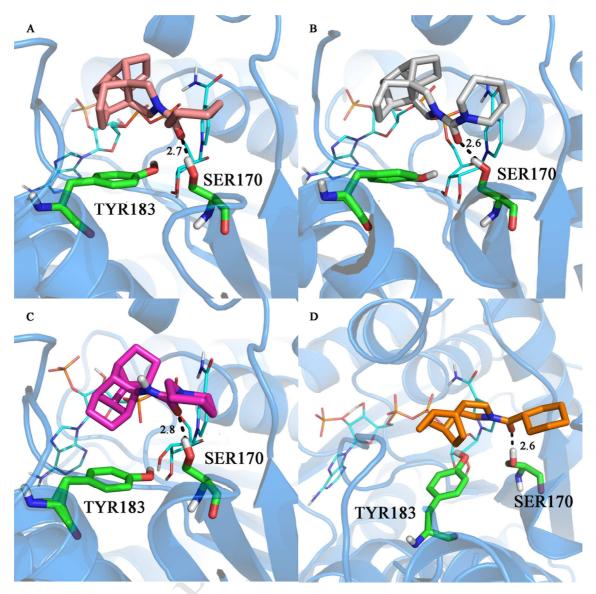


Figure 3. Representative snapshot of the binding mode of compounds 12 (A), 14 (B), 8 (C), and 23 (D) to the human 11β -HSD1 enzyme as determined from the analysis of the MD simulations. The protein backbone is shown as blue carton, the NADP cofactor, residues Tyr183 and Ser170, and the ligands are shown as atom-coloured sticks. The hydrogen bond between the ligand and the hydroxyl group of Ser170 is shown as a dashed line.

Docking studies were combined with molecular dynamics (MD) simulations to shed light into the inhibitory potencies of selected compounds. The structural integrity of the simulated systems was supported by the stability of the positional root-mean square deviation of the residues that define the binding site and the ligand, especially for the most potent compounds (see Figures S1-S3 in Supporting Information). A similar

binding mode was found for compounds **8**, **12** and **14** (see Figure 3). In all cases the carbonyl group of the ligand formed a stable hydrogen bond with Ser170 (distances ranging from 2.6 to 3.0 Å). However, the hydrogen bond of the ligand's carbonyl group with the hydroxyl group of Tyr183, which was retained during the setup of the simulated systems, exhibited larger fluctuations and was eventually disrupted during the trajectories. Docking calculations showed compounds **12** and **14** to have slightly better scores (-9.3 and -9.7 kcal/mol, respectively) than for **6** (-8.4 kcal/mol) and **8** (-9.0 kcal/mol). The higher inhibitory potency of **12** and **14** may also be explained by the fused pyrrolidine ring, which should reduce the contribution of the conformational penalty to the binding affinity of these compounds compared to the more flexible compounds **6** and **8**. Although the results appear to support the ability of the size-expanded hydrophobic cage present in **12** and **14** to occupy the binding pocket, the lower inhibitory potency compared to **1** (IC₅₀ = 86 nM; Figure 2) suggests that the size of the polycyclic substituent in **12** and **14** may be close to the upper limit allowed for ligand binding without triggering significant structural distortions in the pocket

Of note, the *N*-acylpyrrolidine motif contained in **11**, **12**, **15** and **16** has scarcely been explored in the context of the design of 11β -HSD1 inhibitors [49]. However, the pyrrolidine **10** is not an ideal starting compound for a medicinal chemistry program, as its synthesis is tedious and very low-yielding [38]. For this reason, we explored the synthesis of alternative, easily synthesized, pyrrolidine derivatives. To this end, we followed a polycyclic substituent optimization process in which the cyclohexyl was selected as the RHS of the molecule, due to its simplicity (i.e. achiral, easy access) and good performance with both adamantyl (**1**, $IC_{50} = 0.09 \mu M$) and hexacyclic substituents (**12**, $IC_{50} = 0.29 \mu M$). An array of 13 pyrrolidine-based polycyclic amides was prepared from cyclohexanecarboxylic acid, HOBt, EDC, and a series of previously synthesized amines (Figure 4). Our aim was to obtain different pyrrolidine-based polycyclic compounds, some of them simplified analogs of the hexacyclic unit contained in **10** but with higher conformational freedom, in order to find the optimal size and shape to deliver more potent 11β -HSD1 inhibitors.

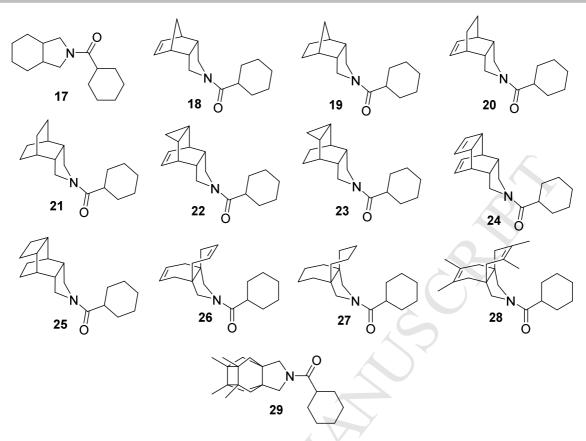


Figure 4. Novel pyrrolidine-based polycyclic amides 17-29.

Following the aforementioned preliminary in vitro microsomal assay, the IC₅₀ values were determined for compounds with an inhibitory activity higher than 50% at 10 µM (Table 2). In general, compounds containing smaller polycyclic rings (i.e. 17-23) were one order of magnitude more potent than our initially best inhibitors 12, 14 and 16, and some of them were also more potent than the adamantyl derivative 1 (IC₅₀ = 0.09 μ M). The most potent inhibitors were 18, 20, 21, 22 and 23 (IC₅₀ values ranging from 0.02 to 0.03 µM). MD simulations of the enzyme complex with compound 23 confirmed the stability of the binding mode (see Figure 3D and S4), which resembled the arrangement of compound 12, and the formation of the hydrogen-bond interaction with Ser170 (average distance of 2.8 ± 0.3 Å). However, there was not a clear trend in terms of activity between the alkene/alkane pairs containing the same polycyclic ring system (compare 20 vs 21, and 22 vs 23). Thus, in the pair 18/19 the alkene derivative presented a slightly higher potency, but the bigger alkene derivatives 24 and 26 were significantly more potent than their alkane analogues 25 and 27 (IC₅₀ = 1.49 μ M vs 27% inhibition at 10 μM, and 0.04 μM vs 1.22 μM, respectively). Finally, the introduction of four methyl groups, either in an extended (28) or compact (29) arrangement, was highly

deleterious to the inhibitory activity (compare **26** *vs* **28** and **29**). Overall, these findings reinforce the assumption that the hexacyclic substituent reaches the upper-limit size to fill the hydrophobic pocket of the binding site.

Compound	hHSD1 % inh at 10 μM	hHSD1 IC ₅₀ (μM)
17	85	0.05
18	89	0.02
19	92	0.09
20	96	0.03
21	95	0.02
22	100	0.02
23	100	0.03
24	83	1.49
25	27	ND
26	82	0.04
27	67	1.22
28	30	ND
29	42	ND

Table 2. 11β-HSD1 inhibition by compounds 17-29. a,b

^a11β-HSD1 inhibition was determined in mixed sex, Human Liver Microsomes (Celsis In-vitro Technologies) by measuring the conversion of ³H-cortisone to ³H-cortisol in a cortisol-Scintillation Proximity Assay. ^bPercentage inhibition was determined relative to a no inhibitor control (see Experimental section for further details). ND, not determined.

3. Biological profiling of the more potent 11β-HSD1 inhibitors

The more potent inhibitors obtained by this polycyclic substituent optimization process have clogP values between 2.68 and 3.99, more desirable than that of the adamantyl-containing analogue 1 (clogP = 4.65). These new compounds were characterized in terms of cellular potency, selectivity over 11β -HSD2, human metabolic stability, cytochromes P450 (CYP) inhibition and predicted brain permeability, in order to select

the best candidate to perform an *in vivo* study in a rodent model of cognitive dysfunction.

The cellular potency was assessed in Human Embryonic Kidney 293 (HEK293) cells stably transfected with the 11 β -HSD1 gene. With the only exception of alkenes **18**, **20** and **26**, which showed a moderate inhibitory activity (55%, 64% and 64%, respectively), all compounds completely inhibited the enzyme at 10 μ M (Table 3).

Selectivity over 11β -HSD2 is required for 11β -HSD1 inhibitors progressing into clinical trials since 11β -HSD2 inhibition in the kidney can lead to sodium retention and increased blood pressure via cortisol stimulation of mineralocorticoid receptors. However, for the purposes of our $in\ vivo$ studiy of cognitive dysfunction in rodents, high selectivity vs 11β -HSD2 was not required. Notwithstanding, 11β -HSD2 inhibition was assessed in a cellular assay with HEK293 cells stably transfected with the 11β -HSD2 gene at 10, 1 and $0.1\ \mu M$ in order to establish the selectivity of our compounds. Ideally, the 11β -HSD2 inhibition at $10\ \mu M$ should be lower than 50% to consider a compound sufficiently selective toward 11β -HSD1. None of our compounds achieved this threshold but some slightly improved the poor selectivity of the adamantyl-containing analogue 1, such as amide $10\ \mu M$, respectively, data not shown). Although $10\ \mu M$ and $10\ \mu M$, so we cannot rigorously consider this compound to be selective against 11β -HSD1, its selectivity index was at least $10\ \mu M$ compared to compounds $10\ \mu M$ and $10\ \mu M$ so we cannot rigorously consider this compound to be selective against 11β -HSD1, its selectivity index was at least $10\ \mu M$ so vecannot rigorously consider this compound to compounds $10\ \mu M$ should displayed selectivities less than $10\ \mu M$ so of $10\ \mu M$ so we cannot rigorously consider this compound to compounds $10\ \mu M$ should displayed selectivities less than $10\ \mu M$ so of $10\ \mu M$ so we cannot rigorously consider this compound to be selective against 11β -HSD1, its selectivity index was at least $10\ \mu M$ so of $10\ \mu M$ so of $10\ \mu M$ so we cannot rigorously consider this compound to be selective against 11β -HSD1, its selectivity index was at least $10\ \mu M$ so of $10\ \mu M$ should be $10\ \mu M$ so we cannot rigorously consider this compound to compounds $10\ \mu M$ should be $10\ \mu$

Microsomal stability was performed in human liver microsomes (HLM), which are widely used to determine the likely degree of primary metabolic clearance in the liver. Compounds **18**, **19** and **20** presented moderate microsomal stabilities between 36 and 60% of remaining parent compound after 30 min incubation, while amides **21** and **22** showed stabilities lower than 28%. Compound **23** displayed a high microsomal stability with 94% of remaining parent compound after the 30 min incubation period.

The active compounds in the 11 β -HSD1 cellular assay (19 and 21-23) were further tested for predicted brain permeation using the widely used *in vitro* PAMPA-BBB model [50]. Unfortunately, the *in vitro* permeabilities (P_e) for compounds 21 and 22 could not be determined due to their lack of UV absorption. Whereas 19 showed an uncertain BBB permeation [CNS \pm with $5.179 > P_e$ (10^{-6} cm s⁻¹) > 2.106], compound 23

had P_e clearly above the threshold established for a high blood-brain barrier (BBB) permeation ($P_e > 30 \text{ x } 10^{-6} \text{ cm s}^{-1}$).

Compound	hHSD1 IC ₅₀ (μM)	HEK hHSD1 % inh at 10 µM ^c	HEK hHSD2 inhibition at 10 μM or IC ₅₀ (μM) ^c	HLM % parent ^d	PAMPA- BBB Pe (10 ⁻⁶ cm s ⁻¹) ^{e,f}
1	0.09	100	88%	79	-
18	0.02	55	$< 0.1~\mu M$	60	-
19	0.099	100	< 0.1 μΜ	37	5.20 ± 0.1 (CNS±)
20	0.03	64	0.1-1 μΜ	44	-
21	0.02	100	0.1-1 μΜ	17	ND^g
22	0.02	100	1-10 μΜ	27	ND
23	0.03	100	< 0.1 μΜ	94	>30 (CNS+)
26	0.04	64	100%	-	-

Table 3. Biological profiling of the most potent compounds. a,b

^aSee Experimental section for further details. ^bPercentage inhibition was determined relative to a no inhibitor control. ^cHEK293 cells stably transfected with the full-length gene coding for human either 11β-HSD1 or 11β-HSD2 were used. ^dThe microsomal stability of each compound was determined using human liver microsomes. ^ePermeability values from PAMPA-BBB assay. Values are expressed as the mean \pm SD of three independent experiments. CNS+, predicted positive brain penetration. ^fCalibration line between 0 and 30 x 10⁻⁶ cm s⁻¹. ^gND, not detected.

4. In vivo study

Recent studies in rodents and humans with brain-penetrant 11β-HSD1 inhibitors have shown that they provide beneficial effects on the cognitive impairment associated with aging [13, 18-23]. SAMP8 has been studied as a non-transgenic murine mouse model of accelerated senescence and late-onset AD [51-52]. These mice exhibit cognitive and emotional disturbances, probably due to early development of brain pathological hallmarks, such as oxidative stress (OS), inflammation, and activation of neuronal death pathways, which mainly affect cerebral cortex and hippocampus [24,53]. To date, this

rodent model has not been used to test 11β -HSD1 inhibitors, being this work the first investigation of the effects of 11β -HSD1 pharmacological inhibition in SAMP8.

The *in vivo* study was performed with amide 23, as this compound had low nanomolar potency against the murine 11 β -HSD1 enzyme (mHSD1, IC₅₀ = 0.08 μ M), high cellular potency, high microsomal stability (both in human and mouse liver microsomes, 94 and 93%, respectively) and positive predicted brain penetration. A pharmacokinetic study of compound 23 was performed in order to assess its oral administration. Although its clearance seems to be rapid, the concentration levels at 30 min post-administration are five fold the IC₅₀ (Table S1, Figure S5 and Table S2). In addition, we could also measure compound concentration in brain tissue at 3 hours post-administration (1.45 ng/mL), hint of *in vivo* BBB permeability. Then, compound 23 was administered to 12-month-old SAMP8 mice in drinking water during four weeks at a concentration of 105 mg/L (average body weight for 48-week-old mice is 25 g; fluid consumption is 5 mL, therefore the dose was 0.105 mg/mL x 5 mL/0.025 kg = 21 mg/kg). Compound 23 was dissolved in polyethylene glycol 400 (PEG400) and then diluted with water to a PEG400 final concentration of 2% (v/v) in drinking fluid. 2% PEG400 in water was given to the remaining mice in drinking fluid as a vehicle control.

Neuroprotective effects were investigated through a behavioural test, the novel object recognition test (NORT), as a common measure of cognition (short-term and long-term memory) [54], and biochemical analysis, which were made through Western blotting and quantitative real-time polymerase chain reaction (qPCR). Brain tissue was analysed upon termination of the study to determine compound levels and *ex vivo* inhibition of 11β-HSD1. Consistent with previous reports, we found that aged SAMP8 mice presented memory impairments in the NORT when compared to young animals [53]. Satisfactorily, treatment with **23** certainly prevented short-term and long-term memory deficits in SAMP8 mice (Figure 5A).

Postsynaptic density 95 (PSD95) protein levels were evaluated as a measure of neuronal synapses, whereas gene expression for interleukin-6 (IL-6), which acts as a proinflammatory cytokine, and for inducible nitric oxide synthase (iNOS), an oxidative stress sensor that catalyzes the production of nitric oxide (NO), were also studied. Treatment with 23 prevented the reduction of PSD95 protein levels, while the oxidative stress and pro-inflammatory gene expression markers, such as iNOS and IL-6, were significantly decreased compared to untreated mice (Figure 5A-B). These observations

indicate a neuroprotective action of 23, whereby reduced cognitive impairment in 23-treated mice is mediated by a reduction of neuroinflammation and oxidative stress as confirmed by in the measurements of pro-inflammatory biomarkers (IL-6 and iNOS).

AD is characterized by the production and deposition of β -amyloid and it has been postulated that its reduction produces beneficial effects [55]. For these reasons, the effect of **23** in modifying amyloid processing pathways was also examined. No changes in amyloid beta A4 precursor (*PreAPP*), β -secretase 1 (*Bace1*), disintegrin and metalloproteinase 10 (*Adam 10*) gene expression levels were found (data not shown). Nevertheless, after treatment with **23**, we found a decrease of APP β -secretase C-terminal fragment (β CTF) protein levels without modification of those of *PreAPP*, together with an increase of the protein levels of insulin-degrading enzyme (IDE) (Figure 1C), a zinc metalloprotease that degrades β -amyloid species. Several *in vitro* and *in vivo* studies have shown correlations between IDE, β -amyloid levels and AD [56].

In an attempt to elucidate the mechanisms underlying the beneficial effects of 23 and the relationship with cognitive amelioration in old SAMP8, we focused our study on amyloid processing and β CTF because of its implication in neurodegeneration and cognitive decline process in this strain [57]. Amide 23 did not alter the proamiloidogenic pathway in SAMP8, as demonstrated by the lack of effect on *Pre-APP*, *Adam10* or *Bace1* gene expression (see Figure S6 in the Supplementary material file). However, the capacity of the brain to remove proamiloidogenic species by activation of specific proteases, such as IDE or neprilysine, appeared greatly increased in treated SAMP8 with higher protein levels of IDE than in control animals and consequently with substantially decreased β CTF protein levels. The results of the behavioural and biochemical studies in SAMP8 mice suggest that compound 23 acts centrally on 11β -HSD1.

Overall, behavioural tests and biochemical analyses confirmed a neuroprotective action of compound **23**, probably by reduction of inflammation and oxidative stress, as measured by reduction of IL-6 and iNOS.

A. COGNITION AND SYNAPTIC DENSITY SP8 young SP8 old SP8 old treated PSD95 95 kDa Summary NORT long-term memory Summary NORT GAPDH 37 kDa short-term memory PSD95 Discrimination Index (DI) %vs.SAMP8 young 5 P8 Old Treated **B. OXIDATIVE STRESS AND INFLAMMATION** iNOS IL-6 Relative mRNA expression Relative mRNA expression 5P8 old SPS old treated C. AMYLOID PROCESSING AND AB CLEARANCE Pre-APP 100kDa SP8 young SP8 old SP8 old treated BCTF 10kDa IDE 118 kDa GAPDH 37kDA GAPDH 37 kDa Ratio βCTF/pre-APP IDE %vs.SAMP8 young %vs.SAMP8 young

Figure 5. Results of *in vivo* study. 2A. -Cognition and synaptic density: -Left panel: Results. Discrimination index of Novel Object recognition test (NORT) obtained in

young SAMP8, old SAMP8 and treated SAMP8, at 2 and 24 h. -Right Panel: Representative Western blot (wb) for PSD95 and quantification. **2B.** -Oxidative stress and inflammation: -Left Panel: Oxidative stress gene expression iNOS. -Right panel: pro-inflammatory gene expression for IL-6. Gene exprexion levels were determined by real-time PCR. **2C.** -Amyloid processing and β -CTF clearance: -Left panel: Representative Western blot (wb) for IDE and quantification. -Right panel: Representative Western blot (wb) for Pre-APP and β -CTF and β -CTF/APP ratio quantification. For Western blot, bars represent mean \pm standard error of the mean (SEM), and values are adjusted to 100% for levels SAMP8 young. For real-time PCR, mean \pm SEM from five independent experiments performed in triplicate are represented. The One-Way ANOVA analysis and Tukey post hoc analysis were conducted. Statistical outliers (Grubbs' test) were removed from the analyses. *p<0.05; **p<0.01; ***p<0.001.

5. Conclusions

We have found that adamantyl, widely used as a lipophilic substituent in 11B-HSD1 inhibitors, may successfully be replaced by other polycyclic hydrocarbons. The previously scarcely explored pyrrolidine-based polycyclic substituents presented here led to potent 11B-HSD1 inhibitors, and potentially these aliphatic ring-systems can serve as alternatives to adamantyl. Of note, the novel nanomolar inhibitors reported are achiral and easily synthesized in maximum four synthetic steps (compounds 19, 21 and 23) from commercially available starting materials. Biological profiling allowed us to select amide 23 for the first in vivo study in SAMP8 aiming to investigate the pharmacological effects of 11β-HSD1 inhibition in this model of cognitive dysfunction. In this study, prevention of cognitive impairment in aged SAMP8 after four-week treatment with 23 was demonstrated in comparison with control animals. The results provide further support for the neuroprotective effect of 11β-HSD1 inhibition, through reduction of neuroinflammation and oxidative stress, in cognitive decline related to the aging process. Due to the promising biological activity of 23, further optimization is currently being carried out, with focus on modifying the RHS of the molecule to improve the selectivity and DMPK properties.

6. Experimental section

6.1. Chemistry

6.1.1. General

Melting points were determined in open capillary tubes with a MFB 595010M Gallenkamp. 400 MHz 1 H/100.6 MHz 13 C NMR spectra, and 500 MHz 1 H NMR spectra were recorded on Varian Mercury 400, and Varian Inova 500 spectrometers, respectively. The chemical shifts are reported in ppm (δ scale) relative to internal tetramethylsilane, and coupling constants are reported in Hertz (Hz). Assignments given for the NMR spectra of the new compounds have been carried out on the basis of DEPT, COSY 1 H/ 1 H (standard procedures), and COSY 1 H/ 1 3C (gHSQC and gHMBC sequences) experiments. IR spectra were run on Perkin-Elmer Spectrum RX I spectrophotometer. Absorption values are expressed as wave-numbers (cm $^{-1}$); only significant absorption bands are given. Column chromatography was performed either on silica gel 60 Å (35–70 mesh) or on aluminium oxide, neutral, 60 Å (50-200 μ m, Brockmann I). Thin-layer chromatography was performed with aluminum-backed

sheets with silica gel 60 F_{254} (Merck, ref 1.05554), and spots were visualized with UV light and 1% aqueous solution of KMnO₄. The analytical samples of all of the new compounds which were subjected to pharmacological evaluation possessed purity \geq 95% as evidenced by their elemental analyses.

6.1.2. General procedures for the synthesis of the compounds

6.1.2.1. General procedure A.

A solution of cyclohexane acyl chloride (1.2 mmol) in anhydrous acetone was added to a solution of the amine hydrochloride (1 mmol) and triethylamine (2 mmol) in anhydrous acetone. The reaction mixture was stirred at 90° C for 3 h. The resulting residue was dissolved with DCM (20 mL) and washed with 1 M aqueous solution of HCl (4 x 25 mL), dried over anh. Na₂SO₄ and filtered. The evaporation *in vacuo* of the organics gave the desired product.

6.1.2.2. General procedure B.

To a solution of the amine hydrochloride (1 mmol) in DCM (10 mL) were added 1-piperidinecarbonyl chloride (1.5 mmol) and triethylamine (4 mmol). The reaction mixture was stirred at rt overnight. To the resulting mixture was added saturated aqueous solution of NaHCO₃ (10 mL) and the phases were separated. The aqueous layer was extracted with further DCM (2 x 10 mL). The organics were washed with 10% Na₂CO₃ solution (30 mL), dried over anh. Na₂SO₄, filtered and concentrated *in vacuo* to give the desired product.

6.1.2.3. General procedure C.

To a solution of amine hydrochloride (1.1 mmol) in EtOAc (15 mL) were added the carboxylic acid (1 mmol), HOBt (1.5 mmol), EDC (1.5 mmol) and triethylamine (4 mmol) and the reaction mixture was stirred at rt overnight. To the resulting suspension was then added water (15 mL) and the phases were separated. The organic phase was washed with saturated aqueous NaHCO₃ solution (15 mL) and brine (15 mL), dried over anh. Na₂SO₄ and filtered. The organic layer was concentrated *in vacuo* to give the desired product.

6.1.2.4. General procedure D.

A solution or suspension of the amide or urea (1 mmol) and 5 wt. % palladium on carbon (50% in water, 10% of the weight) in absolute ethanol (ca 30 mL) was stirred at rt and atmospheric pressure under hydrogen for 3-72 h. The suspension was then

filtered and the solids were washed with EtOH (10 mL). The solvents were removed *in vacuo* to give the desired reduced product.

6.1.3. N-[Pentacyclo[6.4.0.0^{2,10}.0^{3,7}.0^{4,9}]dodec-8-yl] cyclohexanecarboxamide, (**6**).

From cyclohexane acyl chloride (128 mg, 0.88 mmol) in anhydrous acetone (0.6 mL) and amine 4 [38] (152 mg, 0.72 mmol) and triethylamine (0.25 mL, 1.74 mmol) in anhydrous acetone (0.9 mL) and following the general procedure A, amide 6 (215 mg, 95% yield) was obtained as a yellow solid. The analytical sample was obtained by crystallization from EtOAc / pentane (164 mg), mp 228–229 °C; IR (ATR) v: 580, 591, 604, 638, 663, 695, 722, 761, 824, 894, 933, 1143, 1194, 1221, 1277, 1304, 1336, 1386, 1448, 1547, 1639, 2851, 2923, 3271 cm⁻¹; ¹H-NMR (400 MHz, CDCl₃) δ : 1.08 (t, J =2.8 Hz, 1 H, 9'-H), 1.18-1.34 [complex signal, 3 H, 3(5)-H_{ax} and 4-H_{ax}], 1.39-1.55 [complex signal, 10 H, 5'(11')- H_2 , 6'(12')- H_2 and 2(6)- H_{ax}], 1.66 (m, 1 H, 4- H_{eq}), 1.79 $[m, 2 H, 3(5)-H_{eq}], 1.85 [m, 2 H, 2(6)-H_{eq}], 2.08 (tt, J = 11.6 Hz, J' = 3.4 Hz, 1 H, 1-H),$ 2.13 [b. s., 2 H, 4'(10')-H], 2.21 [m, 2 H, 2'(3')-H], 2.81 [b. s., 2 H, 1'(7')-H], 5.47 (s, 1 H, NH); ¹³C-NMR (100.5 MHz, CDCl₃) δ: 21.5 [CH₂, C5'(C11')], 24.4 [CH₂, C6'(C12')], 25.7 (CH₂, C4), 25.8 [CH₂, C3(5)], 30.1 [CH₂, C2(C6)], 45.8 (CH, C1), 46.9 [CH, C2'(3')], 52.8 [CH, C4'(10')], 53.4 [CH, C1'(7')], 54.8 (CH, C9'), 65.4 (C, C8'), 176.7 (C, CO). Anal. Calcd for C₁₉H₂₇NO: C 79.95, H 9.54, N 4.91. Found: C 79.73, H 9.75, N 4.82.

6.1.4. N-[[Pentacyclo[6.4.0.0^{2,10.}0^{3,7}.0^{4,9}]dodec-8-yl]methyl]cyclohexanecarboxamide, (7).

From cyclohexane acyl chloride (119 mg, 0.81 mmol) in anhydrous acetone (0.6 mL) and amine **5** [38] (150 mg, 0.67 mmol) and triethylamine (0.23 mL, 1.62 mmol) in anhydrous acetone (0.8 mL) and following the general procedure A, amide **7** (155 mg, 78% yield) was obtained as a dark solid. The analytical sample was obtained by crystallization from EtOAc / pentane (77 mg), mp 148–149 °C; IR (ATR) v: 576, 589, 604, 621, 660, 672, 711, 800, 894, 956, 979, 1037, 1106, 1123, 1180, 1213, 1252, 1298, 1314, 1380, 1435, 1443, 1550, 1633, 1659, 2855, 2928, 3299 cm⁻¹; ¹H-NMR (400 MHz, CDCl₃) δ : 1.05 (b. s., 1 H, 9'-H), 1.17-1.33 [complex signal, 3 H, 3(5)-H_{ax} and 4-H_{ax}], 1.35-1.55 [complex signal, 8 H, 5'(11')-H_{exo or endo}, 6'(12')-H₂ and 2(6)-H_{ax}], 1.56-1.70 [complex signal, 3 H, 5'(11')-H_{endo or exo} and 4-H_{eq}], 1.74-1.87 [complex signal, 4 H, 2(6)-H_{eq} and 3(5)-H_{eq}], 1.98 [b. s., 2 H, 4'(10')-H], 2.05 (tt, J = 11.6 Hz, J' = 3.2 Hz, 1

H, 1-H), 2.13 [b. s., 2 H, 1'(7')-H], 2.22 [m, 2 H, 2'(3')-H], 3.38 [d, J = 5.6 Hz, 2 H, NCH₂], 5.20 [b. s., 1H, NH]; ¹³C-NMR (100.5 MHz, CDCl₃) δ: 22.3 [CH₂, C5'(C11')], 24.2 [CH₂, C6'(12')], 25.7 (3 CH₂, C3, C4 and C5), 29.8 [CH₂, C2(6)], 36.9 (CH₂, NCH₂), 42.7 (C, C8'), 45.7 (CH, C1), 47.9 [CH, C2'(3')], 51.2 (CH, C9'), 53.1 [CH, C1'(7')], 53.5 [CH, C4'(10')], 175.6 (C, CO). Anal. Calcd for C₂₀H₂₉NO·0.15 EtOAc: C 79.13, H 9.74, N 4.48. Found: C 79.03, H 9.88, N 4.52.

6.1.5. *N-[Pentacyclo[6.4.0.0^{2,10}.0^{3,7}.0^{4,9}]dodec-8-yl]piperidine-1-carboxamide, (8).*

From amine **4** [38] (100 mg, 0.47 mmol), 1-piperidinecarbonyl chloride (0.09 mL, 0.71 mmol) and triethylamine (0.23 mL, 1.88 mmol) in DCM (5 mL) and following the general procedure B, amide **8** (72 mg, 54% yield) was obtained as a white solid. The analytical sample was obtained by crystallization from hot EtOAc (22 mg), mp 225–226 °C; IR (ATR) v: 700, 721, 734, 764, 794, 851, 866, 907, 954, 971, 1003, 1019, 1050, 1128, 1141, 1187, 1232, 1254, 1261, 1276, 1306, 1325, 1357, 1395, 1440, 1479, 1520, 1615, 2024, 2158, 2855, 2930, 3362 cm⁻¹; ¹H-NMR (400 MHz, CDCl₃) δ: 0.99 (broad s, 1 H, 9'-H), 1.43-1.65 complex signal, 14 H, 5'(11')-H₂, 6'(12')-H₂, 3(5)-H₂ and 4-H₂], 2.13 [broad s, 2 H, 4'(10')-H], 2.21 [broad s, 2 H, 2'(3')-H], 2.78 [broad s, 2 H, 1'(7')-H], 3.31 [m, 4 H, 2(6)-H₂], 4.53 (broad s, 1 H, NH); ¹³C-NMR (100.5 MHz, CDCl₃) δ: 21.5 [CH₂, C5'(11')], 24.5 (CH₂, C4), 24.6 [CH₂, C6'(12')], 25.6 [CH₂, C3(5)], 45.2 [CH₂, C2(6)], 46.8 [CH, C2'(3')], 52.8 [CH, C4'(10')], 53.6 [CH, C1'(7')], 55.4 (CH, C9'), 65.8 (C, C8'), 157.4 (C, CO); HRMS-ESI+ m/z [*M*+H]⁺ calcd for [C₁₈H₂₆N₂O+H]⁺: 287.2118, found: 287.2118.

6.1.6. N-[[Pentacyclo[6.4.0.0^{2,10}.0^{3,7}.0^{4,9}]dodec-8-yl]methyl]piperidine-1-carboxamide, (9).

From amine **5** [38] (96 mg, 0.42 mmol), 1-piperidinecarbonyl chloride (0.08 mL, 0.63 mmol) and triethylamine (0.21 mL, 1.68 mmol) in DCM (5 mL) and following the general procedure B, amide **9** (125 mg, quantitative yield) was obtained as a white solid. The analytical sample was obtained by crystallization from hot EtOAc (25 mg), mp 144–145 °C; IR (ATR) v: 554, 572, 623, 637, 678, 734, 763, 851, 869, 900, 908, 945, 969, 992, 1023, 1107, 1155, 1232, 1253, 1261, 1340, 1397, 1438, 1451, 1475, 1524, 1614, 2018, 2158, 2842, 2860, 2930, 3378 cm⁻¹; ¹H-NMR (400 MHz, CDCl₃) δ: 1.06 (broad s, 1 H, 9'-H), 1.40-1.70 [complex signal, 14 H, 5'(11')-H₂, 6'(12')-H₂, 3(5)-H₂ and 4-H₂], 2.00 [broad s, 2 H, 4'(10')], 2.13 [broad s, 2 H, 1'(7')-H], 2.22 [broad s, 2 H, 2'(3')-H], 3.28 [m, 4 H, 2(6)-H₂], 3.37 (d, *J* = 5.2 Hz, 2 H, NCH₂), 4.18 (broad s, 1H, NH); ¹³C-

NMR (100.5 MHz, CDCl₃) δ : 22.4 [CH₂, C5'(11')], 24.3 [CH₂, C6'(12')], 24.4 (CH₂, C4), 25.6 [CH₂, C3(5)], 38.6 (CH₂, NCH₂), 44.5 (C, C8'), 45.0 [CH₂, C2(6)], 47.9 [CH, C2'(3')], 51.2 (CH, C9'), 53.1 [CH, C1'(7')], 53.6 [CH, C4'(10')], 157.8 (C, CO); HRMS-ESI+ m/z [M+H]⁺ calcd for [C₁₉H₂₉N₂O+H]⁺: 301.2274, found: 301.2276. 6.1.7. (3-Azahexacyclo[7.6.0.0^{1,5}.0^{5,12}.0^{6,10}.0^{11,15}]pentadeca-7,13-dien-3-yl)(cyclohexyl) methanone, (II).

From amine 10 [38] (400 mg, 2.03 mmol), cyclohexanecarboxylic acid (237 mg, 1.85 mmol), HOBt (375 mg, 2.78 mmol), EDC (430 mg, 2.78 mmol) and triethylamine (0.570 mL, 4.07 mmol) in ethyl acetate (30 mL) and following the general procedure C, an orange oil (627 mg) was obtained. Column chromatography (Al₂O₃, DCM/methanol) gave amide 11 (260 mg, 42% yield) as a white solid. The analytical sample was obtained by crystallization from tert-butanol, mp 111–113 °C; IR (ATR) v: 2961, 2930, 2853, 1630, 1443, 1425, 1347, 1305, 1223, 1194, 1138, 1067, 998, 897, 878, 830, 780, 749, 734, 696, 659, 646 cm⁻¹; ¹H-NMR (500 MHz, CDCl₃) δ: 1.17-1.24 [complex signal, 3 H, 4'-H_{ax}, 3'(5')-H_{ax}], 1.43 [complex signal, 2 H, 2'(6')-H_{ax}], 1.60-1.66 [complex signal, 3 H, 2'(6')-H_{eq}, 4'-H_{eq}], 1.75-1.78 [complex signal, 2 H, 3'(5')-H_{eq}], 2.13 (tt, J = 12.0 Hz, J' = 3.5 Hz, 1 H, 1'-H), 2.67 [m, 2 H, 10(11)-H], 2.98 [complex signal, 4 H, 6(12)-H, 9(15)-H], 3.18 (s, 2 H, 2-H₂ or 4-H₂), 3.19 (s, 2 H, 4-H₂ or 2-H₂), 6.00 [ddd, J = 6 Hz, J' = 3 Hz, J'' = 1.5 Hz, 2 H, 7(13)-H or 8(14)-H], 6.04 [ddd, J = 6Hz, J' = 3 Hz, J'' = 1 Hz, 2 H, 8(14)-H or 7(13)-H]; ¹³C-NMR (125.7 MHz, CDCl₃) δ : 25.8 (CH₂, C4'), 25.9 [CH₂, C3'(5')], 28.7 [CH₂, C2'(6')], 42.5 (CH, C1'), 45.5 (CH₂, C2 or C4), 46.7 (CH₂, C4 or C2), 62.0 [CH, C6(12) and C9(15)], 62.8 [CH, C10(11)], 69.1 (C, C1 or C5), 70.8 (C, C5 or C1), 132.8 [CH, C7(13) or 8(14)], 134.1 [CH, C8(14) or C7(13)], 174.4 (C, CO); MS (EI), (rt = 25.4 min), m/z (%); significant ions: 308 (20), 307 (M^{+} , 83), 252 (16), 242 (25), 198 (17), 197 (100), 196 [($C_{14}H_{14}N$)⁺, 39], 182 (19), 181 (15), 180 (41), 179 (16), 168 (24), 167 (25), 166 (15), 165 (38), 156 (21), 153 (23), 152 (27), 132 (64), 131 (100), 130 (64), 128 (17), 118 (19), 117 (19), 115 (23), 91 (18), 83 $[(C_6H_{11})^+, 85]$, 77 (15), 55 (72); HRMS-ESI+ m/z $[M+H]^+$ calcd for $[C_{21}H_{25}NO+H]^+$: 308.2009, found: 308.2003.

6.1.8. $(3-Azahexacyclo[7.6.0.0^{1.5}.0^{5.12}.0^{6.10}.0^{11.15}]$ pentadecane-3-yl)(cyclohexyl) methanone, (12).

From amide **11** (118 mg, 0.40 mmol) and Pd/C (13 mg) and following the general procedure D (5 h), amide **12** (94 mg, 78% yield) was obtained as a white solid. The

analytical sample was obtained by crystallization from DCM/diethyl ether, mp 134–135 °C; IR (ATR) v: 2932, 2851, 1621, 1463, 1426, 1357, 1286, 1202, 1120, 1104, 1036, 1013, 969, 925, 886, 860, 825, 768, 732, 702, 657, 627 cm⁻¹; ¹H-NMR (500 MHz, CDCl₃) δ : 1.25 [complex signal, 3 H, 4'-H_{ax}, 3'(5')-H_{ax}], 1.48-1.57 [complex signal, 10 H, 7(13)-H₂, 8(14)-H₂, 17(21)-H_{ax}], 1.68 (m, 1 H, 4'-H_{eq}), 1.75-1.82 [complex signal, 4 H, 2'(6')-H_{eq}, 3'(5')-H_{eq}], 2.08 (broad signal, 4 H, 6(12)-H, 9(15)-H], 2.36 (tt, J = 12 Hz, J' = 3.5 Hz, 1 H, 1'-H), 2.41 [m, 2 H, 10(11)-H], 3.29 [s, 4 H, 2(4)-H₂]; ¹³C-NMR (125.7 MHz, CDCl₃) δ : 21.5 [CH₂, C7(13) or C8(14)], 21.8 [CH₂, C8(14) or C7(13)], 25.8 (CH₂, C4'), 25.9 [CH₂, C3'(5')], 29.0 [CH₂, C2'(6')], 40.7 (CH₂, C2 or C4), 42.0 (CH₂, C4 or C2), 42.9 (CH, C1'), 49.6 [CH, C10(11)], 55.0 [CH, C6(12) or C9(15)], 55.1 [CH, C9(15) or C6(12)], 57.5 (C, C1 or C5], 59.3 (C, C5 or C1], 175.1 (C, CO); MS (EI), (rt = 26.7 min), m/z (%); significant ions: 312 (23), 311 (M⁻⁺, 100), 270 (14), 257 (19), 256 (99), 243 (12), 228 [(C₁₅H₁₈NO)⁺ 13], 202 (12), 201 (30), 184 (29), 129 (15), 128 (15), 91 (16), 83 [(C₆H₁₁)⁺, 24], 55 (25). Anal. Calcd for C₂₁H₂₉NO: C 80.98, H 9.39, N 4.50. Found: 80.76, H 9.61, N 4.33.

6.1.9. $(3-Azahexacyclo[7.6.0.0^{1.5}.0^{5.12}.0^{6.10}.0^{11.15}]$ pentadeca-7,13-diene-3-yl)(piperidin-1-yl)methanone, (13).

From amine **10** [38] (400 mg, 2.03 mmol), 1-piperidinecarbonyl chloride (0.26 mL, 2.13 mmol) and triethylamine (0.56 mL, 4.06 mmol) in DCM and following the general procedure B amide **13** (443 mg, 71% yield) was obtained as a clear oil. Several attempts to crystallize this product met with failure. The product was used in the next step without further purification or characterization; MS (EI), (rt = 24.2 min), m/z (%); significant ions: 308 (M⁻⁺, 46), 196 [(C₁₄H₁₄N)⁺, 17], 165 (14), 130 (15), 112 [(C₆H₁₀NO)⁺, 100], 84 [(C₅H₁₀N)⁺, 17], 69 (41).

 $6.1.10. (3-Azahexacyclo[7.6.0.0^{1.5}.0^{5.12}.0^{6.10}.0^{11.15}]$ pentadeca-3-yl)(piperidin-1-yl) methanone, (14).

From urea **13** (235 mg, 0.76 mmol) and Pd/C (24 mg) and following the general procedure D (3 h), urea **14** (171 mg, 72% yield) was obtained as a white solid, mp 124–126 °C; IR (ATR) v: 3457, 3291, 2936, 2867, 2839, 1615, 1538, 1461, 1415, 1370, 1332, 1304, 1252, 1226, 1202, 1159, 1124, 1110, 1027, 991, 915, 882, 851, 767, 722, 632 cm⁻¹; ¹H-NMR (500 MHz, CDCl₃) δ: 1.47-1.58 [complex signal, 14 H, 7(8,13,14)-H₂, 3'(5')-H₂, 4'-H₂], 2.05 [m, 4 H, 6(9,12,15)-H], 2.38 [m, 2 H, 10(11)-H], 3.19 [m, 8 H, 2(4)-H₂, 2'(6')-H₂]; ¹³C-NMR (125.7 MHz, CDCl₃) δ: 21.7 [CH₂, C7(8, 13, 14)],

24.8 (CH₂, C4'), 25.9 [CH₂, C3'(5')], 43.3 [CH₂, C2(4)], 47.8 [CH₂, C16(20)], 49.6 [CH, C10(11)], 54.8 [CH, C6(9, 12, 15)], 58.8 [C, C1(5)], 163.3 (C, CO); MS (EI), (rt = 25.1 min), m/z (%); significant ions: 312 (M⁻⁺, 100), 201 (15), 200 [(C₁₄H₁₈N)⁺, 82], 184 (33), 129 (36), 112 [(C₆H₁₀NO)⁺, 54], 91 (16), 84 [(C₅H₁₀N)⁺, 59], 69 (28). Anal. Calcd for C₂₀H₂₈N₂O: C 76.88, H 9.03, N 8.97. Found: 76.60, H 9.21, N 8.74.

 $6.1.11.\ (4-Amino-3,5-dichlorophenyl) (3-$

 $azahexacyclo[7.6.0.0^{1.5}.0^{5.12}.0^{6.10}.0^{11.15}]$ pentadeca-7,13-dien-3-yl) methanone, (15).

From amine 10 [38] (400 mg, 2.03 mmol), 4-amino-3,5-dichlorobenzoic acid (380 mg, 1.85 mmol), 1-hydroxybenzotriazole (HOBt) (375 mg, 2.78 mmol), 1-ethyl-3-(3dimethylaminopropyl)carbodiimide (EDC) (430 mg, 2.78 mmol) and triethylamine (0.560 mL, 4.07 mmol) in EtOAc (30 mL) and DMF (2 mL) and following the general procedure C, a yellow solid (677 mg) was obtained. Column chromatography (Al₂O₃, DCM/methanol) furnished 15 (332 mg, 47% yield) as a white solid, mp 236–238 °C; IR (ATR) v: 3449, 3306, 3250, 3204, 2954, 2867, 2150, 1597, 1538, 1501, 1456, 1410, 1384, 1342, 1316, 1295, 1244, 1225, 1192, 1054, 1002, 943, 896, 875, 791, 745, 733, 686, 667, 643 cm⁻¹; ¹H-NMR (500 MHz, CDCl₃) δ: 2.61 [m, 2 H, 10(11)-H], 2.85 [broad s, 2 H, 6(12)-H or 9(15)-H], 2.94 [broad s, 2 H, 9(15)-H or 6(12)-H], 3.13 (broad s, 2 H, 2-H or 4-H), 3.33 (broad s, 2 H, 4-H or 2-H), 5.91 [m, 2 H, 7(13)-H or 8(14)-H], 6.03 [m, 2 H, 8(14)-H or 7(13)-H], 7.19 (s, 2 H, Ar-H); ¹³C-NMR (125.7 MHz, CDCl₃) δ: 45.5 (CH₂, C2 or C4), 49.9 (CH₂, C4 or C2), 61.8 [CH, C6(12), C9(15)], 62.7 [CH, C10(11)], 68.8 (C, C1 or C5), 71.0 (C, C5 or C1), 118.7 (C, Ar-C_{meta}), 126.7 (C, C_{ipso}), 127.1 (CH, Ar-C_{ortho}), 132.9 [CH, C7(13) or 8(14)], 133.9 [CH, C8(14) or C7(13)], 141.3 (C, Ar-C_{para}), 166.8 (C, CO); GC/MS (EI), (rt = 30.7 min), m/z (%); significant $[(C_{21}H_{18}^{37}Cl_2N_2O)^{+}, 2], 386 [(C_{21}H_{18}^{37}Cl_2^{35}Cl_2N_2O)^{+}, 12],$ 388 $[(C_{21}H_{18}^{35}Cl_2N_2O)^{+}, 18], 192 [(C_7H_4^{37}Cl_2NO)^{+}, 10], 190 [(C_7H_4^{37}Cl_3^{35}ClNO)^{+}, 63], 188$ $[(C_7H_4^{35}Cl_2NO)^+, 100]$, 180 (12), 160 (12), 124 (16); HRMS-ESI+ m/z $[M+H]^+$ calcd for $[C_{21}H_{18}Cl_2N_2O+H]^+$: 385.0869, found: 385.0875.

6.1.12. (4-Amino-3,5-dichlorophenyl) (3-azahexacyclo[7.6.0.0^{1,5}.0^{5,12}.0^{6,10}.0^{11,15}] pentadecan-3-yl)methanone, (**16**).

From amide **15** (200 mg, 0.78 mmol) and Pd/C (24 mg) and following the general procedure D (5 h) a yellow solid (207 mg) was obtained. Column chromatography (Al₂O₃, DCM/methanol) gave the desired amide **16** (180 mg, 89% yield) as a white solid, mp 243–244 °C; IR (ATR) v: 3455, 3283, 3238, 3186, 2931, 2865, 1631, 1597,

1545, 1498, 1456, 1425, 1332, 1307, 1269, 1232, 1216, 1118, 1070, 1035, 944, 895, 789, 768, 745, 693, 647, 629 cm⁻¹; ¹H-NMR (500 MHz, CDCl₃) δ : 1.43-1.60 [complex signal, 8 H, 7(13)-H₂, 8(14)-H₂], 2.04 [broad s, 2 H, 6(12)-H or 9(15)-H], 2.12 [broad s, 2 H, 9(15)-H or 6(12)-H], 2.41 [m, 2 H, 10(11)-H], 3.29 (s, 2 H, 2-H₂ or 4-H₂), 3.53 (s, 2 H, 4-H₂ or 2-H₂), 7.40 (s, 2 H, Ar-H); ¹³C-NMR (125.7 MHz, CDCl₃) δ : 21.5 [CH₂, C7(13) or C8(14)], 21.8 [CH₂, C8(14) or C7(13)], 40.8 (CH₂, C2 or C4), 45.5 (CH₂, C4 or C2), 49.6 [CH, C10(11)], 54.8 [broad CH, C6(12) and C9(15)], 57.9 (C, C1 or C5), 59.6 (C, C5 or C1), 118.8 (C, Ar-C_{meta}), 126.9 (C, C_{ipso}), 127.2 (CH, Ar-C_{ortho}), 141.4 (C, Ar-C_{para}), 167.0 (C, CO); MS (EI), (rt = 32.3 min), m/z (%); significant ions: 392 [(C₂₁H₂₂³⁷Cl₂N₂O)⁻⁺, 8], 390 [(C₂₁H₂₂³⁷Cl³⁵ClN₂O)⁻⁺, 42], 388 [(C₂₁H₂₂³⁵Cl₂N₂O)⁻⁺, 63], 200 (18), 192 [(C₇H₄³⁷Cl₂NO)⁺, 11], 190 [(C₇H₄³⁷Cl³⁵ClNO)⁺, 64], 188 [(C₇H₄³⁵Cl₂NO)⁺, 100], 184 (33), 169 (13), 160 (12), 124 (13). Anal. Calcd for C₂₁H₂₂Cl₂N₂O·0.50H₂O: C 63.32, H 5.82, Cl 17.80, N 7.03. Found: C 63.23, H 5.71, Cl 17.82, N 6.77.

6.1.13. (Cyclohexyl)(octahydro-2H-isoindol-2-yl)methanone, (17).

From octahydro-1*H*-isoindole hydrochloride (300)2.40 mmol), mg, cyclohexanecarboxylic acid (279 mg, 2.18 mmol), HOBt (442 mg, 3.27 mmol), EDC (506 mg, 3.27 mmol) and triethylamine (0.7 mL, 4.80 mmol) in EtOAc (10 mL) and following the general procedure C, 17 (458 mg, 89% yield) was obtained as a white solid. The analytical sample was obtained by crystallization from hot EtOAc (82 mg), mp 67 – 68 °C; IR (ATR) v: 734, 890, 973, 1073, 1113, 1136, 1173, 1184, 1307, 1341, 1358, 1443, 1481, 1622, 2851, 2873, 2917 cm⁻¹; ¹H-NMR (400 MHz, CDCl₃) δ: 1.16-1.30 (complex signal, 3 H, 3-H_{ax}, 4-H_{ax} and 5-H_{ax}), 1.31-1.63 (complex signal, 10 H, 2-H_{ax}, 6-H_{ax}, 4'-H₂, 7'-H₂, 5'-H₂, 6'-H₂), 1.64-1.84 (complex signal, 5 H, 2-H_{eq}, 6-H_{eq}, 3-H_{eq}, 4-H_{eq} and 5-H_{eq}), 2.16 (m, 1 H, 3a'-H or 7a'-H), 2.24 (m, 1 H, 7a'-H or 3a'-H), 2.30 [tt, J = 11.6 Hz, J' = 3.6 Hz, 1 H, 1-H], 3.31 (dd, J = 10.0 Hz, J' = 6.0 Hz, 1 H, 1'- H_a or 3'- H_a), 3.36 (dd, J = 12.0 Hz, J' = 6.6 Hz, 1 H, 3'- H_a or 1'- H_a), 3.40 (dd, J = 12.0Hz, J' = 7.8 Hz, 1 H, 3'-H_b or 1'-H_b), 3.45 (dd, J = 10.0 Hz, J' = 7.0 Hz, 1 H, 1'-H_b or 3'-H_b); ¹³C-NMR (100.5 MHz, CDCl₃) δ: 22.5 (CH₂, C5' or C6'), 22.8 (CH₂, C6' or C5'), 25.73 (CH₂), 25.78 (CH₂), 25.80 (CH₂), 25.84 (CH₂) and 25.91 (CH₂) [C4', C7', C3, C4 and C5], 28.8 (CH₂, C2 or C6), 29.0 (CH₂, C6 or C2), 35.8 (CH, C3a' or C7a'), 37.6 (CH, C7a' or C3a'), 42.7 (CH, C1), 49.3 (CH₂, C1' or C3'), 50.4 (CH₂, C3' or

C1'), 175.4 (C, CO). Anal. Calcd for $C_{15}H_{25}NO$: C 76.55, H 10.71, N 5.95. Found: 76.56, H 10.67, N 5.96.

6.1.14 (4-Azatricyclo[5.2.1.0^{2,6}]dec-8-en-4-yl)(cyclohexyl)methanone, (18).

From 4-azatricyclo[5.2.1.0^{2,6}]dec-8-ene hydrochloride [58] (240 mg, 1.40 mmol), cyclohexanecarboxylic acid (163 mg, 1.27 mmol), HOBt (258 mg, 1.91 mmol), EDC (296 mg, 1.91 mmol) and triethylamine (0.8 mL, 5.59 mmol) in EtOAc (8 mL) and following the general procedure C, amide 18 (304 mg, 97% yield) was obtained as a yellowish solid. Column chromatography (hexane/EtOAc) gave 18 as a white solid (209 mg), mp 77–78 °C; IR (ATR) v: 702, 731, 761, 793, 897, 987, 1216, 1256, 1332, 1347, 1428, 1621, 2850, 2920 cm⁻¹; ¹H-NMR (400 MHz, CDCl₃) δ: 1.12-1.29 (complex signal, 3 H, 3-H_{ax}, 4-H_{ax} and 5-H_{ax}), 1.36-1.51 (complex signal, 3 H, 2-H_{ax}, 6-H_{ax} and $10'-H_a$], 1.54 (dt, J = 8.4 Hz, J' = 1.6 Hz, 1 H, $10'-H_b$), 1.60-1.70 [complex signal, 3 H, $2-H_{eq}$, $6-H_{eq}$ and $4-H_{eq}$), 1.71-1.84 (complex signal, 2 H, $3-H_{eq}$ and $5-H_{eq}$), 2.16 (tt, J=11.6 Hz, J' = 3.6 Hz, 1 H, 1-H), 2.82-3.00 (complex signal, 4 H, 1'-H, 7'-H, 2'-H and 6'-H), 3.10 (dd, J = 10.8 Hz, J' = 3.6 Hz, 1 H, 3'-H_a or 5'-H_a), 3.21 (dd, J = 13.5 Hz, J' $= 3.6 \text{ Hz}, 1 \text{ H}, 5' - \text{H}_a \text{ or } 3' - \text{H}_a), 3.29 \text{ (dd, } J = 13.5 \text{ Hz, } J' = 9.2 \text{ Hz, } 1 \text{ H, } 5' - \text{H}_b \text{ or } 3' - \text{H}_b),$ 3.43 (dd, J = 10.8 Hz, J' = 9.2 Hz, 1 H, 3'-H_b or 5'-H_b), 6.14 (dd, J = 6.0 Hz, J' = 3.2Hz, 1 H, 8'-H or 9'-H), 6.19 (dd, J = 6.0 Hz, J' = 3.2 Hz, 1 H, 9'-H or 8'-H); 13 C-NMR (100.5 MHz, CDCl₃) δ: 25.8 [CH₂, C3(5)], 25.9 (CH₂, C4), 28.69 (CH₂, C2 or C6), 28.71 (CH₂, C6 or C2), 42.7 (CH, C1), 43.8 (CH, C2' or C6'), 45.9 (CH, C6' or C2'), 46.66 (CH, C1' or C7'), 46.70 (CH, C7' or C1'), 48.0 (CH₂, C3' or C5'), 48.8 (CH₂, C5' or C3'), 51.9 (CH₂, C10'), 134.7 (CH, C8' or C9'), 136.1 (CH, C9' or C8'), 173.9 (C, CO). Anal. Calcd for C₁₆H₂₃NO: C 78.32, H 9.45, N 5.71. Found: 78.03, H 9.41, N 5.58.

6.1.15.(4-Azatricyclo[5.2.1.0^{2,6}]dec-4-yl)(cyclohexyl)methanone, (**19**).

From amide **18** (177 mg) and Pd/C (36 mg) and following the general procedure D (18 h), amide **19** (156 mg, 88% yield) was obtained as a white solid, mp 83-84 °C; IR (ATR) v: 611, 625, 642, 886, 1002, 1133, 1170, 1187, 1204, 1218, 1290, 1327, 1344, 1357, 1425, 1446, 1623, 2871, 2936, 2945 cm⁻¹; ¹H-NMR (400 MHz, CDCl₃) δ : 1.20–1.94 (complex signal, 16 H, 3-H₂, 4-H₂, 5-H₂, 2-H₂, 6-H₂, 8'-H₂, 9'-H₂ and 10'-H₂), 2.18-2.27 (complex signal, 2 H, 1'-H and 7'-H), 2.37 (tt, J = 11.6 Hz, J' = 3.2 Hz, 1 H, 1-H), 2.52 (m, 1 H, 2'-H or 6'-H), 2.60 (m, 1 H, 6'-H or 2'-H), 3.03 (dd, J = 13.2 Hz, J' = 8.4 Hz, 1 H, 3'-H_a or 5'- H_a), 3.26 (dd, J = 11.6 Hz, J' = 7.6 Hz, 1 H, 5'-H_a or 3'- H_a),

3.57 (dd, J = 11.6 Hz, J' = 1.6 Hz, 1 H, 5'-H_b or 3'- H_b), 3.84 (dd, J = 13.2 Hz, J' = 1.6 Hz, 1 H, 3'-H_b or 5'- H_b); ¹³C-NMR (100.5 MHz, CDCl₃) δ : 22.1 (CH₂, C8' or C9'), 22.8 (CH₂, C9' or C8'), 25.79 (CH₂), 25.86 (CH₂) and 25.88 (CH₂) (C3, C4 and C5), 28.87 (CH₂, C2 or C6), 28.89 (CH₂, C6 or C2), 41.2 (CH, C1' or C7'), 41.4 (CH, C7' or C1'), 42.0 (CH₂, C2' or C6'), 42.1 (CH₂, C10'), 42.9 (CH, C1), 44.0 (CH₂, C6' or C2'), 45.7 (CH₂, C3' or C5'), 46.9 (CH₂, C5' or C3'), 174.4 (C, CO). Anal. Calcd for C₁₆H₂₅NO: 77.68, H 10.19, N 5.66. Found: C 77.55, H 10.05, N 5.54.

6.1.16. (4-Azatricyclo[5.2.2.0^{2,6}]undec-8-en-4-yl)(cyclohexyl)methanone, (**20**).

From 4-azatricyclo[5.2.2.0^{2,6}]undec-8-ene hydrochloride [59] (300 mg, 1.62 mmol), cyclohexanecarboxylic acid (188 mg, 1.47 mmol), HOBt (300 mg, 2.21 mmol), EDC (342 mg, 2.21 mmol) and triethylamine (0.9 mL, 6.47 mmol) in EtOAc (10 mL) and following the general procedure C, amide 20 (323 mg, 85% yield) was obtained as a yellowish solid. Column chromatography (hexane/EtOAc) gave 20 as a white solid (205 mg), mp 86-87 °C; IR (ATR) v: 703, 715, 849, 887, 986, 1044, 1132, 1165, 1216, 1239, 1307, 1347, 1357, 1375, 1431, 1625, 2850, 2921, 3037 cm⁻¹; ¹H-NMR (400 MHz, CDCl₃) δ : 1.15-1.34 (complex signal, 5 H, 10'-H_a, 11'-H_a, 3-H_{ax}, 4-H_{ax} and 5-H_{ax}), 1.37-1.54 (complex signal, 4 H, 2-H_{ax}, 6-H_{ax}, 10'-H_b, 11'-H_b), 1.58-1.86 (complex signal, 5 H, 2-H_{eq}, 6-H_{eq}, 3-H_{eq}, 4-H_{eq} and 5-H_{eq}), 2.23 (tt, J = 11.6 Hz, J' = 3.6 Hz, 1 H, 1-H), 2.42 (m, 1 H, 2'-H or 6'-H), 2.51-2.61 (complex signal, 3 H, 1'-H, 7'-H, 6'-H or 2'-H), 3.09 (d, J = 12.8 Hz, 1 H, 3'-H_a or 5'-H_a), 3.11 (d, J = 12.4 Hz, 1 H, 5'-H_a and 3'- H_a), 3.62 (dd, J = 12.4 Hz, J' = 9.2 Hz, 1 H, 5'- H_b or 3'- H_b), 3.65 (dd, J = 12.8 Hz, J'= 9.6 Hz, 1 H, 3'- H_b and 5'- H_b), 6.15-6.26 (complex signal, 2 H, 8'-H and 9'-H): 13 C-NMR (100.5 MHz, CDCl₃) δ: 24.2 (CH₂, C10' or C11'), 24.3 (CH₂, C11' or C10'), 25.81 (CH₂), 25.84 (CH₂) and 25.9 (CH₂) [C3, C4 and C5], 28.7 (CH₂, C2 or C6), 28.8 (CH₂, C6 or C2), 34.1 (CH, C1' and C7'), 41.9 (CH, C2' or C6'), 42.6 (CH, C1), 44.0 (CH, C6' or C2'), 50.8 (CH₂, C3' or C5'), 51.6 (CH₂, C5' or C3'), 132.9 (CH, C8' or C9'), 134.3 (CH, C9' or C8'), 173.9 (C, CO). Anal. Calcd for C₁₇H₂₅NO: C 78.72, 9.71, N 5.40. Found: C 78.82, H 9.71, N 5.30.

6.1.17. (4-Azatricyclo[5.2.2.0^{2,6}]undec-4-yl)(cyclohexyl)methanone, (21).

From amide **20** (165 mg) and Pd/C (33 mg) and following the general procedure D (18 h) amide **21** (151 mg, 91% yield) was obtained as a white solid, mp 78-79 °C; IR (ATR) v: 622, 725, 868, 976, 1138, 1173, 1204, 1346, 1429, 1443, 1622, 2861, 2901, 2923 cm⁻¹; ¹H-NMR (400 MHz, CDCl₃) δ : 1.18-1.42 (complex signal, 5 H, 3-H_{ax}, 4-

H_{ax}, 5-H_{ax}, 8'-H_a and 9'-H_a), 1.44-1.90 (complex signal, 15 H, 3-H_{eq}, 4-H_{eq}, 5-H_{eq}, 2-H₂, 6-H₂, 1'-H, 7'-H, 8'-H_b, 9'-H_b, 10'-H₂ and 11'-H₂), 2.33 (m, 1 H, 2'-H or 6'-H), 2.37 (tt, J = 11.2 Hz, J' = 3.4 Hz, 1 H, 1-H), 2.43 (m, 1 H, 6'-H and 2'-H), 3.44-3.65 (complex signal, 4 H, 3'-H₂ and 5'-H₂); ¹³C-NMR (100.5 MHz, CDCl₃) δ: 19.8 (CH₂, C9' or C8'), 20.1 (CH₂, C8' or C9'), 25.7 (CH, C1' and C7'), 25.81 (CH₂), 25.86 (CH₂) and 25.91 (CH₂) (C3, C4 and C5), 27.8 (CH₂, C11' or C10'), 28.0 (CH₂, C10' or C11'), 28.86 (CH₂, C2 or C6), 28.92 (CH₂, C6 or C2), 37.8 (CH₂, C2' or C6'), 40.0 (CH₂, C6' or C2'), 42.8 (CH, C1), 49.1 (CH₂, C3' or C5'), 49.9 (CH₂, C5' or C3'), 174.3 (C, CO). Anal. Calcd for C₁₇H₂₇NO: C 78.11, H 10.41, N 5.36. Found: C 78.14, H 10.35, N 5.14. 6.1.18. (4-Azatetracyclo[5.3.2.0^{2,6}.0^{8,10}]dodec-11-en-4-yl)(cyclohexyl)methanone, (22). From 4-azatetracyclo[5.3.2.0^{2,6}.0^{8,10}]dodec-11-ene hydrochloride [40] (2.06 g, 10.6 mmol), cyclohexanecarboxylic acid (1.24 g, 9.67 mmol), HOBt (1.96 g, 14.5 mmol), EDC (2.25 g, 14.5 mmol) and triethylamine (5.9 mL, 42.5 mmol) in EtOAc (150 mL) and following the general procedure C, amide 22 (2.43 g, 94% yield) was obtained as a yellowish solid. The analytical sample was obtained by crystallization from hot EtOAc (2.04 g), mp 96–97 °C; IR (ATR) v: 560, 570, 587, 696, 718, 741, 767, 812, 829, 847, 894, 915, 942, 963, 991, 1036, 1089, 1134, 1167, 1209, 1217, 1242, 1272, 1299, 1361, 1380, 1432, 1624, 2849, 2925, 3002, 3040 cm⁻¹; ¹H-NMR (400 MHz, CDCl₃) δ: 0.12-0.16 (complex signal, 2 H, 9'-H₂), 0.86-0.96 (complex signal, 2 H, 8'-H and 10'-H), 1.14-1.29 (complex signal, 3 H, 3-H_{ax}, 4-H_{ax} and 5-H_{ax}), 1.36-1.53 (complex signal, 2 H, 2-H_{ax} and 6-H_{ax}), 1.60-1.70 (complex signal, 3 H, 2-H_{eq}, 4-H_{eq} and 6-H_{eq}), 1.72-1.80 (complex signal, 2 H, 3-H_{eq} and 5-H_{eq}), 2.21 (tt, J = 11.6 Hz, J' = 3.4 Hz, 1 H, 1-H), 2.55 (dm, J = 12.8 Hz, 1 H, 2'-H or 6'-H), 2.67 (dm, J = 12.8 Hz, 1 H, 6'-H or 2'-H), 2.81-2.87 (complex signal, 2 H, 1'-H and 7'-H), 3.12 (dd, J = 11.0 Hz, J' = 5.0 Hz, 1 H, 3'- H_a or 5'- H_a), 3.15 (dd, J = 13.2 Hz, J' = 5.6 Hz, 1 H, 5'- H_a or 3'- H_a), 3.55 (t, J = 10.0Hz, 1 H, 5'-H_b or 3'-H_b), 3.58 (t, J = 8.8 Hz, 1 H, 3'-H_b or 5'-H_b), 5.73 (ddd, J = 14.4Hz, J' = 8.4 Hz, J'' = 2.0 Hz, 1 H, 11'-H or 12'-H], 5.77 (ddd, J = 14.4 Hz, J' = 8.4 Hz, J'' = 2.0 Hz, 1 H, 12'-H or 11'-H]; ¹³C-NMR (100.5 MHz, CDCl₃) δ : 4.1 (CH₂, C9'), 10.0 (CH, C8' or C10'), 10.2 (CH, C10' or C8'), 25.80 (CH₂), 25.83 (CH₂) and 25.9 (CH₂) [C3, C4 and C5], 28.7 (CH₂, C2 or C6), 28.8 (CH₂, C6 or C2), 35.6 (CH, C1' or C7'), 35.7 (CH, 7' or C1'), 42.6, (CH, C1), 42.7 (CH, C2' or C6'), 44.8 (CH, C6' or C2'), 49.6 (CH₂, C3' or C5'), 50.6 (CH₂, C5' or C3'), 128.1 (CH, C11' or C12'), 129.6

(CH, C12' or C11'), 174.1 (C, CO). Anal. Calcd for C₁₈H₂₅NO: C 79.66, H 9.29, N 5.16. Found: C 79.64, H 9.24, N 5.21.

 $6.1.19.\ (4-Azatetracyclo[5.3.2.0^{2,6}.0^{8,10}]dodec-4-yl)(cyclohexyl)$ methanone, (23).

From 22 (500 mg) and Pd/C (100 mg) and following the general procedure D (72 h) amide 23 (426 mg, 83% yield) was obtained as a white solid, mp 74–75 °C; IR (ATR) v: 651, 679, 707, 729, 748, 789, 805, 826, 839, 865, 885, 950, 961, 988, 1016, 1030, 1082, 1113, 1133, 1171, 1205, 1213, 1234, 1264, 1295, 1325, 1346, 1358, 1427, 1471, 1486, 1623, 2846, 2897, 2928, 3009, cm⁻¹; ¹H-NMR (400 MHz, CDCl₃) δ : 0.45 (dt, J =6.0 Hz, J' = 8.0 Hz, 1 H, $9' \cdot \text{H}_a$), 0.78 (m, J = 6.0 Hz, J' = 3.6 Hz, 1 H, $9' \cdot \text{H}_b$), $0.90 \cdot 0.96 \cdot 0.$ (complex signal, 2 H, 8'-H and 10'-H), 1.00-1.14 (complex signal, 2 H, 11'-H_{ax} and 12'-H_{ax}), 1.17-1.34 [complex signal, 5 H, 11'-H_{eq}, 12'-H_{eq}, 3-H_{ax}, 4-H_{ax} and 5-H_{ax}], 1.42-1.61 (complex signal, 2 H, 2-H_{ax} and 6-H_{ax}), 1.67 (m, 1 H, 4'-H_{eq}), 1.69-1.83 (complex signal, 4 H, 3-H_{eq}, 4-H_{eq}, 5-H_{eq}, 6-H_{eq}), 1.84-1.91 (complex signal, 2 H, 1'-H and 7'-H), 2.38 [tt, J = 11.6 Hz, J' = 3.6 Hz, 1 H, 1-H], 2.50 (dm, J = 12.8 Hz, 1 H, 2'-H or 6'-H), 2.55 (dm, J = 12.8 Hz, 1 H, 6'-H or 2'-H), 3.32 (dd, J = 12.8 Hz, J' = 8.8Hz, 1 H, 3'-H_a or 5'-H_a), 3.52 (m, 2 H, 5'-H₂ or 3'-H₂), 3.76 (dd, J = 12.8 Hz, J' = 3.0Hz, 1 H, 3'-H_b or 5'-H_b); ¹³C-NMR (100.5 MHz, CDCl₃) δ: 4.7 (CH₂, C9'), 14.9 (CH, C8' or C10'), 15.1 (CH, C10' or C8'), 17.3 (CH₂, C11' or C12'), 17.9 (CH₂, C12' or C11'), 25.8 (CH₂, C4), 25.9 [CH₂, C3(5)], 28.87 (CH₂, C2 or C6), 28.92 (CH₂, C6 or C2), 29.0 (CH, C1' or C7'), 29.4 (CH, C7' or C1'), 38.4 (CH, C2' or C6'), 40.5, (CH, C6' or C2'), 42.7 (CH, C1), 48.1 (CH, C3' or C5'), 49.3 (CH₂, C5' or C3'), 174.4 (C, CO). Anal. Calcd for C₁₈H₂₇NO: C 79.07, H 9.95, N 5.12. Found: 79.15, H 9.88, N 5.29.

6.1.20. (4-Azatetracyclo[5.4.2.0^{2,6}.0^{8,11}]trideca-9,12-dien-4-yl)(cyclohexyl) methanone, (24).

From 4-azatetracyclo[5.4.2.0^{2,6}.0^{8,11}]trideca-9,12-diene hydrochloride [40] (139 mg, 0.66 mmol), cyclohexanecarboxylic acid (77 mg, 0.60 mmol), HOBt (122 mg, 0.90 mmol), EDC (139 mg, 0.90 mmol) and triethylamine (0.4 mL, 2.64 mmol) in EtOAc (6 mL) and following the general procedure C, amide **24** (151 mg, 80% yield) was obtained as a yellowish solid. The analytical sample was obtained by crystallization from hot EtOAc (97 mg), mp 120–121 °C; IR (ATR) v: 730, 760 788, 808, 967, 994, 1173, 1187, 1214, 1235, 1294, 1361, 1442, 1463, 1617, 2860, 2901, 2922 cm⁻¹; ¹H-NMR (400 MHz, CDCl₃) δ: 1.15-1.30 [complex signal, 3 H, 3-H_{ax}, 4-H_{ax} and 5-H_{ax}],

1.38-1.54 [complex signal, 2 H, 2- H_{ax} and 6- H_{ax}], 1.60-1.72 (complex signal, 3 H, 2- H_{eq} , 4- H_{eq} and 6- H_{eq}), 1.73-1.82 (complex signal, 2 H, 3- H_{eq} and 5- H_{eq}), 2.24 [tt, J=11.6 Hz, J'=3.6 Hz, 1 H, 1-H], 2.37 (tdd, J=9.6 Hz, J'=5.8 Hz, J''=2.8 Hz, 1 H, 2'-H or 6'-H), 2.50 (tdd, J=8.8 Hz, J'=5.6 Hz, J''=2.8 Hz, 1 H, 6'-H or 2'-H), 2.60-2.66 (complex signal, 2 H, 1'-H and 7'-H), 2.67-2.70 (complex signal, 2 H, 8'-H and 11'-H), 3.17-3.24 (complex signal, 2 H, 3'- H_{a} and 5'- H_{a}), 3.63-3.73 (complex signal, 2 H, 3'- H_{b} and 5'- H_{b}), 5.84-5.87 (complex signal, 2 H, 9'-H and 10'-H), 5.89-5.99 (complex signal, 2 H, 12'-H and 13'-H); 13 C-NMR (100.5 MHz, CDC1₃) &: 25.81 (CH₂), 25.84 (CH₂) and 25.9 (CH₂) (C3, C4 and C5), 28.7 (CH₂, C2 or C6), 28.8 (CH₂, C6 or C2), 39.36 (CH, C1' or C7'), 39.37 (CH, C7' or C1'), 41.1 (CH, C2' or C6'), 42.6 (CH, C1), 43.2 (CH, C6' or C2'), 44.9 (CH, C8' or C11'), 45.1 (CH, C11' or C8'), 50.7 (CH₂, C3' or C5'), 51.5 (CH₂, C5' or C3'), 129.0 (CH, C12' or C13'), 130.3 (CH, C13' or C12'), 137.7 (CH, C9' or C10'), 138.0 (CH, C10' or C9'), 174.0 (C, CO). Anal. Calcd for C₁₉H₂₅NO: C 80.52, H 8.89, N 4.94. Found: 80.31, H 8.81, N 4.97.

 $6.1.21.\ (4-Azatetracyclo[5.4.2.0^{2,6}.0^{8,11}]tridec-4-yl)(cyclohexyl)$ methanone, (25).

From 4-azatetracyclo[5.4.2.0^{2,6}.0^{8,11}]tridecane hydrochloride [40] (235 mg, 1.10 mmol), cyclohexanecarboxylic acid (128 mg, 1.0 mmol), HOBt (203 mg, 1.50 mmol), EDC (232 mg, 1.50 mmol) and triethylamine (0.6 mL, 4.40 mmol) in EtOAc (10 mL) and following the general procedure C, amide 25 (253 mg, 80% yield) was obtained as a yellowish solid. The analytical sample was obtained by crystallization from hot EtOAc (110 mg), mp 115–116 °C; IR (ATR) v: 658, 731, 887, 988, 1133, 1172, 1204, 1236, 1357, 1434, 1440, 1621, 2908, 2925 cm⁻¹; ¹H-NMR (400 MHz, CDCl₃) δ: 1.18-1.34 (complex signal, 3 H, 3-H_{ax}, 4-H_{ax} and 5-H_{ax}), 1.42-1.62 (complex signal, 6 H, 1'-H, 7'-H, 2-H_{ax}, 6-H_{ax}, 12'-H_a and 13'-H_a), 1.64-1.92 (complex signal, 7 H, 2-H_{eq}, 6-H_{eq}, 3-H_{eq}, 4-H_{eq}, 5-H_{eq}, 12'-H_b and 13'-H_b), 2.02-2.20 (complex signal, 5 H, 2'-H or 6'-H, 9'-H_a, 9'-H_b, 10'-H_a and 10'-H_b), 2.27 (m, 1 H, 6'-H or 2'-H), 2.34-2.44 (complex signal, 3 H, 1-H, 8'-H and 11'-H), 3.48-3.70 (complex signal, 4 H, 3'-H₂ and 5'-H₂); ¹³C-NMR (100.5 MHz, CDCl₃) δ: 15.0 (CH₂, C12' or C13'), 15.3 (CH₂, C13' or C12'), 20.8 [CH₂, C9'(10')], 25.82 (CH₂), 25.88 (CH₂) and 25.92 (CH₂) (C3, C4 and C5), 28.88 (CH₂, C2 or C6), 28.95 (CH₂, C6 or C2), 31.2 (CH, C1' or C7'), 31.5 (CH, C7' or C1'), 36.37 (CH, C8' or C11'), 36.44 (CH, C11' or C8'), 37.4 (CH, C2' or C6'), 39.5 (CH, C6' or C2'), 42.8 (CH, C1), 48.8 (CH₂, C3' or C5'), 49.7 (CH₂, C5' or C3'), 174.4 (C, CO).

Anal. Calcd for C₁₉H₂₉NO: C 79.39, H 10.17, N 4.87. Found: C 79.20, H 10.30, N 4.72.

 $6.1.22.\ (12-Azatricyclo[4.4.3.0^{1.6}]trideca-3, 8-dien-12-yl)(cyclohexyl) methanone,\ (\textbf{26}).$

From 12-azatricyclo[4.4.3.0^{1,6}]trideca-3,8-diene hydrochloride [41] (150 mg, 0.71 mmol), cyclohexanecarboxylic acid (82 mg, 0.64 mmol), HOBt (131 mg, 0.97 mmol), EDC (150 mg, 0.97 mmol) and triethylamine (0.4 mL, 2.82 mmol) in EtOAc (6 mL) and following general procedure C, amide 26 (185 mg, 92% yield) was obtained as a yellowish solid. The analytical sample was obtained by crystallization from hot EtOAc (106 mg), mp 116–117 °C; IR (ATR) v: 605, 676, 739, 813, 865, 1004, 1187, 1207, 1224, 1332, 1347, 1427, 1629, 2858, 2918 cm⁻¹; ¹H-NMR (400 MHz, CDCl₃) δ: 1.16-1.31 (complex signal, 3 H, 3(5)-H_{ax} and 4-H_{ax}), 1.51 [m, 2 H, 2(6)-H_{ax}], 1.62-1.84 [complex signal, 5 H, 2(6)-H_{eq}, 3(5)-H_{eq} and 4-H_{eq}], 1.92-2.10 [complex signal, 8 H, $2'(10')-H_2$ and $5'(7')-H_2$, 2.29 [tt, J = 11.6 Hz, J' = 3.6 Hz, 1 H, 1-H], 3.35-3.37 (complex signal, 4 H, 3'-H_a, 3'-H_b, 5'-H_a and 5'-H_b), 5.48-5.57 [complex signal, 4 H, 3'(9')-H and 4'(8')-H]; 13 C-NMR (100.5 MHz, CDCl₃) δ : 25.8 (CH₂, C4), 25.9 [CH₂, C3(5)], 28.9 [CH₂, C2(6)], 32.1 [CH₂, C2'(10') or C5'(7')], 32.2 [CH₂, C5'(7') or C2'(10')], 37.3 (C, C1' or C6'), 39.2 (C, C6' or C1'), 42.6 (CH, C1), 55.3 (CH₂, C11' or C13'), 56.1 (CH₂, C13' or C11'), 123.3 (CH, C3' and C9'), 123.9 (CH, C4' and C8'), 176.3 (C, CO); HRMS-ESI+ m/z $[M+H]^+$ calcd for $[C_{19}H_{27}NO+H]^+$: 286.2165, found: 286.2176.

6.1.23. (12-Azatricyclo[4.4.3.0^{1,6}]tridec-12-yl)(cyclohexyl)methanone, (27).

From **26** (362 mg) and Pd/C (72 mg) and following the general procedure D (24 h), amide **27** (330 mg, 90% yield) was obtained as a white solid. The analytical sample was obtained by crystallization from hot EtOAc (178 mg), mp 160-161 °C; IR (ATR) v: 670, 764, 874, 891, 976, 1157, 1290, 1343, 1360, 1435, 1623, 1635, 2850, 2907, 2921 cm⁻¹; ¹H-NMR (400 MHz, CDCl₃) δ : 1.18-1.29 [complex signal, 3 H, 3(5)-H_{ax} and 4-H_{ax}], 1.31-1.62 [complex signal, 18 H, 2`(6)-H_{ax}, 3'(9')-H₂, 4'(8')-H₂, 5'(7')-H₂, 10'(13')-H₂], 1.64-1.84 [complex signal, 5 H, 2(6)-H_{eq}, 3(5)-H_{eq} and 4-H_{eq}], 2.30 [tt, *J* = 11.6 Hz, *J*' = 3.6 Hz, 1 H, 1-H], 2.70-3.90 (complex signal, 4 H, 3'-H₂ and 5'-H₂); ¹³C-NMR (100.5 MHz, CDCl₃) δ : 21.6 [CH₂, C3'(9') or C4'(8')], 21.8 [CH₂, C4'(8') or C3'(9')], 25.8 (CH₂, C4), 25.9 [CH₂, C3(5)], 28.9 [CH₂, C2(6)], 39.8 (C, C1' and C6'), 41.7 [CH₂, C2'(10') and C5'(7')], 42.6 (CH, C1), 55.1 (CH₂, C11' or C13'), 55.8 (CH₂,

C13' or C11'), 176.0 (C, CO). Anal. Calcd for C₁₉H₃₁NO: C 78.84, H 10.80, N 4.84. Found: 78.83, H 10.74, N 4.75.

6.1.24. (3,4,8,9-Tetramethyl-12-azatricyclo[4.4.3.0^{1,6}]trideca-3,8-dien-12-yl) (cyclohexyl)methanone, (**28**).

From 3,4,8,9-tetramethyl-12-azatricyclo[4.4.3.0^{1,6}]trideca-3,8-diene hydrochloride [41] (136 mg, 0.51 mmol), cyclohexanecarboxylic acid (62 mg, 0.48 mmol), HOBt (111 mg, 0.82 mmol), EDC (127 mg, 0.82 mmol) and triethylamine (0.3 mL, 2.38 mmol) in EtOAc (6 mL) and following the general procedure C, amide 28 (138 mg, 79% yield) was obtained as a yellowish solid. Column chromatography (hexane/EtOAc) gave 28 as a white solid (74 mg), mp 162–163 °C; IR (ATR) v: 748, 785, 848, 973, 1118, 1358, 1434, 1441, 1603, 1613, 2861, 2931 cm⁻¹; ¹H-NMR (400 MHz, CDCl₃) δ: 1.40-1.1.54 [complex signal, 3 H, 3(5)-H_{ax} and 4-H_{ax}], 1.56 (s, 12 H, CH₃), 1.62-1.98 [complex signal, 15 H, 3(5)-H_{eq}, 4-H_{eq}, 2-H₂, 6-H₂, 2'(10')-H₂, 5'(7')-H₂], 2.28 (tt, 1 H, J = 11.6Hz, J' = 3.2 Hz, 1 H, 1-H), 3.28-3.34 (complex signal, 4 H, 11'-H₂ and 13'-H₂); ¹³C-NMR (100.5 MHz, CDCl₃) δ: 18.68 [CH₃, H₃C-C3'(9') or H₃C-C4'(8')], 18.71 [CH₃, H₃C-C4'(8') or H₃C-C3'(9')], 25.8 (CH₂, C4), 25.9 [CH₂, C3(5)], 28.9 [CH₂, C2(6)], 38.6 (CH₂, C1' or C6'), 38.7 [CH₂, C2'(10') or C5'(7')], 38.8 [CH₂, C5'(7') or C2'(10')], 40.4 (CH₂, C6' or C1'), 42.6 (CH, C1), 55.5 (CH₂, C11' or C13'), 56.4 (CH₂, C13' or C11'), 121.6 [C, C3'(9') or C4'(8')], 122.2 (C, C4'(8') or C3'(9')], 176.4 (C, CO); HRMS-ESI+ m/z $[M+H]^+$ calcd for $[C_{23}H_{36}NO+H]^+$: 342.2791, found: 342.2793. (7,8,9,10-Tetramethyl-3-azapentacyclo[7.2.1.1^{5,8}.0^{1,5}.0^{7,10}]tridec-3-yl) 6.1.25. (cyclohexyl) methanone, (29).

From 7,8,9,10-tetramethyl-3-azapentacyclo[7.2.1.1^{5,8}.0^{1,5}.0^{7,10}]tridecane hydrochloride [40] (145 mg, 0.54 mmol), cyclohexanecarboxylic acid (67 mg, 0.52 mmol), HOBt (105 mg, 0.78 mmol), EDC (121 mg, 0.78 mmol) and triethylamine (0.3 mL, 2.29 mmol) in EtOAc (6 mL) and following the general procedure C, amide **29** (143 mg, 78% yield) was obtained as a yellowish solid. Column chromatography (hexane/EtOAc) gave **29** as a white solid (101 mg), mp 100–101 °C; IR (ATR) v: 636, 747, 791, 829, 866, 891, 977, 1026, 1083, 1124, 1207, 1243, 1266, 1346, 1382, 1442, 1625, 2857, 2925 cm⁻¹; 1 H-NMR (400 MHz, CDCl₃) δ : 0.84 [dd, J = 10.8 Hz, J' = 4.8 Hz, 4 H, 6'(13')-H₂ or 11'(12')-H₂], 0.93 (d, J = 3.6 Hz, 12 H, 7'(8')-C \underline{H} ₃ and 9'(10')-C \underline{H} ₃], 1.18-1.34 [complex signal, 3 H, 3(5)-H_{ax} and 4-H_{ax}], 1.52 (m, 2 H, 2(6)-H_{ax}), 1.64-1.84 [complex signal, 9 H, 3(5)-H_{eq}, 4-H_{eq}, 2(6)-H_{ax} and 11'(12')-H₂ or 6'(13')-H₂], 2.35 (tt, 1 H, J =

11.6 Hz, J' = 3.2 Hz, 1 H, 1-H), 3.59 (d, J = 15.6 Hz, 4 H, 2'-H₂ and 4'-H₂); ¹³C-NMR (100.5 MHz, CDCl₃) δ : 15.5 [CH₃, H₃C-C7'(8') or H₃C-C9'(10')], 15.6 [CH₃, H₃C-C9'(10')] or H₃C-C7'(8')], 25.8 (CH₂, C4), 25.9 [CH₂, C3(5)], 29.0 [CH₂, C2(6)], 42.4 (CH, C1), 42,9 [CH₂, C6'(13')] or C11'(12')], 43.0 [CH₂, C11'(12')] or C6'(13')], 45.3 [C, C7'(8') or C9'(10')], 45.4 [C, C9'(10') or C7'(8')], 47.0 (C, C1' or C5'), 48.9 (C, C5' or C1'), 51.6 [CH₂, C2' or C4'], 53.1 [CH₂, C4' or C2'], 174.6 (C, CO). Anal. Calcd for C₂₃H₃₅NO: C 80.88, H 10.33, N 4.10. Found: 80.97, H 10.27, N 4.00.

6.2. Molecular modeling

Docking calculations were carried out using Glide [60], with the X-ray structure of human enzyme 4BB6 [61]. The geometry of each ligand was energy minimized and the centroid of the inhibitor cocrystalised in 4BB6 was used to generate the docking cavity by selecting all the residues located within 20 Å from the ligand. Between 70 and 100 poses were generated for each ligand, and the best-scored poses (and the expected arrangement within the binding pocket) were chosen as starting structures for MD simulations.

For each ligand-protein complex two independent 50ns MD simulations were run to check the consistency of the binding mode. To this end, the ligand-protein complex was immersed in an octahedral box of TIP3P [62] water molecules and sodium ions were added to neutralize the system. The force field ff99SBildn [63-64] was used for the protein parameters, and RESP charges at the HF/6-31G (d) together with the gaff [65] force field were used to for the ligand and NADP parameters. All systems were refined using a three-step energy minimization procedure (involving first hydrogen atoms, then water molecules, and finally the whole system) and a six-step equilibration (heating the system from 0 K to 300 K in 6 steps of 20 ps, the first, 50 ps the next four, and 5 ns the last one).

6.3. Human 11β-HSD1 in vitro enzyme inhibition assay

11β-HSD1 activity was determined in mixed sex, human liver microsomes (Celsis Invitro Technologies) by measuring the conversion of 3 H-cortisone to 3 H-cortisol. Percentage inhibition was determined relative to a no inhibitor control. 5 μg of human liver microsomes were pre-incubated at 37°C for 15 min with inhibitor and 1 mM NADPH in a final volume of 90 μL Krebs buffer. 10 μL of 200 nM 3 H-cortisone was then added followed by incubation at 37°C for a further 30 min. The assay was

terminated by rapid freezing on dry ice and 3 H-cortisone to 3 H-cortisol conversion determined in 50 μ L of the defrosted reaction by capturing liberated 3 H-cortisol on anticortisol (HyTest Ltd)-coated scintillation proximity assay beads (protein A-coated YSi, GE Healthcare). A nanomolar 11 β -HSD1 inhibitor, UE2316, was added as a positive control within in each set of assays. IC₅₀ values for UE2316 were within the normal range across each test occasion [66].

6.4. Mouse 11β-HSD1 in vitro enzyme inhibition assay

11β-HSD1 activity was determined in pooled mouse (CD-1) liver microsomes (Celsis In-vitro Technologies) by measuring the conversion of cortisone to cortisol by LC/MS. Percentage inhibition was determined relative to a no inhibitor control. 5 μg of mouse liver microsomes were pre-incubated at 37°C for 15 min with inhibitor and 1 mM NADPH in a final volume of 90 μL Krebs buffer. 10 μL of 2 μM cortisone was then added followed by incubation at 37°C for a further 30 min. The assay was terminated by rapid freezing on dry ice and subsequent extraction with acetonitrile on thawing. Samples dried down under nitrogen at 65°C and solubilised in 100 μl 70:30 H₂O:ACN and removed to a 96-well V-bottomed plate for LC/MS analysis. Separation was carried out on a sunfire 150 x 2.1 mm, 3.5 μM column using a H₂O:ACN gradient profile. Typical retention times were 2.71 min for cortisol and 2.80 min for cortisone. The peak area was calculated and the concentration of each compound determined from the calibration curve.

6.5. Microsomal stability assay

The microsomal stability of each compound was determined using either human or mouse liver microsomes (Celsis In-vitro Technologies). Microsomes were thawed and diluted to a concentration of 2 mg/mL in 50 mM NaPO₄ buffer pH 7.4. Each compound was diluted in 4 mM NADPH (made in the phosphate buffer above) to a concentration of 10 μ M. Two identical incubation plates were prepared to act as a 0 minute and a 30 minute time point assay. 30 μ L of each compound dilution was added in duplicate to the wells of a U-bottom 96-well plate and warmed at 37°C for approximately 5 min. Verapamil, lidocaine and propranolol at 10 μ M concentration were utilised as reference compounds in this experiment. Microsomes were also pre-warmed at 37°C before the addition of 30 μ L to each well of the plate resulting in a final concentration of 1 mg/mL. The reaction was terminated at the appropriate time point (0 or 30 min) by addition of 60 μ L of ice-cold 0.3 M trichloroacetic acid (TCA) per well. The plates

were centrifuged for 10 min at 112 x g and the supernatant fraction transferred to a fresh U-bottom 96-well plate. Plates were sealed and frozen at -20 °C prior to MS analysis. LC-MS/MS was used to quantify the peak area response of each compound before and after incubation with liver microsomes using MS tune settings established and validated for each compound. These peak intensity measurements were used to calculate the % remaining after incubation with microsomes for each hit compound.

6.6. Cellular 11β-HSD1 enzyme inhibition assay

The cellular 11β-HSD1 enzyme inhibition assay was performed using HEK293 cells stably transfected with the human 11β-HSD1 gene. Cells were incubated with substrate (cortisone) and product (cortisol) was determined by LC/MS. Cells were plated at 2 x 10⁴ cells/well in a 96-well poly-D-lysine coated tissue culture microplate (Greiner Bioone) and incubated overnight at 37 °C in 5% CO₂ 95% O₂. Compounds to be tested were solubilized in 100% DMSO at 10 mM and serially diluted in water and 10% DMSO to final concentration of 10 μM in 10% DMSO. 10 μL of each test dilution and 10 μL of 10% DMSO (for low and high control) were dispensed into the well of a new 96-well microplate (Greiner Bio-one). Medium was removed from the cell assay plate and 100 μL of DMEM solution (containing 1% penicillin, 1% streptomycin and 300 nM cortisone) added to each well. Cells were incubated for 2 h at 37 °C in 5% CO₂ 95% O₂. Following incubation, medium was removed from each well into an eppendorf containing 500 µL of ethyl acetate, mixed by vortex and incubated at rt for 5 min. A calibration curve of known concentrations of cortisol in assay medium was also set up and added to 500 µL of ethyl acetate, vortexed and incubated as above. The supernatant of each eppendorf was removed to a 96-deep-well plate and dried down under liquid nitrogen at 65 °C. Each well was solubilised in 100 µL 70:30 H₂O:ACN and removed to a 96-well V-bottomed plate for LC/MS analysis. Separation was carried out on a sunfire 150 x 2.1 mm, 3.5 μM column using a H₂O:ACN gradient profile. Typical retention times were 2.71 min for cortisol and 2.8 min for cortisone. The peak area was calculated and the concentration of each compound determined from the calibration curve.

6.7. Cellular 11β-HSD2 Enzyme Inhibition Assay

For measurement of inhibition of 11β -HSD2, HEK293 cells stably transfected with the full-length gene coding for human 11β -HSD2 were used. The protocol was the same as for the cellular 11β -HSD1 enzyme inhibition assay, only changing the substrate, this time cortisol, and the concentrations of the tested compounds, 10, 1 and $0.1~\mu M$.

6.8. Parallel Artificial Membrane Permeation Assays- Blood-Brain Barrier (PAMPA-BBB)

To evaluate the brain penetration of the different compounds, a parallel artificial membrane permeation assay for blood-brain barrier was used, following the method described by Di [50]. The in vitro permeability (P_e) of fourteen commercial drugs through lipid extract of porcine brain membrane together with the test compounds were determined. Commercial drugs and assayed compounds were tested using a mixture of PBS:EtOH (70:30). Assay validation was made by comparing the experimental permeability with the reported values of the commercial drugs by bibliography and lineal correlation between experimental and reported permeability of the fourteen commercial drugs using the parallel artificial membrane permeation assay was evaluated (y = 1.5366x - 0.9672; $R_2 = 0.9382$). From this equation and taking into account the limits established by Di et al. for BBB permeation [50], we established the ranges of permeability as compounds of high BBB permeation (CNS +): P_e (10^{-6} cm s⁻¹) > 1.00 cm pounds of low BBB permeation (CNS -): P_e (10^{-6} cm s⁻¹) > 1.00 and compounds of uncertain BBB permeation (CNS ±): 1.00 cm s⁻¹) > 1.00 cm s⁻¹) > 1.00 cm s⁻¹) > 1.00 and compounds of uncertain BBB permeation (CNS ±): 1.00 cm s⁻¹) > 1.00 cm s

6.9. Pharmacokinetic study

All the animal experiments were performed according to the protocols approved by the Animal Experimentation Ethical Committee of Universitat Autònoma de Barcelona and by the Animal Experimentation Commission of the Generalitat de Catalunya (Catalan Government). Male CD-1 mice (20-25 g) purchased from Envigo Laboratories were used. Compound 23 was dissolved in cyclodextrin 10% at 3 mg/mL to give a clear solution. After oral administration (21 mg/kg, 10 mL/Kg), blood (0.6 mL) was collected from cava vein using a syringe (23G needle) rinsed with 5% EDTA(K2) at 0, 0.5, 1, 3, 5 and 24 (3 animals/point). Each blood sample was immediately transferred to a tube containing 40 μ L of water with 5% EDTA. Blood samples were centrifuged at 10000 g for 5 minutes and plasma samples were stored at –20 °C until analysis of compound concentration by UPLC-MS/MS. Brains were transcardially perfused with 10 mL of saline, removed, frozen in liquid N₂ and stored at –80 °C until analysis of the compound concentration by UPLC-MS/MS.

6.10. In vivo study

6.10.1. Animals.

SAMP8 mice 12 months old (n=12) were randomized in 2 experimental groups (control, n=4; treated, n=8), and additional group 2 months old (n=4) were planned as a young population. Mice were used with free access to food and water, under standard temperature conditions ($22 \pm 2^{\circ}$ C) and 12-h:12-h light-dark cycles (300 lx/0 lx). Compound 23 was administered dissolved in tap water and PEG400 (2% final concentration) yielding a dose of 21 mpk for 4 weeks. The dose was selected based on the IC₅₀ value of 23 and our previous expertise in *in vivo* studies with other 11 β -HSD1 inhibitors [20-21]. To maintain the correct dose along the treatment period, once a week the weight of the animals and the quantity of water that they drank were measured. Therefore, we adjusted the concentration (mg/mL) of the compound 23 in the drink bottle to achieve the correct dose of compound (mpk) to be administered to mice. Studies were performed in accordance with the institutional guidelines for the care and use of laboratory animals established by the Ethical Committee for Animal Experimentation at the University of Barcelona.

6.10.2. Novel Object Recognition Test (NORT).

The test was conducted in a 90-degree, two-arm, 25-cm-long, 20-cm-high maze. Light intensity in the middle of the field was 30 lux. The objects to be discriminated were plastic figures (object A, 5.25-cm-high, and object B, 4.75-cm-high). First, mice were individually habituated to the apparatus for 10 min per day during 3 days. On day 4, they were submitted to a 10-min acquisition trial (first trial), during which they were placed in the maze in the presence of two identical novel objects (A+A or B+B) placed at the end of each arm. A 10-min retention trial (second trial) occurred 2 h (short term memory) or 24 h (long term memory) later. During this second trial, objects A and B were placed in the maze, and the times that the animal took to explore the new object (tn) and the old object (to) were recorded. A Discrimination index (DI) was defined as (tn-to)/(tn+to). In order to avoid object preference biases, objects A and B were counterbalanced so that one half of the animals in each experimental group were first exposed to object A and then to object B, whereas the other one half first saw object B and then object A was presented. The maze, the surface, and the objects were cleaned with 96° ethanol between the animals' trials so as to eliminate olfactory cues.

6.10.3. Brain isolation and Western blot analysis.

Mice were euthanized 1 day after the last NORT trial was conducted, and brain quickly removed from the skull. Hippocampus were dissected and frozen in powdered dry ice

and maintained at -80 °C for further use. Tissue samples were homogenized in lysis buffer containing phosphatase and protease inhibitors (Cocktail II, Sigma), and cytosol and nuclear fractions were obtained as described elsewhere. Protein concentration was determined by the Bradford method. 20 µg of protein were separated by Sodium dodecyl sulfate-Polyacrylamide gel electrophoresis (SDS-PAGE) (8-15%) and transferred onto Polyvinylidene diflouride (PVDF) membranes (Millipore). The membranes were blocked in 5% non-fat milk in Tris-buffered saline containing 0.1% Tween 20 (TBS-T) for 1 h at rt, followed by overnight incubation at 4°C with primary antibodies against PSD95 (1:1,000, ab18258/Abcam), IDE (1:1,000, ab32216/Abcam) and APP C-Terminal Fragment (1:1,000, C1/6.1/Covance) diluted in TBS-T and 5% bovine serum albumin (BSA). GAPDH (1:2,000, Millipore) was used as a control protein charge. Membranes were then washed and incubated with secondary antibodies for 1 h at rt. Immunoreactive proteins were visualized utilizing an Enhanced chemiluminescence-based detection kit (ECL kit; Millipore) and digital images were acquired employing a ChemiDoc XRS+System (BioRad). Band intensities were quantified by densitometric analysis using Image Lab software (BioRad) and values were normalized to GAPDH.

6.10.4. RNA extraction and gene expression determination.

Total RNA isolation was carried out by means of Trizol reagent following the manufacturer's instructions. RNA content in the samples was measured at 260 nm, and sample purity was determined by the A260/280 ratio in a NanoDrop™ ND-1000 (Thermo Scientific). Samples were also tested in an Agilent 2100B Bioanalyzer (Agilent Technologies) to determine the RNA integrity number. Reverse transcription-Polymerase chain reaction (RT-PCR) was performed as follows: 2 µg of messenger RNA (mRNA) was reverse-transcribed using the High Capacity complementary DNA (cDNA) Reverse Transcription kit (Applied Biosystems). Real-time quantitative PCR (qPCR) was utilized to quantify the mRNA expression of inflammatory genes Interleukin 6 (IL-6) and inducible nitric oxide synthase (iNOS). Normalization of expression levels was performed with Actin for SYBER Green. The primers were as follows: for IL-6, forward 5'-ATCCAGTTGCCTTCTTGGGACTGA-3' and reverse 5'-TAAGCCTCCGACTTGTGAAGTGGT-3', for iNOS, forward 5'-GGCAGCCTGTGAGACCTTTG-3' and reverse 5'- GAAGCGTTTCGGGATCTGAA-

3', for *Actin*, forward 5'-CAACGAGCGGTTCCGAT-3' and reverse 5'-GCCACAGGTTCCATACCCA-3'.

Real-time PCR was performed on the Step One Plus Detection System (Applied Biosystems) employing the SYBR Green PCR Master Mix (Applied Biosystems). Each reaction mixture contained 7.5 μ L of cDNA, whose concentration was 2 μ g/ μ L, 0.75 μ L of each primer (whose concentration was 100 nM), and 7.5 μ L of SYBR Green PCR Master Mix (2X).

Data were analysed utilizing the comparative Cycle threshold (Ct) method ($\Delta\Delta$ Ct), where the actin transcript level was utilized to normalize differences in sample loading and preparation. Each sample (n = 4-8) was analysed in triplicate, and the results represented the n-fold difference of transcript levels among different samples.

6.10.5. Data analysis.

Data are expressed as the mean \pm Standard Error of the Mean (SEM). Data analysis was conducted using GraphPad Prism[®] ver. 6 statistical software. Means were compared with one-way Analysis of Variance (ANOVA) and Tukey post hoc analysis.

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Appendix A. Supplementary data

Supplementary data related to this article can be found at http://

