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# TITLE

Glucocorticoid receptors signaling impairment potentiates amyloid-β oligomers-induced pathology in an acute model of Alzheimer's disease

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### **Running title**

Glucocorticoid receptors in Alzheimer's disease

# **LIST OF ABBREVIATIONS**

Aβ: amyloid-β peptide

ADAM10: A disintegrin and metalloproteinase Domain-containing protein 10 (*a-secretase*)

- APP: amyloid precursor protein
- AR: androgen receptors
- BACE1:  $\beta$ -APP cleaving enzyme type 1 ( $\beta$ -secretase)
- <mark>β-tub</mark>: β-tubulin
- CDK: cyclin-dependent kinases
- **CORT**: corticosterone
- EPM: elevated plus maze
- GFAP: glial fibrillary acidic protein
- GC: glucocorticoids
- GR: glucocorticoid receptors
- GRE: glucocorticoid response element
- GSK-3β: glycogen synthase kinase-3β
- HPA axis: hypothalamic-pituitary-adrenal axis
- HSP: heat shock protein
- Iba1: ionized calcium-binding adapter molecule 1
- Icv: intracerebroventricular
- IDE: insulin-degrading enzyme
- lp: intraperitoneal
- MR: mineralocorticoid receptors
- NFT: neurofibrillary tangles
- $oA\beta_{25-35}$ : oligomers of A $\beta$  fragment [25-35]
- PDK1: 3-phosphoinositide-dependent kinase
- **PFC:** prefrontal cortex
- PR: progesterone receptors
- PS1: presenilin 1 (*α*-secretase)
- PSD95: postsynaptic density protein 95
- ROCK: Rho-associated coiled-coil kinase
- sAPPα: α-secretase-cleaved soluble APP ectodomain
- sGRm: selective GR modulator
- SYN: synaptotagmine
- TACE: tumor necrosis factor-α-converting enzyme
- TAT: Tyrosine Amino Transferase activity assay

# ABSTRACT (200/200 words)

Dysregulation of the hypothalamic-pituitary-adrenal (HPA) axis occurs early in Alzheimer's disease (AD), associated with elevated circulating glucocorticoids (GC) and glucocorticoid receptors (GR) signaling impairment. However, the precise role of GR in the pathophysiology of AD remains unclear.

Using an acute model of AD induced by the intracerebroventricular injection of amyloid- $\beta$  oligomers (oA $\beta$ ), we analyzed cellular and behavioral hallmarks of AD, GR signaling pathways, processing of amyloid precursor protein, and enzymes involved in Tau phosphorylation. We focused on the prefrontal cortex (PFC), particularly rich in GR, early altered in AD and involved in HPA axis control and cognitive functions.

We found that oAß impaired cognitive and emotional behaviors, increased plasma GC levels, synaptic deficits, apoptosis and neuroinflammatory processes. Moreover, oAß potentiated the amyloidogenic pathway and enzymes involved both in Tau hyperphosphorylation and GR activation. Treatment with a selective GR modulator (sGRm) normalized plasma GC levels and all behavioral and biochemical parameters analyzed.

GR seems to occupy a central position in the pathophysiology of AD. Deregulation of the HPA axis and a feed-forward effect on PFC GR sensitivity could participate in the etiology of AD, in perturbing A $\beta$  and Tau homeostasis. These results also reinforce the therapeutic potential of sGRm in AD.

Key-words: Selective GR modulator; GSK-3β; Cdk5; ROCK; PDK1

### INTRODUCTION

Alzheimer's disease (AD), the most common cause of dementia in the elderly, is characterized by a progressive impairment of cognitive functions and the presence of senile plaques and neurofibrillary tangles (NFT) throughout the brain, including areas particularly involved in memory formation and emotional regulation. Plaques are composed of insoluble extracellular aggregates consisting mainly of amyloid- $\beta$  (A $\beta$ ) peptides, while NFT result from hyper- and abnormal phosphorylation of the microtubule-stabilizing protein Tau (1). There are several forms of AD. Familial forms with known mutations of specific genes represent less than 5% of cases, whereas 95% of patients develop sporadic forms, with unknown mechanisms, but with identified risk factors. The principal risk factor for sporadic AD is aging. But, there is also growing evidence that stressful lifetime events may increase the probability of developing AD (2). This view is particularly supported by the fact that in AD patients, cognitive and psychological symptoms are associated with an early deregulation of the hypothalamic-pituitary-adrenal (HPA) axis, as well as elevated levels of glucocorticoids (GC) in plasma and CSF (3,4).

The HPA axis, highly involved in the stress response, triggers the adrenal cortex to release GC. These steroid hormones readily cross the BBB and bind to low affinity glucocorticoid receptors (GR) and high affinity mineralocorticoid receptors (MR) (5). GC are necessary for normal cellular activity and fundamental for many CNS functions, including learning and memory (6). While MR are localized mainly in the hippocampus, GR are more ubiquitous and are particularly found in several structures of the limbic system (prefrontal cortex (PFC), hippocampus and amygdala), which are strongly involved in cognitive and psychological functions, but also are important components of the neural circuitry modulating HPA axis activity (7).

GC act synergistically with excitatory amino acids (like glutamate) in neurotoxicity. Hence, a deregulation of the HPA axis activity or a modification of GR functioning could be extremely toxic, especially in limbic structures (8), and thus could contribute to the cognitive decline and psychological symptoms that occur in AD. In chronic animal models of AD (transgenic mice), stress and GC administration affect the course of the pathology. Chronic stress accelerates the onset of cognitive deficits, triggers amyloid precursor protein (APP) misprocessing, enhances plaque pathology, reduces  $A\beta$  clearance, increases  $A\beta$  levels, stimulates Tau hyperphosphorylation and its neuronal accumulation (9,10). In the same line of evidence, a recent study showed that early life stress in APP/PS1 mice induced elevated corticosterone levels, associated with enhanced

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hippocampal A $\beta_{1-40/42}$  and BACE1 levels (11). In an acute pathomimetic model of AD obtained after a single intracerebroventricular (icv) injection of an oligomeric solution of A $\beta$  (oA $\beta_{25-35}$ ) (12-14), we demonstrated a strong, long-lasting activation of the HPA axis, associated with a modification of GR and MR expression in brain regions involved in the control of GC secretion (hippocampus, amygdala and hypothalamus) (15), supporting its involvement in the etiology of AD (9,16-19). We also observed that an antagonist and selective modulators of the GR could potently counteract the effects of oA $\beta_{25-35}$  injection in the hippocampus, arguing strongly for a therapeutic potential of modulating GR activity (14).

In the present preclinical study, we focused our attention on the PFC for several reasons. (1) It is a cerebral region highly involved in the control of the HPA axis. Indeed, the PFC is involved in both driving the stress-induced activation of the HPA axis, and in mediating negative feedback regulation in times of stress (20). (2) The different subdivisions of the PFC are particularly involved in cognitive and emotional processing (21). (3) In AD, the PFC is rapidly affected (22). (4) GR levels in the PFC are five-fold higher than MR levels (23), suggesting that the PFC could be particularly sensitive to a dysregulation of the HPA axis activity.

Thus, on the basis of our previous findings (12-14) we took advantage of specific properties of a new selective GR modulator (sGRm) CORT113176 to decipher the role of GR in AD. CORT113176 is representative of a series of novel, selective non-steroidal GR ligands (1H-pyrazolo(3,4-g)hexahydro-isoquinoline sulfonamides) developed by *Corcept Therapeutics* (Fig.1B). It exhibits excellent affinity for GR with no measurable affinity for the other nuclear hormone receptors (progesterone, androgen, mineralocorticoid and estrogen) (14,24-26). CORT113176 demonstrated only partial antagonism and also some agonism in reaction to a viral protein in rat hepatocytes (14). Hence, we refer to it as a sGRm, with the implication of the advantages linked to selective receptor modulation that we previously reported in detail (14). In fact, this family of molecules has the potential to more selectively abrogate pathogenic GR-dependent processes in the brain (*as antagonist*), while retaining beneficial aspects of GR signaling (*as agonist*) (14,27-30).

# **MATERIALS AND METHODS**

#### Animals

Adult male Sprague-Dawley rats (*Janvier Lab., Le Genest-Saint-Isle, France*) weighing 260-280g (8 weeks) at the beginning of the experiments, were housed 1 week before experiments in a standard animal facility (12H/12H light/dark cycle with lights on at

07H00; 21  $\pm$  1°C, food and water *ad libitum*). All experiments, including sacrifices, were performed in conscious rats between 09H00 and 14H00, during the diurnal trough of the HPA axis circadian rhythm.

#### **Ethical considerations**

Animal procedures were conducted in strict adherence to the European Union Directive of 2010 (2010/63/EU). The National French Animal Welfare Committee and the local committee at the University of Montpellier approved all protocols (*authorization: CEEA-LR-12160*). All efforts were made to minimize the number of animals used, potential pain, suffering and distress.

#### *Amyloid-*β peptide

In patients, soluble A $\beta$  oligomers contains mainly the sequences A $\beta_{1.40}$  and A $\beta_{1.42}$  (1). However, they also contain peptides with shorter sequences such as A $\beta_{25\cdot35}$  or A $\beta_{25\cdot35/40}$  (31-33), identical between human and rodent (34). It can be produced in AD patients by enzymatic cleavage of A $\beta_{1-40}$  (31,32). This A $\beta$  peptide includes extracellular and transmembrane residues that have been reported to represent a biologically active region of A $\beta$  (35-37) and to contain the highly hydrophobic region forming stable aggregates (36). Interest in this undecapeptide, which itself shows a  $\beta$ -sheet structure (12,36), has grown over the last decade, mainly because it induces neurite atrophy, neuronal cell death, synaptic loss, as well as synaptic plasticity and memory deficits in a similar way to A $\beta_{1-40}$  and A $\beta_{1-42}$  (37), but with better solubility and efficiency (38,39). A $\beta_{25\cdot35}$  and scrambled A $\beta_{25\cdot35}$  peptides (*PolyPeptide, Strasbourg, France*) were dissolved in sterile water (1 µg/µl) and stored at -20°C. Since soluble A $\beta$  oligomers correlate better with the progression of the disease (40), A $\beta_{25\cdot35}$  and scrambled peptides were aggregated by *in vitro* incubation at 37°C (4 days) to obtain a solution mainly composed (more than 95%) of a mixture of soluble oligomer species (oA $\beta_{25\cdot35}$ ), as previously characterized (13).

#### **Experimental procedures**

To evaluate the impact of  $\alpha A\beta_{25-35}$  (*acute model of AD*), animals were divided into three groups. One group had no surgery (control rats), a second received an icv injection of incubated scrambled peptide (10 µg/rat) and a third received an icv injection of  $\alpha A\beta_{25-35}$ (10 µg/rat). The animals were anesthetized with an intraperitoneal (ip) injection of 1 ml of a mixture of Ketamine and Xylazine (80 and 10 mg/kg b.w., respectively).  $\alpha A\beta_{25-35}$  was injected directly into the lateral ventricles using a David-Kopf stereotaxic apparatus (*Phymep, Paris, France*), (coordinates: AP -1 mm, L ±1.5 mm, DV -3.5 mm) (Fig.1A) (41). Based on a previous study (14) and in order to decipher the role of GR in  $\alpha A\beta$  toxicity, treatment with sGRm was conducted 1 week after the icv injection of  $\alpha A\beta_{25-35}$ . CORT113176 (10 mg/kg b.w. per injection) (*Corcept Therapeutics, Menlo Park, CA, USA*) (Fig.1B) was injected ip twice a day (09H00 and 18H00) for 1 week. The short-term memory or anxiety state of different groups of rats were tested (day 14) in a T-maze or in an elevated plus maze test (EPM), respectively. The following day (day 15) and 30 min after the last ip injection, the unanesthetized animals were sacrificed by decapitation. Blood samples and the PFC were rapidly collected for corticosterone assay and WB analysis. Naive rats received no treatment but were manipulated in the same manner as treated rats. Vehicle rats received only ip injections of sesame oil and served as negative controls for pharmacological treatments.

#### Spatial short-term memory

T-maze test was used to rapidly assess the delayed alternation of rats. This memory behavioral test was used as a non-invasive recurrent readout that we usually perform to make sure we have an appropriate toxicity (12-15). The T-maze consisted of two short arms (A and B), extending from a longer alley and enclosed with high walls. The test involved two trials separated by 1 h. During the training session, one short arm (B) was closed. Rats were placed at the end of the long alley, allowed to visit the maze for 10 min and then returned into their home cage. During the test session, animals were placed in the maze for 2 min, with free access to all arms. The number of visits and time spent in each arm were measured. The results were expressed as ratio of the time spent in the initially closed novel arm, over the time spent in the previous arm and as a ratio of the number of entries into the novel arm over the familiar one. The apparatus was cleaned with diluted ethanol (50%) between animals.

#### Anxiety behavior

The anxiety state of rats was measured using their ability to explore open and enclosed arms of an EPM, as previously described (15). The clear plexiglass apparatus consisted of two open arms (50 x 10 cm) and two enclosed arms (50 x 10 x 45 cm high), extending from a central platform and placed 60 cm above the floor. Each rat was placed at the center of the plus-maze facing the closed arm and its exploration behavior was recorded for 10 min. The results were expressed as total time spent in the open arms and the total number of entries was counted to verify general motor activity. An entry into an arm was recorded if the animal crossed the line that connected that arm with the central platform with all four legs. The apparatus was cleaned with diluted ethanol (50%) between animals.

#### Corticosterone assay

Blood samples were collected at the time of sacrifice (day 15), on 1 mg/ml EDTA (*Sigma-Aldrich, Saint Quentin Fallavier, France*), centrifuged at 4°C, and plasma stored at -20°C until assayed for corticosterone (14). Plasma corticosterone concentrations were assayed using a conventional ELISA kit (*Enzo-Life Sciences, Farmingdale, NY, USA*) in a 10-µl plasma sample diluted (1:40) with the assay buffer. The assay sensitivity was 27 pg/ml. The intra- and inter-assay coefficients were 6.6 and 7.8%, respectively.

#### Aβ<sub>1-42</sub> assay

Rats were sacrificed by decapitation 15 days after  $oA\beta_{25-35}$  injection and brains were rapidly removed, PFC dissected out, weighted, frozen in liquid nitrogen and stored at -80°C until assayed. After thawing, PFC were sonicated (*VibraCell; Sonics & Materials, Newtown, CT, USA*) for 20s in a lysis buffer (42). After centrifugation (14000 rpm for 25 min, 4°C), supernatants were used for  $A\beta_{1-42}$  ELISA assay (*Anaspec, Fremont, CA, USA*), according to the manufacturer's instructions. Absorbance was read at 450 nm (*Tecan icontrol, ThermoFisher Scientific, Illkirch , France*) and sample concentration was calculated using the standard curve (SFig.3A). Results were then expressed in pg of  $A\beta_{1-42}/g$  of tissue. The assay sensitivity was 3.91 pg/ml. The intra- and inter-assay coefficients were 4.3 and 6.4%, respectively.

#### WB analysis

WB were performed as previously described (14) in the whole PFC. All antibodies used are detailed in the Table 1. Briefly, after sacrifice, the PFC was micro-dissected, weighed, immediately frozen on liquid nitrogen and stored at -20°C. Tissues were sonicated (*VibraCell; Sonics & Materials, Newtown, CT, USA*) in a lysis buffer (12) and centrifuged (4°C). Supernatants were collected and the protein concentration was measured using a BCA kit (*ThermoFisher Scientific, Illkirch , France*). Sixty µg from each sample were taken for WB analysis. Samples were separated in SDS-polyacrylamide gel (12%) and transferred to a PVDF membrane (*Merck-Millipore, Dachstein, France*). The membrane was incubated overnight (4°C) with the primary antibody, rinsed and then incubated for 2h with the appropriate horseradish peroxidase-conjugated secondary antibody. Peroxidase activity was revealed by using enhanced-chemiluminescence (ECL) reagents (*Luminata-Crescendo, Merck-Millipore*). The intensity of peroxidase activity was quantified using Image-J software (*NIH, Bethesda, MA, USA*). β-tubulin (β-Tub) was used as a loading control for all immunoblotting experiments.

#### Statistical analysis

Data are presented as mean ± SEM and analyzed using two-way ANOVA followed by a Tukey's multiple comparison test (*GraphPad-Prism 5.0*). P< 0.05 was considered significant. The number of animals in each group is indicated within the columns. Before each analysis of variance, the Gaussian distribution was evaluated and validated by a Kolmogorov-Smirnov test (*GraphPad-Prism 5.0*).

### RESULTS

To characterize the impact of  $\alpha\beta_{25-35}$ , we previously tested over time (*after 1 and 2 weeks*) two different doses (5 and 10 µg/rat) on several parameters previously characterized in this acute model of AD (12) (SupFig.1A). While the scrambled peptide induced no modification in comparison with control naive rats, the dose of 10 µg of  $\alpha\beta_{25-35}$  was more efficient than the dose of 5 µg. Indeed, after 10 µg short-term memory deficit (T-maze) was observed up to 2 weeks post-injection (SupFig.1B) and plasma levels of corticosterone were increased from 1 to 2 weeks (SupFig.1C).

We next evaluated (in the PFC) the role of GR in the oA $\beta$  toxicity using a sGRm (CORT113176). Animals were treated one week after the icv injection of oA $\beta_{25-35}$  with CORT113176 according to a protocol and a dose established in a previous study (14). Control scrambled peptide and vehicle treatment induced no changes in any of the readouts relative to untreated animals (Fig.1-6). All blots of control conditions were presented in a supplementary document (SFig.2) to improve the clarity of figures and to highlight key effects. Two weeks after oA $\beta_{25-35}$ , animals presented anxious behavior (Fig.1C), short-term memory deficits (Fig.1D), and high plasma concentrations of corticosterone (Fig.1E). Treatment with CORT113176 reversed all of these parameters (Fig.1).

Behavioral deficits observed two weeks after  $oA\beta_{25-35}$ , were associated with pre-(SYN) and post-synaptic (PSD95) deficits, increased apoptotic marker expression (Fig.2A-D) and a marked neuroinflammation characterized by the activation of astrocytes (GFAP) (Fig.2A,E) and microglial cells (Iba1) (Fig.2A,F). Treatment with the sGRm normalized Caspase 3 expression, the pre- and post-synaptic deficits (Fig.2A,C,D) and blocked the neuroinflammatory processes (Fig.2A,F), as previously reported at the hippocampus level (14) or at the spinal cord level in an experimental model of amyotrophic lateral sclerosis (30).

In order to determine in the PFC the effects of  $oA\beta_{25-35}$  (and of CORT113176) on glucocorticoid receptor signaling, as part of a potential feed-forward or feedback process, animals were treated as previously detailed (Fig.1A). The icv injection of  $oA\beta_{25-35}$ 

increased the expression of MR and GR (Fig.3A-C). These effects were associated with an over-activation of GR, as characterized by an increase of the phosphorylated form of GR (Ser<sup>211</sup>) (Fig.3A,D). There were opposite changes in the expression of HSP90 and HSP70, the two main chaperones involved in the activity of GR (43), but also involved in the control of A $\beta$  and Tau aggregation (44-46). The HSP90/HSP70 ratio (which reflects GR activation) (43) accordingly was substantially increased (Fig 3A, E). Treatment with CORT113176 reversed the increase of GR and MR, decreased the phosphorylation of GR and normalized the HSP90/HSP70 ratio (Fig.3).

The GR phosphorylation status (47) may constitute an important link between AD and GC. Indeed, GR can be phosphorylated on several serine and threonine residues. Thus, we characterized the impact of  $oA\beta_{25-35}$  on GSK-3 $\beta$  and Cdk5, the two main enzymes involved in both the phosphorylation of GR (48), and the hyperphosphorylation of Tau (49,50). We first confirmed changed expression ratios of p(Ser9)GSK-3β/GSK-3β and p(Tyr216)GSK-3β/GSK-3β (51) (Fig.4A-C), reflecting an increase in GSK-3β activation. Second we observed increased levels of Cdk5 (Fig.4A,D), in association with those of p35 and p25, which are involved in Cdk5 activation (52) (Fig.4A,E). We also measured increased levels of Calpain 1, a member of cysteine proteases family regulated by intracellular calcium and showing aberrant activity in AD (53). Calpain 1 is particularly involved in the activation of GSK-3 $\beta$  (54), and in the maturation of p35 in p25 (55) (Fig.4A,F). Lastly, we observed increases in the levels of Fyn (Fig.4A,G), a Src kinase associated with non-genomic effects of GR (56), involved in the activation of GSK-3β and the phosphorylation of Tau (49,57). This family of enzymes is bound to the inactive form of GR as chaperone, and released when GC bind to their receptors (56). Under physiological conditions, GC inhibit Fyn activity and phosphorylation (58), but under chronic stress, with high levels of GC, Fyn is upregulated especially in the hippocampus of adult rats (59). To summarize, the icv injection of  $oA\beta_{25-35}$  activated both GSK-3 $\beta$  and Cdk5 pathways. This activation was associated with an increase of p25, Calpain 1 and Fyn levels in the PFC (Fig.4). Treatment with CORT113176 inhibited the activation of GSK-3β and Cdk5, the maturation of p35 into p25 and the increase of Calpain 1 and Fyn (Fig.4).

In the next part of this study, we characterized the different pathways of APP maturation, through the assessment of the different cellular elements involved in the mutually exclusive processing pathways of APP, the amyloidogenic and non-amyloidogenic pathways (Figs 5 & 6). APP processing and induction of the amyloidogenic pathway (Fig.5) were evaluated by measuring PFC levels of full-length APP (*precursor of* 

amyloid proteins) (Fig.5A,B), C99 (precursor of Aß peptides) (Fig.5A,C), BACE1 (β-APP cleaving enzyme) (Fig.5A,D), PS1 (presenilin-1, a subunit of the y-secretase) (Fig.5A,E) and IDE (insulin degrading enzyme, involved in the clearance of A $\beta$ ) (Fig.5A,F). Two weeks after the injection of  $oA\beta_{25-35}$ , APP levels were increased and amyloidogenic processing was enhanced. This activation was associated with an increased formation (BACE1 and PS1), and a decreased clearance (IDE). One week of treatment with CORT113176 inhibited the activation of the amyloidogenic pathway. The increase in levels of APP, C99, PS1 and BACE1, but also the IDE down-regulation were fully reversed (Fig.5). To characterize the non-amyloidogenic pathway in the PFC, we evaluated by western blot the levels of the a-secretase-cleaved soluble APP ectodomain (sAPPa) and ADAM10 (A disintegrin and metalloproteinase Domain-containing protein 10, a component of  $\alpha$ -secretase) (Fig.6A-C). Two weeks after the injection of  $\alpha\beta_{25-35}$ , sAPP $\alpha$  and ADAM10 were decreased. This inhibition of the non-amyloidogenic processing of APP was totally reversed by one week of treatment with CORT113176 (Fig.6A-C). Finally, in order to confirm the induction of the amyloidogenic pathway, we assayed the endogenous levels of  $A\beta_{1-42}$  in the PFC (SFig.3). As expected, two weeks after the icv injection of  $oA\beta_{25-35}$ ,  $A\beta_{1-42}$ levels were increased by 18% in the PFC. This upregulation was totally reversed by the treatment with CORT113176 (SFig.3B). Even if the levels of  $A\beta_{1-42}$  assayed in the PFC were relatively low, they are consistent with some previous studies in rats (60,61).

To understand by which mechanisms non-amyloidogenic pathways could be inhibited, we evaluated the involvement of the Rho-kinases system. These Ser/Thr kinases are involved in cell motility, cell proliferation, autophagy and apoptosis (62-64). They have been suggested as potential therapeutic targets for neurodegenerative diseases, including AD (66-68). This effect seems to be mediated by Rho-associated coiled-coil kinases (ROCK)-induced overactivation of the 3-phosphoinositide-dependent kinase 1 (PDK1) activity (65,68). In fact, exacerbated ROCK activity seems to increase the pool of PDK1 molecules physically interacting with and phosphorylated by ROCK (65,68). The overactivation of this system inhibits the non-amyloidogenic pathway (65,68) and affects Tau phosphorylation (69,70). They seem particularly involved in the inhibition of sAPP $\alpha$ synthesis (68) and in the phosphorylation of Tau (69,71). Their activity is modulated by GC (72) via GR activation (73). To characterize the Rho-kinase system, we measured the PFC levels of Rho-associated coiled-coil kinases (ROCK1 & ROCK2) (Fig.6A,D,E) and the 3phosphoinositide-dependent kinase (PDK1) (Fig.6A,F). Here, two weeks after injection of  $oA\beta_{25-35}$ , we observed in the PFC an increase of ROCK1, ROCK2 and PDK1, which was normalized by one week of treatment with the sGRm (Fig.6A, D-F).

#### DISCUSSION

In previous studies, we provided evidence for a vicious cycle between AD and the HPA axis. We showed that the pathology, and especially the amyloid toxicity, rapidly increases GC secretion that in turn, modulates APP processing (14,15). This dysregulation seems to be under the control of GR, since treatment with a new class of selective GR ligands blocks the installation of this cycle in the hippocampus and re-establishes all parameters analyzed and disturbed by the amyloid toxicity (*memory and synaptic deficits, neuroinflammation, apoptosis, APP processing and high levels of GC*) (14). This recent study (14), allowed us to design and validate the experimental protocol of treatment with sGRm. Briefly, we determined treatment timing and duration, doses and specificities of the two sGRm (*CORT108297 & CORT113167*) tested, in comparison to the non-selective antagonist of reference, Mifepristone. It appeared that one week of treatment (*two ip injection per day*) with CORT113176 (*10 mg/kg per injection*) displayed the most effective therapeutic potential against toxicity induced by oA $\beta_{25-35}$ . Owing to its efficacy and selectivity, this sGRm was selected in the present mechanistic study.

Here, on the basis of our precedent finding (14), we aimed to decipher the role of GR in AD and to characterize associated underlying mechanisms. For this purpose, we evaluated the impact of  $oA\beta_{25-35}$  on several intracellular pathways involved in the activation of GR, but also in the pathophysiology of AD. We show the establishment of several intracellular vicious cycles involving GC and GR, providing mechanistic insight to a central role of these receptors in the etiology of AD (Fig.7). The notion of a vicious cycle between GC signaling and pathogenesis is reinforced by the fact that modulation of GR activity with CORT113176 normalized all changes induced by the amyloid toxicity.

In the first part of our study, in order to validate our protocols in comparison with the previous study (14), we confirmed that one week of treatment with a sGRm (CORT113176) at the dose of 10 mg/kg is sufficient to reverse short-term memory deficits and to reestablish plasma concentrations of corticosterone disrupted by the icv injection of  $oA\beta_{25-35}$ . In addition, and for the first time, we show that treatment with sGRm is able to reverse anxious behavior induced by the amyloid toxicity (15). We equally demonstrate that sGRm treatment is able to reverse synaptic deficits, neuroinflammation and apoptosis processes induced by  $oA\beta_{25-35}$  in the PFC, as previously observed in the hippocampus (14).

Then, we showed that the icv injection of  $oA\beta_{25-35}$  increased GR phosphorylation on Ser<sup>211</sup>, a site involved in the activation of GR in rats (48). This activation was associated

with an increase and activation of GSK-3 $\beta$  and Cdk5, with a substantial increase of the HSP90/HSP70 ratio (*important for GR activity* (43)), but also, importantly, in the control of A $\beta$  and Tau aggregation (44-46). The activation of GSK-3 $\beta$  and Cdk5 is under the control of several enzymes (49,50,54,55,57), including Fyn and Calpain 1 that are regulated by GC (58,59), induced by the amyloid toxicity, and increased in AD patients (74). Our data show that the over-activation of GR induced by oA $\beta_{25-35}$  coincided with an increase of these two enzymes (Fig.7).

We also observed that  $\alpha A\beta_{25\cdot35}$  induced a strong increase of Calpain 1. This augmentation could be the reflection of its proteolytic activity, given that it was concomitant with an activation of two of its substrates, GSK-3 $\beta$  and an increase of p25/p35 ratio. These effects were reversed by treatment with the sGRm (CORT113176), providing evidence of an intracellular loop by which pathology increases the activation of GR (*directly via GSK-3\beta and Cdk5 and indirectly via Fyn and Calpain 1*), which in turn increases the activation of these key enzymes and the HSP90/HSP70 ratio, worsening the AD pathogenesis (44-46) (Fig.7).

Our current data show that the seeding and the accumulation of endogenous Aß induced by the icv injection of  $oA\beta_{25-35}$  (43) (SFig.3B), not only results from the activation of amyloid-pathways, but also from the inhibition of non-amyloid-pathways. Indeed, the icv injection of  $oA\beta_{25-35}$  provoked the activation of the amyloidogenic pathway in the PFC, through an increase of A $\beta$  synthesis (APP/C99,  $A\beta_{1-42}$ , BACE1 and PS1 up-regulation) and a decrease of Aβ clearance (*IDE down-regulation*), as previously reported for the hippocampus (14). Besides, Aβ oligomers also inhibit the non-amyloidogenic pathway (sAPPa and ADAM 10 down-regulation) and that this effect could be controlled by the activation of ROCK1 and ROCK2 (Fig.7). We showed that the  $oA\beta_{25-35}$  injection-induced inhibition of sAPPa and ADAM10 is associated with an activation of ROCK/PDK1 pathways. These results are consistent with several studies showing that ROCKs modulate the shedding of sAPPa through an inhibition of tumor necrosis factor-aconverting enzyme (TACE or ADAM) activity, and that ROCKs depletion reduces A<sup>β</sup> levels (68,75,76). ROCK activity seems to be directly up-regulated by GC (72,73,77), and ROCK/PDK1 activation after  $oA\beta_{25-35}$  is reversed by treatment with CORT113176, again constituting a vicious cycle based on feedforward effects on GR signaling (Fig.7).

ROCK also affect Tau hyperphosphorylation (69,70). They activate the two principal enzymes involved in Tau phosphorylation, Cdk5 and GSK-3 $\beta$  (71,78). In addition, ROCK/PDK1 inhibition of sAPP $\alpha$  may affect Tau, since it was shown that sAPP $\alpha$  reduces

GSK-3β-mediated Tau phosphorylation (79). These two mechanisms again link GR and GC to the pathophysiology of AD (Fig.7). Likewise, Tau phosphorylation may also be directly impacted by the other GR-related enzymes we described. Indeed, Fyn can directly phosphorylate Tau on tyrosine residues (80). Fyn can also directly activate GSK-3β to rapidly induce Tau phosphorylation in human neuroblastoma cells (57). In the same line of evidence, it appears that Fyn controls the activity of PDK1 through an up-regulation of ROCK (81,82), and thus could also participate in the inhibition of the non-amyloidogenic pathway. Calpain 1 also activates GSK-3β and Cdk5, promoting Tau phosphorylation and Tau-associated neurodegeneration (49,50). Calpain 1 can cleave the neuron-specific Cdk5 activator p35 to produces p25, which accumulates in the brains of AD patients (52). In fact, it was shown that induction of p25 by Calpain 1 causes prolonged activation and mislocalization of Cdk5 and that the p25/Cdk5 kinase hyperphosphorylates Tau, disrupts the cytoskeleton and promotes the apoptotic death of primary cortical neurons (55). Thus Fyn and Calpain 1 upregulations may be involved in both the increased Tau phosphorylation after  $oA\beta_{25-35}$  injection (13), and in the activity of GSK-3 $\beta$ , Cdk5, and the processing of APP. This highlights the potential link that GR activation could play between A $\beta$  and Tau (Fig.7).

The mechanisms by which GR affects the multiple enzymes identified here is unclear. GR are nuclear receptors that directly interact with specific genes *via* binding to glucocorticoid response element (GRE), interactions with other transcription factors, or *via* non-genomic mechanisms, such as epigenetic modifications (5, 82). A GRE has been described in the promoter regions of APP and BACE1 (84-86). For the other proteins no mechanism of transcriptional regulation *via* GR has been identified. A recent study on hippocampal slices pre-treated with an inhibitor of transcription activity, showed a non-genomic activation of GSK-3 $\beta$  by GC (87). The efficacy of the sGRm suggests that it is able to antagonize also non-genomic GR signaling. Of note, involvement of membrane-localized GR was linked to AD not only for the regulation of GSK-3 $\beta$  (88), but also – surprisingly - in the regulation of BACE1 (89). For the other proteins, further investigations will be needed to decipher which mechanisms are involved in GR regulatory activity. There is also the potential for more indirect mechanisms, since membrane-localized GR can facilitate glutamatergic transmission (90-92), affect some of the factors that we identified here *via* stimulation of excitotoxicity pathways (8).

This new study provides new arguments supporting the development of a vicious cycle based on GR activation in AD, here based on analysis of changes in the PFC. In this

preclinical study, the new sGRm (CORT113176) blocked this cycle and normalized all AD processes analyzed, including extracellular (ADAM10, BACE1 and IDE), intramembrane (PS1) and intracellular enzymes (Cdk5/p25, GSK-3 $\beta$ , Fyn, Calpain 1, ROCKs and PDK1). In addition, we cannot exclude, as previously mentioned in the hippocampus (14), the involvement of MR in the effects observed after sGRm treatment. Indeed, several studies suggested a neuroprotective role of these receptors in a context of GR blockade (93-95). Thus, further investigations are needed to decipher the precise role of MR in the pathophysiology of AD. We moreover demonstrated that the accumulation of endogenous A $\beta$ , induced by the amyloid toxicity and the concomitant dysregulation of HPA axis, resulted from the activation of amyloidogenic and the inhibition of the non-amyloidogenic pathways (Fig.7). All of these data place HPA axis dysregulation and GR in a central and crucial position in the pathophysiology of AD, linking amyloid toxicity and Tau deregulation. This work also highlights the therapeutic potential of sGRm to counteract negative effects induced by the amyloid toxicity and to reestablish the functionality of GR and *a fortiori* to reestablish the primal role of GC in the maintenance of homeostasis.

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Conflict of interest statement: JKB and HH are employees of Corcept Therapeutics (*Menlo Park, CA, USA*), which develops GR ligands for clinical use. Corcept Therapeutics provided compounds and financed part of the experiments. However, they are not involved in the experimental design, results analysis and conclusions of the present study. All other authors declare that they have no competing interests.

# **Author's contributions**

GC, FP, CH and CZ performed experiments. HH and JKB provided glucocorticoid receptor ligand tested in this study. GC, FP, CZ, CH, CD, NC, VP, JT, HH, OM and JKB corrected the manuscript. LG designed the study, wrote the protocol (with the help of GC), performed part of experiments, analyzed the data (with the help of CD, NC, and OM) and wrote the manuscript with GC. All authors read and approved the final manuscript. All funders had no role in data collection, analysis or in the writing of the manuscript.

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# LEGENDS

#### **Figure 1**

Effects of selective GR modulator on, anxious behavior, spatial short-term memory deficits and high levels of corticosterone induced by the icv injection of  $oA\beta_{25-35}$ . Panel A: Experimental protocol - At T0, adult male rats (Sprague Dawley) were injected directly into the lateral ventricles using a stereotaxic apparatus (coordinates: AP -1 mm, L ±1.5 mm, DV -3.5 mm). One group has not undergone surgery (control rats, white column), a second group received an icv injection of scrambled A $\beta_{25-35}$  peptide (negative control - 10  $\mu$ g/rat orange column) and a third group received an icv injection of  $oA\beta_{25-35}$  peptide (A $\beta$  - 10 µg/rat - red column). Animals were weighted daily and one week after the icv injection (at Week 1), animals were treated with vehicle or CORT113176 (10 mg/kg per injection) through two intraperitoneal (ip) injections per day during 7 days. At day 14 (Week 2), the anxious behavior or the spatial short-term memory of each rat were tested in an elevated plus maze (EPM) or in a T-maze, respectively. The following day (day 15), 30 min after the last ip injection, the animals were sacrificed, blood samples and PFC were rapidly collected for corticosterone assay and Western blot analysis, respectively. Panel B: Chemical structure of the selective GR modulator tested in this study, CORT113176 (reproduction with the permission of Corcept Therapeutics). Panel C: Anxious behavior was determined in the EPM paradigm. Each animal was placed at the center of the EPM and allowed to freely explore for 10 min. Data were expressed as time spent in the open arms (OA) in % of total time. Two-way ANOVA:  $F_{2.59}$ = 7.90 for group, p< 0.001,  $F_{2.59}$ = 1.54 for treatment, ns; and  $F_{4,59}$ = 2.95 for interaction, p< 0.05. Panel D: Spatial short-term memory performance was determined in a T-maze test and was expressed as the ratio of the time spent in the initially closed arm (B) over the time spent in the previous arm (A). Two-way ANOVA:  $F_{2.48}$ = 9.10 for group, p< 0.001;  $F_{2.48}$ = 8.51 for treatment, p< 0.001; and F<sub>4.48</sub>= 2.36 for interaction, ns. **Panel E**: Plasma concentrations of corticosterone (CORT) were determined by Elisa and expressed as ng/ml. Two-way ANOVA: F<sub>2.45</sub>= 39.0 for group, p< 0.0001;  $F_{2.45}$ = 15.3 for treatment, p< 0.0001; and  $F_{4,45}$ = 10.8 for interaction, p< 0.0001. \* p< 0.05 and \*\* p< 0.01 vs. respective control (C) group. + p< 0.05 and ++ p< 0.01 vs. respective scrambled (S) group. x p< 0.05 and xx p< 0.01 vs. respective naive rat in each group (C, S or oAβ). The number of animals in each group is indicated within the columns.

#### Figure 2

The effects in the PFC of selective GR modulator on the different cellular markers (**Panel A**) modified by the icv injection of  $oA\beta_{25-35}$ , were evaluated by Western blot. Variations of apoptosis (caspase-3, 19 kDa) (Panels A,B), post-synaptic marker (PSD95, 95 kDa) (Panels A,C), pre-synaptic marker synaptotagmin (SYN, 65 kDa) (Panels A,D), neuroinflammatory markers GFAP (55 kDa, astrocyte cells) (Panels A,E) and Iba1 (17 kDa, microglial cells) (Panels A,F) were evaluated in control (C - white column) and in icv injected rats with 10  $\mu$ g/rat of scrambled peptide (S - orange column) or oA $\beta_{25-35}$  (A $\beta$  - red column), treated or not with vehicle (sesame oil) or selective GR modulator, CORT113176 (10 mg/kg per ip injection). For experimental protocol see Fig.1A. The variations of Caspase-3, PSD95, SYN, Iba1 and GFAP in the PFC were normalized with the variations of β-tubulin (β-tub, 50 kDa) and compared with non-injected rats (control group: C). Twoway ANOVA: **Caspase-3:**  $F_{2.59}$ = 6.75 for group, p< 0.01;  $F_{2.59}$ = 3.08 for treatment, p< 0.05; and  $F_{4.59}$ = 1.73 for interaction, ns; **PSD95**:  $F_{2.67}$ = 4.08 for group, p< 0.05;  $F_{2.67}$ = 0.50 for treatment, ns; and  $F_{4.67}$ = 3.15 for interaction, p< 0.05; **SYN**:  $F_{2.66}$ = 5.97 for group, p< 0.01;  $F_{2,66}$ = 2.36 for treatment, ns; and  $F_{4,66}$ = 3.87 for interaction, p< 0.05; **GFAP**:  $F_{2,66}$ = 31.04 for group, p< 0.0001;  $F_{2.66}$ = 8.52 for treatment, p< 0.001; and  $F_{4.66}$ = 3.49 for interaction, p< 0.05; **Iba1**:  $F_{2.59}$  = 10.1 for group, p< 0.001;  $F_{2.59}$  = 4.28 for treatment, p< 0.05; and  $F_{4.59}$  = 2.08 for interaction, ns.

The variations are expressed as means  $\pm$  SEM in % of control values. \* p< 0.05 and \*\* p< 0.01 *vs.* respective control (C) group. + p< 0.05 and ++ p< 0.01 *vs.* respective scrambled (S) group. x p< 0.05 and xx p< 0.01 *vs.* respective naive rat in each group (C, S or A $\beta$ ).

#### Figure 3

The effects in the PFC of CORT113176 on the activation of GR induced by the icv injection of oA $\beta_{25-35}$ , were evaluated by Western blot. Variations of the expression of MR (100 kDa) (Panels A,B) and GR (95 kDa) (Panels A,C), the phosphorylation of GR (p[Ser211]GR, 95 kDA) (Panels A,D) and the expression ratio of HSP90/HSP70 (90 kDa / 70 kDa) (Panels A,E) were evaluated in control (C - white column) and in icv injected rats with 10 µg/rat of scrambled peptide (S - orange column) or oA $\beta_{25-35}$  (A $\beta$  - red column), treated or not with vehicle (sesame oil) or the selective GR modulator CORT113176 (10 mg/kg per ip injection). For experimental protocol see Fig.1A. The variations of all proteins in the PFC were normalized with the variations of  $\beta$ -tubulin ( $\beta$ -tub, 50 kDa) and compared with non-injected rats (control group: C). Two-way ANOVA: **MR**: F<sub>2,54</sub>= 11.0 for group, p< 0.0001; F<sub>2,54</sub>= 2.96 for treatment, p< 0.05; and F<sub>4,54</sub>= 3.06 for interaction, p< 0.05; **GR**: F<sub>2,51</sub>= 22.1 for group, p< 0.0001; F<sub>2,51</sub>= 8.45 for treatment, p< 0.001; and F<sub>4,51</sub>= 5.97 for

interaction, p< 0.001; **pGR**:  $F_{2,52}$ = 2.99 for group, p< 0.05;  $F_{2,52}$ = 2.56 for treatment, ns; and  $F_{4,52}$ = 4.05 for interaction, p< 0.01; **HSP90/HSP70**:  $F_{2,50}$ = 29.9 for group, p< 0.0001;  $F_{2,50}$ = 6.00 for treatment, p< 0.01; and  $F_{4,50}$ = 6.55 for interaction, p< 0.001.

The variations are expressed as means  $\pm$  SEM in % of control values. \* p< 0.05 and \*\* p< 0.01 *vs.* respective control (C) group. + p< 0.05 and ++ p< 0.01 *vs.* respective scrambled (S) group. x p< 0.05 and xx p< 0.01 *vs.* respective naive rat in each group (C, S or A $\beta$ ).

#### **Figure 4**

The effects in the PFC of CORT113176 on the activation of GSK-3ß and Cdk5 pathways induced by the icv injection of  $oA\beta_{25-35}$ , were evaluated by Western blot. The activation of GSK-3ß (ratio of p[Tyr216]GSK-3ß/GSK-3ß total & ratio of p[Ser9]GSK-3β/GSK-3β total, 46kDa each) (Panels A-C) and Cdk5 (30 kDa) (Panels A,D) pathways, the expression ratio of p25/p35 (25 & 35 kDa) (Panels A,E), Calpain 1 (80 kDa) (Panels A,F) and FYN (59 kDa) (Panels A,G) were evaluated in control (C - white column) and in icv injected rats with 10  $\mu$ g/rat of scrambled peptide (S - orange column) or oA $\beta_{25-35}$  (A $\beta$  red column), treated or not with vehicle (sesame oil) or the selective GR modulator CORT113176 (10 mg/kg per ip injection). For experimental protocol see Fig.1A. The variations of all proteins in the PFC were normalized with the variations of  $\beta$ -tubulin ( $\beta$ -tub, 50 kDa) and compared with non-injected rats (control group: C). Two-way ANOVA: p[Tyr216]**GSK-3**β/**GSK-3**β: F<sub>2,51</sub>= 17.0 for group, p< 0.0001; F<sub>2,51</sub>= 6.07 for treatment, p< 0.01; and  $F_{4.51}$ = 2.49 for interaction, p< 0.05; p[Ser9]**GSK-3** $\beta$ /**GSK-3** $\beta$ :  $F_{2.51}$ = 11.9 for group, p< 0.0001;  $F_{2.51}$ = 6.99 for treatment, p< 0.01; and  $F_{4.51}$ = 2.81 for interaction, p< 0.05; Cdk5:  $F_{2,48}$ = 12.1 for group, p< 0.0001;  $F_{2,48}$ = 0.87 for treatment, ns; and  $F_{4,48}$ = 3.73 for interaction, p< 0.01; **p25/p35**:  $F_{2,47}$ = 13. 2 for group, p< 0.0001;  $F_{2,47}$ = 12.1 for treatment, p< 0.0001; and  $F_{4,47}$ = 16.5 for interaction, p< 0.0001; **Calpain 1**:  $F_{2,49}$ = 15.5 for group, p< 0.0001;  $F_{2,49}$ = 5.54 for treatment, p< 0.01; and  $F_{4,49}$ = 4.10 for interaction, p< 0.01; **Fyn**:  $F_{2.50}$ = 16.1 for group, p< 0.0001;  $F_{2.50}$ = 2.82 for treatment, p< 0.05; and  $F_{4.50}$ = 4.70 for interaction, p < 0.01.

The variations are expressed as means  $\pm$  SEM in % of control values. \* p< 0.05 and \*\* p< 0.01 *vs.* respective control (C) group. + p< 0.05 and ++ p< 0.01 *vs.* respective scrambled (S) group. x p< 0.05 and xx p< 0.01 *vs.* respective naive rat in each group (C, S or A $\beta$ ). The number of animals in each group is indicated within the columns.

#### Figure 5

The effects in the PFC of selective GR modulator on the APP processing and the induction of the amyloidogenic pathway (Panel A) induced by the icv injection of  $oA\beta_{25-35}$ , were evaluated by Western blot. Variations of full-length APP (precursor of amyloid peptides, 125 kDa) (Panels A,B), C99 (precursor of amyloid-β peptides, 13 kDa) (Panels A,C), β-APP cleaving enzyme (BACE1, 70 kDa) (Panels A,D), Presenilin 1 (PS1, 20 kDa) (Panels A,E) and Insulin Degrading Enzyme (IDE, 110 kDa) (Panels A,F) were evaluated in control (C - white column) and in icv injected rats with 10 µg/rat of scrambled peptide (S - orange column) or  $oA\beta_{25-35}$  (A $\beta$  - red column), treated or not with vehicle (sesame oil) or selective GR modulator, CORT113176 (10 mg/kg per ip injection). For experimental protocol see Fig.1A. The variations of APP, C99, BACE1, PS1 and IDE in the PFC were normalized with the variations of β-tubulin (β-tub, 50 kDa) and compared with non-injected rats (control group: C). Two-way ANOVA: **APP**:  $F_{2.66}$ = 7.76 for group, p< 0.001;  $F_{2.66}$ = 3.20 for treatment, p< 0.05; and  $F_{4.66}$ = 4.60 for interaction, p< 0.01; C99:  $F_{2.56}$ = 21.1 for group, p< 0.0001;  $F_{2,56}$ = 11.5 for treatment, p< 0.0001; and  $F_{4,56}$ = 7.12 for interaction, p< 0.001; **BACE1**:  $F_{2.58}$ = 5.59 for group, p< 0.01;  $F_{2.58}$ = 3.99 for treatment, p< 0.05; and  $F_{4.58}$ = 2.92 for interaction, p< 0.05; **PS1**:  $F_{2,51}$ = 29.4 for group, p< 0.0001;  $F_{2,51}$ = 1.61 for treatment, ns; and  $F_{4.51}$  = 4.89 for interaction, p< 0.01; **IDE**:  $F_{2.56}$  = 6.47 for group, p< 0.01;  $F_{2.56}$  = 4.06 for treatment, p< 0.05; and  $F_{4.56}$ = 1.99 for interaction, ns.

The variations are expressed as means  $\pm$  SEM in % of control values. \* p< 0.05 and \*\* p< 0.01 *vs.* respective control (C) group. + p< 0.05 and ++ p< 0.01 *vs.* respective scrambled (S) group. x p< 0.05 and xx p< 0.01 *vs.* respective naive rat in each group (C, S or A $\beta$ ).

#### Figure 6

The effects in the PFC of selective GR modulator on the non-amyloidogenic pathways (**Panel A**) modulated by the icv injection of oA $\beta_{25-35}$ , were evaluated by Western blot. Variations of sAPP $\alpha$  (100 kDa) (**Panels A,B**), ADAM10 ( $\alpha$ -secretase, 70 kDa) (**Panels A,C**), Rho-associated coiled-coil kinases (ROCK1 & ROCK2, 160 kDa) (**Panels A,D,E**) and 3-phosphoinositide–dependent kinase (PDK1, 60 kDa) (**Panels A,F**) were evaluated in control (C - white column) and in icv injected rats with 10 µg/rat of scrambled peptide (S - orange column) or oA $\beta_{25-35}$  (A $\beta$  - red column), treated or not with vehicle (sesame oil) or selective GR modulator, CORT113176 (10 mg/kg per ip injection). For experimental protocol see Fig.1A. The variations of sAPP $\alpha$ , ADAM10, ROCK1, ROCK2 and PDK1 in the PFC were normalized with the variations of  $\beta$ -tubulin ( $\beta$ -tub, 50 kDa) and compared with non-injected rats (control group: C). Two-way ANOVA: **sAPP\alpha**: F<sub>2,49</sub>= 8.65 for group, p< 0.001; F<sub>2,49</sub>= 3.92 for treatment, p< 0.05; and F<sub>4,49</sub>= 4.25 for interaction, p<

0.01; **ADAM10**:  $F_{2,48}$ = 6.56 for group, p< 0.01;  $F_{2,48}$ = 7.07 for treatment, p< 0.01; and  $F_{4,48}$ = 7.08 for interaction, p< 0.001; **ROCK1**:  $F_{2,31}$ = 16.1 for group, p< 0.0001;  $F_{2,31}$ = 4.11 for treatment, p< 0.05; and  $F_{4,31}$ = 3.30 for interaction, p< 0.05; **ROCK2**:  $F_{2,51}$ = 10.5 for group, p< 0.001;  $F_{2,51}$ = 7.23 for treatment, p< 0.01; and  $F_{4,51}$ = 3.77 for interaction, p< 0.01; **PDK1**:  $F_{2,50}$ = 15.9 for group, p< 0.0001;  $F_{2,50}$ = 1.55 for treatment, ns; and  $F_{4,50}$ = 4.76 for interaction, p< 0.01.

The variations are expressed as means  $\pm$  SEM in % of control values. \* p< 0.05 and \*\* p< 0.01 *vs.* respective control (C) group. + p< 0.05 and ++ p< 0.01 *vs.* respective scrambled (S) group. x p< 0.05 and xx p< 0.01 *vs.* respective naive rat in each group (C, S or A $\beta$ ).

#### **Figure 7**

Schematic figure recapitulating the central role of GR and the therapeutic potential of selective GR modulators in AD. The icv injection of Aβ oligomers increases APP, C99 PS1 and BACE-1 contents, concomitantly with a decrease of IDE, evidencing the induction of the amyloidogenic pathway and, as previously observed,  $A\beta_{1-42}$  production (42) and Tau hyperphosphorylation (13). In parallel, the icv injection of A $\beta$  oligomers inhibits sAPP $\alpha$  and ADAM10, evidencing the inhibition of non amyloidogenic pathway. Aß oligomers induce, in addition to an excess of circulating GC, an overactivation of GR, which is associated with an increase of the two main chaperones (HSP90 & HSP70) particularly involved in the activity of GR (43), but also in the control of A $\beta$  and Tau aggregation (44-46). Thus, active GR translocate to the nucleus where they exert their genomic effects through GRE, inducing APP and BACE-1 gene transcription, and potentiating A $\beta$  oligomers production. Therefore, it evidences a first intracellular vicious cycle by which pathology increases circulating GC, which in turn increase pathology. Aß oligomers also enhance contents and activity of key enzymes involved directly (Cdk5 and GSK-3β) or indirectly (Calpain 1 and Fyn) in the activation of GR, but also in the hyperphosphorylation of Tau. Enzyme inductions which are regulated by GR, as evidenced after treatment with the selective GR modulator (CORT113176), demonstrate non-genomic effects of GR and thus a second intracellular vicious cycle. Indeed, pathology increases the activation of GR via several keys enzymes (Cdk5, GSK-3β, Calpain 1 and Fyn), which in turn increase the activation of these enzymes involved in the pathophysiology of AD. Finally, it appears that amyloid toxicity inhibits also the non-amyloidogenic pathways, reinforcing the displacement of the equilibrium in favor of endogenous amyloid seeding and evidencing another intracellular loop by which pathology increases the activation of GR, which in turn increases the activation of ROCKs/PDK1 pathways, as evidenced after treatment with the selective GR

modulator (*CORT113176*). This activation accentuates the pathology through the inhibition of  $\alpha$ -secretase (ADAM10) and sAPP $\alpha$  synthesis, as previously reported (79), but also by increasing the phosphorylation of Tau (69,70,71,78).

Red arrow: showed in this study. Purple arrow: showed in our previous studies (13,65). Black arrow: known in the literature (*See discussion for references*). Blue arrow: hypothesis to assess. Red cross: Schematic effects of sGRm. AD: Alzheimer's Disease; ADAM10: A disintegrin and metalloproteinase domain-containing protein 10 ( $\alpha$ -secretase); APP: amyloid precursor protein; A $\beta$ : amyloid- $\beta$  peptide; BACE-1:  $\beta$ -APP cleaving enzyme ( $\beta$ -secretase); Cdk5: cyclin-dependent kinase-5; GC: glucocorticoids; GR: glucocorticoid receptors; GRE: glucocorticoid responsive element; GSK-3 $\beta$ : glycogen synthase kinase 3- $\beta$ ; HSP: heat-shock protein; IDE: insulin degrading enzyme; PDK1: 3-phosphoinositide–dependent kinase; PS1: Presenilin 1 ( $\gamma$ -secretase); ROCKs: Rho-associated coiled-coil kinases; sGRm: selective GR modulator (CORT113176).

# Figure 1









Ε.



D.



# Figure 2













lba1

# Figure 3





# Figure 4



p(Ser9)GSK-3β/GSK-3β

C SoAβ Vehicle





F.

Cdk5



C S οΑβ C-113176

Calpain 1





C.

140

120

100

80

60

40

12

C SoAβ Naive

Prefrontal cortex content (% of control)



G.



Fyn

# Figure 5



C.







F.

IDE

C S οΑβ C-113176





C S oAβ Vehicle

C SoAβ Naive

# Figure 6



Figure 7





# Table 1: Antibodies used in Western blot experiments

| Protein                         | Nol.<br>weight                           | Antibody  | Dilutio<br>n | Ref.      | Supplier  |
|---------------------------------|--|---|--------------|-----------|---|
| Primary antibodies              | Hoight                                   |   |              |           |   |
| ADAM10 <sup>(85)</sup>          | 72 kDa                                   | Rabbit anti-<br>ADAM10                              | 1/1000       | AB19026   | Merck-Millipore, France                           |
| APP/C99 <sup>(14)</sup>         | 125/13<br>kDa                            | Rabbit anti-<br>APP/C99                             | 1/750        | PA1-84165 | Thermo-Fisher<br>Scientific, France               |
| BACE1 <sup>(14)</sup>           | 70 kDa                                   | Rabbit anti-<br>BACE                                | 1/1000       | #5606     | Cell Signaling/Ozyme, ,<br>St Cyr-l'Ecole, France |
| Calpain 1 <sup>(86)</sup>       | 80 kDa                                   | Rabbit anti-<br>calpain 1 large<br>subunit (µ-type) | 1/1000       | #2556     | Cell Signaling/Ozyme,<br>France                   |
| Caspase 3 <sup>(14)</sup>       | 19 kDa                                   | Rabbit anti-<br>caspase 3                           | 1/500        | #9665     | Cell Signaling/Ozyme,<br>France                   |
| Cdk5 <sup>(87)</sup>            | 30 kDa                                   | Rabbit anti-Cdk5                                    | 1/500        | #2506     | Cell Signaling/Ozyme,<br>France                   |
| Fyn <sup>(88)</sup>             | 59 kDa                                   | Rabbit anti-Fyn                                     | 1/500        | #4023     | Cell Signaling/Ozyme,<br>France                   |
| GFAP <sup>(14)</sup>            | 55 kDa                                   | Mouse anti-<br>GFAP                                 | 1/2000       | G3893     | Sigma-Aldrich, France                             |
| GR <sup>(15)</sup>              | 95 kDa                                   | Rabbit anti-GR                                      | 1/1000       | #3660     | Cell Signaling/Ozyme,<br>France                   |
| GSK-3β <sup>(39)</sup>          | 46 kDa                                   | Mouse anti-<br>GSK-3β                               | 1/2000       | 610202    | BD-Biosciences,<br>Rungis, France                 |
| HSP70 <sup>(34)</sup>           | 70 kDa                                   | Rabbit anti-<br>HSP70                               | 1/500        | #4872     | Cell Signaling/Ozyme,<br>France                   |
| HSP90 <sup>(34)</sup>           | 90 kDa                                   | Rabbit anti-<br>HSP90                               | 1/1000       | #4877     | Cell Signaling/Ozyme,<br>France                   |
| lba1 <sup>(14)</sup>            | 17 kDa                                   | Rabbit anti-Iba1                                    | 1/750        | 013-19741 | Wako Chem,,Osaka,<br>Japan                        |
| IDE (14)                        | 110 kDa                                  | Rabbit anti-IDE                                     | 1/3000       | AB9210    | Merck-Millipore, France                           |
| MR <sup>(15)</sup>              | 100 kDa                                  | Rabbit anti-MR                                      | 1/100        | SC11-412  | Dallas, TX, USA                                   |
| p[Ser9]GSK-3β <sup>(39)</sup>   | 46 kDa                                   | Mouse anti-<br>p[Ser9]GSK-3β                        | 1/1000       | #9336     | Cell Signaling/Ozyme,<br>France                   |
| p[Tyr211]GSK-3β <sup>(39)</sup> | 46 kDa                                   | Mouse anti-<br>p[Tyr216]GSK-<br>3β                  | 1/2000       | 612313    | BD-Biosciences,<br>France                         |
| p35/p25 <sup>(87)</sup>         | 35/25<br>kDa                             | Rabbit anti-<br>p35/p25                             | 1/500        | #2680     | Cell Signaling/Ozyme,<br>France                   |
| PDK1 <sup>(58)</sup>            | 56-68<br>kDa                             | Rabbit anti-<br>PDK1                                | 1/1000       | #5662     | Cell Signaling/Ozyme,<br>France                   |
| pGR <sup>(89)</sup>             | 95 kDa                                   | Rabbit anti-<br>p[Ser211]GR                         | 1/1000       | #4161     | Cell Signaling/Ozyme,<br>France                   |
| PS1 <sup>(90)</sup>             | 22 kDa                                   | Rabbit anti-PS1                                     | 1/1000       | #5643     | Cell Signaling/Ozyme,<br>France                   |
| PSD95 <sup>(14)</sup>           | 95 kDa                                   | Rabbit anti-<br>PSD95                               | 1/2000       | #3450     | Cell Signaling/Ozyme,<br>France                   |
| ROCK1 <sup>(58)</sup>           | 160 kDa                                  | Rabbit anti-<br>ROCK1                               | 1/500        | #4035     | Cell Signaling/Ozyme,<br>France                   |
| ROCK2 <sup>(58)</sup>           | 160 kDa                                  | Rabbit anti-<br>ROCK2                               | 1/500        | #8236     | Cell Signaling/Ozyme,<br>France                   |
| sAPPα <sup>(58)</sup>           | 100 kDa                                  | Mouse anti-<br>sAPP $\alpha$                        | 1/50         | 11098     | IBL, Hamburg,<br>Germany                          |
| SYN <sup>(14)</sup>             | 65 kDa                                   | Mouse anti-<br>synaptotagmine                       | 1/1000       | MAB5200   | Merck-Millipore, France                           |
| β-Tub                           | 50 kDa                                   | Mouse anti-β-<br>Tubulin                            | 1/7500       | T4026     | Sigma-Aldrich, France                             |
| Secondary antibodies            |  |   |              |           |   |
| IgG                             | Goat anti-rabbit IgG                     |   | 1/2000       | A61-54    | Sigma-Aldrich, France                             |
| IgG                             | Goat anti-mouse IgG peroxidase conjugate |   | 1/2000       | A67-82    | Sigma-Aldrich, France                             |





### Sup. Figure 2

A.  $\frac{N}{C S} \frac{Veh}{C S} \frac{C-11}{C S}$ Caspase 3- $\beta$ -Tub- $\frac{N}{C S} \frac{Veh}{C S} \frac{C-11}{C S}$ B.  $\frac{N}{C S} \frac{Veh}{C S} \frac{C-11}{C S}$   $\frac{N}{C S} \frac{Veh}{C S} \frac{C-11}{C S}$   $\frac{PSD95}{SYN} - \frac{-55 \text{ kDa}}{-55 \text{ kDa}}$   $\frac{154}{154} - \frac{-17 \text{ kDa}}{-55 \text{ kDa}}$ 

β-Tub – – – – 50 kDa

- C. N Veh C-11 C S C S C S MR pGR -GR -HSP90 -HSP70 β-Tub -- 50 kDa
- D.  $\frac{N}{C} \frac{Veh}{S} \frac{Veh}{C} \frac{C-11}{C} \frac{S}{S}$ Calpain 1 –  $\frac{N}{C} \frac{Veh}{S} \frac{C-11}{C} \frac{S}{S}$  $Fyn - \frac{S}{S} \frac{S}{$

- F. <u>N</u> <u>Veh</u> <u>C-11</u> <u>C S</u> <u>C S</u> <u>C S</u> ADAM10 <u>- - - - - - - - - - - 125 kDa</u> ROCK1 <u>- - - - - - - - - - - - - - 160 kDa</u> ROCK2 <u>- - - - - - - - - - - 160 kDa</u>

# Sup. Figure 3

Α.



Β.



#### SUPPLEMENTARY LEGEND

#### S-Figure 1

Time-course effect of a single icv injection of an oligometric solution of A $\beta_{25-35}$ (oAB25-35) on spatial short term memory (T-maze test) and plasma levels of corticosterone (CORT). Panel A: Experimental protocol - At T0, adult male rats (Sprague Dawley) were injected directly into the lateral ventricles using a stereotaxic apparatus (coordinates: AP -1 mm, L ±1.5 mm, DV -3.5 mm). One group has not undergone surgery (control rats - white column), a second group received an icv injection of scrambled A $\beta_{25-35}$  peptide (negative control - 10  $\mu$ g/rat - light blue) and a third group received an icv injection of  $oA\beta_{25-35}$  peptide ( $oA\beta_{25-35}$  - 5 or 10 µg/rat medium and dark blue, respectively). Animals were tested or sacrificed 1 and 2 weeks after the icv injection. Panel B: As detailed in Materials and Methods, shortterm memory performance (T-maze test) was expressed as the ratio of the time spent in the initially closed arm (B) over the time spent in the previous arm (A). Twoway ANOVA:  $F_{3.54} = 15.3$ , p< 0.0001 for Group;  $F_{1.54} = 0.02$ , p> 0.05 for Time and  $F_{3,54} = 0.42$ , p> 0.05 for Interaction. Panel C: Plasma concentrations of corticosterone (CORT) were determined by Elisa and expressed as ng/ml. Two-way ANOVA:  $F_{3,47} = 24.5$ , p< 0.0001 for Group;  $F_{1,47} = 1.18$ , p> 0.05 for Time and  $F_{3,47} =$ 1.11, p> 0.05 for Interaction.

Data were expressed as Mean  $\pm$  SEM. \* p< 0.05 and \*\* p< 0.01 *vs.* control non-injected rats (control group, C) and + p< 0.05 and ++ p< 0.01 *vs.* scrambled injected rats (negative control group, S).  $\emptyset \emptyset$  p< 0.01 *vs.* selected group. The number of animals in each group is indicated within the columns.

#### S-Figure 2

The lack of effects in the PFC of scrambled Aβ<sub>25-35</sub> peptide in the different groups of animals tested in this study (naive (N), treated with vehicle (V) or with CORT113176 (C-11)) were resumed in this Figure in order to improve the clarity of main figures and to highlight key effects. The different markers were evaluated in these control groups by Western blot in the same time as treated groups. **Panel A:** Apoptosis (caspase 3); **Panel B:** Synaptic (PSD95 & SYN) and Neuroinflammatory (GFAP & Iba1) markers; **Panel C:** GR system (GR, p[Ser211]GR, MR, HSP90, HSP70); **Panel D:** enzymes involved both in GR activation and Tau phosphorylation (Cdk5, p35/p25, GSK-3β, p[Ser9]GSK-3β, p[Tyr216]GSK-3β, Fyn, Calpain 1); **Panel E-F:** amyloidogenic (APP/C99) and non-amyloidogenic (sAPPα) pathways, focusing in particular on the extracellular (ADAM10, BACE1 and IDE), intramembrane (PS1) and intracellular enzymes (ROCK1, ROCK2 and PDK1).

#### S-Figure 3

The effects in the PFC of selective GR modulator on the endogenous  $A\beta_{1-42}$  levels induced by the icv injection of  $oA\beta_{25-35}$ , were evaluated by ELISA.  $A\beta_{1-42}$  levels were evaluated in control (C - white column) and in icv injected rats with 10 µg/rat of scrambled peptide (S - orange column) or  $oA\beta_{25-35}$  (A $\beta$  - red column), treated or not with selective GR modulator, CORT113176 (10 mg/kg per ip injection). For experimental protocol see Fig.1A. **Panel A:**  $A\beta_{1-42}$  ELISA standard curve obtained at 450 nm. **Panel B:** The endogenous concentrations of  $A\beta_{1-42}$  in the PFC were expressed as means ± SEM in pg/g tissue. Two-way ANOVA:  $F_{2,24} = 6.22$  for Group, p< 0.01;  $F_{1,24} = 5.09$  for Treatment, p< 0.05; and  $F_{2,24} = 2.21$  for Interaction, p> 0.05. \* p< 0.05 vs. respective control (C) group, + p< 0.05 vs. respective scrambled (S) group and x p< 0.05 vs. respective naive rat in each group. The number of animals in each group is indicated within the columns.