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Decreased striatal adenosine A_{2A} -dopamine D_2 receptor heteromerization in schizophrenia

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According to the adenosine hypothesis of schizophrenia, the classically associated hyperdopaminergic state may be secondary to a loss of function of the adenosinergic system. Such a hypoadenosinergic state might either be due to a reduction of the extracellular levels of adenosine or alterations in the density of adenosine A_{2A} receptors ($A_{2A}Rs$) or their degree of functional heteromerization with dopamine D_2 receptors (D_2R). In the present study, we provide preclinical and clinical evidences for this latter mechanism. Two animal models for the study of schizophrenia endophenotypes, namely the phencyclidine (PCP) mouse model and the $A_{2A}R$ knockout mice, were used to establish correlations between behavioural and molecular studies. In addition, a new AlphaLISA-based method was implemented to detect native $A_{2A}R$ - D_2R heteromers in mouse and human brain. First, we observed a reduction of prepulse inhibition in $A_{2A}R$ knockout mice, similar to that observed in the PCP animal model of sensory gating impairment of schizophrenia, as well as a significant upregulation of striatal D_2R without changes in $A_{2A}R$ expression in PCP-treated animals. In addition, PCP-treated animals showed a significant reduction of striatal $A_{2A}R$ - D_2R heteromers, as demonstrated by the AlphaLISA-based method. A significant and pronounced reduction of $A_{2A}R$ - D_2R heteromers was next demonstrated in postmortem caudate nucleus from schizophrenic subjects, even though both D_2R and $A_{2A}R$ were upregulated. Finally, in PCP-treated animals, sub-chronic administration of haloperidol or clozapine counteracted the reduction of striatal $A_{2A}R$ - D_2R heteromers. The degree of $A_{2A}R$ - D_2R heteromer formation in schizophrenia might constitute a hallmark of the illness, which indeed should be further studied to establish possible correlations with chronic antipsychotic treatments.

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INTRODUCTION

Schizophrenia is a complex psychiatric disorder with a heterogeneous genetic and neurobiological background, which influence early brain development [1]. The most accepted mechanistic hypothesis of schizophrenia is based on a simultaneous deregulation of glutamatergic and dopaminergic systems [1, 2]. In 1967, Van Rossum [3] proposed a dopaminergic basis of schizophrenia, an increase in dopaminergic neurotransmission. Indeed, most individuals with schizophrenia are behaviourally supersensitive to drugs inducing activation of dopamine receptors [4] and the positive symptoms are thought to be associated with hyperactivation of the dopaminergic system. Early work analyzing post-mortem brains from patients with schizophrenia revealed a higher striatal density of dopamine D_2 receptors (D_2Rs) [5]. Recent neuroimaging studies indicate that striatal D_2R are augmented by about $5.8 \pm 2.7\%$ in schizophrenic subjects [6] and it has been

postulated that the D_2R population with higher affinity for dopamine are preferentially involved [7], although with some considerations [8]. Thus, a meta-analysis of imaging studies supported the elevation in the D_2R availability but gave more significance to a presynaptic component of dopamine release [8]. In agreement with the involvement of dopamine and D_2R in schizophrenia, classical antipsychotics such as haloperidol [9] are D_2R antagonists or weak partial agonists [10]. However, although efficacious against positive symptoms, haloperidol and related compounds induce severe side effects, including motor impairments [11].

The purinergic system may also be involved in the biology of psychiatric disorders, including schizophrenia [12]. Indeed, the adenosine hypothesis of schizophrenia [13, 14] posits that dysregulation of both glutamatergic and dopaminergic signalling are secondary to a hypoadenosinergic state. The adenosinergic

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control of the striatal dopaminergic system largely depends on the existence of complexes (i.e., heteromers) formed by adenosine and dopamine receptors, more specifically heteromers of adenosine A_{2A} receptors (A_{2A}R) and D₂R [15]. Accordingly, a hypoadenosinergic state, with a decline in extracellular adenosine concentration or a relative downregulation of A_{2A}R vs. D₂R, would reduce the well-established tonic A_{2A}R-mediated inhibition of D₂R function in the A_{2A}R–D₂R heteromer [15]. It has therefore been postulated that increasing A_{2A}R-mediated allosteric inhibition of D₂R within the striatopallidal GABA pathway might represent a promising strategy for schizophrenia management [15–17]. In line with this, administration of a selective A_{2A}R agonist effectively antagonized phencyclidine (PCP)-induced locomotor activity in rats [18], an animal model of schizophrenic symptomatology [19]. Similarly, the non-selective adenosine receptor antagonist caffeine has been described to exacerbate psychosis in schizophrenic patients [20].

The present study aimed at exploring potential alterations of A_{2A}R and D₂R striatal expression and their degree of heteromerization in mice treated sub-chronically with PCP and in postmortem striatal tissue from schizophrenic subjects, using our recently developed AlphaLISA-based approach [21]. Furthermore, we evaluated whether sub-chronic treatment with the typical antipsychotic haloperidol and the atypical antipsychotic clozapine could modify these alterations in PCP-treated mice.

METHODS AND MATERIALS

Human brain samples

Postmortem human brain samples were obtained at autopsy in the Basque Institute of Legal Medicine, Bilbao, Spain. The study was carried out in compliance with the policies of research and ethical review boards for postmortem brain studies at the moment of sample collection (Basque Institute of Legal Medicine, Bilbao). After a retrospective search for antemortem medical information, ten brain samples of subjects diagnosed with schizophrenia according to DSM-IV, DSM-IV-R or CIE-10 criteria were matched to ten control subjects in a paired design. Mean age, postmortem interval, RNA integrity number, brain pH or tissue storage time did not differ significantly between controls and schizophrenic subjects (Supplementary Table S1). Samples from the caudate nucleus were dissected at autopsy following standard procedures and were immediately stored at –70 °C until assay. For every experimental procedure, each subject ascribed to the schizophrenia group was processed in parallel to a matched control subject.

Animals

A_{2A}R^{–/–} and D₂R^{–/–} mice generated on a CD-1 genetic background [22, 23] and the corresponding littermates weighing 20–25 g were used. The animal protocol (#7085) was approved by the University of Barcelona Committee on Animal Use and Care. Animals were housed and tested in compliance with the guidelines provided by the Guide for the Care and Use of Laboratory Animals [24] and following the European Union directives (2010/63/EU). Mice were housed in groups of five in standard cages with ad libitum access to food and water and maintained under a 12 h dark/light cycle (starting at 7:30 a.m.), 22 °C temperature and 66% humidity (standard conditions).

Drug administration

PCP (10 mg/kg, subcutaneously; Tocris, Bristol, UK), haloperidol (0.1 mg/kg, intraperitoneally (i.p.); Tocris) and clozapine (10 mg/kg, i.p.; Tocris) were dissolved in physiological saline (NaCl 0.9%) and administered for 5 consecutive days. At the end of treatment (2–3 h after the final drug administration) animals were used for behavioural assessments and biochemical experiments.

Prepulse inhibition of the acoustic startle response

Mice aged 2–3 months were first handled for 5 days prior to behavioural tests. All the tests were performed during the light phase of the circadian cycle (between 8:30 and 14:00 h) by a researcher blind to drug treatments. At the end of the trial, mice were briefly returned to their home cage before being used in biochemical experiments. Prepulse inhibition (PPI) test was performed using the StartFear System (Panlab, Cornellà de Llobregat, Spain), as described in detailed elsewhere [25] and in the Supplementary Information.

Determination of ribonucleosides and monoamines in mouse striatum

Mice were rapidly killed by cervical dislocation and 400 µm brain coronal sections were obtained in carbogen (95% O₂ and 5% CO₂) bubbled artificial cerebrospinal fluid buffer (127 mM NaCl, 1 mM KCl, 1.2 mM KH₂PO₄, 26 mM NaHCO₃, 10 mM D-glucose, 2.4 mM CaCl₂ and 1.3 mM MgCl₂, pH 7.4). Next, the striatum was dissected from the slice and homogenized in 200 µl of acetonitrile (Sigma-Aldrich) containing 2% CH₃COOH (Sigma-Aldrich) and dC-d₃ (1 ng/µl) at 4 °C in a 1 ml Potter–Elvehjem glass tube using a homogenizer stirrer HS-30E (Witeg Labor Technik GmbH, Wertheim, Germany) with 10 strokes at 700–900 rotations per min. Finally, samples were centrifuged at 13,200 r.p.m. at 4 °C for 20 min and the supernatants were stored at –80 °C until analysis. Quantitative analysis of ribonucleosides (i.e., adenosine, cytidine, guanosine and uridine) and monoamines (i.e., dopamine and serotonin) was carried out by liquid chromatography technique coupled with tandem mass spectrometry and is described in detail in the Supplementary Information.

Immunohistofluorescence analysis

Mice were anaesthetized and perfused intracardially with 100–200 ml ice-cold 4% paraformaldehyde (PFA) in phosphate-buffered saline (PBS; 8.07 mM Na₂HPO₄, 1.47 mM KH₂PO₄, 137 mM NaCl, 0.27 mM KCl, pH 7.2). Brains were post-fixed in the same solution of PFA at 4 °C during 12 h. Coronal sections (50 µm) were obtained using a vibratome (Leica Lasertechnik GmbH, Heidelberg, Germany). Slices were collected in Walter's Antifreezing solution (30% glycerol, 30% ethylene glycol in PBS, pH 7.2) and kept at –20 °C until processing. For immunohistofluorescence analysis, the slices were washed three times with PBS, permeabilized with 0.5% Triton X-100 in PBS for 2 h and rinsed again three times with washing solution (0.05% Triton X-100 in PBS). The slices were then incubated with washing solution containing 10% normal donkey serum (NDS; Jackson ImmunoResearch Laboratories, Inc., West Grove, PA, USA) for 2 h at room temperature. Subsequently, slices were incubated with goat anti-A_{2A}R (3 µg/ml; Santa Cruz Biotechnology, Inc., Dallas, TX, USA), rabbit anti-D₂R (3 µg/ml; Frontier Institute Co. Ltd, Hokkaido, Japan) or goat anti-DAT (0.5 µg/ml; Santa Cruz Biotechnology, Inc.) in washing solution containing 10% NDS for 24 h at 4 °C. Next, slices were washed with washing solution containing 1% NDS before the incubation with Cy3-conjugated donkey anti-goat or Cy2-conjugated donkey anti-rabbit IgG antibody (1:200; Jackson ImmunoResearch Laboratories, West Grove, PA, USA) in washing solution for 2 h at room temperature. Finally, slices were washed twice with washing solution containing 10% NDS and then mounted with Vectashield immunofluorescence medium (Vector Laboratories, Peterborough, UK) in glass slides. Fluorescence striatal images were captured using a Leica TCS 4D confocal scanning laser microscope (Leica Lasertechnik GmbH).

Gel electrophoresis and immunoblotting

Mouse striatum or frozen human caudate were homogenized in ice-cold 10 mM Tris HCl, pH 7.4, 1 mM EDTA, 300 mM KCl buffer containing a protease inhibitor cocktail (Roche Molecular Systems, USA) using a Polytron for three periods of 10 s each. The

homogenate was centrifuged for 10 min at 1000 × *g*. The resulting supernatant was centrifuged for 30 min at 12,000 × *g*. The membranes were dispersed in 50 mM Tris HCl (pH 7.4) and 10 mM MgCl₂, washed, and resuspended in the same medium as described previously [26]. Protein concentration was determined using the BCA protein assay kit (Thermo Fisher Scientific, Inc., Rockford, IL, USA) and 50–80 µg of protein was used for immunoblotting.

Sodium dodecyl sulfate-polyacrylamide gel electrophoresis was performed using 10% polyacrylamide gels. Proteins were transferred to Hybond[®]-LFP polyvinylidene difluoride (PVDF) membranes (GE Healthcare, Chicago, IL, USA) using a Trans-Blot[®] SD Semi-Dry Transfer Cell (Bio-Rad, Hercules, CA, USA). PVDF membranes were blocked with 5% (wt/vol) dry non-fat milk in PBS containing 0.05% Tween-20 (PBS-T) during 45 min and immunoblotted using goat polyclonal anti-A_{2A}R (0.5 µg/ml; Santa Cruz Biotechnology, Inc.), rabbit polyclonal anti-D₂R (1 µg/ml; Frontier Institute Co. Ltd), guinea pig polyclonal anti-A_{2A}R (1 µg/ml; Frontier Institute Co. Ltd), goat anti-DAT (0.5 µg/ml; Santa Cruz Biotechnology, Inc.) and rabbit polyclonal anti-α-actinin (0.5 µg/ml; Santa Cruz Biotechnology, Inc.) antibodies in blocking solution overnight at 4 °C. PVDF membranes were washed with PBS-T three times (5 min each) before incubation with either a horseradish peroxidase (HRP)-conjugated goat anti-rabbit IgG (1/30,000; Pierce Biotechnology, Rockford, IL, USA), HRP-conjugated rabbit anti-goat IgG (1/10,000; Pierce Biotechnology) or HRP-conjugated rabbit anti-guinea pig IgG (1/3,000; Pierce Biotechnology) in blocking solution at 20 °C during 2 h. After washing the PVDF membranes with PBS-T three times (5 min each), the immunoreactive bands were developed using a chemiluminescent detection kit (Thermo Fisher Scientific, Waltham, MA, USA) and were detected with an Amersham Imager 600 (GE Healthcare Europe GmbH, Barcelona, Spain).

AlphaLISA method

Mouse striatal and human caudate membranes were homogenized in AlphaLISA buffer (AlphaLISA HiBlock Buffer, PerkinElmer, Waltham, MA, EEUU) and protein concentration determined. Thus, 5 µg of protein/well was placed in white 384 well-plates (384 Well Small Volume™ HiBase Microplates, Greiner Bio-one, Kremsmünster, Austria). Subsequently, membranes were incubated with 5 µl of 10 nM donor primary antibody plus 5 µl of 10 nM acceptor primary antibody (rabbit anti-D₂R and guinea pig anti-A_{2A}R, respectively) overnight at 4 °C. As a negative control, samples were also incubated in the absence of either the acceptor or donor primary antibody (Supplementary Fig. S1a). In addition, striatal membranes from A_{2A}R^{-/-} and D₂R^{-/-} mice were used negative controls to further validate the new AlphaLISA method implemented here for the detection of A_{2A}R–D₂R heteromers in native tissue (Supplementary Fig. S1b). All reagents were mixed by pipetting up and down before incubation. After the overnight incubation, plates were tempered at 22 °C before acceptor beads were added. Anti-guinea pig conjugated acceptor beads were generated by covalent binding of AffiniPure F(ab)₂ Fragment Donkey Anti-Guinea Pig IgG (H + L) (Jackson ImmunoResearch Laboratories, Inc., West Grove, PA, USA) to acceptor beads (AlphaLISA acceptor beads, 6772001; PerkinElmer) following the manufacturer's instructions. Subsequently, acceptor beads were diluted in AlphaLISA buffer and 5 µL added to each well (final concentration 40 µg/mL). The reagents were mixed by pipetting. After 1 h incubation (in darkness at 22 °C), donor beads (anti-rabbit IgG AlphaLISA Donor beads, AS105D, PerkinElmer) were diluted in AlphaLISA buffer and 5 µL of diluted beads added to each well (final concentration 10 µg/mL). The final reaction volume was 30 µL. The reagents were mixed by pipetting. Finally, after 24 h incubation (in darkness at 4 °C), the microplate was read using a CLARIOstar plate-reader (BMG Labtech, Durham, NC, USA). The plate was left in the dark at room temperature and read again at 48 h. A background signal (B) was determined in wells where all

reagents were added without sample. For each membrane brain sample a negative control (C_n) was determined by incubating in the absence of acceptor primary antibody (i.e., guinea pig anti-A_{2A}R antibody). Therefore, for each sample the specific AlphaLISA signal (i.e., ΔAlphaLISA) was calculated as follows:

$$\Delta\text{AlphaLISA}_{(n)} = (\text{AlphaLISA signal}_{(n)} - B) - (C_{(n)} - B)$$

Statistics

Data are represented as mean ± SEM. The number of samples/animals (*n*) in each experimental condition is indicated in the corresponding figure legend. Data normality was assessed by the Shapiro–Wilk normality test. Outliers were assessed by the Grubbs' test; thus, data from any animal found to be an outlier was excluded. Comparisons among experimental groups were performed by Student's *t*-test, one-way analysis of variance (ANOVA) with Dunnett's post hoc test or two-way ANOVA with Sidak's post hoc test using GraphPad Prism 6.01 (San Diego, CA, USA), as indicated. Statistical difference was accepted when *P* < 0.05.

RESULTS

Alterations in sensorimotor gating in PCP-treated animals and in mice lacking A_{2A}R

The adenosine hypothesis of schizophrenia [13, 14] was recently bolstered by the observation that A_{2A}R deletion (A_{2A}R^{-/-}) prompts behaviours that mimic symptoms of psychosis and molecular/anatomical alterations in mice, resembling the relevant phenotype features of the human disorder [23, 27, 28]. We first validated the A_{2A}R^{-/-}-based animal model of psychotic symptoms by comparing it with the well-established PCP animal model of sensory gating impairment of schizophrenia [19, 29, 30]. A_{2A}R^{-/-} and A_{2A}R^{+/+} saline- and PCP-treated mice were evaluated with the PPI test, a classically accepted behavioural assessment of sensorimotor processing altered in schizophrenic patients and in PCP-treated rodents [19, 29]. As previously reported [23, 28], A_{2A}R^{-/-} mice showed a significant reduction in basal PPI at 70 dB (*P* = 0.0156), 75 dB (*P* = 0.0023) and 80 dB (*P* = 0.0001) (Fig. 1). PCP administration to A_{2A}R^{+/+} mice also induced a significant PPI reduction, up to half of basal values, at 70 dB (*P* = 0.0068), 75 dB (*P* = 0.0025) and 80 dB (*P* = 0.0119) (Fig. 1). These results demonstrated that sensorimotor processing alterations in A_{2A}R^{-/-} mice were comparable to those observed in PCP-treated wild-type (WT) animals.

Striatal A_{2A}R and D₂R expression profiles in A_{2A}R^{-/-} and PCP-treated mice

Next, we aimed at determining possible alterations in striatal A_{2A}R content in the PCP animal model, as well as possible alterations in striatal D₂R densities in PCP-treated and in A_{2A}R^{-/-} mice groups, by performing immunoblotting and immunocytochemistry experiments. First, we assessed A_{2A}R and D₂R expression by immunoblot analysis. A_{2A}R and D₂R expression was ascertained by the presence of protein bands of molecular weight ~45 kDa and ~70–80 kDa, respectively (Fig. 2a, d), which were absent in A_{2A}R- and D₂R-knockout mice samples as previously demonstrated [23]. In addition, we assessed the dopamine transporter (DAT), as alterations in DAT expression have been recently described in another rodent model of psychosis [31]. Importantly, although A_{2A}R and DAT expression were not altered in the striatum of PCP-treated animals, the expression of D₂R was significantly increased (*P* = 0.0006) (Fig. 2a, b). Immunohistochemistry analysis of A_{2A}R, D₂R and DAT in coronal slices at the level of the anterior commissure from PCP-treated mice also revealed higher D₂R immunoreactivity in the striatum (Fig. 2c). Overall, these results indicate that PCP treatment raised D₂R expression in the striatum, while A_{2A}R and DAT were unaffected.

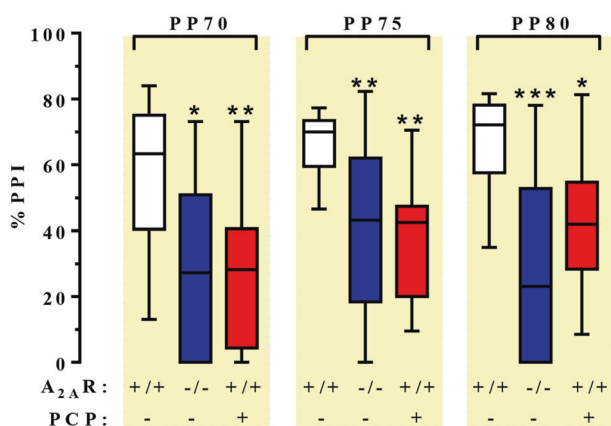


Fig. 1 Sensorimotor processing in WT, PCP-treated and A_{2A}R^{-/-} mice. The prepulse inhibition (PPI) impairment of the acoustic startle response (ASR) was assessed and expressed as percentage mean ± SEM ($n = 10-15$ animals) of inhibition of ASR at the indicated prepulse acoustic stimulus amplitudes (i.e., 70, 75 and 80 dB). * $P < 0.05$, ** $P < 0.01$ and *** $P < 0.001$, one-way ANOVA followed by Dunnett's post hoc test compared with WT (A_{2A}R^{+/+}) group. Two-way ANOVA (PP × phenotype/PCP) revealed a significant main effect of phenotype/PCP ($F_{(2,117)} = 22.65$, $P < 0.0001$) and a nonsignificant effect of PP ($F_{(2,117)} = 2.616$, $P = 0.0774$) or the interaction between both factors ($F_{(4,117)} = 0.5762$, $P = 0.6804$).

Subsequently, we analyzed A_{2A}R, D₂R and DAT expression in WT and A_{2A}R^{-/-} mice. Similar to what was observed in PCP-treated mice, immunoblotting analysis revealed that deletion of A_{2A}R induced a significant increase in D₂R ($P = 0.0433$) but also in DAT ($P = 0.0014$) expression in the striatum (Fig. 2d, e). As expected, A_{2A}R expression was abolished in A_{2A}R^{-/-} (Fig. 2d, e). In addition, immunohistochemistry detection of A_{2A}R, D₂R and DAT in WT mice also revealed increased D₂R and DAT immunoreactivity in the striatum (Fig. 2f). As expected, A_{2A}R immunoreactivity could not be detected in A_{2A}R^{-/-} mice (Fig. 2f). Collectively, these results indicate that the expression of D₂R in the striatum of A_{2A}R^{-/-} mice is increased, similar to what was observed in the PCP-based animal model of schizophrenic symptomatology.

Striatal levels of ribonucleosides and monoamines in A_{2A}R^{-/-} and PCP-treated mice

To determine a possible presynaptic imbalance, striatal levels of adenosine and dopamine were determined in A_{2A}R^{+/+}, A_{2A}R^{-/-} and PCP-treated mice. The levels of the ribonucleosides cytidine, guanosine and uridine, and the monoamine serotonin were also determined (Supplementary Table S2). Interestingly, although the levels of dopamine and serotonin were not altered in A_{2A}R^{-/-} or in PCP-treated animals, the amounts of adenosine and cytidine were significantly increased in A_{2A}R^{-/-} mice (Supplementary Table S2). Also, the levels of uridine were significantly raised both in A_{2A}R^{-/-} and PCP-treated animals (Supplementary Table S2). These results indicate that purine metabolism is altered in the A_{2A}R^{-/-} mouse model, which displays a striatal neurochemical pattern different from that observed in the PCP model of schizophrenic symptomatology. Altogether, the results (i.e., density of striatal A_{2A}R and D₂R, and neurotransmitter levels) obtained from PCP-treated and A_{2A}R^{-/-} mouse models do not support a presynaptic adenosine–dopamine imbalance, but rather support a postsynaptic A_{2A}R–D₂R density imbalance in schizophrenia.

A_{2A}R, D₂R and DAT expression profiles in the caudate nucleus from postmortem schizophrenic subjects

As mentioned in the introductory section, previous postmortem and imaging studies suggested the presence of an increased D₂R expression in the striatum of schizophrenic subjects [5, 6].

Conversely, the results with DAT and A_{2A}R are inconsistent. A recent meta-analysis indicates that there is a significantly greater interindividual variability of striatal DAT availability in patients with schizophrenia, compared to control subjects. These data suggest that altered DAT expression may occur only in a group of patients [32]. Inconsistent results have also been obtained in relation to striatal A_{2A}R expression in schizophrenia, with reports of either upregulation or downregulation [33, 34]. Therefore, we aimed at determining whether the relative expression of D₂R, A_{2A}R and DAT were altered in postmortem caudate from a group of schizophrenic subjects (Supplementary Table S1). From immunoblot analysis of postmortem caudate nucleus membranes obtained from a cohort of ten healthy controls and ten subjects diagnosed with paranoid schizophrenia (Supplementary Table S1), we found that D₂R, DAT and A_{2A}R expression was significantly ($P < 0.05$) increased in the caudate nucleus from subjects with schizophrenia (Fig. 3a, b).

Decrease in A_{2A}R–D₂R heteromerization in the caudate nucleus of schizophrenic subjects and the striatum of PCP-treated mice

We next interrogated whether the postulated hypoadenosinergic state in schizophrenia could be dependent on a significant change in the interactions between A_{2A}R and D₂R, i.e., in their well-established ability to form functional heteromers [15]. To this end, we used our recently engineered AlphaLISA-based method to assess A_{2A}R–D₂R interactions in human postmortem brain [21] (Fig. 4a). The AlphaLISA approach revealed a strong and specific energy transfer between the donor and acceptor beads coupled to specific receptor antibodies (Supplementary Fig. 1). This energy transfer confirmed that the intermolecular interaction between A_{2A}R and D₂R, i.e., A_{2A}R–D₂R heteromers, can be detected in postmortem membrane extracts (Fig. 4b), as recently reported [21]. A significant and pronounced reduction of $59 \pm 11\%$ ($P = 0.0076$) in A_{2A}R–D₂R heteromerization was observed in the human caudate membrane extracts from schizophrenic subjects as compared to controls (Fig. 4b, right panel). Similarly, when striatal membrane extracts from PCP-treated animals were assessed, a comparable significant reduction of $75 \pm 7\%$ ($P = 0.0001$) in A_{2A}R–D₂R heteromerization was detected as compared to controls (Fig. 4b, left panel). Overall, these results indicate that striatal A_{2A}R–D₂R heteromerization was reduced both in schizophrenic subjects and in the PCP-based animal model of schizophrenic symptoms, thus predicting a role for alterations in the formation of the striatal A_{2A}R–D₂R heteromers in the pathophysiology of schizophrenia.

Counteracting effects of sub-chronic antipsychotic treatment on PCP-induced reduction of A_{2A}R–D₂R heteromerization

In view that schizophrenic patients showed a reduced A_{2A}R–D₂R heteromer formation, we next aimed at determining whether antipsychotic medications may prevent PCP-mediated A_{2A}R–D₂R heteromer downregulation in mice. Interestingly, previous pre-clinical experiments evaluating the effects of neuroleptics on A_{2A}R–D₂R interactions found that chronic haloperidol treatment induced an increase in functional A_{2A}R–D₂R interactions. More specifically, chronic treatment with haloperidol (administered in drinking water for 30 consecutive days, adjusted to 1.5 mg/kg) in rats led to an increased potency of A_{2A}R agonists in reducing the affinity of D₂R agonists in striatal membrane preparations and to an increase in the locomotor-activating effects of the adenosine receptor antagonist theophylline [35]. However, we could not evaluate a possible effect of antipsychotics on striatal A_{2A}R–D₂R heteromerization in our cohort sample, since it was limited to ten subjects with paranoid schizophrenia treated with different compounds (Supplementary Table S1). Therefore, we aimed at ascertaining the impact of neuroleptic treatment on A_{2A}R–D₂R heteromer formation in PCP-treated mice. We evaluated the possible modification of striatal A_{2A}R–D₂R heteromerization upon

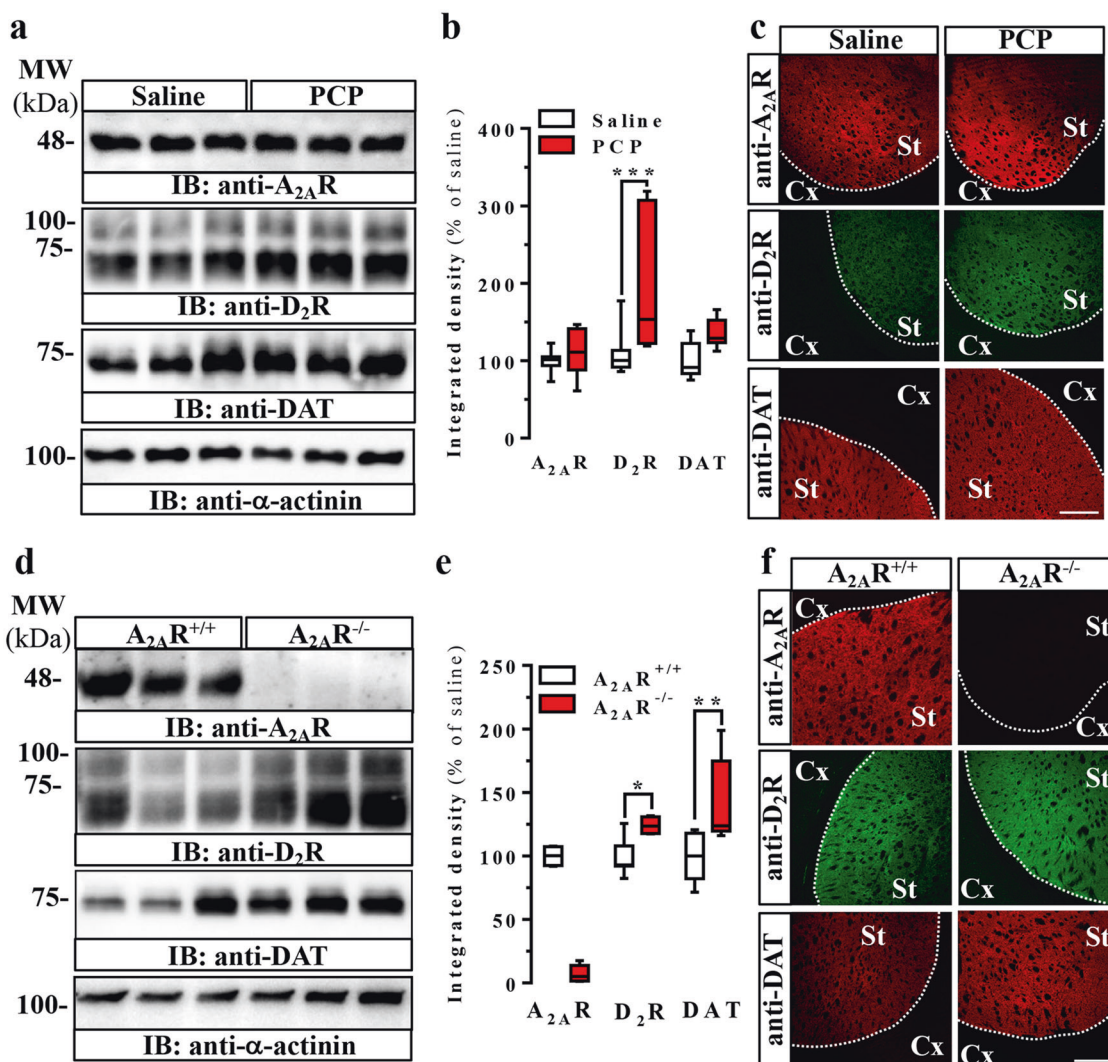


Fig. 2 Expression of A_{2A}R, D₂R and DAT in the striatum of PCP-treated and A_{2A}R^{-/-} mice. **a, d** Representative immunoblots showing the expression of A_{2A}R, D₂R and DAT in striatal membranes from saline and PCP-treated mice (**a**) and striatal membranes from WT (A_{2A}R^{+/+}) and A_{2A}R^{-/-} mice (**d**); striatal membranes were analyzed by SDS-PAGE (50 μg of protein/lane) and immunoblotted using goat anti-A_{2A}R, rabbit anti-D₂R, goat anti-DAT and rabbit anti-α-actinin antibodies (see Methods). **b, e** Relative quantification of A_{2A}R, D₂R and DAT expression. The immunoblot protein bands corresponding to A_{2A}R, D₂R, DAT and α-actinin from saline and PCP-treated mice (**b**; n = 6–9) and from WT and A_{2A}R^{-/-} mice (**e**; n = 5–7) were quantified by densitometric scanning; values were normalized to the respective amount of α-actinin in each lane to correct for protein loading; results are expressed as percentage (mean ± SEM) of the corresponding saline-treated or WT animals; *P < 0.05, **P < 0.01 and ***P < 0.001 two-way ANOVA with Sidak's post hoc test. **c, f** Representative images of A_{2A}R, D₂R and DAT immunoreactivities in the dorsal striatum of saline and PCP-treated mice (**c**) or WT and A_{2A}R^{-/-} mice (**f**); scale bar: 350 μm. Cx, cortex; St, striatum.

administration of haloperidol or clozapine in mice repeatedly administered with PCP. We could first confirm that sub-chronic treatment with haloperidol or clozapine did not modify PPI in control mice. Conversely, they both counteracted PCP-induced disruption of PPI (Fig. 5a). On the other hand, the AlphaLISA approach revealed that neither haloperidol nor clozapine modified A_{2A}R–D₂R heteromerization in control mice but prevented the reduction of A_{2A}R–D₂R heteromerization in PCP-treated animals (Fig. 5b). Finally, when the density of striatal A_{2A}R and D₂R was assessed by immunoblotting, we confirmed that PCP produced a significant increase in the density of D₂R (Fig. 5c, d). This effect was not modified by haloperidol or clozapine treatment (Fig. 5c, d). Similarly, in the absence of PCP, neither haloperidol nor clozapine treatment significantly modified A_{2A}R and D₂R densities (Fig. 5c, d). These results imply that sub-chronic treatment with haloperidol or clozapine does not lead to an increase in A_{2A}R–D₂R heteromerization in healthy subjects. However, antipsychotics are

able to reverse PCP-induced reduction in striatal A_{2A}R–D₂R heteromerization and, as previously reported [36], PCP-induced disruption of PPI.

DISCUSSION

Targeting D₂R-containing heteromers (i.e., A_{2A}R–D₂R) represents a new strategy for psychosis management [16]. Our results indicate that the presence of sensitized D₂R in schizophrenia may be concomitant to a reduction in A_{2A}R activity secondary to a deficiency in A_{2A}R–D₂R heteromer formation. Indeed, we observed a reduction of PPI in A_{2A}R^{-/-} mice, similar to the PPI deficits in PCP-treated mice. In addition, these animals showed a significant upregulation of striatal D₂R without concomitant changes in the expression of A_{2A}R (a relative downregulation of A_{2A}R vs. D₂R). More importantly, we could establish a correlation between striatal A_{2A}R–D₂R heteromer downregulation and PCP-induced

sensory gating impairment in mice. Finally, in schizophrenic subjects, A_{2A}R–D₂R heteromers were also reduced, even though the expression of both A_{2A}R and D₂R was increased.

Apart from the already supported antipsychotic potential for A_{2A}R agonists [18], drugs promoting A_{2A}R–D₂R heteromer formation might constitute an alternative strategy for the treatment of schizophrenia. In fact, we could demonstrate that, in the PCP animal model of schizophrenic symptomatology, sub-chronic treatment with either the typical antipsychotic, haloperidol, or the atypical antipsychotic, clozapine, counteracted PCP-induced

PPI disruption and PCP-induced reduction in striatal A_{2A}R–D₂R heteromerization. Conversely, antipsychotics did not change the degree of A_{2A}R–D₂R heteromerization in control animals, without PCP treatment. A more sustained treatment with antipsychotics could therefore lead to further increase in striatal A_{2A}R–D₂R heteromerization, as suggested from previous experiments in rats chronically treated with haloperidol [35]. Upregulation of striatal A_{2A}R–D₂R heteromers could therefore represent a protective mechanism associated with chronic neuroleptic treatment.

Schizophrenia has a strong heritable component and some genetic risk factors have been already established. Accordingly, genetic animal models are important tools to study the mechanisms underlying this disease. Based on the evidence that striatal D₂R are upregulated in schizophrenic patients [6], an animal model of schizophrenia based on D₂R overexpression in the striatum (i.e., D₂R-OE mice) was created using an artificial transcription factor system [37]. However, although D₂R-OE mice showed deficits in working memory tasks and behavioural flexibility, they did not present deficits in PPI [37]. Several gene knockouts, including those of the metabotropic glutamate mGlu₂ and mGlu₃ receptors, PSD95 (postsynaptic density protein 95), GPRK6 (G-protein receptor kinase 6), the Trace Amine-1 receptor, and RGS9-2 (regulator of G-protein signalling 9-2), have also been proposed as animal model of psychosis, which may represent different aspects of the disease [38, 39]. Our results support the validity of a novel genetic model; A_{2A}R^{-/-} mice. Thus, similar to what is observed in the widely used PCP animal model schizophrenic symptomatology, A_{2A}R^{-/-} mice display PPI deficits and increased striatal D₂R and DAT densities.

Altogether, the results of the present study provide strong support to the adenosine hypothesis of schizophrenia [13, 14]. However, rather than pointing to adenosine itself, our data highlight A_{2A}Rs and their fundamental role in controlling striatal D₂R function [15, 16, 40]. Overall, apart from presynaptic mechanisms [8, 32], the pathophysiology of schizophrenia seems to involve postsynaptic mechanisms which, rather than be related to D₂R upregulation, it involves the loss of the control of D₂R by A_{2A}R in the striatum. Changes in the density of striatal A_{2A}R–D₂R heteromers should then be expected to contribute to the changes observed in positron emission tomography (PET) imaging studies that use D₂R ligands in patients with schizophrenia. Those changes would then depend on the complex allosteric

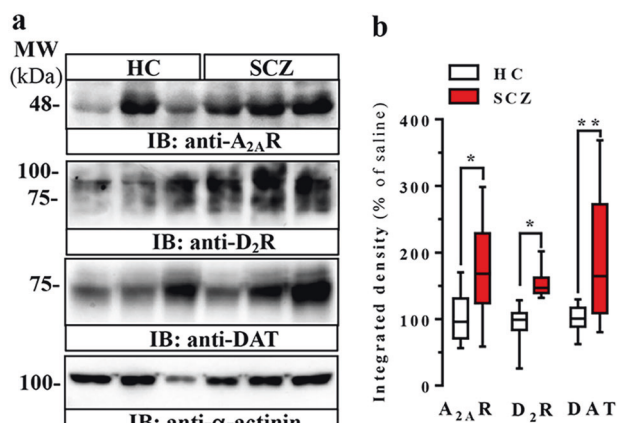


Fig. 3 Expression of A_{2A}R, D₂R and DAT in schizophrenic subjects. **a** Representative immunoblot showing the expression of A_{2A}R, D₂R and DAT in postmortem caudate membranes from healthy control (HC) and schizophrenic (SCZ) subjects; membranes from human postmortem caudate were analyzed by SDS-PAGE (50 μg of protein/lane) and immunoblotted using goat anti-A_{2A}R, rabbit anti-D₂R, goat anti-DAT and rabbit anti-α-actinin antibodies (see ‘Methods’). **b** Relative quantification of A_{2A}R, D₂R and DAT expression; the immunoblot protein bands corresponding to A_{2A}R, D₂R, DAT and α-actinin from healthy control (*n* = 10) and schizophrenic (*n* = 10) individuals were quantified by densitometric scanning; values were normalized to the respective amount of α-actinin in each lane to correct for protein loading. Results are expressed as percentage (mean ± SEM) of the control. **P* < 0.05, ***P* < 0.01, two-way ANOVA with Sidak’s post hoc test.

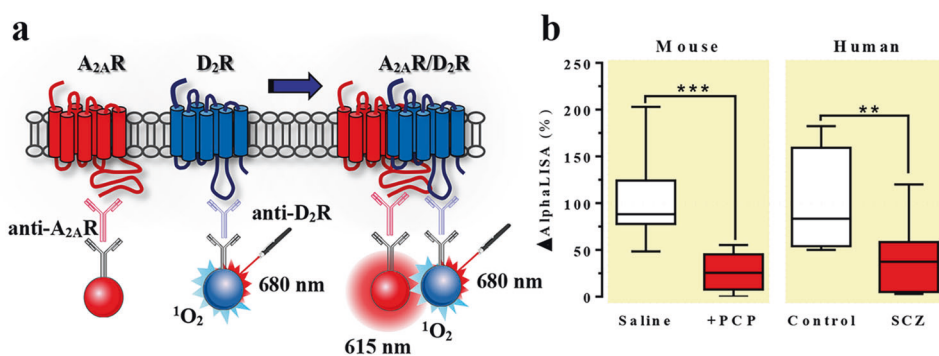


Fig. 4 A_{2A}R–D₂R heteromer assessment in mouse striatum and human caudate using an AlphaLISA approach. **a** Illustration of the specific AlphaLISA protein–protein interaction assay designed for A_{2A}R–D₂R heteromer identification and quantification in native tissue; anti-guinea pig-coated acceptor beads (red sphere) were generated to capture a guinea pig anti-A_{2A}R antibodies bound to the receptors within the membrane extract; anti-rabbit coated donor beads (blue sphere) capture the immune complexes between the rabbit anti-D₂R antibodies and the receptors within the membrane extract; A_{2A}R–D₂R heteromerization brings donor beads into close proximity (<200 nm) to the acceptor beads. The excitation of the donor beads at 680 nm generates singlet oxygen (¹O₂) molecules triggering a chemical reaction within the acceptor beads, which results in a sharp peak of fluorescent emission at 615 nm (figure designed using image templates from Servier Medical Art <https://smart.servier.com/image-set-download/>). **b** The A_{2A}R–D₂R interaction capacity in membranes from saline- (*n* = 10) and PCP- (*n* = 10) treated mouse striatum (left panel) or from postmortem control (*n* = 10) and schizophrenic (*n* = 10; SCZ) caudate (right panel) was determined by AlphaLISA method (see ‘Methods’); the specific AlphaLISA signal (i.e., ΔAlphaLISA) was calculated as described in the ‘Methods’ section and expressed as percentage (mean ± SEM) of either the saline-treated mice or control subjects. ***P* < 0.01 and ****P* < 0.001, Student’s *t*-test.

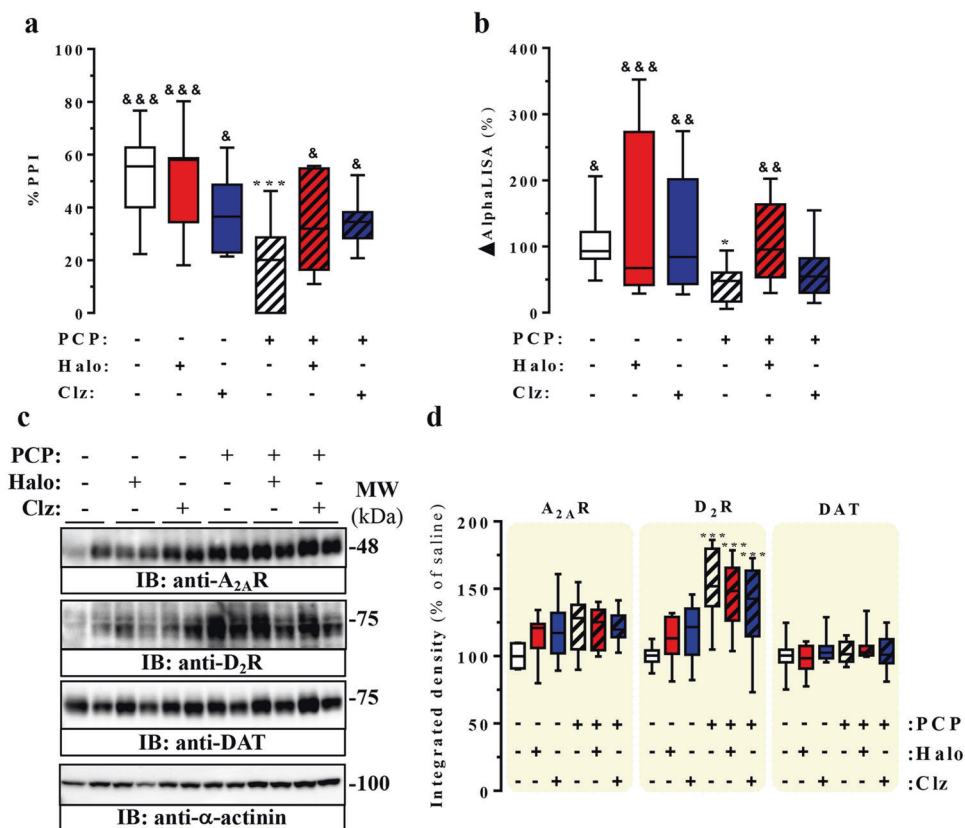


Fig. 5 Effect of antipsychotics on A_{2A}R and D₂R density and A_{2A}R–D₂R heteromerization in the PCP animal model. **a** PPI impairment in saline- (*n* = 10) and PCP-treated animals (*n* = 10) with or without chronic treatment with haloperidol (Halo, 0.1 mg/kg/day for 5 days) or clozapine (Clz, 10 mg/kg/day for 5 days); results are expressed as percentage (mean ± SEM) of inhibition of ASR at the 75 dB amplitude prepulse acoustic stimulus; * *P* < 0.05, one-way ANOVA followed by Dunnett's post hoc test when compared to saline-treated mice. **b** A_{2A}R–D₂R heteromerization in striatal membranes from the same animals shown in **a** determined by AlphaLISA method (see 'Methods'); the specific AlphaLISA signal (i.e., ΔAlphaLISA) was calculated as described in the 'Methods' section and expressed as percentage (mean ± SEM) of the saline-treated mice. **P* < 0.05 one-way ANOVA with Dunnett's post hoc test when compared to saline-treated mice. **c** Representative immunoblot showing the expression of A_{2A}R, D₂R and DAT in striatal membranes from animals from the same groups shown in **a**; striatal membranes from PCP-treated mice were analyzed by SDS-PAGE (50 μg of protein/lane) and immunoblotted using guinea pig anti-A_{2A}R, rabbit anti-D₂R, goat anti-DAT and rabbit anti-α-actinin antibodies (see 'Methods'). **d** Relative quantification of A_{2A}R, D₂R and DAT expression; the immunoblot protein bands corresponding to A_{2A}R, D₂R, DAT and α-actinin from the same animals shown in **a** were quantified by densitometric scanning; values were normalized by the respective amount α-actinin in each lane to correct for protein loading. Results are expressed as percentage (mean ± SEM) of the corresponding saline-treated animal; ****P* < 0.001, two-way ANOVA with Sidak's post hoc test.

interactions between A_{2A}R and D₂R agonists and antagonists that occur in the A_{2A}R–D₂R heteromer, where any orthosteric A_{2A}R ligand, agonist or antagonist, exerts a negative allosteric modulation on the affinity of any orthosteric D₂R ligand, agonist or antagonist [41]. Those allosteric interactions provided a plausible mechanism for the reported increase of [¹¹C]raclopride binding in humans induced by caffeine [42], due to its ability to antagonize the effect of endogenous adenosine on the binding of the exogenous D₂R antagonist [41]. It can thus be predicted that a decrease in A_{2A}R–D₂R heteromerization should lead to a reduced effect of endogenous adenosine on exogenous antagonist binding, which might be mainly responsible for the increased striatal D₂R antagonist binding of schizophrenic patients.

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AUTHOR CONTRIBUTIONS

Substantial contributions to the conception or design of the work (all authors), or the acquisition (MV-L, MMC-M and C.S.), analysis (MV-L, LFC and ML-C), or interpretation of data for the work (LFC and KS). Drafting the work or revising it critically for important intellectual content (all authors). Final approval of the version to be published (all authors). Agree to be accountable for all aspects of the work in ensuring that questions related to the accuracy or integrity of any part of the work are appropriately investigated and resolved (all authors).

ADDITIONAL INFORMATION

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