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# **A<sub>1</sub> receptor deficiency causes increased insulin and glucagon secretion in mice**

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## ABSTRACT

Adenosine influences metabolism and the adenosine receptor antagonist caffeine decreases the risk of type 2 diabetes. In this study the metabolic role of one adenosine receptor subtype, the adenosine A<sub>1</sub>R, was evaluated in mice lacking this receptor [A<sub>1</sub>R (–/–)]. The HbA1c levels and body weight were not significantly different between wild type [A<sub>1</sub>R (+/+)] and A<sub>1</sub>R (–/–) mice (3–4 months) fed normal lab chow. At rest, plasma levels of glucose, insulin and glucagon were similar in both genotypes. Following glucose injection, glucose tolerance was not appreciably altered in A<sub>1</sub>R (–/–) mice. Glucose injection induced sustained increases in plasma insulin and glucagon levels in A<sub>1</sub>R (–/–) mice, whereas A<sub>1</sub>R (+/+) control mice reacted with the expected transient increase in insulin and decrease in glucagon levels. Pancreas perfusion experiments showed that A<sub>1</sub>R (–/–) mice had a slightly higher basal insulin secretion than A<sub>1</sub>R (+/+) mice. The first phase insulin secretion (initiated with 16.7 mM glucose) was of the same magnitude in both genotypes, but the second phase was significantly enhanced in the A<sub>1</sub>R (–/–) pancreata compared with A<sub>1</sub>R (+/+) mice. Insulin- and contraction-mediated glucose uptake in skeletal muscle were not significantly different between in A<sub>1</sub>R (–/–) and A<sub>1</sub>R (+/+) mice. All adenosine receptors were expressed at mRNA level in skeletal muscle in A<sub>1</sub>R (+/+) mice and the mRNA A<sub>2A</sub>R, A<sub>2B</sub>R and A<sub>3</sub>R levels were similar in A<sub>1</sub>R (–/–) and A<sub>1</sub>R (+/+) mice. In conclusion, the A<sub>1</sub>R minimally affects muscle glucose uptake, but is important in regulating pancreatic islet function.

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## 1. Introduction

Adenosine is the endogenous ligand for four pharmacologically well defined G protein-coupled adenosine receptors, the A<sub>1</sub>, A<sub>2A</sub>, A<sub>2B</sub> and A<sub>3</sub> receptors [1]. Caffeine can block all the adenosine receptors, although the affinity of caffeine is much higher for the A<sub>1</sub>, A<sub>2A</sub> and A<sub>2B</sub> receptors than for the A<sub>3</sub> receptor [2]. Several, but not all, epidemiological studies have

concluded that coffee consumption decreases the risk of developing non-insulin-dependent diabetes (type 2 diabetes) [3,4]. Coffee contains several thousand active components including chlorogenic acid and magnesium [2]. Nevertheless it is tempting to speculate that the inverse association between coffee and type 2 diabetes may be due to the effects of caffeine, and hence that adenosine acting on one or more of the caffeine sensitive receptors is important in the regulation of glucose

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homeostasis. It is not certain which receptor is involved and how it transmits its signal although the results of previous studies using pharmacological approaches have suggested that the  $A_1R$  is the adenosine receptor most involved in metabolism [5–8].

Adenosine and adenosine agonists have also been shown to decrease insulin secretion [9–11] and increase glucagon secretion [12–14] in pancreas. Regarding skeletal muscle, some studies suggest that adenosine has positive effects on glucose uptake [15–18], whereas others indicate negative effects [19–22]. It is still unclear which adenosine receptor underlies these changes.

In the present study, we used  $A_1$  knock out [ $A_1R$  (–/–)] mice to examine the role of  $A_1$  receptors in glucose homeostasis. The results indicate that the  $A_1R$  is not critical for regulating muscle glucose uptake, but strongly influences pancreatic islet function.

## 2. Materials and methods

### 2.1. Materials

Midazolam was from Hoffmann-La Roche (Nutley, NJ). Fentanyl was from Janssen Pharmaceuticals (Neuss, Germany). Trasylol was from Bayer (Leverkusen, Germany). 2-Deoxy-D-[1,2- $^3H$ ]glucose (2-DG) and carboxy-[ $^{14}C$ ]inulin were from Amersham Bioscience (Buckinghamshire, UK). Human insulin (Actrapid) was from Novo Nordisk (Bagsvaerd, Denmark). 2-Chloro- $N^6$ -cyclopentyladenosine (CCPA) and tribromoethanol were from Sigma (St. Louis, MO). Scintillation liquid Ultima Gold was from Packard (Meriden, CT). Qiagen RNeasy kit was from Qiagen GmbH (Hilden, Germany). High-Capacity cDNA Archive Kit, primers and probes for real time RT-PCR and TaqMan Universal PCR Master Mix No AmpErase UNG were from Applied Biosystems (Foster City, CA). The kit for determination of insulin was obtained from Diagnostica (Falkenberg, Sweden) and the kit for glucagon determination was from Eurodiagnostica (Malmö, Sweden). Glucose reagent strips were from Medisense, Baxter Travenol (Deerfield, IL). Hemoglobin A1c (HbA1c) kit was from Roche Diagnostics GmbH (Mannheim, Germany). All other reagents and enzymes were from Sigma (St. Louis, MO), Boehringer Mannheim, GmbH (Mannheim, Germany) or Merck (Darmstadt, Germany).

### 2.2. Animals

Wild type  $A_1R$  (+/+) and knock out  $A_1R$  (–/–) mice with a C57BL/6 background and wild type mice (NMRI strain) were used. Male mice were used in all experiments except in the real-time RT-PCR experiments. In these experiments, both male and female mice were used. The  $A_1R$  (–/–) mice were generated as previously described [23] and back-crossed to a C57BL/6 congenic strain by Jackson Laboratory (Bar Harbor, ME) according to their general procedures for back-crossing, until the mice were determined to be congenic by 140 genomic markers. PCR-based genotyping was used to identify  $A_1R$  (–/–) offspring [24]. The mice (3–4 months old) were housed at a constant temperature (22–23 °C) and 12-h light/dark cycles with free access to standard pellet food and tap water ad libitum. All

the experimental protocols were evaluated and approved by the local ethical committees in Stockholm, Uppsala or Malmö/Lund, Sweden.

### 2.3. In vivo glucose challenges

For the in vivo studies, glucose (1 g/kg body weight) was dissolved in 0.9% NaCl and delivered to freely fed mice by an intraperitoneal (i.p.) injection. Blood samples were taken by the retrobulbar approach as previously described [25]. In other experiments in freely fed mice, glucose (3 g/kg body weight) was delivered by intravenous injection (i.v.) into the tail vein and thereafter blood samples were taken at different time-points from the cut tip of the tail [26].

### 2.4. In situ pancreatic perfusion

The pancreatic perfusion experiments were performed as described previously [27]. Briefly, the mice were anesthetized with midazolam (0.4 mg/25 g body weight) and fentanyl (0.02 mg/25 g body weight) and kept on a heating pad during the entire experiment. The abdominal cavity was opened and arteries (renal, hepatic and splenic) were ligated and the aorta was tied off above the level of pancreatic artery. The pancreas was perfused with Krebs-Ringer HEPES buffer (1 ml/min) supplemented with glucose (3.3 mM) and 0.20% bovine serum albumin via a silicone catheter placed in aorta. After 10 min of infusion, the medium glucose concentration was changed to 16.7 mM. The perfusate was collected via a silicone catheter from the portal vein in tubes at different time points for further measurements of insulin and glucose levels.

### 2.5. Muscle stimulation

Skeletal muscles from  $A_1R$  (+/+) and  $A_1R$  (–/–) mice were used in all experiments except those with the  $A_1$  agonist, CCPA, where muscles from NMRI mice were used. The mice were sacrificed and both extensor digitorum longus (EDL; glycolytic) and soleus (oxidative) muscles were rapidly removed. The tendons of the muscles were tied with nylon thread to stainless steel hooks and each muscle was transferred to a stimulation chamber (World Precision Instruments), which contained a Tyrode solution consisting of (in mM): NaCl, 121; KCl, 5;  $CaCl_2$ , 1.8;  $NaH_2PO_4$ , 0.4;  $MgCl_2$ , 0.5;  $NaHCO_3$ , 24; EDTA, 0.1; glucose, 5.5; 0.1% fetal calf serum, gassed continuously with 5%  $CO_2$ /95%  $O_2$ , yielding a pH of 7.4 [28]. The temperature was set to 25 °C. In the chamber, the muscles were mounted between a force transducer and an adjustable holder. The muscles were stretched to the length where maximum tetanic force was obtained. After a 25 min recovery period, the muscles were stimulated with current pulses (0.5 ms duration; ~150% of the voltage required for maximum force response) via plate electrodes lying parallel to the fibers. The muscles were stimulated at 50 Hz (tetanic duration 100 ms, 2 trains/s) for 10 min and immediately transferred to vials for glucose uptake measurements. In experiments where insulin-mediated glucose uptake was measured, insulin was added to the vials after a 30-min preincubation period. When CCPA (100 nM) was used, it was included in the Tyrode solution prior to addition of the muscles.

## 2.6. 2-Deoxy-D-[1,2-<sup>3</sup>H]glucose uptake

The 2-Deoxy-D-[1,2-<sup>3</sup>H]glucose (2-DG) uptake was measured both in contraction- and insulin-stimulated muscles as described elsewhere [28,29]. Briefly, the muscles were incubated in continuously gassed (as above) vials containing 1.5 ml Tyrode solution without glucose but supplemented with 2 mM pyruvate in a shaking water bath (100 oscillations/min) at 35 °C for a total of 80 (insulin, 2 or 20 mU/ml) or 40 min (contraction). 2-DG (final concentration 1 mM; 1 mCi/mmol) and carboxy-[<sup>14</sup>C]inulin (0.2 µCi/ml medium, for assessment of extracellular space) were present during the last 20 min. Thereafter, muscles were blotted, frozen and added to tubes containing 0.5 ml of 1N NaOH. The muscles were weighed, digested by heating at 70 °C for 15 min, cooled, centrifuged (23,000 × *g* for 5 min) and duplicate 200 µl aliquots of the supernatant were added to 4 ml scintillation liquid (Ultima Gold) and counted for <sup>3</sup>H and <sup>14</sup>C with a scintillation counter.

## 2.7. Real time reverse transcription polymerase chain reaction (RT-PCR)

The muscles were dissected out and rapidly frozen. They were homogenized in a lysis buffer (Qiagen) and RNA was isolated from the muscles with Qiagen RNeasy kit according to the manufacturer's protocol (Qiagen GmbH). The cDNA synthesis was carried out with a High-Capacity cDNA Archive Kit with random primers and multiscribe reverse transcriptase enzyme according to the manufacturer's instructions (Applied Biosystems). The Applied Biosystem 2720 Terminal cycler was used for the reverse transcription and the incubation conditions were set to 25 °C for 10 min followed by 37 °C for 2 h. Detection of A<sub>1</sub>R, A<sub>2A</sub>R, A<sub>2B</sub>R and A<sub>3</sub>R mRNA was performed by using real time RT-PCR [30]. The real time RT-PCR reactions were run in an ABI Prism 7500 Sequence Detector System (Applied Biosystems, Foster City, CA). Each run consisted of 50 °C for 2 min and 95 °C for 10 min, followed by 40 cycles of 95 °C for 15 s and 60 °C for 1 min. Primers (900 nM of each primer per reaction) and probes (200 nM per reaction) used for the adenosine receptors are described elsewhere [31]. All the reactions were performed in triplicate. TaqMan Universal PCR No AmpErase UNG master mix was used in all reactions. Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) and β-actin were used as endogenous controls. The difference in the number of cycles needed to reach the detection threshold (ct = cycle at threshold) using GAPDH and β-actin as reference ( $\Delta\text{ct} = \text{ct}_{\text{adenosine receptor}} - \text{ct}_{\text{endogenous control}}$ ) was calculated.  $\Delta\Delta\text{ct}$  was calculated as ( $\Delta\Delta\text{ct} = (\Delta\text{ct}_{\text{adenosine receptor}})_{\text{sample}} - (\Delta\text{ct}_{\text{adenosine receptor}})_{\text{calibrator}}$  (A<sub>1</sub>R (+/+) mouse)) in the experiments where the mRNA expression between A<sub>1</sub>R (+/+) and A<sub>1</sub>R (–/–) mice was compared. The  $\Delta\Delta\text{ct}$ -value was finally expressed as relative fold change in gene expression (mean ± 95% confidence interval) by using the  $2^{-(\Delta\Delta\text{ct})}$  formula.

## 2.8. Body weight and hemoglobin A1c (HbA1c)

The body weights of the mice were measured before experiments were performed. For the HbA1c determination, the mice were decapitated and blood samples were collected from the site of decapitation and heparinized.

## 2.9. Analyses of blood samples

Plasma glucose (i.p. glucose tolerance test) was determined enzymatically and insulin and glucagon with a radioimmunoassay (RIA) as described previously [32–34]. Blood glucose concentrations (i.v. glucose tolerance test) were measured with glucose reagent strips. HbA1c levels were determined with a kit according to the manufacturer's instructions (Roche Diagnostics GmbH).

## 2.10. Statistics

Levels of significance ( $p < 0.05$ ) between means were assessed using Student's *t*-test for unpaired data or an ANOVA followed by Tukey–Kramer's multiple comparisons test where appropriate. Values are presented as mean ± S.E.M. unless otherwise stated.

# 3. Results

## 3.1. Body weight and HbA1c levels

Body weight and HbA1c values were not significantly different between A<sub>1</sub>R (+/+) and A<sub>1</sub>R (–/–) mice (Table 1). The HbA1c data thus do not indicate an abnormal blood glucose status over a longer period of time in the A<sub>1</sub>R (–/–) mice. The A<sub>1</sub>R (–/–) mice fed normal lab chow had a normal body weight despite the fact that a major antilipolytic factor (the A<sub>1</sub>R) has been eliminated [5,35,36]. The weight of the abdominal adipose tissue was also not significantly different between genotypes (data not shown).

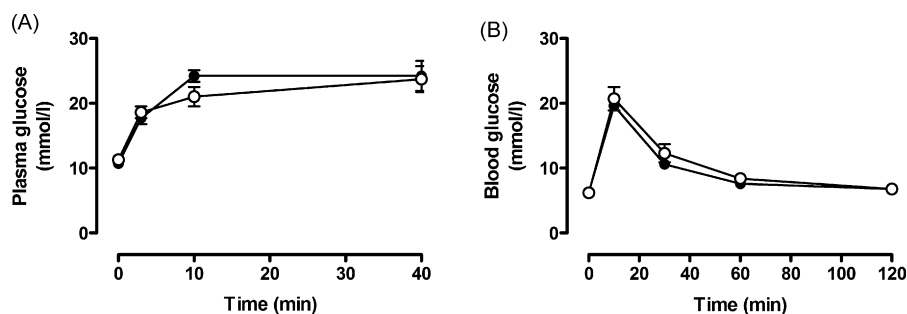
## 3.2. In vivo glucose tolerance and pancreatic hormone response after glucose challenge

In vivo glucose tolerance was investigated by a glucose challenge in freely fed mice. There were no significant differences in blood/plasma glucose concentrations in the basal state. The changes in the glucose concentrations measured at different time-points after glucose injection in A<sub>1</sub>R (–/–) mice (i.p., Fig. 1A or i.v., Fig. 1B) were not appreciably different from those measured in A<sub>1</sub>R (+/+) animals. Thus glucose tolerance was not significantly altered in A<sub>1</sub>R (–/–) mice.

The pancreatic hormone responses were measured when glucose was administered i.p. to freely fed mice. The basal plasma insulin levels did not differ between the A<sub>1</sub>R (+/+) and A<sub>1</sub>R (–/–) mice. In A<sub>1</sub>R (+/+) mice, glucose injection resulted in a rapid increase of plasma insulin concentration. However, by 10 min after the glucose challenge the insulin levels had

**Table 1 – Body weight and HbA1c levels in A<sub>1</sub>R (+/+) and A<sub>1</sub>R (–/–) mice. Values are mean ± S.E.M. for 8–11 mice of each genotype**

Genotype	A <sub>1</sub> R (+/+)	A <sub>1</sub> R (–/–)
Body weight (g)	30.9 ± 0.2	31.3 ± 1.5
HbA1c (%)	2.7 ± 0.1	2.7 ± 0.1



**Fig. 1 – Circulating glucose levels in A<sub>1</sub>R (+/+) (●) and A<sub>1</sub>R (-/-) (○) freely fed mice at different time points after i.p. glucose injection (1 g/kg body weight) (A) or i.v. glucose injection (3 g/kg body weight) (B). Values are mean ± S.E.M., n = 8–10 mice of each genotype.**

returned to basal values. In A<sub>1</sub>R (-/-) mice, the glucose-induced increase in plasma insulin was maintained until the end of the 40 min period of measurement (Fig. 2A). The basal glucagon levels were also similar in A<sub>1</sub>R (+/+) and A<sub>1</sub>R (-/-) mice. In A<sub>1</sub>R (+/+) mice, as expected, the glucose load suppressed plasma glucagon levels, but paradoxically glucagon levels increased continuously in A<sub>1</sub>R (-/-) mice (Fig. 2B).

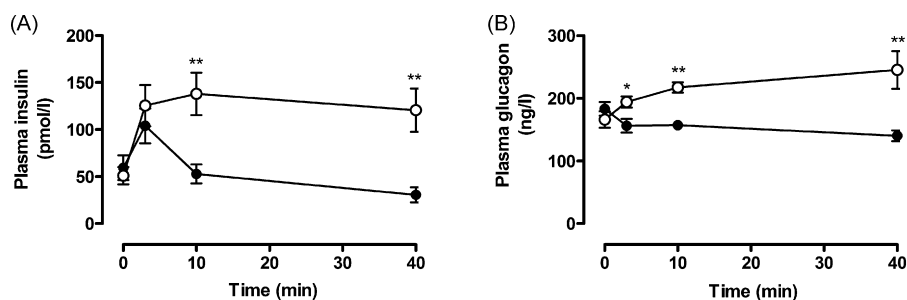
### 3.3. Phasic insulin release from adenosine A<sub>1</sub>R (-/-) mice measured by *in situ* pancreatic perfusion

To assess the role of A<sub>1</sub>R in the dynamics of insulin release, we performed *in situ* pancreatic perfusions with fractionated sampling. Fig. 3 illustrates that basal insulin release (first 10 min with 3.3 mM glucose) from perfused pancreas of A<sub>1</sub>R (-/-) mice was slightly higher compared to A<sub>1</sub>R (+/+) controls. Glucose stimulated insulin release showed a typical biphasic pattern when the glucose concentration in perfusate was raised from 3.3 to 16.7 mM in the A<sub>1</sub>R (+/+) mice. Thus after the glucose concentration increased to 16.7 mM (Fig. 3B), first-phase insulin release was initiated and it lasted for 2–3 min. When the same experiment was repeated in adenosine A<sub>1</sub>R (-/-) pancreata (Fig. 3), the peak in first-phase insulin secretion was of the same magnitude as that seen in A<sub>1</sub>R (+/+) mice (n = 4). However, ablation of A<sub>1</sub>R brought about a significant marked increase in second-phase insulin secretion representing a 100% increase compared to the A<sub>1</sub>R (+/+) mice (P < 0.005). There were no differences in the response to a high potassium pulse (results not shown).

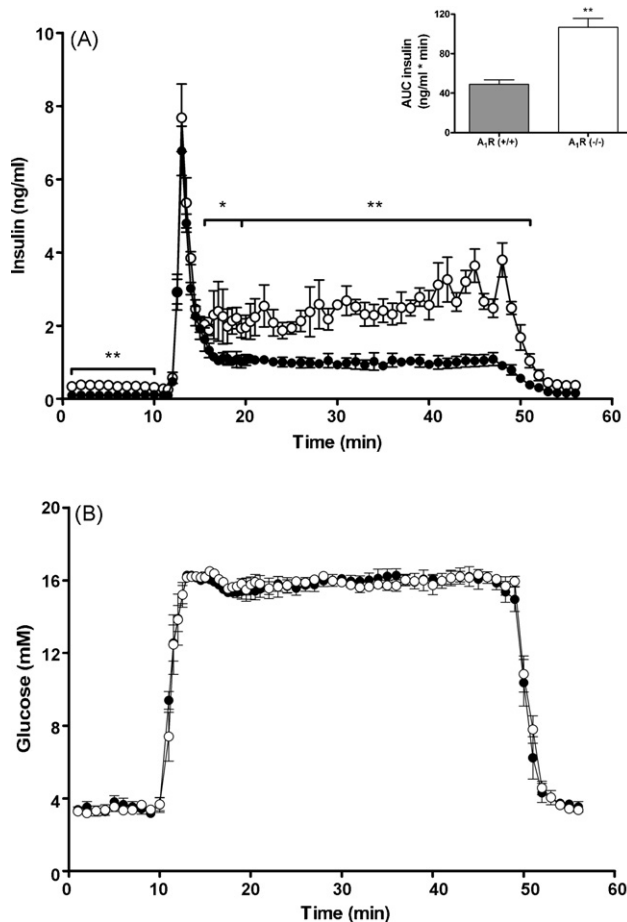
### 3.4. Adenosine receptor expression and glucose uptake in muscle tissue

The expression of the adenosine receptors in skeletal muscles was measured. All adenosine receptors were present at mRNA level in both EDL and soleus muscles in A<sub>1</sub>R (+/+) mice. The  $\Delta$ ct values for the different receptors in the A<sub>1</sub>R (+/+) mouse were: 11.2 (A<sub>1</sub>R), 9.1 (A<sub>2A</sub>R), 10.9 (A<sub>2B</sub>R) and 14.9 (A<sub>3</sub>R) in soleus and 11.6 (A<sub>1</sub>R), 9.2 (A<sub>2A</sub>R), 10.6 (A<sub>2B</sub>R) and 13.4 (A<sub>3</sub>R) in EDL. The A<sub>1</sub>R was not present in A<sub>1</sub>R (-/-) mice (ct value—not detectable). When the mRNA levels of the adenosine receptors that were present in both the A<sub>1</sub>R (+/+) and A<sub>1</sub>R (-/-) mice were compared, no significant differences in mRNA levels of the A<sub>2A</sub>R, A<sub>2B</sub>R and A<sub>3</sub>R in EDL and soleus between the A<sub>1</sub>R (+/+) and A<sub>1</sub>R (-/-) mice were found (Fig. 4). These results are in agreement with previous studies, where it was shown that our A<sub>1</sub>R (-/-) mice did not have an altered expression of the other adenosine receptors in heart tissue [38].

The observation that the glucose tolerance test (GTT) elicited a normal blood glucose response, but an increased plasma insulin response in A<sub>1</sub>R (-/-) mice suggested a peripheral insulin resistance in these mice. Therefore, glucose uptake measurements were performed on isolated muscles. Glucose uptake in soleus and EDL muscles was similar in A<sub>1</sub>R (-/-) and A<sub>1</sub>R (+/+) mice in the basal state and after exposure to insulin (2 or 20 mU/ml) (Fig. 5). There was also no difference in contraction-mediated glucose uptake between groups (Fig. 5). The effect of the A<sub>1</sub>R agonist CCPA (100 nM) on glucose uptake in the presence of saturating insulin was also studied in wild type (NMRI) mice. CCPA did not significantly



**Fig. 2 – Plasma insulin (A) and glucagon (B) response to i.p. glucose injection (1 g/kg body weight). Values are mean ± S.E.M. for A<sub>1</sub>R (+/+) (●) and A<sub>1</sub>R (-/-) (○) mice. n = 8–10 mice of each genotype. \*p < 0.05, \*\*p < 0.01 between groups.**

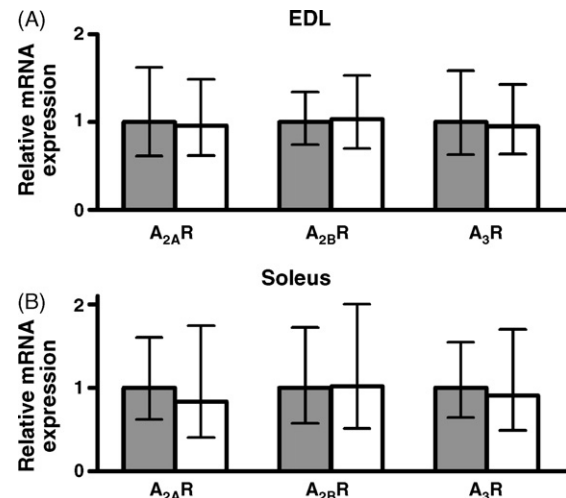


**Fig. 3 – Dynamics of insulin release from perfused pancreata in  $A_{1R} (+/+)$  and  $A_{1R} (-/-)$  mice. (A)** Insulin release measured in adenosine  $A_{1R} (+/+)$  and  $A_{1R} (-/-)$  pancreata before and after the glucose concentration in the perfusate was increased from 3.3 to 16.7 mM. Samples were taken at 60-s intervals, except during the first 10 min after increasing the glucose concentration ( $t = 11$ –21 min, when the sample interval was 30 s). Values are mean  $\pm$  S.E.M. for  $A_{1R} (+/+)$  ( $\bullet$ ) and  $A_{1R} (-/-)$  ( $\circ$ ) mice.  $n = 4$  mice of each genotype. The insert shows AUC of the data. **(B)** An illustration of glucose levels in the same samples shown in A. Statistical significance is provided for comparisons between  $A_{1R} (+/+)$  and  $A_{1R} (-/-)$  mice are denoted by asterisks (\*)  $p < 0.05$ , \*\* $p < 0.01$ .

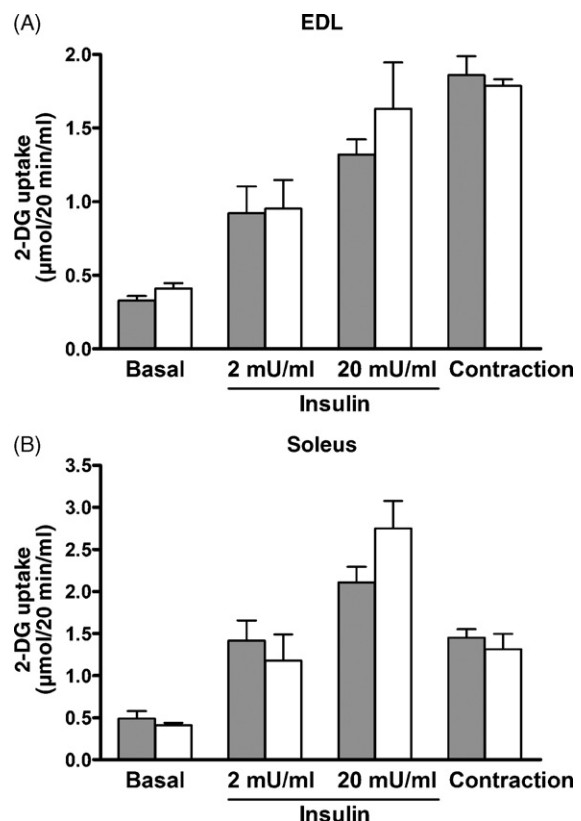
affect insulin-mediated 2-DG uptake (see [Supplemental material, Fig. S1](#)). Nor did CCPA have a significant effect when the muscles were stimulated with intermediate concentrations of insulin (0.2 and 0.6 mU/ml) (data not shown).

#### 4. Discussion

The major findings of the present study are: (1) an essentially normal glucose homeostasis and (2) altered pancreatic hormone responses to a glucose challenge in  $A_{1R} (-/-)$  mice fed normal lab chow. That glucose levels were normal despite elevated plasma insulin responses to the glucose challenge in



**Fig. 4 – Relative mRNA levels of the adenosine  $A_1$ ,  $A_{2A}$ ,  $A_{2B}$  and  $A_3$  receptors in EDL (A) and soleus (B) muscles in  $A_{1R} (+/+)$  ( $\blacksquare$ ) mice compared to  $A_{1R} (-/-)$  ( $\square$ ) mice. Values are means (mean values in  $A_{1R} (+/+)$  mice are set to 1) and the error bars represent 95% confidence intervals,  $n = 3$ –8 mice in each group.**



**Fig. 5 – 2-DG uptake in EDL and soleus muscles. Values are mean  $\pm$  S.E.M.,  $n = 6$  mice of each genotype. Data were obtained in the basal state, after exposure to insulin or after repeated contractions in  $A_{1R} (+/+)$  ( $\blacksquare$ ) and  $A_{1R} (-/-)$  ( $\square$ ) mice.**



the  $A_1R$  ( $-/-$ ) mice suggested the presence of peripheral insulin resistance. Since skeletal muscle is quantitatively the most important organ for glucose disposal during hyperinsulinemia [39], glucose uptake experiments were performed on isolated skeletal muscles. However, the  $A_1R$  agonist, CCPA, did not affect insulin-mediated glucose uptake. This is in apparent conflict with several previous studies on rat muscle [16,17], and could perhaps be due to high endogenous levels of adenosine. However, this possibility is less likely, as there were no significant differences between  $A_1R$  ( $-/-$ ) and  $A_1R$  ( $+/+$ ) mice in terms of basal or insulin-mediated glucose uptake in either EDL or soleus muscles. Nor were there any significant differences between the genotypes in contraction-mediated glucose uptake. Thus both genetic and pharmacological approaches yielded consistent results to indicate that  $A_1Rs$  do not play a significant role in the control of muscle glucose uptake under several physiological conditions (basal, exercise and insulin) in muscle from young fed mice.

Pharmacological studies indicate that adenosine receptors are present in skeletal muscles [15,16,18,37], although the expression of the different adenosine receptors in skeletal muscle (EDL and soleus) in mice at mRNA level has not been clarified. Therefore the expression of the adenosine receptors was measured and we verified that mRNA for all adenosine receptors were present in EDL and soleus muscles of mice. This suggests that the receptor proteins are also present. We do not know, however, if all receptors are present in the same type of cell.

If the muscles are not insulin resistant, then the higher plasma insulin levels in the  $A_1R$  ( $-/-$ ) mice during the GTT should be associated with higher rates of peripheral (i.e. muscle) glucose utilization. This should result in lower blood glucose concentrations during the GTT. One possible reason why this was not observed could be a compensatory release of glucose from the liver. There are two potential mechanisms whereby this could have occurred: the elevated glucagon levels in the  $A_1R$  ( $-/-$ ) mice stimulate glycogenolysis in the liver; or insulin-mediated inhibition of hepatic gluconeogenesis [40] is defective in the  $A_1R$  ( $-/-$ ) mice. These explanations are not mutually exclusive and thus both mechanisms may be involved. The net effect would be a maintained blood glucose level in the face of a higher glucose turnover in the  $A_1R$  ( $-/-$ ) mice.

These data do not rule out the possibility that  $A_1Rs$  play a significant role in glucose transport in tissues other than skeletal muscle. It has for example been shown that adenosine stimulates and adenosine deaminase inhibits glucose uptake in adipose tissue [41,42] and heart [43,44].

The increased glucose-stimulated insulin release and the paradoxical glucagon response after glucose injection in the  $A_1R$  ( $-/-$ ) mice suggested a role for the  $A_1R$  in pancreatic islet function. Therefore we examined insulin secretion in the perfused pancreas. Previous studies have shown that adenosine and its analogues can affect insulin secretion [45]. For example, it has been shown that the adenosine analogue phenylisopropyladenosine (PIA) can inhibit glucose-stimulated insulin secretion [10,46] and the  $A_1R$  has been postulated to be the adenosine receptor involved [11,46]. One hypothesis that has been proposed to explain the adenosine-mediated inhibition of insulin secretion is that the  $A_1R$  is coupled to the

inhibitory  $G_i$  protein and a stimulation of  $A_1R$  thereby will inhibit adenylyl cyclase, which will result in lower cellular cAMP levels. cAMP stimulates the release of insulin [47]. Thus deletion of the  $A_1R$  would be expected to result in an increased insulin secretion. Recently, however, Rüsing et al. [48] showed that another second messenger system must also be involved in the adenosine-mediated inhibition of insulin secretion since their results showed that agonists to both  $G_i$ -coupled adenosine receptors ( $A_1R$  and  $A_3R$ ) and  $G_s$ -coupled adenosine receptor ( $A_{2A}R$ ) were able to decrease the plasma insulin levels in rats and reduce insulin secretion in INS-1 cells. Antagonists to the  $G_s$ -coupled  $A_{2B}R$  were also able to counteract the inhibitory effect of the adenosine analogue, 5'-N-ethylcarboxamidoadenosine (NECA). Our current findings are consistent with the idea that the  $A_1R$  is involved in insulin secretion. The function of the other adenosine receptors and the intracellular signal transduction pathways behind the involvement of the adenosine receptors in insulin secretion await further evaluation.

Glucose-induced insulin release follows a biphasic time course. A transient first phase is followed by a more or less pronounced second phase depending on the amount of glucose given. Our present results both in vivo and in situ show that the  $A_1R$  is involved in the second phase, but not in the first phase of insulin secretion. The mechanisms behind the two phases of release are still unclear but it has been suggested that the rapid first phase reflects the release of a limited pool of readily releasable granules in close proximity to the L-type  $Ca^{2+}$  channels in the plasma membrane, while the second phase is regulated by replenishment of this pool by granules originating from a "reserve pool" situated at a greater distance from the channels [49]. Hence the  $A_1Rs$  seem to influence the mobilization of the granules from this "reserve pool".

Glucagon is known as an amplifier of insulin secretion [50]. It is thus possible that the elevated glucagon levels after the initial phase of the GTT (i.e. >5 min) enhanced insulin secretion by the  $\beta$  cells. High cAMP levels in the  $\alpha$  cells can increase the secretion of glucagon and therefore we also consider a role for the  $A_1R$  in glucagon release. Previous studies [12,13] have shown that adenosine can stimulate glucagon release, but this effect is probably mediated by  $A_2$  receptors (unknown whether  $A_{2A}$  or  $A_{2B}$ ) that couple to stimulatory G proteins [14]. If  $A_1Rs$  are present in  $\alpha$  cells, stimulation of these receptors could be expected to inhibit glucagon release. Conversely, if the  $A_1R$  is lacking, glucagon release would be expected to be stimulated.

These studies thus suggest that although the basal plasma glucose and insulin levels are essentially normal in the  $A_1R$  ( $-/-$ ) mice, there are major perturbations that could influence the animal's ability to cope with metabolic stress. Future studies with animals that are older than our mice (>3–4 months) or stressed with a high fat diet would be a good complement to this study. There is also good reason to perform studies under fasting conditions. Even an overnight fast represents a major metabolic stress in mice and there is evidence (H. Edlund, Umeå University, personal communication) that 12 h fasting leads to lower basal glucose levels and improved glucose tolerance in  $A_1R$  ( $-/-$ ) mice. Furthermore, investigation of the mechanisms behind the pancreatic

hormone response in the A<sub>1</sub>R (–/–) mice would also give more information about the role of the A<sub>1</sub>R in metabolism. These studies should also be of value to assess the feasibility of using a drug acting on the A<sub>1</sub>R in treatment of metabolic disorders such as type 2 diabetes mellitus.

In conclusion, this study suggests that A<sub>1</sub>Rs are important in regulating pancreatic islet function, but have less influence on muscle glucose transport. Furthermore, our results raise the question whether altered function of A<sub>1</sub>Rs in the pancreas is involved in the pathogenesis of type 2 diabetes.

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## Appendix A. Supplementary data

Supplementary data associated with this article can be found, in the online version, at [doi:10.1016/j.bcp.2007.08.006](https://doi.org/10.1016/j.bcp.2007.08.006).

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