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Key amino acids for differential coupling of α 1-adrenergic receptor subtypes to Gs[☆]

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Abstract

We have established that differing effects of α 1-adrenergic receptor (AR) subtypes on cell proliferation are due to differential coupling to the Gs/cAMP pathway; thus, both α 1A- and α 1B-ARs couple to Gs, while α 1D-AR does not. To identify the region responsible for this difference in subtype-specific Gs coupling, we constructed a series of chimeric and a set of point-mutated human α 1A- and α 1D-ARs, and examined their signaling ability. Here, we show that the amino acid residues Thr 136 and Val138 in the intracellular loop II of the human α 1A-AR are intimately involved with Gs coupling.

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α 1-adrenergic receptors (ARs) are members of the G protein-coupled receptor superfamily, which mediate the effects of the sympathetic nervous system, including smooth muscle contraction, cell proliferation and hypertrophy, and hepatic glucose metabolism [1–6]. The α 1-ARs mediate their physiological effects by activating phospholipase C, resulting in production of inositol 1,4,5-triphosphate, mobilization of intracellular calcium and diacylglycerol, and finally activation of protein kinase C [7,8]. More recently, it has been demonstrated that the α 1-ARs can also influence a variety of other effectors, such as the mitogen-activated protein kinase (MAPK) pathway, cAMP metabolism, and activation of

phospholipases D and A2 in different cells [9–14]. α 1-ARs comprise a heterogeneous family and molecular cloning studies have identified three distinct cDNAs encoding α 1-AR subtypes (α 1a, α 1b, and α 1d) [15–18]. Although the pharmacological properties of three distinct α 1-AR subtypes (α 1A-AR, α 1B-AR, and α 1D-AR) are well documented, little is known regarding differences in their physiological roles and signaling pathways.

We have previously reported that α 1B-AR couples not only to the Gq/Ca^{2+} signaling pathway, but also the Gs/cAMP pathway via direct interaction with Gs [19]. Furthermore, we have recently observed that the activation of α 1A- or α 1B-AR inhibits serum-promoted cell proliferation, whereas the activation of α 1D-AR has a potent growth-promoting effect. This subtype-dependent anti-proliferative effect might be caused by cAMP-dependent inhibition of serum-promoted down-regulation of $\text{p}27^{\text{kip1}}$ (submitted for publication).

In the present study, we have constructed a series of human chimeric α 1A and α 1D-ARs (α 1A/ α 1D-CRs) to identify the region responsible for the difference between α 1A-AR and α 1D-AR in serum-stimulated cell proliferation and signaling pathways.

[☆] Abbreviations: AR, α 1-adrenergic receptor; CHO, Chinese hamster ovary; CHO α 1A, CHO cells stably expressing cloned α 1A-AR; CHO α 1B, CHO cells stably expressing cloned α 1B-AR; CHO α 1D, CHO cells stably expressing cloned α 1D-AR; NA, noradrenaline; α 1A/ α 1D-CR, chimeric α 1A and α 1D-AR; CHO α 1A/ α 1D-CR, CHO cells stably expressing α 1A/ α 1D-CR; HEK, human embryonic kidney; TMD, transmembrane domain; ICL, intracellular loop; MAPK, mitogen-activated protein kinase; FCS, fetal calf serum.

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Materials and methods

Materials. The following drugs and reagents were used: 2-[2-(4-hydroxy-3- 125 I]iodo-phenyl)ethylaminomethyl]- α -tetralone (125 I]HEAT) (specific activity, 2200 Ci mmol $^{-1}$; New England Nuclear, Boston, MA, USA); [methyl- 3 H]thymidine (specific activity, 2 Ci mmol $^{-1}$; New England Nuclear, Boston, MA, USA); phentolamine mesylate (Novartis Pharmaceutical, Summit, NJ, USA); (–)-noradrenaline bitartrate, phenylephrine, DL-propranolol hydrochloride, and 3-isobutyl-1-methylxanthine (IBMX) (Sigma Chemical, St. Louis, MO, USA); and Ham's F-12 medium, G418 (Gibco Life Technologies, Gaithersburg, MD, USA). All other chemicals were of reagent grade.

Construction of human $\alpha 1A/\alpha 1D$ -CRs and amino acid-substituted $\alpha 1A$ -AR mutants. The cDNAs encoding human $\alpha 1A/\alpha 1D$ -CRs were constructed by digestion at common restriction sites (the *Sna*BI and *Afl*II sites created by oligonucleotide-directed mutagenesis within the wild-type human $\alpha 1A$ - and $\alpha 1D$ -AR cDNAs, Fig. 1) and then exchanging the corresponding regions of each. The restriction sites and their positions in the deduced amino acid sequence of the $\alpha 1A/\alpha 1D$ -ARs, respectively, are as follows: *Sna*BI, at Y208/Y278 in the transmembrane domain (TMD) V; *Afl*II, at Leu262/Leu338 in the intracellular loop (ICL) III (Fig. 1). The structures of the chimeras are shown in Fig. 2. The positions of the junctions for individual human $\alpha 1A/\alpha 1D$ -CRs and their component amino acids are as follows: AAD, $\alpha 1A$ 1–262/ $\alpha 1D$ 339–569; ADA, $\alpha 1A$ 1–208/ $\alpha 1D$ 279–338/ $\alpha 1A$ 263–438; ADD, $\alpha 1A$ 1–208/ $\alpha 1D$ 279–569; DDA, $\alpha 1D$ 1–338/ $\alpha 1A$ 263–438; DAD, $\alpha 1D$ 1–278/ $\alpha 1A$ 209–262/ $\alpha 1D$ 339–569; DAA, $\alpha 1D$ 1–278/ $\alpha 1A$ 209–438. Amino acid-substituted mutants of $\alpha 1A$ -AR were constructed by PCR using site-directed mutagenesis. Comparing amino acid sequences of human $\alpha 1A$ -AR, $\alpha 1B$ -AR, and $\alpha 1D$ -AR showed that homology was highly conserved. In the region extending from the amino-terminal extracellular tail to TMD V of the human $\alpha 1D$ -AR, we found 31 amino acids that were different from those common to $\alpha 1A$ -AR and $\alpha 1B$ -AR (shaded amino acids in Fig. 1). In the ICLII of $\alpha 1D$ -AR, only two amino acids, Ala-206 and Met-208, were replaced by Thr and Val in that region of $\alpha 1A$ -AR and $\alpha 1B$ -AR (Thr-136 and Val-138) (Fig. 1). Either Thr-136 or Val-138 or both in the $\alpha 1A$ -AR was changed to the amino acid corresponding to the $\alpha 1D$ -AR sequence. Mutations generated were confirmed by sequencing. The wild-type human $\alpha 1A$ -AR, $\alpha 1D$ -AR and all of the mutant receptors were inserted into *Eco*RI and *Xho*I sites of the mammalian expression vector pMEK19.

Cell transfection. Cell transfection was performed as described previously [20]. For stable expression of the human $\alpha 1A$ -AR, $\alpha 1D$ -AR and all $\alpha 1A/\alpha 1D$ -CRs, CHO-K1 cells were seeded at a density of 2×10^4 cells into 35-mm tissue culture dishes. On the next day, F-12 medium was removed and 1 ml of serum-free F-12 medium containing 13.8 μ g of LipofectAMINE reagent (Gibco Life Technologies, Gaithersburg, MD, USA) and 9.2 μ g of the recombinant expression plasmid was added to the cells. After 24 h, 1 ml F-12 medium containing 20% fetal calf serum (FCS) was added. The cells were passaged at a low density 72 h later. Single colonies resistant to the antibiotic G418 (600 μ g/ml) were isolated and maintained in F-12 medium with 10% fetal calf serum (FCS) and G418 (200 μ g/ml) for four weeks.

In some experiments, the human $\alpha 1A$ -AR, $\alpha 1D$ -AR and the amino acid-substituted mutants were transiently expressed in HEK293 cells by electroporation (240 V, 975 μ F, Bio-Rad gene Pulser II electroporator, Bio-Rad, Hercules, CA, USA) according to manufacturer's instructions. Cells were harvested 48–72 h after transfection.

Membrane preparations. The transfected cells were collected and homogenized by a Branson sonicator (model SONIFIER 250, Branson, Danbury, CT, USA; setting 5 for 8 s) in 2 ml ice-cold buffer A (250 mM sucrose, 5 mM Tris–HCl, and 1 mM MgCl $_2$, pH 7.4). The mixture was then centrifuged at 1000g at 4°C for 10 min to remove nuclei. The supernatant was centrifuged at 35,000g for 20 min at 4°C and the pellet was resuspended in the binding buffer (50 mM Tris–HCl,

10 mM MgCl $_2$, and 10 mM EGTA, pH 7.4) [20]. The protein concentration was measured using the bicinchoninic acid protein assay kit (Pierce, Rockford, IL, USA).

Radioligand binding assay. Radioligand binding studies were performed as described previously [18,20]. Briefly, membrane aliquots (~ 0.1 – 0.5 μ g protein) were incubated for 60 min at 25°C with 125 I]HEAT, and either with or without a competing drug, in a final volume of 150 μ l of the binding buffer. After dilution with ice-cold buffer, samples were immediately filtered through Whatman GF/C glass fiber filters with a Brandel cell harvester (Model-30, Gaithersburg, MD, USA). Each filter was collected and the radioactivity was measured. Binding assays were performed in duplicate at all times, and specific 125 I]HEAT binding was experimentally determined from the difference between counts in the presence or absence of 10 μ M phentolamine. Data were analyzed by the iterative nonlinear regression program, LIGAND [21].

3 H]Thymidine incorporation. We measured 3 H]thymidine incorporation as previously reported [22]. CHO cells stably expressing the cloned $\alpha 1A$ -AR (CHO $\alpha 1A$), $\alpha 1D$ -AR (CHO $\alpha 1D$), or the $\alpha 1A/\alpha 1D$ -CRs (CHO $\alpha 1A/\alpha 1D$ -CRs), were grown to confluence in 96-well plates and rendered quiescent by serum deprivation for 24 h. These cells were treated with 10 μ M phenylephrine or left untreated; they were also pretreated or not pretreated with 10 μ M phentolamine and then incubated with 10% FCS for 18 h. Next, cells were incubated for the following 12 h with 3 H]thymidine (1 μ Ci/ml). The cells were rinsed twice with ice-cold phosphate-buffered saline and precipitated with cold 5% trichloroacetic acid, then washed with cold ethanol, and solubilized with 0.5 N NaOH. DNA radioactivity was measured by scintillation counting.

Measurement of cAMP production. cAMP production in transiently transfected HEK293 cells was determined as described previously [19]. Briefly, these cells were seeded in 6-well plates at a density of 5×10^5 cells/well and cultured for 12–16 h. The cells were washed twice with phosphate-buffered saline and incubated at 37°C for 30 min in a buffered salt solution (140 mM NaCl, 4 mM KCl, 1 mM MgCl $_2$, 1.25 mM CaCl $_2$, 1 mM NaHPO $_4$, 5 mM HEPES, and 11 mM glucose, pH 7.4) and 1 mM IBMX. Propranolol (1 μ M) was included to block endogenous β_1 -ARs present in the cells [6]. The reaction was started by adding NA (10 μ M) with or without phentolamine (10 μ M). After incubation for 10 min at 37°C, the medium was aspirated and the reaction was stopped with 100 mM HCl. Aliquots were taken and assayed for cAMP by radioimmunoassay (Yamasa cAMP Assay kit, Yamasa Shoyu, Chiba, Japan).

Results and discussion

Chimeric receptors

To determine the region responsible for the $\alpha 1$ -AR subtype-specific differential effects on cell proliferation, we first constructed a series of $\alpha 1A/\alpha 1D$ -CRs. The structures of these chimeras (AAD, ADA, ADD, DDA, DAD, and DAA) are shown in Fig. 2. They consist of three components that belong to either the wild-type human $\alpha 1A$ -AR or $\alpha 1D$ -AR. The first component is from the amino-terminal extracellular tail to TMD V. The second component is the ICL III and the third component is from TMD III to the intracellular carboxy-terminus. Three components of the wild-type human $\alpha 1A$ -AR or $\alpha 1D$ -AR were substituted by one another. Radioligand binding studies using 125 I]HEAT

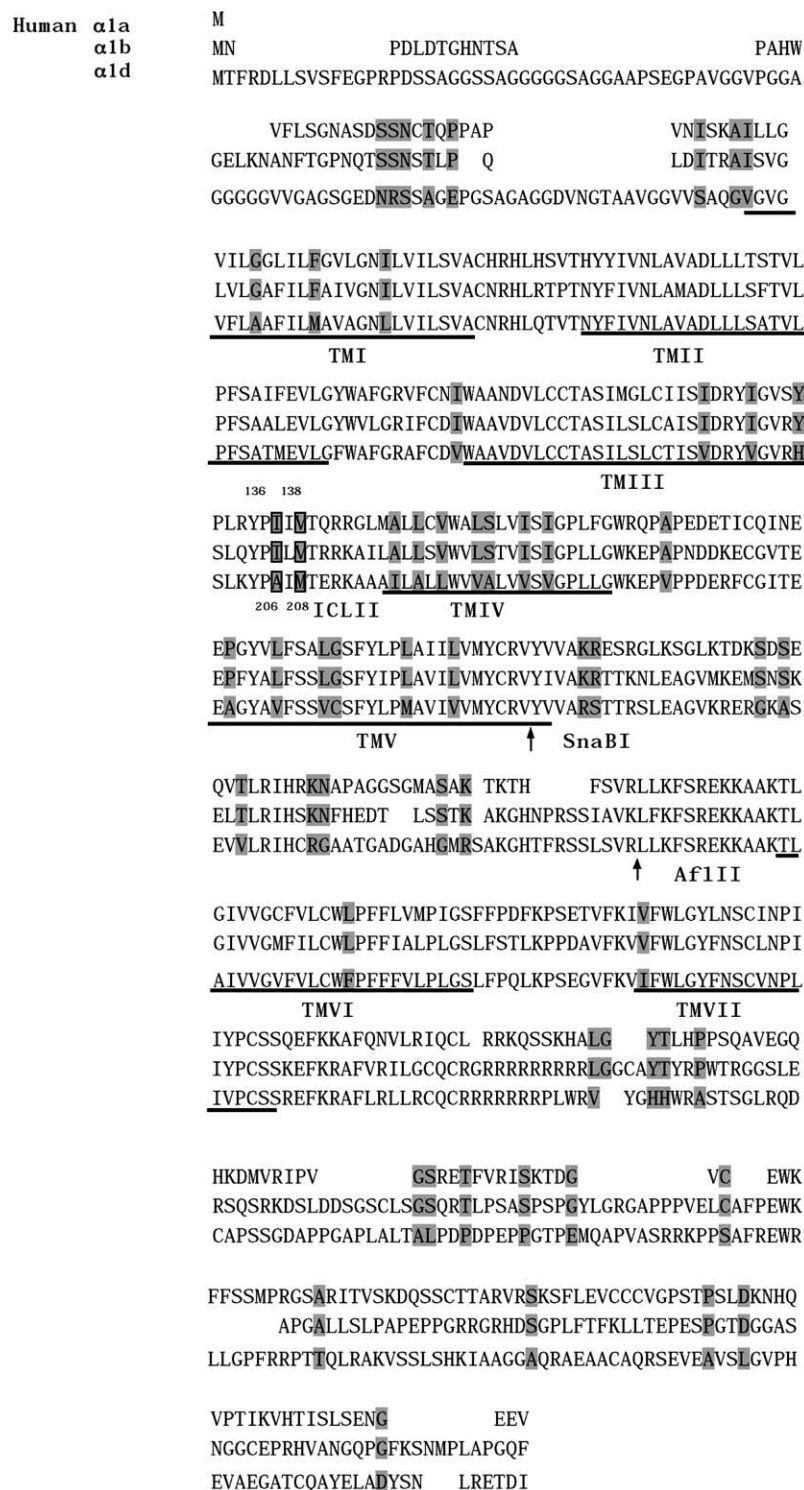


Fig. 1. Alignment of the wild-type human $\alpha 1A$ -AR, $\alpha 1B$ -AR, and $\alpha 1D$ -AR amino acid sequences. Positions of the restriction sites used to construct chimeric receptors are shown by arrowheads with the enzymes (*Sna*BI, *Af*III). The shaded amino acids indicate the observed differences in $\alpha 1D$ -AR, whereas they are identical between the aligned human $\alpha 1A$ -AR and $\alpha 1B$ -AR sequences. The shaded box indicates two amino acids, Thr-136 and Val-138 in ICL II of $\alpha 1A$ -AR, which differ from those of $\alpha 1D$ -AR. These amino acids were changed individually to the corresponding amino acids of the $\alpha 1D$ -AR, Ala-206, and Met-208, respectively, by PCR using site-directed mutagenesis.

confirmed that all the cell lines examined are exhibiting similar receptor expression; CHO $\alpha 1A$ (K_d = 110 pM, B_{max} = 1.3 pmol/mg protein), CHO $\alpha 1D$ (K_d = 300 pM,

B_{max} = 1.1 pmol/mg protein), and CHO $\alpha 1A/\alpha 1D$ -CRs (K_d = 180–350 pM, B_{max} = 0.50–1.3 pmol/mg protein) (Table 1).

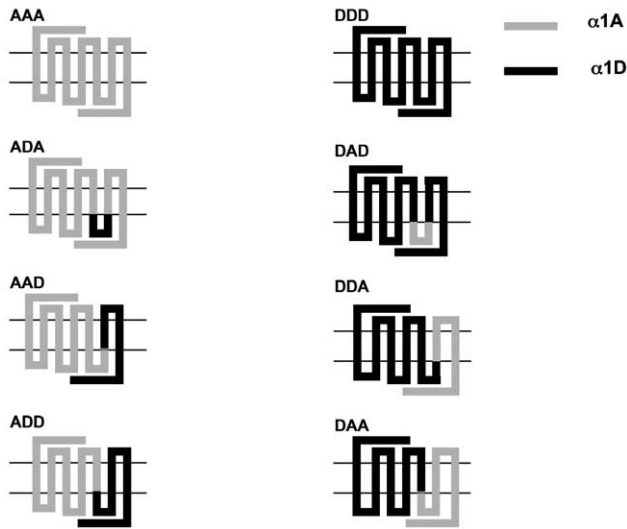


Fig. 2. Structure of chimeric human $\alpha 1A/\alpha 1D$ -ARs ($\alpha 1A/\alpha 1D$ -CRs). Filled black bars, sequences of the wild-type human $\alpha 1A$ -AR; filled blue bars, sequences of the wild-type human $\alpha 1D$ -AR. The position of junctional amino acids in these chimeric receptors and the methods used for their construction are described in Materials and methods.

Table 1
Ligand-binding characteristics of the wild-type $\alpha 1A$, $\alpha 1D$, and chimeric $\alpha 1A$ and $\alpha 1D$ adrenergic receptor

| | B_{\max} (pmol/mg protein) | K_d (nM) |
|-----------------|------------------------------|--------------|
| CHO $\alpha 1A$ | 1.3 ± 0.2 | 110 ± 21 |
| AAD | 0.65 ± 0.1 | 200 ± 10 |
| ADA | 1.3 ± 0.15 | 350 ± 23 |
| ADD | 0.5 ± 0.05 | 180 ± 16 |
| CHO $\alpha 1D$ | 1.1 ± 0.1 | 300 ± 26 |
| DDA | 0.56 ± 0.22 | 200 ± 32 |
| DAD | 0.63 ± 0.18 | 210 ± 18 |
| DAA | 0.84 ± 0.11 | 250 ± 36 |

The specific binding of various concentrations of [125 I]HEAT to membranes of stable clones expressing the wild-type $\alpha 1A$, $\alpha 1D$, and $\alpha 1A/\alpha 1D$ -CRs were measured, and B_{\max} and K_d values were estimated. Data are presented as means \pm SEM from three independent experiments.

Cell proliferation

Proliferation of CHO $\alpha 1A$, CHO $\alpha 1D$, and CHO $\alpha 1A/\alpha 1D$ -CRs was measured by [3 H]thymidine incorporation after serum deprivation. No significant differences were observed between these cells regarding the basal level of [3 H]thymidine incorporation. When they were exposed to 10% serum, in CHO $\alpha 1A$, CHO $\alpha 1D$, and CHO $\alpha 1A/\alpha 1D$ -CRs, a 3.2–5.8-fold increase in [3 H]thymidine incorporation compared to serum-deprived cells was observed (Fig. 3A). Activation of $\alpha 1$ -AR by phenylephrine (10 μ M) inhibited this serum-promoted [3 H]thymidine incorporation to 31–49% in CHO $\alpha 1A$, AAD, ADA, and ADD (Fig. 3A). All of these receptors contained the region stretching from the amino-terminal extracellular tail to TMD V of $\alpha 1A$ -AR. Pretreatment

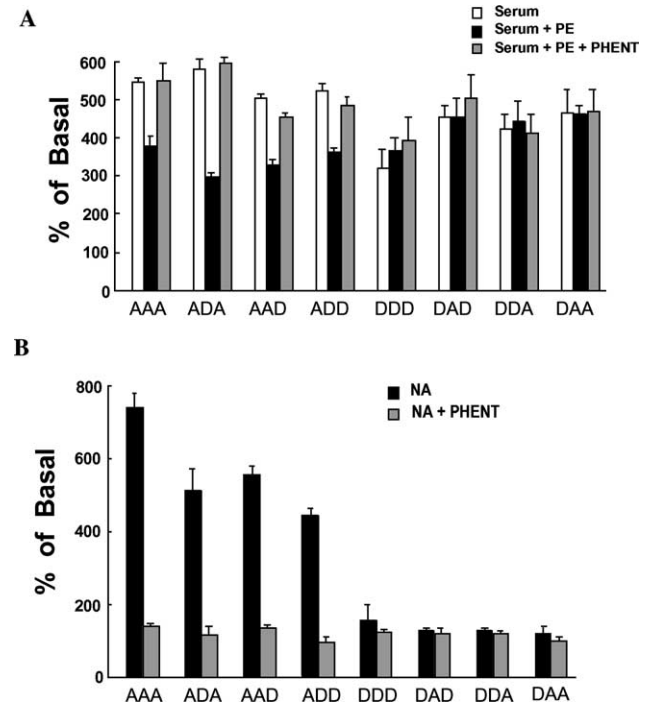


Fig. 3. (A) Effect of phenylephrine on serum-stimulated [3 H]thymidine incorporation. [3 H]Thymidine incorporation was measured in each CHO cell line stably expressing the wild-type $\alpha 1A$ -AR, $\alpha 1D$ -AR or the $\alpha 1A/\alpha 1D$ -CRs. (B) Effect of $\alpha 1$ -AR subtype activation on intracellular cAMP production in each CHO cell line stably expressing the wild-type $\alpha 1A$ -AR, $\alpha 1D$ -AR or the $\alpha 1A/\alpha 1D$ -CRs. In all experiments, phentolamine (10 μ M) treatment was started 30 min prior to stimulation by phenylephrine or noradrenaline. The results are expressed as means \pm SEM of 3–4 different experiments each performed in duplicate. PE, phenylephrine; PHENT, phentolamine; NA, noradrenaline.

with the α -antagonist phentolamine (10 μ M) almost completely blocked the phenylephrine-induced inhibitory effects on these cells, showing that inhibition is mediated through $\alpha 1$ -AR (Fig. 3A). However, in CHO $\alpha 1D$, DDA, DAD, and DAA, which had the region encompassing the amino-terminal extracellular tail to TMD V of $\alpha 1D$ -AR, such an inhibitory effect of phenylephrine was not observed (Fig. 3A). Taken together, these results suggest that the domains within $\alpha 1A$ -AR and $\alpha 1D$ -AR which are responsible for the different subtype-specific effects on cell proliferation may reside in the region extending from the amino-terminal extracellular tail to TMD V.

cAMP production

As described above, we have recently observed that all of the $\alpha 1$ -AR subtypes share the inositol 1,4,5-triphosphate/ Ca^{2+} signaling pathway, but only $\alpha 1A$ - and $\alpha 1B$ -ARs are coupled to the cAMP signaling pathway, and thereby inhibiting serum-promoted cell proliferation (submitted for publication). We first confirmed that all the chimeric $\alpha 1$ -ARs couple to Ca^{2+} signaling pathway; hence, the [Ca^{2+}] $_i$ elevation response induced by NA

(10 μ M) in CHO α 1A, CHO α 1D, and CHO α 1A/ α 1D-CRs (data not shown). Next, we measured α 1-AR-mediated cAMP production. In CHO α 1A, AAD, ADA, and ADD, NA (10 μ M) caused a 4.4–7.4-fold increase in cAMP production over basal levels. On the other hand, α 1-AR activation did not promote cAMP synthesis in CHO α 1D, DDA, DAD, and DAA (Fig. 3B). Furthermore, pretreatment with the α -AR antagonist phentolamine (10 μ M) abolished this α 1-AR-mediated cAMP production in CHO α 1A, AAD, ADA, and ADD. These results indicate that the domains responsible for the difference between α 1A-AR and α 1D-AR in the outcome of α 1-AR-mediated cAMP production may reside in the region stretching from the amino-terminal extracellular tail to TMD V. This region, which is important for the cAMP production signaling pathway, is the same as that responsible for the inhibition of serum-promoted cell proliferation.

cAMP production in the amino acid-substituted mutant receptors

To identify the amino acid(s) responsible for α 1-AR subtype-specific cAMP signaling pathways, we first compared the sequences of wild-type human α 1A-AR and α 1B-AR with α 1D-AR. In the region from the amino-terminal extracellular tail to TMD V of α 1D-AR, we found 31 amino acids different from those common to the human α 1A-AR and α 1B-AR (Fig. 1). Among ICLs, which are considered to be important for coupling to G-proteins, although there are no such amino acid differences in ICL I of α 1D-AR, we identified two such amino acids in ICL II, Thr-136, and Val-138. These two amino acids of α 1A-AR were individually, as well as in combination, changed to the corresponding amino acids of α 1D-AR in the different mutants. The positions of the mutated amino acids are indicated in Fig. 1. The K_d values of [125 I]HEAT for the amino acid-substituted mutant receptors were not significantly different from those of the wild-type human α 1A- and α 1D-ARs (K_d = 100–220 pM, B_{max} = 0.9–2.1 pmol/mg protein). The mutation of Thr-136 to Ala (T136A) resulted in NA (10 μ M) stimulation causing a 3.6-fold increase in cAMP production over the level without stimulation (Fig. 4). This increased level was almost 80% of the stimulation of α 1A-AR-induced cAMP production. However, mutation of Val-138 to Met (V138M) showed only a 2-fold increase in cAMP production that was less than 50% the stimulation of α 1A-AR induced cAMP production (Fig. 4). Furthermore, the double mutant (T136A/V138M) showed a 2.5-fold increase in cAMP production over basal levels, almost the same as V138M (Fig. 4). Pretreatment with phentolamine (10 μ M) almost completely blocked NA-stimulated responses in the wild-type α 1A-, α 1D-AR, and the amino acid-substituted mutants (Fig. 4). Taken together, these results may indicate that the region extending from the

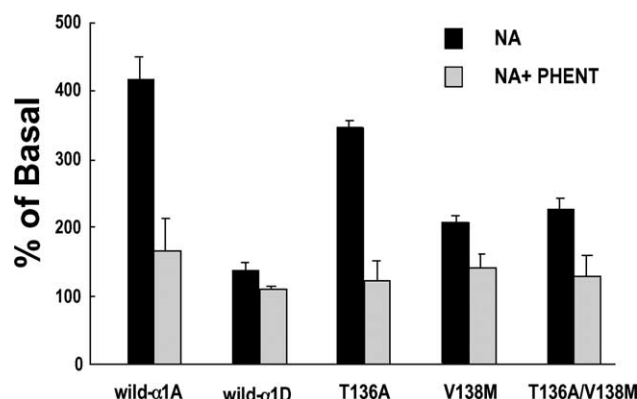


Fig. 4. Effect of α 1-AR subtype activation on intracellular cAMP production in each HEK293 cell transiently expressing the wild-type α 1A-AR, α 1D-AR, T136A, V138M, or T136A/V138M. Phentolamine (10 μ M) treatment was started 30 min prior to stimulation by noradrenaline. Values are means \pm SEM of at least three different experiments each performed in duplicate. NA, noradrenaline; PHENT, phentolamine.

amino-terminal extracellular tail to TMD V, and in particular, residue Val138 in ICL II of the α 1A-AR appears to be intimately connected to the Gs/cAMP pathway.

As described in the introductory material, α 1-ARs mediate the effects of the sympathetic nervous system, especially those effects related to the regulation of cellular hypertrophy and proliferation. As the three different α 1-AR subtypes are generally co-expressed, little is known regarding the physiological role of each receptor subtype at a cellular level and in the regulation of cell proliferation. Using a heterologous expression system, we have recently found that these highly homologous receptor subtypes have distinct signal transduction coupling properties and have differential effects on cell proliferation; thus, activation of α 1A- or α 1B-AR inhibits serum-promoted cell proliferation, whereas the activation of α 1D-AR has a growth-promoting effect (submitted for publication). The present study further demonstrates that certain different amino acids between each α 1-AR subtype may influence and differentiate functional responses mediated by α 1-AR subtypes. This study provides valuable information on the α 1-AR subtype and G-protein coupling.

Acknowledgments

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