Sarcosine¹, Glycine⁸ Angiotensin II Is a Functional AT₁ Angiotensin Receptor Antagonist

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Sarcosine¹, glycine⁸ angiotensin II (SG Ang II) displayed unusual characteristics in early pharmacological studies. It was a potent antagonist of the dipsogenic actions of intracerebroventricularly administered Ang II in the rat, but showed low affinity for bovine cerebellar Ang II receptors. It has recently been shown that SG Ang II binds preferentially to the AT₁ receptor. To determine if SG Ang II is a functional antagonist of the AT₁ receptor-mediated calcium signaling, CHO cells stably transfected with AT₁ receptors were exposed to Ang II in the presence and absence of SG Ang II. At concentrations of 10–100 nM, SG Ang II completely inhibited Ang II-stimulated intracellular Ca²⁺ mobilization in AT_{1A} and AT_{1B} receptor-transfected cells. SG Ang II and ¹²⁵I-SG Ang II bound to AT_{1A} and AT_{1B} receptor-transfected cells with K_D and K_I values of 2–4 nM. In contrast, SG Ang II bound to AT_2 transfected cells with a K_1 value of 7.86 µM. These results demonstrate that SG Ang II is a selective and functional peptide antagonist of the AT_1 angiotensin II receptor subtype.

Key Words: AT₁ receptors; calcium signaling; sarcosine¹, glycine⁸ angiotensin II; antagonist; CHO cells; transfection.

Introduction

The actions of angiotensin II (Ang II) are mediated by two receptor subtypes: AT_1 and AT_2 (1). This discovery arose from the development of peptidomimetic and nonpeptide antagonists selective for the two major Ang II receptor subtypes (2). Although some analogs of Ang II have been demonstrated to have AT_2 selectivity (3,4), it is only recently that an Ang II analog selective for the AT_1 receptor has been observed (5).

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Based on functional studies in the brain (6), SG Ang II was classified as a potent Ang II receptor antagonist based on its ability to block the dipsogenic actions of intracerebroventricular (ICV) Ang II. However, the inability of SG Ang II to compete for ¹²⁵I-Ang II binding sites in the bovine cerebellum (7) raised questions about the mechanism whereby SG Ang II blocked the dipsogenic response to Ang II. Subsequent discovery that the predominant Ang II receptor subtype in the bovine cerebellum was the AT₂ subtype (8), and that the predominant receptor mediating the dipsogenic response to Ang II is the AT₁ subtype (9–11) led to the suggestion that SG Ang II was an AT₁ selective ligand (5).

The studies described herein demonstrate that SG Ang II is a functional antagonist of AT₁ receptor-mediated stimulation of intracellular Ca²⁺ mobilization and confirm the AT₁ receptor selectivity of this Ang II analog.

Results

The ability of SG Ang II and sarcosine¹, isoleucine⁸ Ang II (SI Ang II) to compete for ¹²⁵I-SI Ang II binding to cells stably transfected with Ang II receptors is demonstrated in Fig. 1 and Table 1. SG Ang II bound with high affinity to both AT_{1A} and AT_{1B} transfected CHO cells. In contrast, the K_I of SG Ang II for AT₂ transfected cells was more than 1000-fold lower than for the AT₁ receptor transfected cells. The K_I for SI Ang II was about threefold higher affinity than that for SG Ang II in AT_{1A} and AT_{1B} transfected cells and about 900-fold higher than that for SG Ang II in AT₂ transfected cells.

The affinity of 125 I-SG Ang II and 125 I-SI Ang II for the Ang II receptor transfected CHO cells was similar to that of the corresponding un-iodinated peptides (Table 1). The K_D for 125 I-SG Ang II binding in AT $_{1A}$ and AT $_{1B}$ transfected CHO cells was 4 and 2.2 nM, respectively. It was not possible to determine a K_D for 125 I-SG Ang II binding in the AT $_2$ transfected cells. The K_D for 125 I-SI Ang II binding in AT $_{1A}$ and AT $_{1B}$ transfected CHO cells was 0.32 and 0.73 nM, respectively, while the K_D of 125 I-SI Ang II for AT $_2$ transfected cells was 16.6 nM. The $B_{\rm max}$ values for 125 I-SG Ang II and 125 I-SI Ang II were similar (Table 1).

Table 1
Comparison of Binding of Sar¹,Gly⁸ Ang II and Sar¹,Ile⁸ Ang II and ¹²⁵I-labeled Sar¹, Gly⁸ Ang II, and Sar¹,Ile⁸ Ang II in CHO Cells Stably Transfected with AT_{1A}, AT_{1B}, or AT₂ Receptors

	Sar ¹ ,Gly ⁸	Sar ¹ ,Ile ⁸	¹²⁵ I-Sar ¹ ,Gly ⁸	¹²⁵ I-Sar ¹ ,IIe ⁸	¹²⁵ I-Sar ¹ ,Gly ⁸	¹²⁵ I-Sar ¹ ,Ile ⁸
Receptor	Ang II	Ang II	Ang II	Ang II	Ang II	Ang II
subtype	K_I	K_I	K_D	K_D	B_{\max}	B_{max}
AT_{1A}	2.16 ± 0.83 (2)	0.60 ± 0.33 (2)	4.00 ± 1.28 (2)	0.317 ± 0.069 (2)	532 ± 216 (2)	$579 \pm 118 (2)$
AT_{1B}	3.12 ± 1.00 (3)	1.10 ± 0.09 (3)	2.19 ± 0.69 (3)	0.734 ± 0.206 (3)	$151 \pm 67 (3)$	$150 \pm 25 (3)$
AT_2	7856 ± 2398 (4)	8.92 ± 3.14 (4)	_	$16.6 \pm 2.2 (4)$	_	$2437 \pm 97 (4)$

Values are mean \pm SEM (n). Units for K_I and K_D are nM.

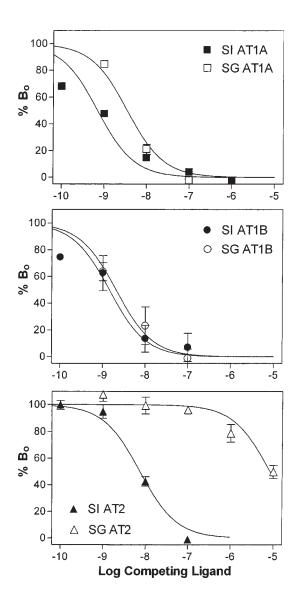


Fig. 1. Representative competition curves of Sar¹,Gly⁸ Ang II (SG) and Sar¹,Ile⁸ Ang II (SI) for specific ¹²⁵I-Sar¹,Ile⁸ Ang II binding in CHO cells stably transfected with either AT_{1A}, AT_{1B}, or AT₂ receptors. ¹²⁵I- Sar¹,Ile⁸ Ang II was present at a concentration of 0.4 n*M*. IC₅₀ values for this graph are SI AT_{1A} = 0.70 n*M*, SI AT_{1B} = 1.40 n*M*, SI AT₂ = 7.49 n*M*, SG AT_{1A} = 3.48 n*M*, SG AT_{1B} = 2.02 n*M*, and SG AT₂ = 8355 n*M*. Specific binding was defined as binding that was displaceable by 3 μ *M* Ang II. Error bars indicate average \pm SEM of triplicate measurements.

To test the functional effect of SG Ang II on Ang II receptors, CHO cells stably transfected with AT_{1A}, AT_{1B}, or AT₂ receptors were loaded with Fura-2 and placed in a perfusion chamber. The Ca²⁺ mobilization response to Ang II was documented in both AT_{1A} and AT_{1B} transfected cells (Fig. 2, Panels A–E), but was absent in AT₂ receptor transfected cells (Fig. 2, Panel F). Perfusion of the cells with 100 nM SG Ang II immediately prior to and during perfusion with Ang II completely blocked the Ca²⁺ mobilization response to 10 nM Ang II in both AT_{1A} and AT_{1B} transfected cells (Fig. 2, Panels A, B, D, E). SG Ang II was completely devoid of any agonistic effect on Ca²⁺ mobilization in AT_{1A} and AT_{1B} transfected cells (Fig. 2, Panels B and E). In AT_{1A} transfected cells, 10 nM SG Ang II was able to completely block the Ca²⁺ mobilization response to 1 nM Ang II (Fig. 2, Panel C). The inhibition of the Ca²⁺ mobilization response to SG Ang II was reversible. When Ang II was perfused 16–20 min following washout of SG Ang II, it again caused an intracellular Ca²⁺ mobilization response (Fig. 2, Panels A,B,C,E).

Discussion

The ability of Ang II receptors to cause an intracellular Ca²⁺ mobilization response is well established (12–15). Stimulation of both AT_{1A} and AT_{1B} (16,17) receptors increases intracellular Ca²⁺. Consistent with the absence of an AT₂ receptor-mediated Ca²⁺ mobilization response, it is generally recognized that the Ca²⁺ mobilization responses to Ang II are exclusively mediated by AT₁ receptors (16–19). However, some studies using the AT₂ receptor-selective antagonist PD123319 suggest that the AT₂ receptor can participate in Ca²⁺ mobilization responses induced by Ang II (20,21).

The observation that nanomolar concentrations of SG Ang II are capable of completely inhibiting the Ca^{2+} mobilization to a 10-fold lower concentration of Ang II suggests that SG Ang II is a potent antagonist of the AT_1 receptor. This confirms observations of Fitzsimons et al. (6), made prior to the discovery of multiple Ang II receptor subtypes, that SG Ang II is a potent inhibitor of the dipsogenic response. It is now well established that the dipsogenic response is mediated by the AT_1 receptor subtype (9,10,22). Of note,

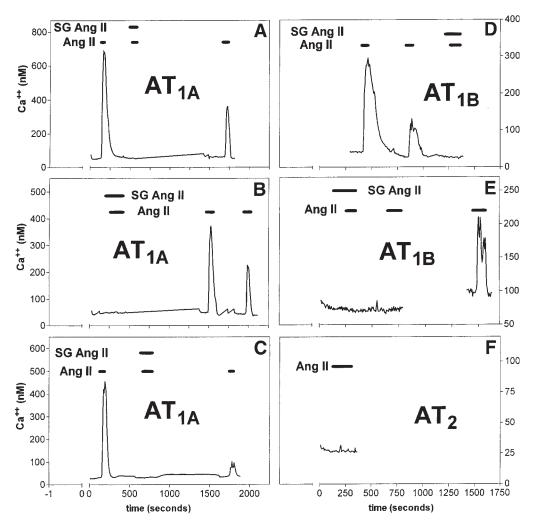


Fig. 2. Inhibition of Ang II-induced calcium signaling by Sar¹,Gly⁸ Ang II (SG). (A) The initial response generated by 10 nM Ang II in AT_{1A} transfected cells was $689 \pm 80 \text{ nM}$ (mean $\pm \text{ SEM}$, n = 20). 5 min later 100 nM of Sar^1 , Gly⁸ Ang II was applied to the cells; 30 s later 10 nM Ang II was applied to the cells in the continued presence of Sar¹, Gly⁸ Ang II. The Ang II was applied for 1 min; 20 min later Ang II (10 nM) was again applied to the cells to cause the second, smaller response of 356 ± 33 nM. (B) Sar¹, Gly⁸ Ang II (100 nM) was applied initially to AT_{1A} transfected cells, followed 1 min later by Ang II (10 nM) in the continued presence of Sar¹, Gly⁸ Ang II for 3 min; 17 min later Ang II (10 nM) was applied to the cells for 90 s, generating the first calcium response of 371 ± 52 nM. Six minutes later Ang II (10 nM) was again applied to the cells for 90 s generating the second, smaller response of 225 ± 40 nM. (C) The protocol was similar to that in A except that Ang II was applied to the AT_{1A} transfected cells at a concentration of 1 nM giving a response of 643 ± 41 nM and Sar¹,Gly⁸ Ang II was applied at a concentration of 10 nM completely blocking the Ang II response and largely blocking the subsequent response to Ang II applied 15 min later. (D) Initial response generated by 10 nM Ang II applied to AT_{IR} transfected cells for 1 min was $296 \pm 69 \text{ nM}$. Six minutes later Ang II (10 nM) was again applied to the cells, generating the second, smaller response, $118 \pm 35 \text{ nM}$. Six minutes later 100 nM Sar¹,Gly⁸ Ang II was applied to the cells followed by Ang II (10 nM) 45 s later in the continued presence of Sar¹,Gly⁸ Ang II. The Ang II was applied for 90 s with no response. (E) Sar¹, Gly⁸ Ang II (100 nM) was applied to AT_{1B} transfected cells followed 2 min later by Ang II (10 nM) in the continued presence of Sar¹,Gly⁸ Ang II for 90 s. Ang II (10 nM) was again applied to the cells 5 min later with no effect. Eleven minutes later Ang II (10 nM) was applied a third time to the cells generating an intracellular calcium response in a single cell. (F) Ang II (10 nM) was applied to the AT, transfected cells for 3 min without causing any cytoplasmic calcium response.

Zhu and Herbert (23) have shown that the dipsogenic response to Ang II in rats involves mobilization of Ca²⁺ via an L-type calcium channel.

There was a profound AT_1 receptor selectivity observed for SG Ang II in the present study: more than a 1000-fold difference in binding affinity. This is considerably more than that previously observed in studies of naturally occurring Ang II receptor subtypes in a variety of rat tissues (5).

However, SI Ang II also displayed a small selectivity for the AT $_1$ receptor in this study, ranging from 8-fold (AT $_2$ vs AT $_{1B}$) to more than 50-fold (125 I-SI Ang II binding to AT $_2$ vs AT $_{1A}$). This suggests that the AT $_2$ receptor transfected into these CHO cells displayed a lower Ang II binding affinity than the AT $_1$ transfected cells. Possible explanations for this difference could be different membrane properties of CHO cells, different receptor-associated proteins in the CHO

membranes, or differences in post-translational modifications of the AT₂ receptor in CHO cells. When the difference in affinity of SI Ang II for the AT₁ and AT₂ is factored in, the relative selectivity of SG Ang II for the AT₁ receptor is 244- and 311-fold for the AT_{1A} and AT_{1B} receptors, respectively, compared to the AT₂ receptor. This selectivity is closer, but still substantially larger than SG Ang II's 40- and 82-fold higher affinity for AT_{1A} and AT_{1B}, respectively, compared to the AT2 in naturally occurring Ang II receptors reported previously (5). Of note, the B_{max} values for ¹²⁵I-SG Ang II and ¹²⁵I-SI Ang II were similar in the AT₁ receptor transfected cells. This also differs from the previous study of ¹²⁵I-SG Ang II and ¹²⁵I-SI Ang II binding to naturally occurring AT₁ receptors in rat tissues in which there is an unresolved reduction in $B_{\rm max}$ for $^{125}\text{I-SG}$ Ang II relative to that of ¹²⁵I-SI Ang II (5).

To date, SG Ang II is the only Ang II analog identified as having a substantial AT_1 Ang II receptor subtype selectivity. However, many of the analyses of Ang II congeners were carried out prior to the discovery of different Ang II receptor subtypes. Thus, it is possible that other Ang II analogs might also possess AT_1 receptor selectivity. Discovery of such analogs might further aid in our understanding of ligand-docking characteristics of Ang II receptors as well as their signal transduction mechanisms.

In summary, these results confirm and extend previous studies of SG Ang II which suggest that it is a potent and subtype-selective AT₁ receptor antagonist.

Materials and Methods

Generation of Stably Transfected AT_{IA} , AT_{IB} , and AT_2 Receptor-Expressing Cells

The generation of the rat AT_{1A} and rat AT_{1B} receptor-expressing, stably transfected CHO cell lines is described previously (24). Briefly, amplimer sets that were specific for either the rAT1AR or the rAT1BR subtype were used to amplify the coding region of each receptor. The amplified products were subcloned into the eukaryotic expression vector pRc/CMV (Invitrogen, San Diego, CA) and sequenced to ensure authenticity and proper orientation.

An amplimer set specific for the rAT₂R was also synthesized. The sense primer (5'-ATGAAGGACAACTTCAGT TTTGCT-3') corresponds to the sequence beginning at the start codon of the open reading frame of the rat AT₂R. The antisense primer (5'-TTAAGACACAAAGGTGTCCATT TC-3') corresponds to the sequence that encompasses the stop codon of the open reading frame of the rat AT₂R. The receptor-specific amplimer sets were utilized in PCR experiments with rat genomic DNA as template and the proofreading DNA polymerase PFU (Stratagene, La Jolla, CA). The amplified products were subcloned into the eukaryotic expression vector pCR3 (Invitrogen) and sequenced to ensure authenticity and proper orientation.

The rAT_{1A}R-pRc/CMV, rAT_{1B}R-pRc/CMV, and rAT2-pCR3 constructs were transfected into CHO cells utilizing the calcium phosphate precipitation procedure (Stratagene). Pure clones, expressing either the rAT_{1A}R, rAT_{1B}R, or rAT₂ subtype, were selected by the antibiotic G418 (Gibco BRL) and purified by the limited dilution technique.

Cell Culture

The AT_{1A}, AT_{1B}, and AT₂ stably transfected CHO cells were grown in six-well plates (Costar) on round microscope cover slips for the Ca²⁺ imaging experiments, or 75 mm culture flasks for the radioligand binding assays at 37°C in a humidified atmosphere of 5% CO₂/95% air (v/v). The medium was Dulbecco's modified Eagle's medium, high glucose (DMEM-Hi, Gibco BRL, Gaithersburg, MD), supplemented with 10% fetal calf serum (FCS, Hyclone Laboratories, Ogden, UT), 80 units/mL penicillin, 80 µg/mL streptomycin, 1.6 mM L-glutamine, and 0.175 g/mL L-proline for untransfected cells. For transfected cells, the medium also included 1000 µg/mL geneticin (G418, Gibco BRL). Cells were grown to approx 50–75% confluency and were used between three and six passages post-thawing for these experiments.

Intracellular Ca²⁺ Imaging

All Ca²⁺ imaging studies were performed in a standard physiological saline solution (PSS; 140 mM NaCl, 5 mM KCl, 2 mM CaCl₂, 1 mM MgCl₂, 10 mM Hepes, and 6 mM glucose, pH adjusted to 7.4 with NaOH) at room temperature. Coverslips with attached cells were washed in PSS and then loaded with 1 µM Fura-2-AM in PSS for 30 min. The cells were then rinsed with the PSS and allowed to de-esterify the dye for an additional 30 min. The coverslip was then inserted into a sealed chamber (0.5 mL volume) mounted on the stage of an inverted microscope (Nikon Diaphot; 40× objective) and the cells were constantly perfused with PSS (2 mL/min). Exposure of the cells to the various test compounds (all in PSS) was achieved by switching inflow lines through a common manifold mounted just upstream from the sealed chamber. Test compounds would reach the cells within 30 s of switching. Paired images (excitation at 340 nm and 380 nm, emissions monitored at 510 nm) were collected with a CCD camera every 6 s, corrected for background fluorescence, and then the ratio of fluorescence intensity (340/380) for a region of interest was calculated and converted to a calcium value based on a standard curve. In each of the experimental conditions, approximately 20 cells were targeted for fluorescence monitoring. The results from all the cells were averaged to obtain the final value portrayed in the figures. The experiments were carried out two times using different passages of cells. The standard Ca^{2+} curve was generated with 10 μM Fura-2 in a Hepes-buffered (10 mM, pH 7.4) 130 mM KCl, 10 mM EGTA solution with varying amounts of CaCl₂ added to generate specific Ca²⁺ levels. Free Ca²⁺ levels in the standard bath were calculated with EQCAL software (Biosoft, Ferguson, MO). Control of light exposures, image collection, and data manipulation was done with MetaFluor imaging software (Universal Imaging Corp., West Chester, PA). Fura-2 and Fura-2-AM were obtained from Molecular Probes (Eugene, OR). All salts and buffer components were from Sigma. The results from all the cells were averaged on an Excel (Microsoft Corp., Redmond, WA) spreadsheet and translated into XY coordinate graphs using PRISM (Graphpad Software, San Diego, CA).

Radioligand Binding Assays

Measurement of Ang II receptor binding in cultured cells was carried out using procedures described previously (4). Briefly, the culture medium was aspirated and replaced with a hypotonic, 20 mM NaPO₄ 5 mM EDTA, pH 7.2 solution. The cells were scraped from the culture flasks with a rubber policeman and mechanically homogenized in this buffer. The homogenate was centrifuged (48,000g for 20 min at 4°C) and the supernatant decanted. The membrane pellet was resuspended by homogenization in assay buffer (150 mM NaCl, 5 mM EDTA, 0.1 mM bacitracin, 50 mM NaPO₄, pH 7.1–7.2). The homogenate was centrifuged as before. The supernatant was again decanted and the membrane pellet was resuspended in 5 mL of assay buffer by mechanical homogenization.

To measure Ang II receptor binding, cell membranes were incubated with ¹²⁵I-SI Ang II or ¹²⁵I-SG Ang II prepared by the chloramine T procedure (25). The radioligands were purified as the mono-radioiodinated peptides by HPLC using a C18 reverse phase column eluted isocratically with a triethylamine phosphate (pH 3.0):acetonitrile mobile phase (26).

For saturation isotherm analysis, membranes were incubated with increasing concentrations of $^{125}\text{I-SI}$ Ang II or $^{125}\text{I-SG}$ Ang II ranging from 0.1 to 4 nM for 2 h at 22–24°C. To assess nonspecific binding, one of three tubes containing each concentration of radioligand also contained 3 μM Ang II to saturate the Ang II receptors. Values presented represent specific (total minus nonspecific) binding.

For competition binding studies, membranes were incubated with 0.4 nM $^{125}\text{I-SI}$ Ang II in the presence of varying concentrations of SG Ang II or SI Ang II ranging from 0.1 nM to $10 \text{ }\mu\text{M}$. Non-specific binding was again determined in tubes containing $3 \text{ }\mu\text{M}$ Ang II and the non-specific binding was subtracted to determine specific Ang II receptor binding.

Determination of K_D and K_I values for ¹²⁵I-SG Ang II, ¹²⁵I-SI Ang II, SG Ang II, and SI Ang II were made using PRISM One-Site Saturation and One-Site Competition equations: $Y = B_{\text{max}} / (K_D + L)$ and $Y = B_0 - B_0 \times I / (I + \text{IC}_{50})$, respectively, where Y = amount of bound radioligand, L = radioligand concentration, $B_0 =$ radioligand bound in the

absence of competing ligand, and I = concentration of competing ligand.

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