# SHORT COMMUNICATION

# **Tripeptides with C-Terminal Arginine as Potential Inhibitors** of Urokinase

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**Abstract** We synthesized and tested ten peptides with the molecular structure being H–D-Ser–AA–Arg–OH for their effect on the amidolytic activities against urokinase, thrombin, trypsin, plasmin, tissue plasminogen activator and kallikrein. The inserted amino acid in each peptide was either leucine, norleucine, izoleucine, valine, norvaline,  $\alpha$ -metyloalanine,  $\alpha$ -aminobutanoic acid, homoleucine, tert-leucine or neoglycine. H–D-Ser–NVal–Arg–OH (compound 4) was the most active inhibitor of urokinase plasminogen activator with a K<sub>i</sub> value of 0.85 µM. Compound 4 showed cytotoxic effect against MDA-MB-231 and DLD cell lines, respectively, with IC<sub>50</sub> values of 25 and 19 µM. Synthesised compounds did not have activity against MCF-7 cancer cells. These peptides were nontoxic against pig's erythrocytes in vitro.

**Keywords** Urokinase inhibitor · Anti-amidolytic activity · Low molecular weight peptides

# Introduction

Urokinase plasminogen activator (uPA), a trypsin-like serine protease, plays an important role in several biological processes including tissue remodeling, cell migration and matrix degradation (Irigoyen et al. 1999; Syrovets and Simmet 2004). The primary role of uPA is converting

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plasminogen into its active form plasmin, a broad spectrum serine protease, which participates in the fibrinolytic cascade (Collen 2001; Rijken and Sakharov 2001). uPA and plasmin are strongly associated with tumor cells by influencing tumor initiation, proliferation, migration, invasion, metastasis and apoptosis (Andreasen et al. 1997; Duffy 2004; Mazar et al. 1999). Both these enzymes directly promote proteolysis. However, plasmin acts also indirectly through the activation of matrix metaloproteases. Overexpression of uPA has been found in various malignant tumors, especially in the gastrointestinal, respiratory, genitourinary systems, and in bones, bone marrow, skin, breast and brain (Dass et al. 2008). High levels of urokinase are correlated with enhanced tumor invasiveness, metastasis and poor prognosis in certain cancers (Sidenius and Blasi 2003).

The inhibition of plasminogen activation by uPA appears to be an attractive approach to cancer therapy, especially against tumor growth and metastasis (Rockway et al. 2002; Rockway 2003; Schweinitz et al. 2004). In 1997, Ke et al. analyzed a peptide library and selected the primary peptide sequence SGRSA (from position P3 to P2) as the substrate of uPA and tPA. Zeslawska et al. (2000, 2003) investigated the structure-function relationship of certain peptides such as phenethyl-sufonamidino-D-seryl-Lalanyl-L-argininal, and peptidomimetic inhibitors of uPA (e.g., benzamidine, amiloride, 2,4,6-triisopropyl-phenylsulfonyl-L-(3-amidino)-phenylalanine-piperazine-N- $\beta$ alanine). Studies have demonstrated that the S1 pocket of uPA is the critical site for interaction with inhibitors. The P1 positively charged residue of an inhibitor (arginine, benzamidine or phenylguanidine) makes a salt bridge with Asp189 in the S1 pocket of uPA. The P2 residue of the potential inhibitor, which should be hydrophobic, is accommodated in S2 side. When the P3 position is

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occupied by an unnatural D-amino acid, the side-chain projects into the S4 pocket. The residue of D-Ser could form a favorable interaction with His-99 in the S4 site of uPA.

Recently, we described a series of peptide analogs containing arginine as inhibitors of urokinase. The first series was H(Ac)-D-Ser-Ala-(Gly)-Arg-OH(NH<sub>2</sub>) (Markowska et al. 2008) and the most active inhibitor of urokinase was H-D-Ser-Gly-Arg-OH with an IC50 value of 1 mM. The second series of peptide analogs of arginine we synthesized were the peptides of the general formula H-D-Ser-Ala-Arg-NH-X (Markowska et al. 2010), where  $X = (CH_2)_n - NH_2$ , n = 2-9,  $(CH_2)_m - OH$ , m = 2-4. H-D-Ser-Ala-Arg-NH-(CH<sub>2</sub>)<sub>5</sub>-NH<sub>2</sub> inhibited urokinase with a  $K_i$  value of 6.3  $\mu$ M. The Lineweaver–Burke analysis of the H-D-Ser-Ala-Arg-NH-(CH2)5-NH2 compound proved this compound to be a competitive inhibitor of urokinase. The third series of compounds with D-Ser-Ala-Arg sequence were N-sulfonylamides peptides (Markowska et al. 2010). 2,4,6-triisopropylphenyl-SO<sub>2</sub>-D-Ser-Ala-Arg-OH was the most selective inhibitor of urokinase and α-tolyl-SO<sub>2</sub>-D-Ser-Ala-Arg-OH was the most active inhibitor of uPA with a  $K_i$  value of 24  $\mu$ M.

The purpose of this paper was the synthesis and the amidolytic activity of 10 novel peptides against urokinase, thrombin, trypsin, plasmin, tissue plasminogen activator (t-PA) and kallikrein in vitro. We also tested the hemolytic activity of the peptides against swine erythrocytes and the anti-tumor activity against the following human breast cancer cells: standard MCF-7, estrogen-independent MDA-MB-231 and colorectal adenocarcinoma tumor DLD cell line.

# **Materials and Methods**

The inserted amino acid (AA) in each of the novel peptides was either leucine, norleucine, izoleucine, valine, norvaline,  $\alpha$ -metyloalanine,  $\alpha$ -aminobutanoic acid, homoleucine, tert-leucine or neoglycine. Table 1 represents the identity of the inserted amino acid in each of the 10 peptides synthesized from X–D-Ser–AA–Arg–OH.

### Source of Reagents

Fmoc-Arg(Pbf)-OH (Fmoc = 9-fluorenylmethyloxycarbonyl, Pbf = pentamethyl-dihydrobenzofuran), chloranil, acetaldehyde, HOBt = 1-hydroxybenzotriazole were purchased from Fluka (Schnelldorf, Germany). Fmoc-D-Ser (t-Bu)-OH (t-Bu = t-butyl) and 2-chlorotrityl chloride resin were purchased from Merck (Novabiochem, Darmstadt, Germany). TFA = trifluoroacetic acid, DIPEA = diisopropylethylamine, piperidine, TBTU = tetrafluoroborate salt of

Table 1 Amino acid insertion in novel peptides

Peptide	Inserted amino acid
1	α-Metyloalanine
2	α-Aminobutanoic acid
3	Valine
4	Norvaline
5	Leucine
6	Norleucine
7	Izoleucine
8	tert-Leucine
9	Homoleucine
10	Neoglycine

the O-(benzotriazol-1-yl)-N,N,N',N'-tetramethyluronium tetrafluoroborate, DIC = diisopropylcarbodiimide, NMP = 1-methyl-2-pirrolidon, Fmoc-L-Leu-OH, Fmoc-L-NLe-OH, Fmoc-L-Ile-OH, Fmoc-L-Val-OH, Fmoc-L-NVa-OH, Fmoc-Aib-OH, Fmoc-L-2Abu-OH, Fmoc-L-HLeu-OH, Fmoc-L-TLe-OH, Fmoc-L-NptGly-OH were obtained from Iris Biotech GmbH (Marktrewitz, Germany). DCM = dichloromethane and DMF = dimethylformamide were the products of Chempur (Piekary Slaskie, Poland) DCM was used without further purification. DMF was distillated over ninhidrin and stored under molecular sieves 4A. HPLC solvent acetonitrile was purchased from Merck (Darmstadt, Germany). Urokinase, trypsin, kallikrein, amiloride and  $Bzl-L-Arg-pNA\cdot HCl$  (Bzl = benzyl) were purchased from Sigma (Schnelldorf, Germany). Plasmin, S-2444 (pyro-Glu-Gly-Arg-pNA·HCl), S-2238 (H-D-Phe-Pip-Arg-pNA), S-2251 (H-D-Val-Leu-Lys-pNA), S-2266 (H-D-Val-Leu-Arg-pNA·2HCl and S-2288 (H-D-Ile-Pro-Arg-pNA) were obtained from Chromogenix (Milano, Italy). Thrombin and phosphate buffered saline (PBS) were purchased from Lubelska Wytwórnia Szczepionek (Lublin, Poland). t-PA was obtained from Boehringer Ingelheim GmbH (Ingelheim, Germany).

### Peptide Synthesis

The peptides shown in Table 1 were synthesized manually using the standard Fmoc-based strategy (Chan and White 2000). Fmoc deprotection steps were carried out with 20% (v/v) piperidine in DMF/NMP (1:1) for 15 min. The coupling reactions of Fmoc amino acids were performed in DMF/NMP/DCM (1:1:1) using a molar ratio of amino acid/DIC/HOBt/resin 3:3:3:1. In the case of the coupling of Fmoc–D-Ser(t-Bu)–OH molar ratio of amino acid/TBTU/HOBt/DIPEA/resin was 2:2:2:4:1. The reactions were monitored with the Steward chloranil test. The cleavage from the resin was carried out with TFA/water (95/5). After 2.5 h stirring, the resin was filtered and washed with TFA.

The combined filtrates were concentrated under reduced pressure. The crude peptide was washed with cold diethyl ether, filtered, dissolved in glacial acetic acid and lyophilized.

The Shimadzu LC-10A system (Smith and Hanly 1997; Hartmann et al. 2009) (Shimadzu Europa GmbH, Duisburg, Germany) was used for analytical and semipreparatory HPLC (Phenomenex C18, Jupiter 90A, 4  $\mu$ m, 250 × 10 mm; Phenomenex C18, Jupiter 300A, 5  $\mu$ m, 250 × 4 mm; solvents: A, 0.1% aqueous TFA; B, 0.1% TFA in acetonitrile, gradient 0% B to 100% B in A in 30 min, flow rate 1 ml/min, monitored at 220 nm). The major peak fraction was pooled and lyophilized. The molecular weight determination was performed by mass spectrometry using a Bruker Daltonics Esquire 6000 (Bruker Daltonik GmbH, Leipzig, Germany) with electrospray ionization (ESI), (Table 2).

### **Enzymatic Investigations**

Determination of amidolytic activity was performed as previously described (Okada et al. 1988). A detailed description of the method is given below. Buffer and  $0.1 \text{ cm}^3$  of enzyme solution was added to  $0.2 \text{ cm}^3$  of examined compound (1–10) (as control 0.15 M NaCl). The buffer and the enzyme solution included the following constituents:

- a. Tris buffer: 0.6 cm<sup>3</sup> (pH 8.8), enzyme: urokinase (50 units/cm<sup>3</sup>), synthetic substrate: S-2444 (0.1 cm<sup>3</sup>, 3 mM/dm<sup>3</sup>);
- b. Tris buffer: 0.5 cm<sup>3</sup> (pH 8.4), enzyme: thrombin (1 units/cm<sup>3</sup>), synthetic substrate: S-2238 (0.2 cm<sup>3</sup>, 0.75 mM/dm<sup>3</sup>);
- c. Tris buffer: 0.5 cm<sup>3</sup> (pH 7.4), enzyme: plasmin (0.4 units/cm<sup>3</sup>), synthetic substrate: S-2251 (0.2 cm<sup>3</sup>, 3 mM/dm<sup>3</sup>);

- d. borane buffer: 0.5 cm<sup>3</sup> (pH 7.5), enzyme: trypsin (0.4 units/cm<sup>3</sup>), synthetic substrate: Bzl-L-Arg-pNA·HCl (0.2 cm<sup>3</sup>, 8 mM/dm<sup>3</sup>);
- e. Tris buffer: 0.6 cm<sup>3</sup> (pH 9.0), enzyme: kallikrein (3 units/cm<sup>3</sup>), synthetic substrate: S-2266 (0.1 cm<sup>3</sup>, 7.5 mM/dm<sup>3</sup>);
- f. Tris buffer: 0.6 cm<sup>3</sup> (pH 8.4), enzyme: t-PA (1.67 mg/cm<sup>3</sup>), synthetic substrate: S-2288 (0.1 cm3, 10 mM/dm<sup>3</sup>).

The mixture was incubated for 3 min at 37°C, followed by the addition of synthetic substrate solution in the same buffer. After 20 min of incubation, the reaction was stopped by adding 0.1 cm<sup>3</sup> of 50% acetic acid and the absorbance of the released p-nitroaniline was measured at 405 nm. As shown in Table 3, each value represents the average of a triplicate determination. The IC<sub>50</sub> value for each peptide was considered as the concentration of the inhibitor, which decreased the absorbance at 405 nm by 50%, compared with the absorbance measured under the same conditions without the inhibitor. Also, K<sub>i</sub> was calculated for each peptide from IC<sub>50</sub> based on Cheng–Prusoff equation (Cheng and Prusoff 1973). The results are shown in Table 3.

Our results were compared with the data obtained by Tamura et al. (2000) for 2-phenethyl–SO<sub>2</sub>–D-Ser–Ala–Arg–al, the irreversible urokinase inhibitor with the same tripeptide sequence. The determination methods were identical.

# Tissue Culture

All studies were performed on MCF-7, MDA-MB-231 and DLD cells lines were purchased from American Type Culture Collection, (Rockville, MD). The cells were maintained in DMEM supplemented with 5% fetal bovine serum (FBS), 2 mmol/ml glutamine, 50 U/ml penicillin, 50 mg/ml streptomycin at 37°C in a 5% CO<sub>2</sub> incubator.

Peptide no.	Yield (%)	Retention time (min)	Melting point (°C)	MW	$[M^+H]^+$
1	46	8.93	223-225	346.4	347.7
2	47	8.89	231-234	346.4	347.6
3	45	9.72	211-212	360.4	361.7
4	45	9.99	243-245	360.4	361.3
5	47	11.32	219-220	374.4	375.6
6	56	11.27	232–233	374.4	375.5
7	43	10.99	228-229	374.4	375.6
8	59	10.93	197–199	374.4	375.6
9	45	13.08	221-225	388.5	389.4
10	58	12.10	138–139	388.4	389.5

**Table 2** Analytical data of thesynthesized peptides

Table 3 Inhibition of amidolytic activity of enzymes against synthetic peptides with the general H-D-Ser-AA-Arg-OH formula

No.	AA	$K_i (\mu M)$				
		Urokinase S-2444	Thrombin S-2238	Trypsin Bzl–L-Arg– pNA·HCl	Plasmin S-2251	t-PA S-2288
1.	α-Metyloalanine	n.i.	n.i.	n.i.	$59.09 \pm 4.73$	n.i.
2.	α-Aminobutanoic acid	n.i.	$1.19\pm0.09$	$4.73\pm0.38$	$2.54\pm0.20$	n.i.
3.	Valine	$182.92 \pm 14.63$	n.i.	$4.48 \pm 0.36$	$163.63 \pm 13.09$	n.i.
4.	Norvaline	$0.85\pm0.07$	n.i.	n.i.	$4.36\pm0.35$	n.i.
5.	Leucine	$219.51 \pm 17.56$	$1.1\pm0.09$	$4.73\pm0.38$	$27.27\pm2.18$	$154.54 \pm 12.36$
6.	Norleucine	$4.45\pm0.34$	n.i.	n.i.	$40.9\pm3.27$	n.i.
7.	Izoleucine	$158.53 \pm 12.68$	n.i.	$0.77\pm0.06$	$1.81\pm0.15$	$19.09\pm1.53$
8.	tert-Leucine	$26.62\pm2.15$	$1.43 \pm 0.11$	n.i.	$31.81\pm2.55$	n.i.
9.	Homoleucine	n.i.	n.i.	$3.74\pm0.30$	n.i.	n.i.
10.	Neoglycine	n.i.	n.i.	n.i.	n.i.	$163.63 \pm 13.09$
	2-Ph-(CH <sub>2</sub> ) <sub>2</sub> -SO <sub>2</sub> -D-Ser-Ala- Arg-al	3 nM (Tamura et al. 2000)	-	-	_	_
	Benzyl–SO <sub>2</sub> –D-Ser–4- amidinobenzylamid	7.7 nM (Zeslawska et al. 2003)	-	-	-	-
	Amiloride	7 μM (Zeslawska et al. 2003)				

n.i. No inhibition was observed at maximum concentration (0.02 M)

# Cytotoxicity Assay

The toxicity of the evaluated compounds was determined by the method of Plumb et al. (Plumb et al. 1989). MCF-7, MDA-MB-231 and DLD cells were maintained as described above. After 48 h of incubation of the cells with synthesized peptides, the medium was discarded and the cells were rinsed three times with PBS. The cells were then incubated for 4 h in 2 ml of PBS with 50 ml of MTT (5 mg/ml). After removal of the medium, the cells were lysed in 200 ml of DMSO with 20 ml of Sorensen's buffer (0.1 M glycine with 0.1 M NaCl, pH 10.5). The absorbance was measured at 570 nm. The cytotoxic activity towards MDA-MB-231 and DLD of synthesized peptides was calculated as percentage of nonviable cells and the IC<sub>50</sub> value was estimated from logarithm curves as shown in Table 4.

# Hemolytic Activity

Pig's fresh red blood cells (p-RBC) were washed three times with PBS (35 M phosphate buffer/0.15 mM NaCl, pH 7.4) and were centrifuged at  $1000 \times g$  for 10 min to remove plasma and the buffy coat. Erythrocytes in suspension were incubated with various concentrations of peptides for 1 h at 37°C. The final erythrocyte concentration was 5% v/v. After the centrifugation ( $1000 \times g$  for 10 min), 100 µl of the supernatant was transferred into sterilized 96-well plates, where hemoglobin release was monitored, using Infinite M200 plate reader and measuring

 Table 4
 The nonviability of DLD and MDA-MB-231 cells treated for 24 h with different concentrations of the synthesized inhibitors

Compound no.	IC <sub>50</sub> ( $\mu$ M) (% of control $\pm$ 2)		
	DLD	MDA-MB-231	
1	32	54	
2	43	76	
3	55	48	
4	19	25	
5	56	119	
6	52	84	
7	59	80	
8	57	81	
9	n.e.	n.e.	
10	n.e.	n.e.	
Amiloride	62.1	126	

n.e. No cytotoxic effect was observed in maximum concentration (250  $\mu$ M)

 $^{\rm a}$  Mean values  $\pm$  SD from three independent experiments done in duplicate

the absorbance at 414 nm. Zero hemolysis (blank), hemolysis with amiloride as reference compound for synthesised peptides and 100% hemolysis, which consisted of p-RBC suspended in PBS and 0.1% Triton-X-100, were determined respectively. The percentage of hemolysis was calculated with the following formula: %hemolysis = (Abs<sub>414nm</sub> in the peptide solution in PBS/Abs<sub>414nm</sub> in 0.1% Triton-X-100 in PBS)  $\times$  100.

# **Results and Discussion**

We obtained ten new peptides as potential inhibitors of urokinase by the manual solid phase synthesis and tested them for the effect on the amidolytic activities of urokinase, thrombin, trypsin, plasmin, t-PA and kallikrein. These peptides did not influence the enzymatic activity of kallikrein.

The results indicated that the concentration up to 1000  $\mu$ g/ml of the synthesized peptides did not lyse erythrocytes.

The synthesized peptides were not selective inhibitors of urokinase. Peptides 1 (H–D-Ser–Aib–Arg–OH) and 2 (H–D-Ser–Abu–Arg–OH), containing two carbon atoms in their amino acid side chains inhibited the activity of plasmin. Peptides with three and four carbon atoms in the amino acid side chains (H–D-Ser–Val–Arg–OH 3, H–D-Ser–NVal–Arg– OH 4, H–D-Ser–Leu–Arg–OH 5, H–D-Ser–NLeu–Arg–OH 6, H–D-Ser–ILeu–Arg–OH 7, H–D-Ser–tLeu–Arg–OH 8) inhibited plasmin and urokinase. Tripeptides 9 and 10, containing homoleucine and neoglycine respectively as P2 residue, were non active inhibitors of plasmin and urokinase. The most active peptide against urokinase was H–D-Ser– NVal–Arg–OH 4 with a K<sub>i</sub> value 0.85  $\mu$ M.

The obtained values of  $K_i$  were higher than  $K_i$  of the standard inhibitors. However, 2-phenethyl–SO<sub>2</sub>–D-Ser–Ala–Arg–H was an alkylating agent and irreversibly inhibited urokinase by forming a covalent adduct with an active site of the enzyme (Zeslawska et al. 2003; Tamura et al. 2000). Benzyl–SO<sub>2</sub>–D-Ser–4-amidinobenzylamid with  $K_i = 0.0077 \mu$ M inhibitory activity was a non-peptidic inhibitor of urokinase. 4-Amidinobenzylamine was a decarboxylated arginine mimetic, which has been widely used for the development of urokinase inhibitors (Rockway 2003; Schweinitz et al. 2004). The previous analysis of the effect of H–D-Ser–Ala–Arg–NH–(CH<sub>2</sub>)<sub>5</sub>–NH<sub>2</sub> on the activity of urokinase showed that the derivatives with these kinds of sequence reversibly and competitively inhibited urokinase.

The peptides **1–10** were tested in vitro for their antitumour activity in human breast cancer cells: standard MCF-7, estrogen-independent MDA-MB-231 and colorectal adenocarcinoma tumor DLD line. Peptides **9** and **10** were not active in vivo. The most interesting peptide was compound **4** with an IC<sub>50</sub> values of 19  $\mu$ M for DLD cells and 25  $\mu$ M for MDA-MB-231 cells, which was the most selective inhibitor of urokinase. All potential inhibitors were insignificantly less cytotoxic to MDA-MB-231 and DLD cells than amiloride. Amiloride, which is a potassium-sparing diuretic drug, has been shown to inhibit uPA with K<sub>i</sub> = 7  $\mu$ M and to have antitumor activity in vivo (Zeslawska et al. 2003).

The synthesized peptides did not influence MCF-7 cancer cells. It was found that the proteolytic activity of

uPA is closely related to cell-surface events when incubated with the breast cancer cells. MCF-7 had low uPAR/ uPA-expressing and low plasminogen-binding, whereas MDA-MB-231 had high uPAR/uPA expressing and high plasminogen-binding (Stillfried et al. 2007). Thus, the influence of the synthetic peptides on the cytotoxic effect of MCF-7 cells could be insignificant.

We hope that our studies provided information about the structure–activity and may be used for synthesizing more active enzyme inhibitors and potential antitumor agents.

### Conclusion

Our investigation of the structure–activity relationship of a series of new peptide sequences as potential urokinase inhibitors reveals that the obtained H–D-Ser–NVal–Arg–OH was the most active inhibitor of uPA with K<sub>i</sub> 0.85  $\mu$ M value. The compound showed cytotoxic effects against MDA-MB-231 cell lines with a IC<sub>50</sub> 25  $\mu$ M value and with a IC<sub>50</sub> 19  $\mu$ M value against DLD cell lines. The examined compound did not influence MCF-7 cancer cells.

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