

***N*-*t*-Boc-amino acid esters of isomannide Potential inhibitors of serine proteases**

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Summary. Hepatitis C, Dengue and West Nile virus are some of the most important flaviviruses, that share one important serine protease enzyme. Serine proteases are the most studied class of proteolytic enzyme and, in these cases, a primary target for drug discovery. In this paper, we describe the synthesis and preliminary molecular modeling studies of a novel class of *N*-*t*-Boc amino acid esters derived of isomannide as potential serine proteases inhibitors.

Keywords: Isomannide derivatives – Serine protease – Dengue virus

Introduction

The family *Flaviviridae* comprises more than 60 viruses, many of which are important human pathogens. Hepatitis C virus (HCV), West Nile virus (WN) and Dengue virus are among the most important flaviviruses. HCV is the major cause of blood-borne chronic hepatitis, with nearly 200 million infected people worldwide (Houghton et al., 1991). Chronic HCV infection is associated with liver cirrhosis and hepatocellular carcinoma (Bruix et al., 1989). Current therapeutic regimens based on alpha interferon and the nucleoside analog ribavirin are only partially effective and are limited by the adverse effects of both agents (Wright et al., 2001).

For many years, WN has been recognized as one of the most widely distributed flaviviruses, with a geographic range which includes Africa, Australia, Europe, the Middle East and West Asia. In 1999, WN first established itself in the U.S. In acute severe illness, WN virus can cause hepatitis, meningitis, and encephalitis, lead-

ing to paralysis and coma (Hayes, 1989; Pletnev et al., 2002).

Dengue viruses infect up to 20 million people annually, and are a significant cause of morbidity and mortality in tropical and sub-tropical regions throughout the world (World Health Organization, 1996). There are four serotypes (types 1 to 4) that cause widespread human diseases such as dengue fever, dengue hemorrhagic fever and dengue shock syndrome (Yusof, 2000). There is currently no approved vaccine or effective antiviral therapy for these flaviviruses.

All flaviviruses have a positive-sense nonsegmented RNA genome that encodes a single long polyprotein processed to yield three structural proteins (C, prM, and E) and seven nonstructural proteins (NS1, NS2A, NS2B, NS3, NS4A, NS4B, and NS5) (Leung et al., 2001). A single virus-encoded protease comprising 180 amino acids of NS3 (NS3pro) is responsible for the cleavage of both in *cis* and in *trans* that generates viral proteins that are essential for viral replication and maturation of infectious virions. The presence of a trypsin-like serine protease within the N-terminal one-third of the flavivirus NS3 protein was first proposed by Bazan and Fletterick (1989, 1990) and Gorbalenya and co-workers (Gorbalenya et al., 1989). Their analysis of virus sequence alignments revealed that structural motifs as well as the characteristic catalytic triad (His⁵¹, Asp⁷⁵, and Ser¹³⁵) of mammalian serine proteases were conserved

in all flaviviruses. As NS3pro activity is essential for viral replication, it represents a suitable target for the development of chemotherapeutic approaches for the treatment of flaviviruses. As part of our antiviral program for flaviviruses, we describe in this paper the synthesis and preliminary molecular modeling study of a series of *N*-*t*-Boc amino acid esters of isomannide, designed as a potential inhibitors of the catalytic triad of serine proteases. Isomannide was designed as the center of the peptide mimetic compounds due to its structural analogy with cyclic rigid dipeptides (Bencsik et al., 2003; Dietrich and Lubell, 2003) and its C_2 symmetry, thus providing a rigid scaffold for the peptide mimetic compounds described in the present work.

Material and methods

Computational methods. Computer graphics, structural manipulations, energy minimization and docking calculations were carried out with a Silicon Graphics O2 Workstation (CPU MIPS R10000, processor speed 150 MHz, and main memory 128 MB) using the Insight II 97.0 (Discover) software package, under the operating system IRIX 6.3. Energy minimization and docking calculations were carried out with the Discover 2.9.7 program (Costi and Ferrari, 2001) and Docking module (Kuntz et al., 1982), respectively, available within Insight II, using the molecular mechanics CVFF force field. The enzyme-ligand reference structure is the Dengue virus NS3-serine-protease complexed with Mung-bean Bowman-Birk inhibitor (MbBBI) (PDB code: 1DF9) (Berman et al., 2000; Murthy et al., 2000).

Chemistry. All solvents were purchased as a reagent grade, dried using standard conditions, and stored over molecular sieves. Purification of products was accomplished using silica gel flash chromatography (Whatman 60, 230–400 mesh). Routine NMR analyses were accomplished on a Bruker Advance DPX-400. Melting points were obtained on Thomas-Hoover capillary melting point apparatus and are uncorrected. Mass spectral measurements were made on a Waters 2690 mass spectrometer. Analytical results are within $\pm 0.40\%$ of the theoretical values and were determined by QTI, Whitehouse, NJ.s.

1,4:3,6-Dianhydro-D-mannitol (2). D-mannitol (25 g), HOAc (37 mL) and H_2SO_4 conc. (5 mL) were mixed in a round bottom flask. The mixture was refluxed for 24 h and basified until pH 10 with Na_2CO_3 saturated solution. The mixture was again refluxed for 48 h. Then, it was concentrated in vacuum until a precipitate was formed. Water was added and the solution was extracted with ethyl acetate (continuous extraction/24 h). The organic phases were washed with brine and dried. The solvent was evaporated and the product was obtained as an oil which was distilled under vacuum to afford the product as a white solid in 40% yield.

General procedure for *N*-*tert*-Butoxycarbonyl protection

***N*-*tert*-Butoxycarbonyl-L-alanine (3).** The procedure is similar to the general procedure for **7**. 1H NMR δ ($CDCl_3$, 400 MHz): 9.38 (br s, 1H, –OH), 5.13 (br s, 1H, –NH), 4.31–4.15 (m, 1H, –CH), 1.43 (s, 9H, – CH_3), 1.41 (s, 3H, – CH_3). mp 77–78°C. $\alpha_D^{25} = -24.5$ (c, 2.0) AcOH.

***N*-*tert*-Butoxycarbonyl-L-valine (4).** The procedure is similar to the general procedure for **7**. 1H NMR δ ($CDCl_3$, 400 MHz): 10.25 (br s, 1H, –OH), 5.12 (br s, 1H, –NH), 4.21–4.10 (m, 1H, –CH), 2.16–2.00 (m, 1H, –CH), 1.42 (s, 9H, – CH_3), 0.97–0.91 (m, 6H, – CH_3). mp 75–76°C. $\alpha_D^{25} = +12.9$ (c, 1.0) $CHCl_3$.

***N*-*tert*-Butoxycarbonyl-L-phenylalanine (5).** The procedure is similar to the general procedure for **7**. 1H NMR δ ($CDCl_3$, 400 MHz): (1:2 mixture of rotamers) 8.10 (br s, 1H, –NH), 7.27–7.19 (m, 5H, –CH), 6.21 (br s, 1H, –NH), 4.98 (br s, 1H, –NH), 4.60 (m, 1H, –CH), 3.17–2.90 (m, 2H, – CH_2), 1.40 (s, 6H, – CH_3), 1.30 (s, 3H, – CH_3). mp 68–69°C. $\alpha_D^{27} = +23.2$ (c, 1.0) EtOH.

***N*-1-Bis[*tert*-Butoxycarbonyl]-L-histidine (6).** To a suspension of L-histidine (6.44 mmol, 1 g) and triethylamine (19.33 mmol, 2.7 mL) in water (3.2 mL), a solution of di-*t*-butyldicarbonate (16.1 mol, 3.54 g) in dioxane (3.2 mL) was slowly added. After the mixture had been stirred overnight at room temperature, water (6 mL) was added and the aqueous phase was washed with diethyl ether (3×50 mL) and ethyl acetate (3×50 mL). The total volume was increased to 200 mL with ethyl acetate and the solution acidified with 10% chloridric acid until it reached pH 3. The aqueous phase was poured off and extracted with ethyl acetate (100 mL). The organic phases were washed with brine and dried. The solvent was evaporated and the product was obtained (1.65 g). 1H NMR δ ($CDCl_3$, 400 MHz): 8.15 (s, 1H, –CH), 7.19 (s, 1H, –CH), 5.47 (br s, 1H, –NH), 4.48 (s, 1H, –CH), 3.20–3.08 (m, 2H, – CH_2), 1.59 (s, 9H, – CH_3), 1.46 (s, 9H, – CH_3). mp 165–166°C. $\alpha_D^{27} = +95.5$ (c, 1.0) $CHCl_3$.

***N*-*tert*-Butoxycarbonyl-L-proline (7).** L-Proline (17.39 mmol, 2 g) and sodium hydroxide (20.1 mmol, 0.807 g) were dissolved in *t*-butanol-water mixture (30 mL, 1:1, v/v). To this solution, di-*t*-butyldicarbonate (21.1 mmol, 4.36 g) was added in one portion. The reaction mixture was stirred at rt overnight. The aqueous layer was treated with aqueous acetic acid (2 mL in 11 mL H_2O) and, then, extracted with ethyl acetate (3×70 mL). The combined extracts were washed with water and dried. The crude product was recrystallised from chloroform-hexane at 0°C to afford the compound (2.27 g). 1H NMR δ ($CDCl_3$, 400 MHz): 9.24 (br s, 1H, –OH), 4.27–4.20 (m, 1H, –CH), 3.50–3.48 (m, 2H, – CH_2), 2.12–1.84 (m, 4H, – CH_2), 1.43 (s, 9H, – CH_3). mp 120–122°C. $\alpha_D^{27} = -85.5$ (c, 1.0) $CHCl_3$.

***N*-*tert*-Butoxycarbonyl-L-glutamic acid 5-methyl ester (8).** To a stirred solution of L-glutamic acid 5-methyl ester (6.20 mmol, 1 g) in water (18 mL) $NaHCO_3$ was added (13.6 mmol, 1.14 g) at 0°C, followed by di-*t*-butyl dicarbonate (6.57 mmol, 1.42 g) in dioxane (18 mL). The reaction mixture was stirred at rt overnight. The solution was extracted with ethyl ether (3×20 mL). The aqueous phase was acidified with 10% chloridric acid until it reached pH 3. The product was extracted by ethyl acetate and the organic phase washed with brine and dried. The solvent was evaporated and the product was obtained (1.56 g). 1H NMR δ ($CDCl_3$, 400 MHz): 8.74 (br s, 1H, –NH), 4.33–4.22 (m, 1H, –CH), 3.70 (s, 3H, – CH_3), 2.48–2.03 (m, 4H, – CH_2), 1.43 (s, 9H, – CH_3). mp 75–76°C. $\alpha_D^{27} = +8.4$ (c, 1.0) $CHCl_3$.

O-Benzyl-L-Pyroglutamic acid (9a). To a solution of L-Pyroglutamic acid (1 g, 7.74 mmol) in benzene (20 mL), benzyl alcohol (1.12 mL, 10.83 mmol), and *p*-TsOH (cat.) were added. The mixture was refluxed in a Dean-Stark apparatus overnight. Ethyl acetate was added to the organic layer and washed with 5% $NaHCO_3$ (3×50 mL). The solvent was evaporated under vacuum to obtain 1.66 g of the crude product. Purification by column chromatography gives the product (0.9 g) as oil in 51% yield. 1H NMR δ ($CDCl_3$, 400 MHz): 7.35–7.33 (m, 5H, –CH), 6.76 (br s, 1H, –NH), 5.17 (s, 2H, – CH_2), 4.27–4.24 (m, 1H, –CH), 2.48–2.25 (m, 4H, – CH_2). ^{13}C NMR: 178.2 (–C=O), 172.0 (C=O), 135.1 (–Cq), 128.7, 128.6, 128.3 (–CH), 67.2 (– CH_2), 55.6 (–CH), 29.3 (– CH_2), 24.7 (– CH_2). ESMS m/z 218 ($M - 1$). $\alpha_D^{27} = -1.6$ (c, 1.0) $CHCl_3$.

O-Benzyl-N-*tert*-Butoxycarbonyl-L-Pyroglutamic acid (9b). To a solution of O-Benzyl-Pyroglutamic acid (**9a**) (5.07 mmol, 1.17 g) in dry THF (25 mL), at –78°C, a NaHMDS 1M solution in THF (5.31 mmol, 5.31 mL) was added. The mixture was stirred for 20 min. A solution of di-*t*-butyldicarbonate (7.6 mmol, 1.66 g) in THF (10 mL) was added, and the mixture was allowed to stir for 20 min more at low temperature. The solvent was evaporated, ethyl acetate was added and the organic phase washed with saturated ammonium chloride. The solvent was evaporated

under vacuum to obtain 1.86 g of the crude product. Purification by column chromatography gives the product (1.36 g) in 81.4% yield. ^1H NMR δ (CDCl_3 , 400 MHz): 7.38–7.32 (m, 5H, $-\text{CH}$), 5.20 (s, 2H, $-\text{CH}_2$), 4.65 (dd, $J = 2.8\text{ Hz}$, $J = 9.6\text{ Hz}$, 1H, $-\text{CH}$), 2.65–2.35 (m, 4H, $-\text{CH}_2$), 1.40 (s, 9H, $-\text{CH}_3$). ^{13}C NMR: 173.3 ($-\text{C}=\text{O}$), 171.1 ($-\text{C}=\text{O}$), 149.2 ($-\text{C}=\text{O}$), 135.0 ($-\text{Cq}$), 128.7, 128.6, 128.5 ($-\text{CH}$), 83.6 ($-\text{Cq}$), 67.3 ($-\text{CH}_2$), 58.9 ($-\text{CH}$), 31.1 ($-\text{CH}_2$), 27.8 ($-\text{CH}_3$), 21.5 ($-\text{CH}_2$). ESMS m/z 318 ($\text{M} - 1$). $\alpha_{\text{D}}^{26} = -34.8$ (c, 1.0) CH_2Cl_2 , $\alpha_{\text{D}}^{\text{lit}} = -31.9$ (c, 1.0) CH_2Cl_2 .

L-*N*-tert-Butoxycarbonylpyroglutamic acid (9). A solution of *O*-Benzyl-*N*-tert-Butoxycarbonyl-L-Pyroglutamic acid (**9b**) (4.23 mmol, 1.35 g) in ethyl acetate (30 mL) and 10% Pd/C (0.310 g) was vigorously shaken under an atmosphere of H_2 (40 psi) for 1 h. The catalyst was filtered off and the solution was concentrated affording the crude product. Purification by column chromatography gives the product (0.872 g) in 90% yield. ^1H NMR δ (CDCl_3 , 400 MHz): 8.11 (br s, 1H), 4.66–4.63 (m, 1H, $-\text{CH}$), 2.70–2.36 (m, 4H, $-\text{CH}_2$), 1.49 (s, 9H, $-\text{CH}_3$). ^{13}C NMR: 175.8 ($-\text{C}=\text{O}$), 174.0 ($-\text{C}=\text{O}$), 149.3 ($-\text{C}=\text{O}$), 84.0 ($-\text{Cq}$), 58.7 ($-\text{CH}$), 31.1 ($-\text{CH}_2$), 27.8 ($-\text{CH}_3$), 21.4 ($-\text{CH}_2$). ESMS m/z 228 ($\text{M} - 1$). mp 111–112°C. $\alpha_{\text{D}}^{29} = -21.3$ (c, 1.0) CHCl_3 .

General procedure for DCC-mediated condensation: To a solution of isomannide (2 mmol) and the corresponding protected amino acid (4 mmol) in dry CH_2Cl_2 (10 mL) DCC (4 mmol, 0.605 g) was added at 0°C. The mixture was stirred at room temperature overnight, and the solvent was evaporated. Purification by column chromatography gives the corresponding products in around 40% yield.

(3*R*,6*R*)-bis-*O*-(*N*-tert-Butoxycarbonyl-L-alanine)-1,4-dioxabicyclo[3.3.0]octane (10a). ^1H NMR δ (CDCl_3 , 400 MHz): 5.10 (br s, 2H, $-\text{CH}$), 4.69 (s, 2H, $-\text{CH}$), 4.37 (br s, 2H, $-\text{CH}$), 4.03–4.01 (m, 2H, $-\text{CH}_2$), 3.81–3.77 (m, 2H, $-\text{CH}_2$), 1.43 (s, 18H, $-\text{CH}_3$), 1.39 (d, 6H, $J = 7.18\text{ Hz}$, $-\text{CH}_3$). ^{13}C NMR: 173.0 ($-\text{C}=\text{O}$), 155.3 ($-\text{C}=\text{O}$), 80.6 ($-\text{CH}$), 80.2 ($-\text{Cq}$), 74.6 ($-\text{CH}$), 70.8 ($-\text{CH}_2$), 49.6 ($-\text{CH}$), 28.7 ($-\text{CH}_3$), 19.0 ($-\text{CH}_3$). ESMS m/z 489 ($\text{M} + 1$). IR (KBr) $\nu\text{ cm}^{-1}$: 3347, 2979, 2888, 1747, 1683, 1517, 1303, 1252, 1160, 1074, 978. Anal. Calcd. for $\text{C}_{22}\text{H}_{36}\text{N}_2\text{O}_{10}$: C, 54.09; H, 7.43; N, 5.73. Found: C, 53.96; H, 7.56; N, 5.36. $\alpha_{\text{D}}^{27} = +89.0$ (c, 1.0) CHCl_3 , mp = 105–106°C.

(3*R*,6*R*)-bis-*O*-(*N*-tert-Butoxycarbonyl-L-phenylalanine)-1,4-dioxabicyclo[3.3.0]octane (10b). ^1H NMR δ (CDCl_3 , 400 MHz): 7.30–7.13 (m, 10H, $-\text{CH}$), 5.04–5.01 (m, 2H, $-\text{CH}$), 4.65 (s, 2H, $-\text{CH}$), 4.18 (m, 2H, $-\text{CH}$), 3.94–3.90 (m, 2H, $-\text{CH}_2$), 3.64–3.62 (m, 2H, $-\text{CH}_2$), 3.10 (m, 4H, $-\text{CH}_2$), 1.41 (s, 18H, $-\text{CH}_3$). ^{13}C NMR: 171.7 ($-\text{C}=\text{O}$), 155.3 ($-\text{C}=\text{O}$), 136.3 ($-\text{Cq}$), 129.7 ($-\text{CH}$), 128.9 ($-\text{CH}$), 127.4 ($-\text{CH}$), 80.6 ($-\text{CH}$), 80.3 ($-\text{Cq}$), 74.6 ($-\text{CH}$), 70.5 ($-\text{CH}_2$), 54.8 ($-\text{CH}$), 38.8 ($-\text{CH}_2$), 28.7 ($-\text{CH}_3$). ESMS m/z 641 ($\text{M} + 1$). IR (KBr) $\nu\text{ cm}^{-1}$: 2976, 2931, 1750, 1710, 1624, 1500, 1451, 1365, 1247, 1163, 1055. Anal. Calcd. for $\text{C}_{34}\text{H}_{44}\text{N}_2\text{O}_{10}$: C, 63.74; H, 6.92; N, 4.37. Found: C, 63.39; H, 7.01; N, 4.22. $\alpha_{\text{D}}^{27} = +105.3$ (c, 1.0) CHCl_3 , mp = 55–56°C.

(3*R*,6*R*)-bis-*O*-(*N*-tert-Butoxycarbonyl-L-valine)-1,4-dioxabicyclo[3.3.0]octane (10c). ^1H NMR δ (CDCl_3 , 400 MHz): 5.04–5.00 (m, 2H, $-\text{CH}$), 4.66 (s, 2H, $-\text{CH}$), 4.25 (br s, 2H, $-\text{CH}$), 3.99–3.95 (m, 2H, $-\text{CH}_2$), 3.76–3.72 (m, 2H, $-\text{CH}_2$), 2.15–1.98 (m, 2H, $-\text{CH}$), 1.44 (s, 18H, $-\text{CH}_3$), 0.90 (dd, 12H, $J = 6.7\text{ Hz}$, $J = 28.3\text{ Hz}$, $-\text{CH}_3$). ^{13}C NMR: 172.1 ($-\text{C}=\text{O}$), 155.9 ($-\text{C}=\text{O}$), 80.6 ($-\text{CH}$), 80.1 ($-\text{Cq}$), 74.6 ($-\text{CH}$), 70.7 ($-\text{CH}_2$), 58.9 ($-\text{CH}$), 31.6 ($-\text{CH}$), 28.7 ($-\text{CH}_3$), 19.3 ($-\text{CH}_3$), 17.9 ($-\text{CH}_3$). ESMS m/z 545 ($\text{M} + 1$). IR (CH_2Cl_2) $\nu\text{ cm}^{-1}$: 2970, 2932, 1742, 1710, 1504, 1365, 1308, 1246, 1162, 1095, 1017. Anal. Calcd. for $\text{C}_{26}\text{H}_{44}\text{N}_2\text{O}_{10}$: C, 57.34; H, 8.14; N, 5.14. Found: C, 57.33; H, 8.20; N, 5.42. $\alpha_{\text{D}}^{27} = +85.1$ (c, 1.0) CHCl_3 . The product is a colorless oil.

(3*R*,6*R*)-bis-*O*-(*N*-tert-Butoxycarbonyl-L-proline)-1,4-dioxabicyclo[3.3.0]octane (10d). ^1H NMR δ (CDCl_3 , 400 MHz): 5.12–5.10 (m, 2H, $-\text{CH}$), 4.68–4.67 (s, 2H, $-\text{CH}$), 4.28–4.27 (m, 2H, $-\text{CH}$), 4.08–4.07 (m, 2H, $-\text{CH}_2$), 3.94–3.92 (m, 2H, $-\text{CH}_2$), 3.54–3.45 (m, 4H, $-\text{CH}_2$), 2.21–1.83 (m, 8H, $-\text{CH}_2$), 1.43 (s, 18H, $-\text{CH}_3$). ^{13}C NMR: 172.8 ($-\text{C}=\text{O}$), 154.2 ($-\text{C}=\text{O}$), 82.1 ($-\text{CH}$), 80.6 ($-\text{Cq}$), 74.6 ($-\text{CH}$), 71.1 ($-\text{CH}_2$), 59.2 ($-\text{CH}$), 46.9 ($-\text{CH}_2$), 28.8 ($-\text{CH}_3$), 25.3 ($-\text{CH}_2$), 23.8 ($-\text{CH}_2$). ESMS m/z 539 ($\text{M} - 1$). Anal. Calcd. for $\text{C}_{26}\text{H}_{40}\text{N}_2\text{O}_{10}$: C, 57.76; H, 7.46; N, 5.18. Found:

C, 57.40; H, 7.08; N, 4.89. $\alpha_{\text{D}}^{27} = +16.3$ (c, 1.0) CHCl_3 . The product is colorless oil.

(3*R*,6*R*)-bis-*O*-(*N*-1-Bis[*tert*-Butoxycarbonyl]-L-histidine)-1,4-dioxabicyclo[3.3.0]octane (10e). ^1H NMR δ (CDCl_3 , 400 MHz): δ 7.96 (s, 2H, $-\text{CH}$), 7.15 (s, 2H, $-\text{CH}$), 5.68–5.66 (m, 2H, $-\text{CH}$), 5.08 (s, 2H, $-\text{CH}$), 4.65 (s, 2H, $-\text{CH}$), 3.99–3.98 (m, 2H, $-\text{CH}_2$), 3.80–3.75 (m, 2H, $-\text{CH}_2$), 3.08 (br s, 4H, $-\text{CH}_2$), 1.59 (s, 18H, $-\text{CH}_3$), 1.42 (s, 18H, $-\text{CH}_3$). ^{13}C NMR: 171.6 ($-\text{C}=\text{O}$), 155.6 ($-\text{C}=\text{O}$), 147.2 ($-\text{C}=\text{O}$), 138.8 (Cq), 137.2 ($-\text{CH}$), 115.1 ($-\text{CH}$), 85.9 ($-\text{Cq}$), 80.6 ($-\text{CH}$), 80.1 ($-\text{Cq}$), 74.7 ($-\text{CH}$), 70.6 ($-\text{CH}_2$), 53.5 ($-\text{CH}$), 30.6 ($-\text{CH}_2$), 28.6 ($-\text{CH}_3$), 28.2 ($-\text{CH}_3$). ESMS m/z 821 ($\text{M} + 1$). IR (KBr) $\nu\text{ cm}^{-1}$: 3388, 2979, 2932, 1755, 1712, 1490, 1391, 1289, 1254, 1159, 1055, 1011, 843. Anal. Calcd. for $\text{C}_{38}\text{H}_{56}\text{N}_6\text{O}_{14}$: C, 55.60; H, 6.88; N, 10.24. Found: C, 55.25; H, 7.08; N, 10.01. $\alpha_{\text{D}}^{21} = +72.2$ (c, 1.0) CHCl_3 , mp = 74–75°C.

(3*R*,6*R*)-bis-*O*-(*N*-tert-Butoxycarbonyl-L-glutamic acid 5-methyl ester)-1,4-dioxabicyclo[3.3.0]octane (10f). ^1H NMR δ (CDCl_3 , 400 MHz): δ 5.15–5.07 (m, 2H, $-\text{CH}$), 4.70 (s, 2H, $-\text{CH}$), 4.40 (br s, 2H, $-\text{CH}$), 4.03 (m, 2H, $-\text{CH}_2$), 3.81 (m, 2H, $-\text{CH}_2$), 3.67 (s, 6H, $-\text{OCH}_3$), 2.46–2.36 (m, 4H, $-\text{CH}_2$), 2.21–2.18 (m, 2H, $-\text{CH}_2$), 2.02–1.95 (m, 2H, $-\text{CH}_2$), 1.43 (s, 18H, $-\text{CH}_3$). ^{13}C NMR: δ 173.5 ($-\text{C}=\text{O}$), 171.8 ($-\text{C}=\text{O}$), 155.6 ($-\text{C}=\text{O}$), 80.5 ($-\text{CH}$), 80.2 ($-\text{Cq}$), 74.6 ($-\text{CH}$), 70.7 ($-\text{CH}_2$), 53.2 ($-\text{CH}$), 52.0 ($-\text{OCH}_3$), 30.2 ($-\text{CH}_2$), 28.5 ($-\text{CH}_3$), 27.8 ($-\text{CH}_2$). ESMS m/z 633 ($\text{M} + 1$). IR (CH_2Cl_2) $\nu\text{ cm}^{-1}$: 3349, 2977, 2931, 1738, 1713, 1632, 1583, 1515, 1445, 1366, 1251, 1166. Anal. Calcd. for $\text{C}_{28}\text{H}_{44}\text{N}_2\text{O}_{14}$: C, 51.65; H, 6.67; N, 4.63. Found: C, 52.01; H, 6.98; N, 4.66. $\alpha_{\text{D}}^{21} = +91.9$ (c, 1.0) CHCl_3 . The product is a colorless oil.

(3*R*,6*R*)-bis-*O*-(*N*-tert-Butoxycarbonyl-L-pyroglutamic acid)-1,4-dioxabicyclo[3.3.0]octane (10g). ^1H NMR δ (CDCl_3 , 400 MHz): 5.13–5.11 (m, 2H, $-\text{CH}$), 4.74–4.73 (m, 2H, $-\text{CH}$), 4.68–4.64 (m, 2H, $-\text{CH}$), 4.03–3.98 (m, 2H, $-\text{CH}_2$), 3.76–3.74 (m, 2H, $-\text{CH}_2$), 2.66–2.34 (m, 8H, $-\text{CH}_2$), 1.48 (s, 18H, $-\text{CH}_3$). ^{13}C NMR: 173.5 ($-\text{C}=\text{O}$), 170.6 ($-\text{C}=\text{O}$), 149.2 ($-\text{C}=\text{O}$), 83.8 ($-\text{Cq}$), 80.2 ($-\text{CH}$), 74.4 ($-\text{CH}$), 70.1 ($-\text{CH}_2$), 58.9 ($-\text{CH}$), 31.1 ($-\text{CH}_2$), 27.8 ($-\text{CH}_3$), 21.8 ($-\text{CH}_2$). ESMS m/z 569 ($\text{M} + 1$). Anal. Calcd. for $\text{C}_{26}\text{H}_{36}\text{N}_2\text{O}_{12}$: C, 53.45; H, 6.38; N, 4.93. Found: C, 53.16; H, 6.40; N, 5.25. $\alpha_{\text{D}}^{27} = +21.3$ (c, 1.0) CHCl_3 .

Results and discussion

Chemistry

The start material for the synthesis of the isomannide esters was D-mannitol (**1**). Ready dehydration of naturally occurring **1** provided the symmetrical molecule, 1,4:3,6-dianhydro-D-mannitol, the isomannide (**2**) (Wiggins, 1945). Its structure was resolved by X-ray diffraction analysis to assure that no isomerization took place in the acidic cyclization step. Ester formation proved to be very difficult to achieve. Initial attempts to obtain the isomannide esters with the Boc-protected L-amino acids, **3–9** (described in Table 1), in the presence of *p*-toluenesulfonic acid in benzene were unsuccessful (Gomurashvili, 2000). The L-amino acids were protected according to literature (Benoiton et al., 1993; Ye and McKerver, 1992; Gołębowski et al., 1987; Barcelo et al., 1986; Le Nguyen et al., 1985; Keller et al., 1985; Ookawa and Soai, 1987; Feng and Edstrom, 1999; El Marini et al., 1992; Li et al., 1995; Yoshifuji et al., 1986). In view of these results, other amino acid esterification

Table 1. Boc-amino acids prepared by acylation with Di-*t*-Butyl dicarbonate

No.	Boc-amino acids	Conditions	mp, °C	α_D^{25}
3	Boc-Ala	NaOH, <i>t</i> BuOH/H ₂ O	77–78	–24.5 (AcOH, <i>c</i> 2.0)
4	Boc-Val	NaOH, <i>t</i> BuOH/H ₂ O	75–76	+12.9 (CHCl ₃ , <i>c</i> 1.0)
5	Boc-Phe	NaOH, <i>t</i> BuOH/H ₂ O	68–69	+23.2 (EtOH, <i>c</i> 1.0)
6	Boc-His	Et ₃ N, dioxane/H ₂ O	165–166	+95.5 (CHCl ₃ , <i>c</i> 1.0)
7	Boc-Pro	NaOH, <i>t</i> BuOH/H ₂ O	120–122	–85.5 (CHCl ₃ , <i>c</i> 1.0)
8	Boc-OMeGlu	NaHCO ₃ , dioxane/H ₂ O	75–76	+8.4 (CHCl ₃ , <i>c</i> 1.0)
9	Boc-PyroGlu	NaHMDS, THF	111–112	–21.3 (CHCl ₃ , <i>c</i> 1.0)

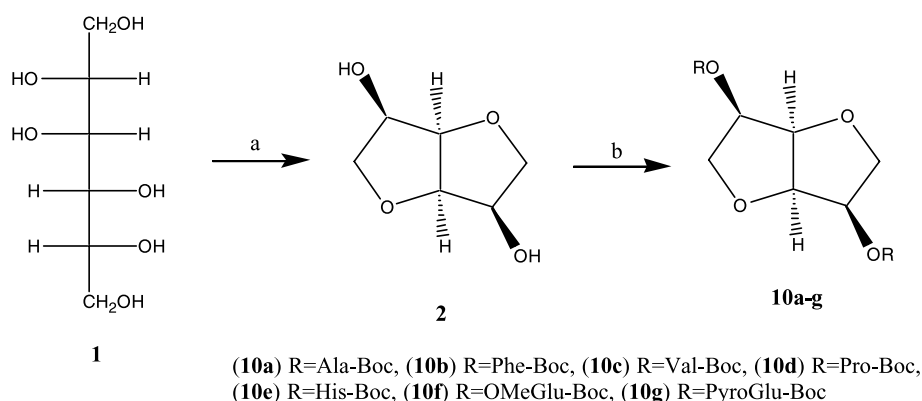
methods were attempted, i.e., as the activation of the carboxylic acid by formation of a mixed anhydride with isobutyl chloroformate (Kim et al., 1985; Muri et al., 2000), and the treatment of amino acids with *N,N*-bis-(2-oxooxazolidin-3-yl)phosphorodiamidic chloride (BOP-Cl) in pyridine (Herranz et al., 1991). Unfortunately, these methodologies led to negative results. The best result one was achieved by the *N,N'*-dicyclohexylcarbodiimide (DCC)-mediated condensation of Boc-protected amino acids with isomannide (Benoiton et al., 1993; Barany and Albericio, 1995). In this manner, compounds **10a–g** were obtained free of racemization (Scheme 1). The structural assignment of all the esters of amino acids was based on analytical and spectroscopic data.

Molecular modeling

The Mung-bean Bowman-Birk inhibitor (MbBBI) fragment (nine residues from Cys⁵¹⁸ to Cys⁵²⁶), (Murthy, 2000), was docked into the active site of the Dengue virus NS3 protease showing six hydrogen bond interactions between the NS3 protease and the MbBBI fragment (Fig. 1A/B). The observed hydrogen bonds are between the following pairs of residues, Gln³⁵...Gln⁵²⁵ (2.17 Å), His⁵¹...Thr⁵¹⁹ (2.41 Å), Gly¹³³...Lys⁵²⁰ (2.06 Å), Ser¹³⁵...Lys⁵²⁰ (1.77 Å),

Asn¹⁵²...Lys⁵²⁰ (1.91 Å), and Gly¹⁵³...Cys⁵¹⁸ (2.41 Å), from NS3 protease and MbBBI fragment, respectively. It is to be noted that, as in the crystallographic structure of NS3-MbBBI complex, the most extensive interactions occur at the P1 site, where the Lys⁵²⁰ makes three hydrogen bond interactions. Although these interactions do not present the same pattern as in the crystallographic structure, they are closer to the same residues. The residues around (5 Å), the oxygen atom of the Lys⁵²⁰ carbonyl group, are Pro¹³², Gly¹³³, Thr¹³⁴, and Ser¹³⁵; and the residues around (5 Å), the nitrogen atom of the Lys⁵²⁰ amine group, are Tyr¹⁵⁰, Gly¹⁵¹, Gly¹⁵³, and Ser¹⁶³. Overall, comparing the crystallographic NS3-MbBBI complex with the docked structure of NS3 and MbBBI fragment, it may be pointed, therefore, that the fragment of the MbBBI was able to maintain the same orientation in the active site of NS3 protease and the principal interactions in the docked model, therefore validating the proposed model of the NS3 docked with our compound.

The most simple compound of our series, the Ala-Boc ester **10a**, was docked into the active site of the NS3 protease, showing three hydrogen bond interactions with the NS3 protease (Fig. 2A/B). One hydrogen bond is between the backbone NH group of Gly¹³³ and one oxygen atom of the bicycle group (1.88 Å), that corresponds

**Scheme 1.** a HOAc/H₂SO₄; b Boc-amino acids (3–9), DCC, CH₂Cl₂

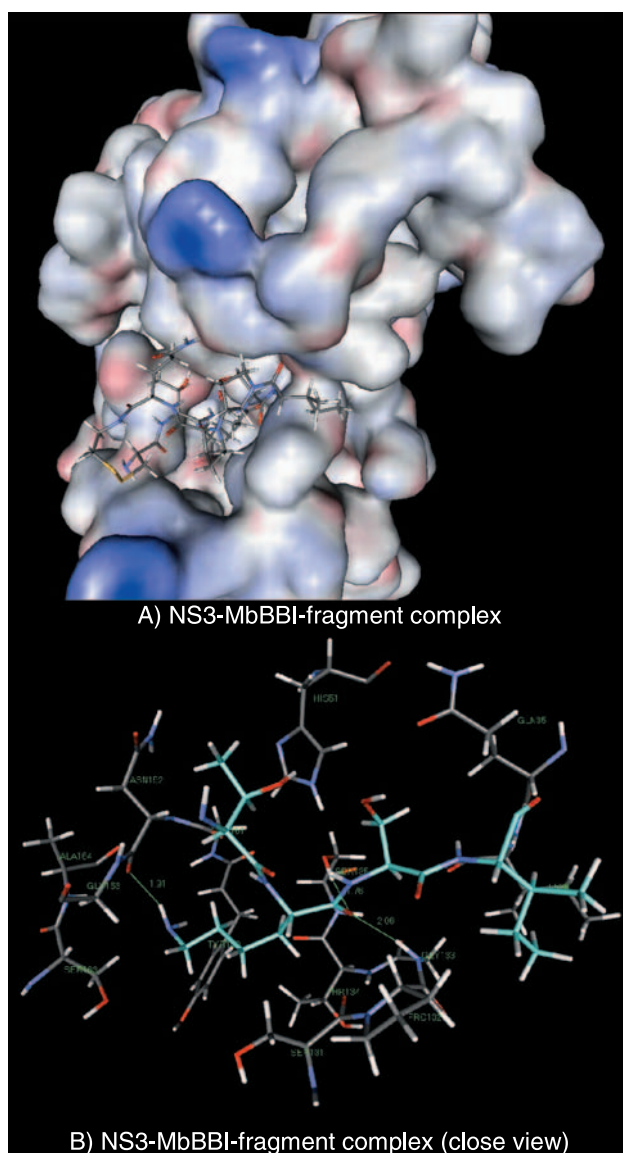


Fig. 1. **A** Docked structure of MbBBI-fragment into the NS3-protease active site, where the enzyme is represented with a solid surface color by electrostatic potential. **B** Close view of this complex representing only the residues P2'-P1'-P1-P2 of MbBBI-fragment and the aminoacids residues no more distant than 5 Å from this tetrapeptide fragment. The carbon atoms of the MbBBI tetrapeptide are colored in light blue

to the P1' position of compound **10a**, according to Fig. 3. The other two hydrogen bonds correspond to the P1 position: i.e., the Ala residue from compound **10a**, where one of them includes the hydroxyl group of Ser¹³⁵ (catalytic residue) and the carbonyl oxygen atom of Ala residue (1.68 Å), while the other includes the carbonyl oxygen atom of Asn¹⁵² and NH group of Ala residue (1.96 Å). There are three residues, namely, Ile³⁶, Leu¹⁰⁰, and Pro¹³², potentially capable of providing specific van der Waals interactions stabilizing the complex NS3 protease-com-

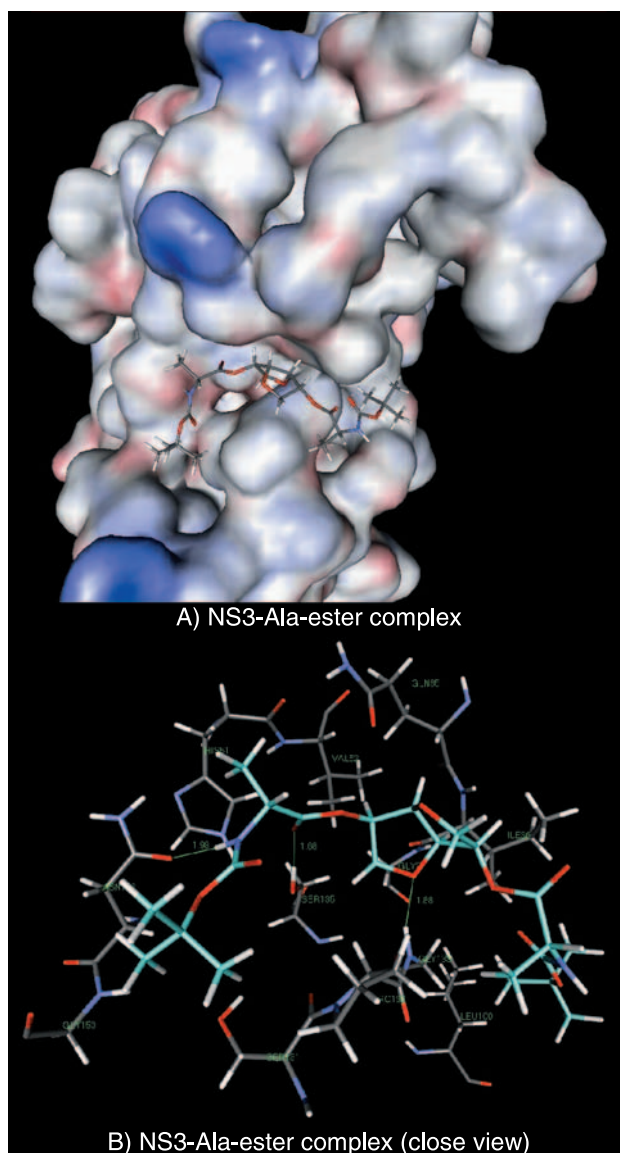


Fig. 2. **A** Docked structure of compound **10a** into the NS3-protease active site, where the enzyme is represented with a solid surface color by electrostatic potential. **B** Close view of this complex representing only the aminoacids residues no more distant than 5 Å from this tetrapeptide fragment. The carbon atoms of the compound **10a** are colored in light blue

pound **10a**. In addition, the catalytic His⁵¹ and residues Gln³⁵, Ser¹³¹, and Gly¹⁵³ make nonspecific interactions. It is clear that the putative ligand, the Ala-Boc ester compound, is well accommodated into the active site of the NS3 protease, although it should be pointed out that it does not have at P1 position a basic residue: e.g., Arg or Lys. Therefore, we are planning to obtain a new series of compounds showing basic residues in order to reach Tyr¹⁵⁰ residue, that stays at the bottom of the cleft at S1 position, as well as polypeptide derivatives.

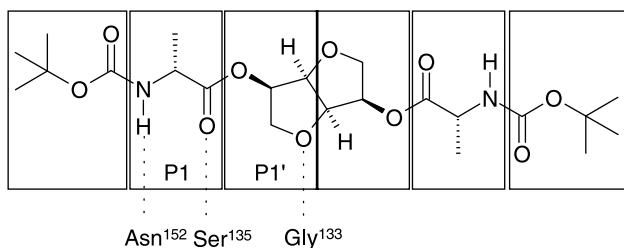


Fig. 3. Schematic hydrogen bonding representation between the compound **10a** and the residues of NS3-protease after docking calculations

As part of our antiviral program (Peçanha et al., 2003), we described here the synthesis of a new series of *N*-*t*-Boc amino acid esters of isomannide, designed as potential inhibitors of serine proteases. In order to have a model to validate the proposed model of NS3 protease docked with our compound, a preliminary molecular modeling study was developed using a peptide inhibitor (MbBBI) from the literature, the bioassay of the new compounds described here in the dengue virus test are underway.

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