

Article

Structures of Benzenesulfonylamino-3-(4-benzenesulfonyloxy-phenyl)-propionic acid and 2-(toluene-4-sulfonylamino)-3-[4-(toluene-4-sulponyl-oxy)-phenyl]-propionic acid: Variations in L-tyrosine Backbone Conformation, Intramolecular Aromatic π - π Stacking and Short C-H···O Interactions

Muneeb Hayat Khan 1,* , Islam Ullah Khan 1,* , Muhammad Nadeem Arshad 2 , H. M. Rafique 2 and William T. A. Harrison 3,*

- ¹ Materials Chemistry laboratory, Department of Chemistry, Government College University, Lahore-54000, Pakistan
- ² X-ray Diffraction and Crystallography Laboratory, Department of Physics, School of Physical Sciences, University of the Punjab, Quaid-e-Azam Campus, Lahore-54590, Pakistan
- ³ Department of Chemistry, University of Aberdeen, Aberdeen AB24 3UE, Scotland, UK
- * Authors to whom correspondence should be addressed; E-Mails: koolmuneeb@yahoo.com (M.H.K.); iukhan@gcu.edu.pk (I.U.K.); w.harrison@abdn.ac.uk (W.T.A.H.); Tel.: +92-331-4248092 (M.H.K.); +92-42-111-000-010 Ext. 262 (I.U.K.); +44-1224-272897 (W.T.A.H.).

Received: 26 April 2011; in revised form: 8 June 2011 / Accepted: 10 June 2011 /

Published: 14 June 2011

Abstract: The syntheses and crystal structures of benzenesulfonylamino-3-(4-benzenesulfonyloxy-phenyl)-propionic acid (1) and 2-(toluene-4-sulfonylamino)-3-[4-(toluene-4-sulponyloxy)-phenyl]-propionic acid (2) are described. The L-tyrosine cores of the molecules show significant conformational differences. In 1, both organic molecules show intramolecular aromatic π - π stacking and in 2 a very short intermolecular C^{α} -H···O interaction is seen. The structures of 1 and 2 are compared with those of related materials. Crystal data: 1₂·H₂O·MeOH [2(C₂₁H₁₉NO7S₂)·H₂O·CH₄O], M_r = 973.04, monoclinic, P2₁ (No. 4), a = 8.0078 (4) Å, b = 34.0704 (16) Å, c = 8.5506 (3) Å, β = 94.239 (3)°, V = 2326.47 (18) ų, Z = 2, T = 296 K, R(F) = 0.062, $wR(F^2)$ = 0.157, 2·H₂O (C₂₃H₂₅NO7S₂·H₂O), M_r = 507.56, monoclinic, P2₁ (No. 4), a = 5.7171 (7) Å, b = 24.359 (3) Å, c = 9.1043 (10) Å, β = 104.563 (6)°, V = 1227.2 (2) ų, Z = 2, T = 296 K, R(F) = 0.055, $wR(F^2)$ = 0.092.

Keywords: L-tyrosine; green chemistry; torsion angles; C–H···O interactions

1. Introduction

Tyrosine ($C_9H_{11}NO_3$) is one of the non-essential amino acids, which has been found as a constituent of naturally occurring proteins [1,2]. The crystal structures of chiral L-tyrosine and racemic DL-tyrosine were reported nearly 40 years ago, with both found to crystallize in their zwitterionic forms. The chiral form (space group $P2_12_12_1$) [3] features intermolecular N–H···O_c and N–H···O_h (c = carboxyl, h = hydroxyl) hydrogen bonds as well as O–H···O_h links, which generate a three-dimensional network in the crystal. Although it was not identified at the time, a very short intermolecular C–H···O_c link (H···O = 2.36 Å, compared to the van der Waals' contact distance of about 2.72 Å) arising from the stereogenic (chiral) α carbon atom also occurs [3]. This is now recognized as an important general interaction in amino acids, peptides and even proteins [4,5]. There is no possibility of aromatic π – π stacking in L-tyrosine, as the minimum separation of aromatic ring centroids is greater than 4.51 Å. In the racemic DL-tyrosine crystal structure [6] (space group $Pna2_1$), the zwitterions are linked by the same types of hydrogen bond, although the N–H···O_h bond is notably longer (H···O = 2.31 Å). Again, a three-dimensional array of molecules is generated by the hydrogen bonds and an intermolecular C–H···O_c link also occurs: the equivalent α C atom is again involved, although the H···O separation of 2.54 Å is significantly longer than that in the chiral form.

The crystal structures of various acylated L-tyrosine derivatives (containing C–N–C bonds) have been reported, including N-acetyl-L-tyrosine [7], N-acetyl-L-phenylalanyl-L-tyrosine [8] and (R)-2-(2-(5-fluoro-2,4-dioxo-3,4-dihydropyrimidin-1(2H)-yl)acetamido)-3-(4-hydroxyphenyl)-propanoic acid dimethylformamide solvate [9]. It is notable that intermolecular C–H···O interactions from the α -carbon atom occur in the first and third of these, with H···O = 2.57 and 2.30 Å, respectively.

In this paper, we report the syntheses and crystal structures of benzenesulfonylamino-3-(4-benzenesulfonyloxy-phenyl)-propionic acid (1) (see Figure 1 below) and 2-(toluene-4-sulfonylamino)-3-[4-(toluene-4-sulponyloxy)-phenyl]-propionic acid (2), in which the L-tyrosine core is substituted at both its N and O_h atoms. Their structures are compared to those of the related compounds noted above.

Figure 1. Structures of 1 and 2

(1)

2. Results and Discussion

2.1. Structure of 1

The asymmetric unit of $\mathbf{1}_2$ ·H₂O·MeOH consists of two organic molecules (**1A** containing N1 and **1B** containing N2), accompanied by one molecule of water and one molecule of methanol, which is consistent with the mixed solvent used for recrystallization (see Experimental section). **1** has arisen from the reaction of L-tyrosine with two molecules of benzene sulfonyl chloride to form new S–N and S–O bonds in the product [10]. It is notable that there are no other compounds containing L-tyrosine with an equivalent S–N bond in the Cambridge Structural Database (version 5.32 of November 2010 with two updates) [11]. The transferrable hydrogen atoms (acidic protons) are clearly located on atoms O4 and O12, *i.e.* the molecules of **1** are not zwitterions. The absolute structure of $\mathbf{1}_2$ ·H₂O·MeOH is well established based on the refined value of -0.03 (8) of the Flack absolute structure parameter [12] and the *S* configurations of the stereogenic atoms in **1** (C6 in molecule A and C27 in molecule B) are consistent with that of the equivalent carbon atom in L-tyrosine itself [3]. In order to simplify the comparison of **1** with related structures, we will designate atom-sets C13, C6, C5 and C4 (molecule A) and C34, C27, C26 and C25 (molecule B) as Ca, C α , C β and C γ , respectively.

S–N–C α –C β or C–N–C α –C β					
1a	140.5 (4)	1b	146.9 (4)		
2	129.7 (3)	3	-176.6		
4	-171.1	5	-68.6		
Ν–Cα–Cβ–Cγ					
1a	-67.6 (6)	1b	-59.7 (6)		
2	-61.1 (5)	3	-71.5		
4	-179.8	5	-60.5		
6	69.3	7	-69.7		
Са–Сα–Сβ–Сγ					
1a	170.2 (5)	1b	178.3 (5)		
2	174.4 (4)	3	165.6		
4	58.2	5	64.6		
6	-53.1	7	52.5		
N–Cα–Ca–OH					
1a	166.0 (5)	1b	-37.4 (6)		
2	165.3 (3)	3	-28.7		
4	-46.9	5	170.0		

Table 1. Key torsion angles (°) in **1**, **2** and related compounds.

Key torsion angles for the amino-acid cores of the molecules are as follows: in **1A**, the conformation of the S–N–C α –C β bond is substantially twisted from nominal *gauche* [torsion angle =

³ = N-acetyl-L-tyrosine (ref. 7). Standard uncertainty (s.u.) for angles $\approx 0.2^\circ$; **4** = N-acetyl-L-phenylalanyl-L-tyrosine (ref. 8). S.u. for angles $\approx 0.4^\circ$; **5** = (*R*)-2-(2-(5-fluoro-2,4-dioxo-3,4-dihydropyrimidin-1(2H)-yl)acetamido)-3-(4-hydroxyphenyl)-propanoic acid dimethylformamide solvate (ref. 9). S.u. for angles $\approx 0.2^\circ$; **6** = L-tyrosine (ref. 3). S.u. for angles $\approx 0.5^\circ$; **7** = DL-tyrosine (ref. 6). S.u. for angles $\approx 0.5^\circ$.

140.5 (4)°] whilst the N–C α –C β –C γ [–67.6 (6)°] series is closer to being *gauche*. The Ca–C α –C β –C γ torsion angle is close to *anti* [170.2 (5)°] as is the N–C α –Ca–O(H) torsion angle of 166.0 (5)°. In **1B**, the conformation of the S–N–C α –C β bond is even more twisted from *gauche* [torsion angle = 146.9 (4)°] whilst N–C α –C β –C γ [–59.7 (6)°] is almost ideally *gauche*. The Ca–C α –C β –C γ torsion angle is close to *anti* [178.3 (5)°], and compares well to the equivalent value for molecule A. However, the N–C α –Ca–O(H) torsion angle of –37.4 (6)° indicates a completely different conformation, as compared to molecule A. This is backed up by the dihedral angles of 80.4 (5)° between C34/O11/12 and C26/C27/N2 in molecule B and 65.2 (5)° for the equivalent atoms in molecule A. The torsion angles in **1A** are similar to those in **2** (*vide infra*) but quite different to those in related N-acylated L-tyrosine derivatives (Table 1) and in L-tyrosine and DL-tyrosine.

The dihedral angles between the mean planes of the central C1/C2/C3/C4/C14/C15 aromatic ring and the terminal C7–C12 and C16–C21 rings in **1A** are 7.9 (2)° and 60.1 (2)°, respectively. The dihedral angle between the terminal rings is 66.9 (2)°. Equivalent values for **1B** for the C22/C23/C24/C25/C35/C36, C28–C33 and C37–C42 rings are 6.1 (2)°, 53.5 (2)° and 58.33 (19)°, respectively. The molecular conformation of **1A** (Figure 2) leads to a striking intramolecular aromatic π–π stacking interaction between the C1/C2/C3/C4/C14/C15 and C7–C12 rings, with a centroid–centroid separation of 3.807 (3) Å. In **1B**, which has a broadly similar overall shape (Figure 3), the separation between the equivalent rings is significantly longer at 4.067 (3) Å, which must indicate a much weaker interaction. A similar situation has been seen in a series of isostructural 5-phenyl-2-(benzalhydrazonyl)-1,3,4-thiadiazoles [13], in which some compounds show much shorter centroid–centroid separations than others.

Figure 2. The molecular structure of molecule **1A** showing 40% displacement ellipsoids. The intramolecular π – π stacking interaction is indicated by a double-dashed line.

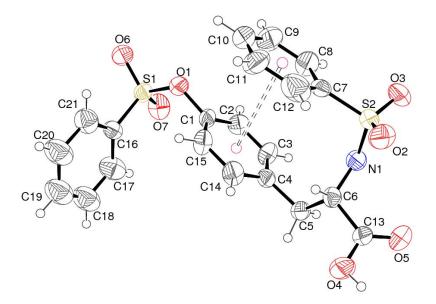
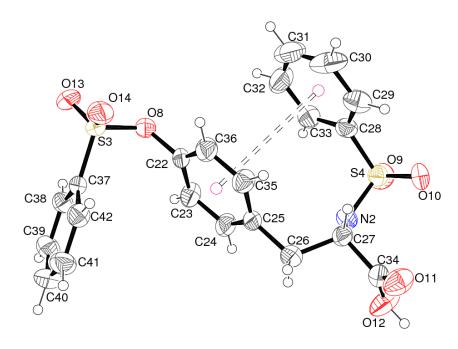


Figure 3. The molecular structure of molecule **1B** showing 50% displacement ellipsoids. The intramolecular π – π stacking interaction is indicated by a double-dashed line.



In the crystal of **1**, the components are linked by N–H···O and O–H···O hydrogen bonds (Table 2). One of the N–H···O links is to a C=O (carboxylic acid) acceptor oxygen atom and the other is to a sulfonyl O atom. Both –OH groups of the carboxylic acids form hydrogen bonds to the solvent molecules (one to a water O atom and one to a methanol O atom). In terms of the C6–H6 and C27–H27 groupings (*i.e.*, the α carbon atoms), the first of these has a distant H···O contact to a sulfonyl O atom (2.63 Å) whereas the second does not have any possible acceptor atoms within 3.5 Å.

Finally, the solvent molecules form further O–H···O links and overall a two-dimensional network lying parallel to (010) arises. With this diversity of hydrogen-bond types observed in the structure, it is hard to draw conclusions as to the preferred hydrogen bonding modes of the organic molecule in 1, although the very short C^{α} –H···O link noted in the introduction is not present. Any intermolecular aromatic π – π stacking must be very weak, with a minimum centroid–centroid separation of greater than 4.06 Å.

N1-H1N···O11 0.87(4)2.06(4)2.922 (6) 168 (5) N2-H2N···O14i 0.87(4)2.10(4)2.925 (6) 160(4)O4-H4O···O15ⁱⁱ 0.82 1.95 2.702 (6) 153 O12-H12O···O1W 0.82 1.75 2.547(7)164 O15-H15A...O10 151 0.82 2.07 2.817 (6) O1W-H1WA···O5 2.08 (6) 2.847(7)146 (6) 0.88(5)O1W-H1WB···O15¹¹¹ 0.87(5)2.14(5)2.998(7)172 (7)

Table 2. Hydrogen bonds in **1**.

The four values refer to the *D*–H, H···A, *D*···A separations (Å) and the D–H···A angle (°), respectively (also in Table 3). Symmetry codes: (i) x, y, z–1; (ii) x, y, z+1; (iii) x+1, y, z.

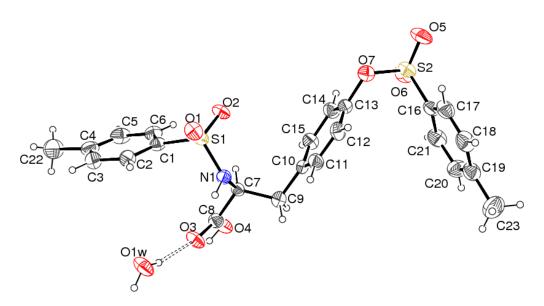
2.2. Structure of 2

The crystal structure of $2 \cdot H_2O$ (Figure 4) shows that L-tyrosine has reacted with two equivalents of 4-toluenesulfonyl chloride to form the S1–N1 and S2–O7 bonds in the product and that a water molecule of crystallization is present. The S-configuration of the C7 stereogenic center [refined value of the Flack absolute structure parameter = -0.01 (7)] is consistent with that of the L-tyrosine starting material and the bond lengths and angles in 2 may be regarded as normal. The molecule of 2 is non-zwitterionic, *i.e.* the proton is clearly attached to O4 and not to N1, as also seen for related molecules [5-7]. In order to simplify the comparison with related structures, atoms C8, C7 C9 and C10 in 2 are referred to as Ca, C α , C β and C γ , respectively.

The conformation of the S1–N1–C α –C β bond in **2** is close to *gauche* [torsion angle = 129.7 (3)°] as is N1–C α –C β –C γ [–61.1 (5)°] whereas that of Ca–C α –C β –C γ is close to anti [174.4 (4)°]. The N1–C α –Ca–O4 torsion angle of 165.3 (3)° indicates that the –OH group of the carboxylic acid is close to *anti* with respect to the NH group. These torsion angles are broadly similar to the situation in molecule **1A** but different to those of other L-tyrosine derivatives (Table 1).

The dihedral angles between the mean planes of the central C10–C15 aromatic ring and the terminal C1–C6 and C16–C21 rings are 44.8 (2)° and 56.1 (2)°, respectively. The dihedral angle between the terminal rings is 81.4 (2)°. Their relative orientations, which are completely different to the conformations of molecules **1A** and **1B** in **1** indicate that no intramolecular aromatic π – π stacking interactions can occur in **2**.

Figure 4. The molecular structure of $2 \cdot H_2O$ showing 40% displacement ellipsoids. The O-H···O hydrogen bond is indicated by a double-dashed line.



In the crystal, the molecules are linked by N–H···O and O–H···O hydrogen bonds. The N–H···O (see Table 3 for atom labels and symmetry codes) hydrogen bond links the organic molecules into C(5) chains propagating in the [100] direction. The carboxylic acid forms an O–H···O bond to the water molecule and the water molecule forms two further O–H···O links; one of the acceptors is the C=O bond of the carboxylic acid, the other is part of a sulfonyl group. Together, these bonds generate a three-dimensional network. An extremely short intermolecular C–H···O bond (2.34 Å) from the α -

carbon atom occurs in **2**, as does a longer C–H···O interaction from C9 (the β carbon atom). There are no aromatic π – π stacking interactions in the crystal of **2**, the shortest centroid–centroid separation being longer than 4.5 Å.

N1-H1N···O11	0.87 (4)	2.06 (4)	2.922 (6)	168 (5)
N1–H1N···O4 ⁱ	0.89 (4)	2.41 (4)	3.241 (5)	156 (3)
O4–H4O···O1W ⁱⁱ	0.82 (5)	1.80 (5)	2.557 (4)	153 (6)
O1W-H1WA···O2 ⁱⁱⁱ	0.818 (18)	2.04(2)	2.842 (3)	166 (5)
O1W-H1WB···O3	0.82 (4)	2.06 (5)	2.864 (4)	165 (5)
C7–H7····O1 ⁱⁱ	0.98	2.34	3.260 (5)	156
C9–H9A···O6 ⁱⁱⁱ	0.97	2.59	3.468 (5)	150

Table 3. Hydrogen bonds in **2**.

Symmetry codes: (i) x-1, y, z; (ii) x+1, y, z; (iii) x, y, z+1.

3. Experimental Section

3.1. Syntheses

All reagents were purchased from commercial sources and used without further purification and compound **1** was synthesized following the "green" literature method [10]. L-Tyrosine (0.50 g, 5.52 mmol) was dissolved in 15 mL of 1 M Na₂CO₃ solution. Benzene sulfonyl chloride (1.41 mL, 11.04 mmol) was suspended in the solution with stirring at room temperature. The pH was maintained at 8–9 until a clear solution resulted. Then, the pH was adjusted to 1–2, using 1 M HCl solution. The white precipitate obtained was filtered, washed with distilled water, dried and recrystallized from methanol and water (98:2 v/v) to yield colorless blocks. Calc. (obs.) analysis (%) for **1** (C₂₁H₁₉NO₇S₂): C 54.65 (54.65), H 4.15 (4.15), N 3.04 (3.03), S 13.90 (13.90).

To prepare compound **2**, L-tyrosine (0.50 g, 5.52 mmol) was dissolved in 15 ml of 1 M Na₂CO₃ solution and 4-toluenesulfonyl chloride (2.1 g, 11.03 mmol) was suspended in the solution with stirring at room temperature. The pH was maintained at 8–9 until the consumption of suspended 4-toluenesulfonyl chloride, to yield a clear solution and the pH was adjusted to 1–2, using 1 M HCl solution. The white precipitate obtained was filtered, washed with distilled water, dried and recrystallized from methanol and water (98:2 v/v) to yield colorless blocks. Calc. (obs.) analysis (%) for **2** ($C_{23}H_{25}NO_7S_2$): C 56.20 (56.43), H 5.13 (4.75), N 2.85 (2.86), S 13.05 (13.10).

3.2. Data collections and refinements

Intensity data for the solvated crystals of **1** and **2** were collected at 296 K using a Bruker Kappa APEXII CCD diffractometer. The unit-cell refinements and data reductions were carried out with SAINT [14]. The structures were solved by direct methods with SHELXS-97 and the atomic models refined against $|F|^2$ with SHELXL-97 [15]. The "observed data" criterion for calculating the R(F) residuals was set as $I > 2\sigma(I)$. The carbon-bound H atoms were geometrically placed and refined as riding atoms with $U_{iso}(H) = 1.2-1.5U_{eq}(C)$. The methyl groups were allowed to rotate, but not to tip, to best fit the electron density. The O- and N-bond H atoms were located in difference maps and refined as riding atoms in their as-found relative positions with $U_{iso}(H) = 1.2U_{eq}(carrier)$. Because of the poor

data to parameter ratio for **1**, some of the benzene rings were modeled as rigid hexagons. The molecular graphics were generated with ORTEP-3 for Windows [16] and the molecular geometries were analyzed with the aid of PLATON [17].

Crystal data for 1_2 ·H₂O·MeOH [2(C₂₁H₁₉NO₇S₂)·H₂O·CH₄O]: colorless block, $0.47 \times 0.28 \times 0.21$ mm, $M_r = 973.04$, monoclinic, $P2_1$ (No. 4), a = 8.0078 (4) Å, b = 34.0704 (16) Å, c = 8.5506 (3) Å, $\beta = 94.239$ (3)°, V = 2326.47 (18) Å³, Z = 2, T = 296 K, $\rho_{calc} = 1.389$ g cm⁻³, $\mu = 0.276$ mm⁻¹, $T_{min} = 0.881$, $T_{max} = 0.944$, 22507 reflections measured ($-10 \le h \le 9$, $-45 \le k \le 27$, $-8 \le l \le 11$; $5.98^{\circ} \le 20 \le 57.00^{\circ}$), $R_{Int} = 0.048$, 8822 merged reflections, 5884 with $I > 2\sigma(I)$, 536 parameters, R(F) = 0.062, $wR(F^2) = 0.157$, S (goodness of fit) = 1.031, Flack parameter = -0.03 (8), $w = 1/[\sigma^2(F_o^2) + 0.0669P^2 + 0.9056P]$, where $P = (F_o^2 + 2 F_c^2)/3$, min./max. $\Delta \rho = -0.29$, +0.57 e Å⁻³, Cambridge Database deposition number: CCDC-822154.

Crystal data for $2 \cdot \text{H}_2\text{O}$ (C₂₃H₂₅NO7S₂·H₂O): colorless block, $0.19 \times 0.18 \times 0.09$ mm, $M_r = 507.56$, monoclinic, $P2_1$ (No. 4), a = 5.7171 (7) Å, b = 24.359 (3) Å, c = 9.1043 (10) Å, $\beta = 104.563$ (6)°, V = 1227.2 (2) Å³, Z = 2, T = 296 K, $\rho_{\text{calc}} = 1.374$ g cm⁻³, $\mu = 0.265$ mm⁻¹, $T_{\text{min}} = 0.00$, $T_{\text{max}} = 0.00$, 13467 reflections measured ($-7 \le h \le 7$, $-32 \le k \le 32$, $-12 \le l \le 12$; $4.62^{\circ} \le 20 \le 56.68^{\circ}$), $R_{\text{Int}} = 0.062$, 5594 merged reflections, 2508 with $I > 2\sigma(I)$, 323 parameters, R(F) = 0.055, $wR(F^2) = 0.092$), S (goodness of fit) = 0.953, Flack parameter = -0.01 (7), $w = 1/[\sigma^2(F_0^2) + 0.0218P^2]$, where $P = (F_0^2 + 2F_0^2)/3$, min./max. $\Delta \rho = -0.17$, +0.18 e Å⁻³, Cambridge Database deposition number: CCDC-822155.

4. Conclusions

We have structurally characterized two doubly-substituted L-tyrosine derivatives, prepared by a "green" synthesis. The change from benzene-sulfonyl to toluene-sulfonyl substituents leads to totally different molecular conformations in the respective crystals: the benzene compound (1) displays a folded conformation stabilized by intramolecular π - π stacking in both asymmetric molecules, whereas the toluene-sulfonyl compound adopts an "extended" conformation. The toluene sulfonyl (2) derivative displays a very short intermolecular C^{α} - $H\cdots$ O interaction in its crystal structure, in common with L-tyrosine and DL-tyrosine, whereas the benzene derivative does not. Both crystal structures contain N- $H\cdots$ O and O- $H\cdots$ O hydrogen bonds linking the tyrosine and the solvent molecules.

Acknowledgements

The authors thank Sohail Anjum Shahzad for his discussion regarding this article and also acknowledge the Higher Education Commission of Pakistan for providing a grant under the project strengthening the Materials Chemistry Laboratory at GC University, Lahore.

References and Notes

- 1. Eagle, H.; Piez, K.A.; Fleishman, R. The utilization of phenylalanine and tyrosine for protein synthesis by human cells in tissue culture. *J. Biol. Chem.* **1957**, 228, 847–861.
- 2. Ip, C.; Harper, A.E. Protein-synthesis in liver, muscle and brain of rats fed a high tyrosine low protein diet. *J. Nutr.* **1975**, *105*, 885–893.

3. Mostad, A.; Nissen, H.M.; Romming, C. Crystal structure of L-tyrosine. *Acta Chem. Scand.* **1972**, 26, 3819–3833.

- 4. Jeffrey, G.A.; Maluszynska, H. A survey of hydrogen bond geometries in the crystal structures of amino acids. *Int. J. Biol. Macromol.* **1982**, *4*, 173–185.
- 5. Scheiner, S.; Kar, T.; Gu, Y. Strength of the $C^{\alpha}H\cdots O$ hydrogen bond of amino acid residues. *J. Biol. Chem.* **2001**, 276, 9832–9837.
- 6. Mostad, A.; Romming, C. Crystal-structure of DL-tyrosine. *Acta Chem. Scand.* **1973**, 27, 401–410.
- 7. Koszelak, S.N.; van der Helm, D. *N*-acetyl-L-tyrosine. *Acta Cryst.* **1981**, *B37*, 1122–1124.
- 8. Stenkamp, R.E.; Jensen, L.H. The crystal structure of pepsin substrate: *N*-acetyl-L-phenyl-alanyl-L-tyrosine. *Acta Cryst.* **1973**, *B29*, 2872–2878.
- 9. Yan, X.; Hu, M.; Miao, Q.; Wang, S.; Zhao, K. The synthesis and anticancer activities of peptide 5-fluorouracil derivatives. *J. Chem. Res.* **2009**, 261–264.
- 10. Deng, X.; Mani, N.S. A facile, environmentally benign sulfonamide synthesis in water. *Green Chem.* **2006**, *8*, 835–838.
- 11. Allen, F.H.; Motherwell, W.D.S. Applications of the Cambridge Structural Database in organic chemistry and crystal chemistry. *Acta Cryst.* **2002**, *B58*, 407–422.
- 12. Flack, H.D. On enantiomorph–polarity estimation. Acta Cryst. 1983, A39, 876–881.
- 13. Carvalho, S.M.; Harrison, W.T.A.; Fraga, C.A.M.; da Silva, E.F.; Wardell, J.L.; Wardell, S.M.S.V. 5-Phenyl-2-(benzalhydrazonyl)-1,3,4-thiadiazoles, potential trypanocidal agents: consistent dimer formation *via* N–H···N intermolecular hydrogen bonds. *Zeit Krist.* **2009**, *224*, 598–606.
- 14. SAINT data reduction software. Bruker AXS Inc.: Madison, Wisconsin, USA, 1997.
- 15. Sheldrick, G.M. A short history of SHELX. Acta Cryst. 2008, A64, 112–122.
- 16. Farrugia, L.J. *ORTEP*-3 for Windows a version of *ORTEP*-III with a Graphical User Interface (GUI). *J. Appl Cryst.* **1997**, *30*, 565.
- 17. Spek, A.L. Structure validation in chemical crystallography. Acta Cryst. 2009, D65, 148–155.
- © 2011 by the authors; licensee MDPI, Basel, Switzerland. This article is an open access article distributed under the terms and conditions of the Creative Commons Attribution license (http://creativecommons.org/licenses/by/3.0/).