

# NEW THIAZOLYLHYDRAZONE DERIVATIVES AS POTENT MONOAMINE OXIDASE AND AROMATASE INHIBITORS

Asaf Evrim Evren<sup>1,2</sup>, Demokrat Nuha<sup>1,3</sup>, Sam Dawbaa<sup>1,4</sup>, Begüm Nurpelin Sağlık<sup>1</sup>, Leyla Yurttaş<sup>1,\*</sup>

<sup>1</sup>Anadolu University, Faculty of Pharmacy, Department of Pharmaceutical Chemistry, 26470, Eskişehir, Turkey

<sup>2</sup>Vocational School of Health Services, Pharmacy Services, Bilecik Seyh Edebali University, Bilecik, Turkey

<sup>3</sup>Eskişehir Technical University, Faculty of Science, Department of Chemistry, 26555, Eskişehir, Turkey

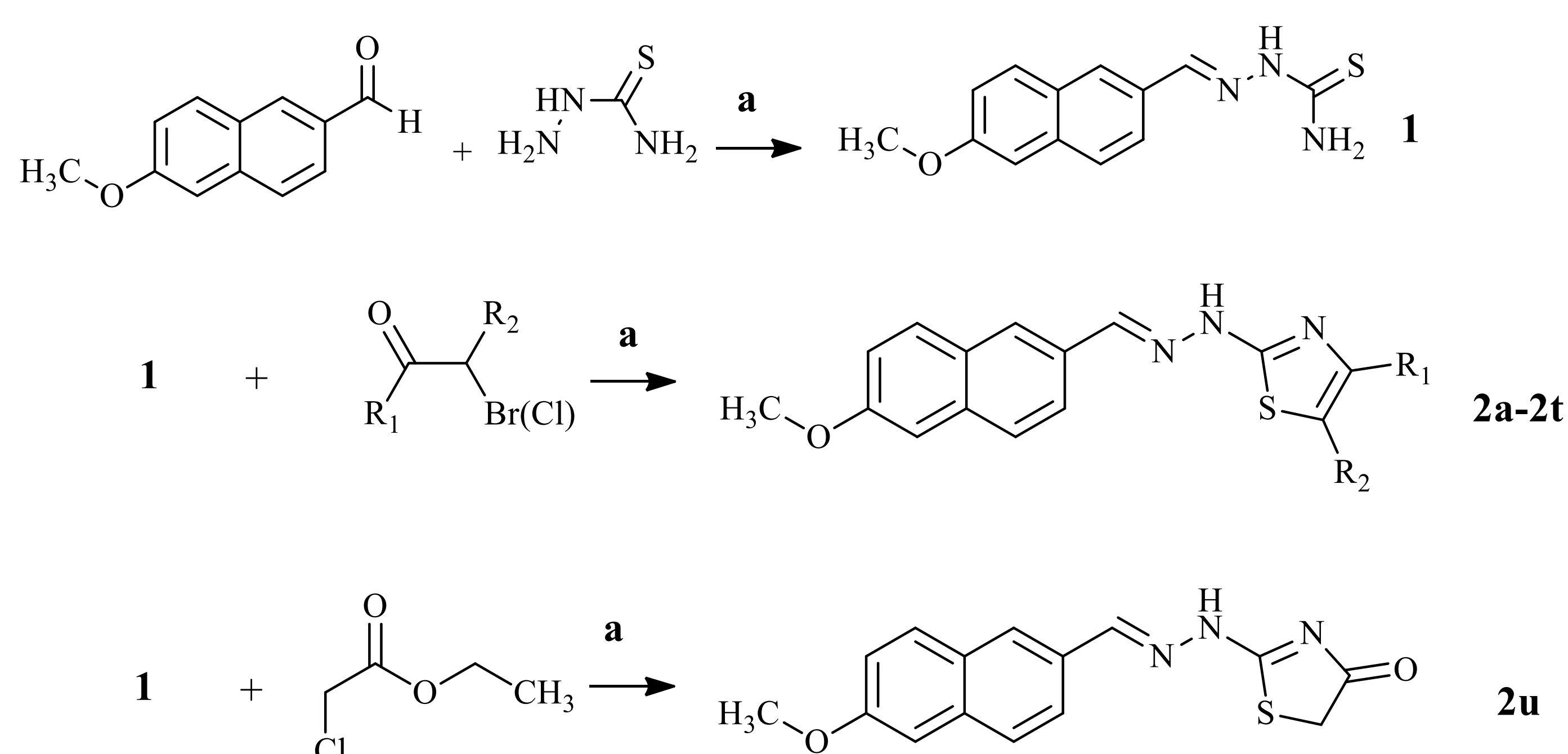
<sup>4</sup>Department of Pharmacy, Faculty of Medicine and Health Sciences, Tamar University, Dhamar, Yemen

**Abstract:** 2-Thiazolylhydrazone nucleus carrying various structures have been known with inhibitory effects on monoamine oxidase (MAO) enzymes. In this study, twenty-one novel 2-((2-((6-methoxynaphthalen-2-yl)methylene)hydrazinyl)thiazole derivatives (**2a-u**) were synthesized and investigated for their MAO and aromatase inhibitory effects. As a result of the study, compound **2j** carrying 3-nitrophenyl residue on the thiazole ring extremely inhibited MAO-A, and compound **2t** carrying phenyl and methyl on thiazole ring was found to inhibit MAO-A both very strongly and selectively. Compounds **2k** and **2q** exhibited selective and high inhibitory potential on MAO-B. Compounds **2q** and **2u** showed satisfying inhibition on aromatase enzyme. Molecular docking and molecular dynamic simulation studies were carried out with the aforementioned compounds and MAO and aromatase enzymes, and findings were correlated with the experimental results.

**Keywords :** 2-Thiazolylhydrazones; aromatase inhibition; molecular docking; molecular dynamic simulation; monoamine oxidase inhibition

## Introduction:

2-Thiazolylhydrazones were defined as a new pharmacophore group in terms of monoamine oxidase inhibition activity by Chimenti and his working group, and derivatives containing different groups were reported in many studies [1-3]. In addition, thiazole-bearing compounds are in a group of non-steroidal aromatase inhibitor drugs since they are inazole structure [4,5]. In the light of reported data, in this study, new thiazolylhydrazone derivatives bearing 6-methoxynaphthalene ring were synthesized and their enzyme inhibitory effects were investigated on MAO-A, MAO-B and aromatase. The synthetic protocol was given below in **Scheme 1**. The structures of all twenty-one compounds were elucidated with spectroscopic data.

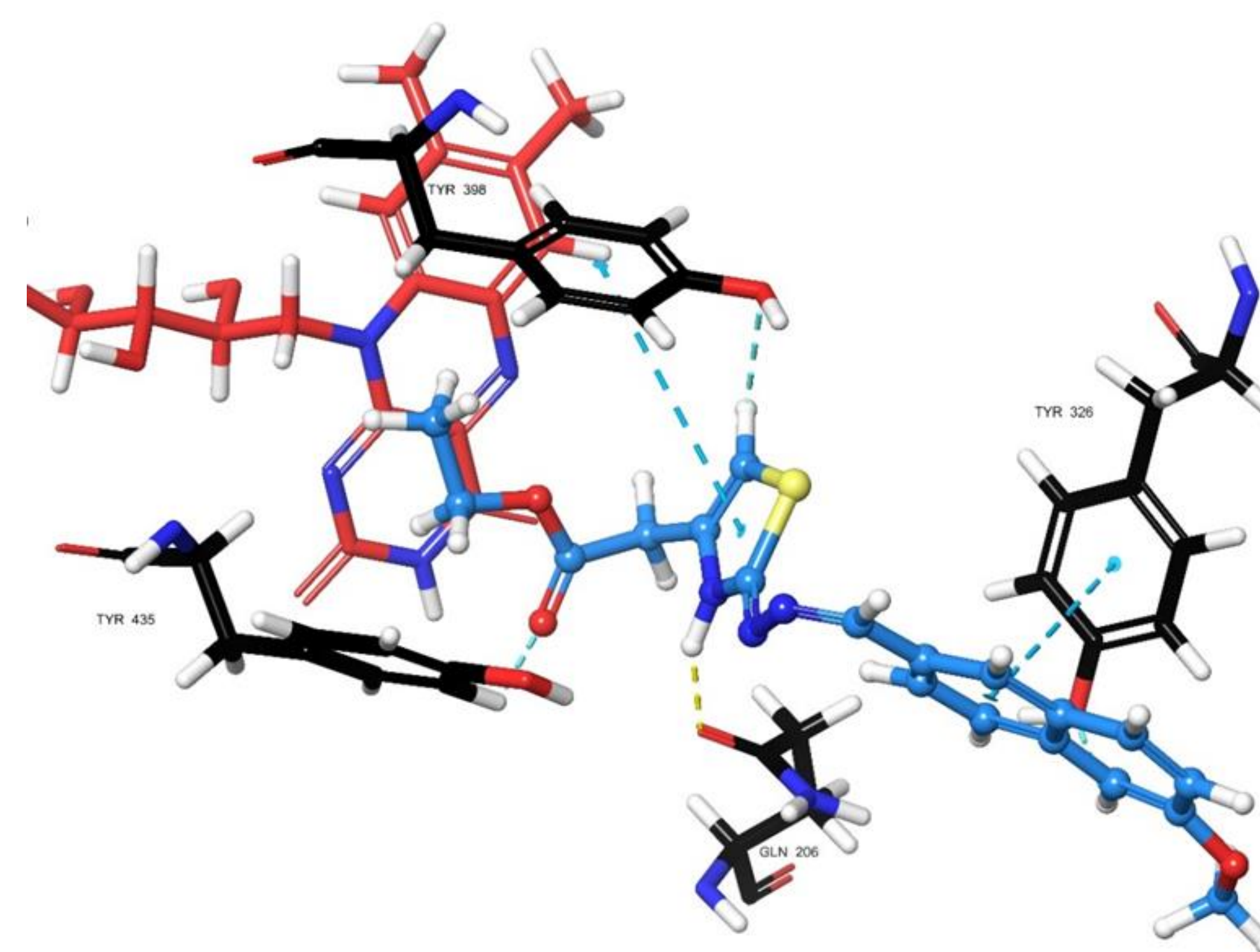


Molecule code	R <sub>1</sub>	R <sub>2</sub>
2a	phenyl	H
2b	4-methylphenyl	H
2c	3-methoxyphenyl	H
2d	4-methoxyphenyl	H
2e	3-chlorophenyl	H
2f	4-chlorophenyl	H
2g	3-fluorophenyl	H
2h	4-fluorophenyl	H
2i	4-cyanophenyl	H
2j	3-nitrophenyl	H
2k	4-nitrophenyl	H
2l	4-(methylsulfonyl)phenyl	H
2m	(1,1'-biphenyl)-4-yl	H
2n	naphthalen-2-yl	H
2o	3,4-dichlorophenyl	H
2p	benzofuran-2-yl	H
2q	-CH <sub>2</sub> COOEt	H
2r	CH <sub>3</sub>	CH <sub>3</sub>
2s	CH <sub>3</sub>	-COOEt
2t	phenyl	CH <sub>3</sub>
2u	-	-

**Biological results :** The enzyme inhibition potencies of the compounds on monoamine oxidase A and B enzyme were studied at 10<sup>-3</sup> and 10<sup>-4</sup> M concentrations. The inhibition concentrations (IC<sub>50</sub>) were calculated for appropriate compounds and were represented in **Table 1**. Compounds **2j** and **2t** exhibited magnificent inhibitory activity on the MAO-A enzyme, nearly 100 times higher than standard drug. Compounds **2j**, **2k** and **2q** acquired to inhibit MAO-B, with very close or half the potency of selegiline. Compounds **2q** and **2u** displayed high aromatase inhibition which was very close to letrozole. Molecular docking and molecular dynamic simulations studies were done for active compounds. Only, **2q**-MAO-B docking pose was given in this poster as **Figure 1**. The interaction with Gln206 was observed for compound **2q**, which is an important residue for the MAO-B selectivity that supports in-vitro results. Compounds **2q** and **2u** were determined to interact HEM protein with π-π bonds which is a necessary for aromatase inhibition. Molecular dynamic simulation and docking studies were determined to support *in vitro* activity results.

**Table 1.** IC<sub>50</sub> values of compounds **2j**, **2k**, **2q**, **2t**, **2u**, moclobemide, selegiline and letrozole against MAO-A, MAO-B, and aromatase enzymes.

Compounds	IC <sub>50</sub> (μM)		
	MAO-A	MAO-B	Aromatase
<b>2j</b>	0.068±0.002	0.046±0.002	-
<b>2k</b>	-	0.082±0.003	-
<b>2t</b>	0.072±0.003	-	-
<b>2q</b>	-	0.039±0.001	0.031±0.001
<b>2u</b>	-	-	0.042±0.001
Moclobemide	6.061±0.262	-	-
Selegiline	-	0.036±0.001	-
Letrozole	-	-	0.026±0.001



**Figure 1.** The aspect of **2q**-MAO-B enzyme complex as 3D (PDBID: 2V5Z)

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**Scheme 1.** The synthesis and derivatives of the compounds (**2a-2u**)



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