## Accepted Manuscript

Title: Conventional and microwave irradiated synthesis, biological activity evaluation and molecular docking studies of highly substituted piperazine-azole hybrids



Author: Arif Mermer Serpil Demirci Serap Basoglu Ozdemir Ahmet Demirbas Serdar Ulker Faik Ahmet Ayaz Fatma Aksakal Neslihan Demirbas

| PII:           | S1001-8417(16)30451-X                             |
|----------------|---|
| DOI:           | http://dx.doi.org/doi:10.1016/j.cclet.2016.12.012 |
| Reference:     | CCLET 3918  |
| To appear in:  | Chinese Chemical Letters                          |
| Received date: | 1-9-2016  |
| Revised date:  | 31-10-2016  |
| Accepted date: | 18-11-2016  |

Please cite this article as: Arif Mermer, Serpil Demirci, Serap Basoglu Ozdemir, Ahmet Demirbas, Serdar Ulker, Faik Ahmet Ayaz, Fatma Aksakal, Neslihan Demirbas, Conventional and microwave irradiated synthesis, biological activity evaluation and molecular docking studies of highly substituted piperazine-azole hybrids, Chinese Chemical Letters http://dx.doi.org/10.1016/j.cclet.2016.12.012

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## ACCEPTED MANUSCRIPT

#### Original article

## Conventional and microwave irradiated synthesis, biological activity evaluation and molecular docking studies of highly substituted piperazine-azole hybrids

Arif Mermer<sup>a</sup>, Serpil Demirci<sup>b</sup>, Serap Basoglu Ozdemir<sup>a</sup>, Ahmet Demirbas<sup>a</sup>, Serdar Ulker<sup>c</sup>, Faik Ahmet Ayaz<sup>d</sup>, Fatma Aksakal<sup>e</sup>, Neslihan Demirbas<sup>a\*</sup>

<sup>a</sup> Karadeniz Technical University, Department of Chemistry, Trabzon 61080, Turkey

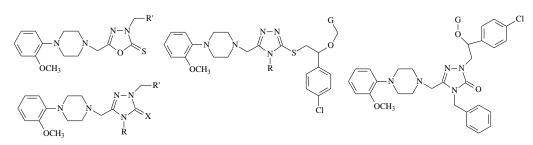
<sup>b</sup> Giresun University, school of Applied Science, Department of Crop Production and Technology, Giresun 28000, Turkey

<sup>d</sup> Karadeniz Technical University, Department of Biology, Trabzon 61080, Turkey

<sup>e</sup> Department of Chemistry, Faculty of Science, Gebze Technical University, Kocaeli 41400, Turkey

\* Corresponding author. *E-mail address*; neslihan@ktu.edu.tr (N. Demirbas)

### **Graphical Abstract**



New hybrid molecules consisting of fluoroquinolone, methoxyphenylpiperazine and azole moieties were synthesized by microwave irradiated and conventional methods. The newly synthesized compounds were screened for their antimicrobial, antiurease, antiglucosidase and antioxidant activities. Also, molecular docking studies were performed.

#### ABSTRACT

Azole derivatives (3, 6) obtained starting from 1-(2-methoxyphenyl)piperazine were converted to the corresponding Mannich bases containing  $\beta$ -lactame or flouroquinolone core *via* a one pot three component reaction. The synthesis of conazole analogues was carried out starting from triazoles by three steps. Reactions were carried out under conventional and microwave mediated conditions. All the newly synthesized compounds were screened for their antimicrobial, enzyme inhibition and antioxidant activity, and most of them displayed good-moderate activity. Binding affinities and non-covalent interactions between enzyme-ligand complexes were predicted with molecular docking method at molecular level. Docking results complemented well the experimental results on  $\alpha$ -glucosidase and urease inhibitory effects of the compounds. Higher binding affinities and much more interaction networks were observed for active compounds in contrary to inactive ones. It was predicted with the docking studies that triazole and anisole moieties in the structure of the synthesized compounds contributed to the stabilization of corresponding enzymes through noncovalent interactions.

Keywords: Fluoroquinolone 1,2,4-Triazole Microwave Mannich reaction Biological activity Molecular docking

<sup>&</sup>lt;sup>c</sup> RecepTayyipErdoğan University, Department of Biology, Rize 53100, Turkey

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