### SUPPORTING INFORMATION

# Design, Synthesis, and Structure–Activity Relationships of Thiazole Analogs as Anticholinesterase Agents for Alzheimer's Disease

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Compounds	AChE % Inhibition							
	10 <sup>-3</sup> M	10-4 M	10 <sup>-5</sup> M	10 <sup>-6</sup> M	10-7 M	10 <sup>-8</sup> M	10-9 M	(μΜ)
2a	93.425	90.465	81.239	74.155	60.975	41.275	20.388	0.063
	±1.652	±1.322	±1.421	±1.518	±1.182	±0.954	±0.711	±0.003
26	90.285	82.151	74.565	63.758	57.965	43.948	21.208	0.056
20	±1.451	±1.478	±1.279	±1.230	±1.023	±0.963	±0.839	±0.002
24	89.462	82.445	78.103	67.321	55.922	38.162	29.751	0.147
20	±2.041	±1.695	±1.766	±1.159	±1.004	±0.847	±0.855	±0.006
20	90.611	85.387	81.828	74.797	62.356	44.275	21.599	0.040
26	±1.815	±1.730	±1.311	±1.488	±1.103	±0.904	±0.631	±0.001
2~	93.461	88.347	81.465	76.911	69.188	43.104	24.903	0.031
2g	±1.632	±1.604	±1.752	±1.326	±1.055	±0.952	±0.789	±0.001
2:	95.207	92.130	86.602	80.964	71.840	42.651	22.155	0.028
21	±1.502	±1.798	±1.214	±1.124	±1.270	±0.866	±0.830	±0.001
2:	91.326	84.945	76.841	62.108	57.365	40.715	25.575	0.138
<b>∠</b> J	±2.107	±1.369	±1.830	±1.411	±1.036	±0.949	±0.899	±0.005
Dononozil	99.254	97.426	92.258	90.318	81.365	43.875	21.418	0.021
Donepezh	±2.104	±1.890	±1.510	±1.104	±1.104	±0.601	±0.548	±0.001

Table S1. IC50 values of compounds 2a, 2b, 2d, 2e, 2g, 2i, 2j and donepezil against AChE.



**Figure S1.** The superimposition pose of selected compounds in the enzyme active site (AChE PDB Code: 4EY7).



**Figure S2.** The two- (**A**) and three-dimensional (**B**) interacting mode of compound **2a** in the active region of AChE. The inhibitor and important residues in the active site of enzyme are presented by tube model and colored with grey and aquamarine, respectively (AChE PDB Code: 4EY7).



**Figure S3.** The two- (**A**) and three-dimensional (**B**) interacting mode of compound **2b** in the active region of AChE. The inhibitor and important residues in the active site of enzyme are presented by tube model and colored with blue and aquamarine, respectively (AChE PDB Code: 4EY7).



**Figure S4.** The two- (**A**) and three-dimensional (**B**) interacting mode of compound **2e** in the active region of AChE. The inhibitor and important residues in the active site of enzyme are presented by tube model and colored with orange and aquamarine, respectively (AChE PDB Code: 4EY7).



**Figure S5.** The two- (**A**) and three-dimensional (**B**) interacting mode of compound **2g** in the active region of AChE. The inhibitor and important residues in the active site of enzyme are presented by tube model and colored with dark green and aquamarine, respectively (AChE PDB Code: 4EY7).



**Figure S6.** The two- (**A**) and three-dimensional (**B**) interacting mode of compound **2d** in the active region of AChE. The inhibitor and important residues in the active site of enzyme are presented by tube model and colored with pink and aquamarine, respectively (AChE PDB Code: 4EY7).



Figure S7. The thin-layer chromatography of the synthesized compounds.

## ==== Shimadzu LCMSsolution Analysis Report ====

: Admin
: tf-14
:
: 32
: 0.3 uL
: tf-14_43.lcd
: isocratic.lcm
: batch.lcb
: DefaultLCMS.lcr
: 12.03.2020 01:13:26
: 12.03.2020 01:23:28

#### <Chromatogram>



PDA Ch1 25	4nm 4nm	F Cak I dole			
Peak#	Ret. Time	Area	Height	Area %	Height %
1	0.985	7787	705	3.059	3.608
2	1.871	3512	321	1.380	1.643
3	3.412	232017	17619	91.149	90.187
4	4.911	6455	509	2.536	2.603
5	7.601	4776	383	1.876	1.959
Total		254546	19537	100.000	100.000

Figure S8. Purity of compound 2i.



Figure S9. <sup>1</sup>H NMR spectra of the compound 2i.



Figure S10. <sup>13</sup>C NMR spectra of the compound 2i.

Data File: C:\LabSolutions\Data\Analiz\derya\tf-14\_43.lcd



Figure S11. HRMS spectra of the compound 2i.