

Trisubstituted 1,3,5-Triazines and Their Effect on BACE1[†]

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Abstract: Alzheimer's disease (AD) is a multifactorial neurological disease of unknown etiology that is associated with various risk factors. Various pharmacological approaches targeting distinct mechanisms have been investigated; however, they have not yet achieved disease-modifying effects. A series of nine trisubstituted 1,3,5-triazine-based derivatives was investigated as potential inhibitors of the β -secretase enzyme (beta-site amyloid precursor protein-cleaving enzyme 1, BACE1), one of the key enzymes in the pathogenesis of AD. Although the triazine-based derivatives are reported to be potent BACE1 inhibitors, the compounds discussed in this contribution, at a concentration of 10 μ M, demonstrated completely insignificant activity. It is worth noting that methyl 4-(4-[(2,3-dihydroxypropyl)amino]-6-[(4-sulfamoylbenzyl)amino]-1,3,5-triazin-2-yl)piperazin-1-yl)- acetate and 4-((4-chloro-6-[(3-hydroxypropyl)amino]-1,3,5-triazin-2-yl)amino)benzenesulfonamide showed an approximately 9% and 2% inhibition of BACE1 activity, respectively.

Keywords: triazinylaminobenzenesulfonamides; Alzheimer's disease; BACE1; modulation



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1. Introduction

The triazine structure consists of a heterocyclic six-membered ring containing three nitrogen atoms; thus, three isomers can be found: 1,2,3-triazine, 1,2,4-triazine and 1,3,5-triazine. Triazine molecules are basic in nature but are weaker bases than pyridine. Although triazines are aromatic compounds, their resonance nature is much lower than that of benzene. Electrophilic substitution is difficult, but nucleophilic aromatic substitution is quite easy [1,2].

The best-known derivative of 1,3,5-triazine is melamine (2,4,6-triamino-1,3,5-triazine), which has found industrial use in the production of resins. Other widely used 1,3,5-triazines are, e.g., cyanuric chloride (2,4,6-trichloro-1,3,5-triazine) and 6-alkyl/aryl-1,3,5-triazine-2,4-diamines. Triazine-based agents such as atrazine, ametryn, prometryn, cyanazine, propazine, simazine, terbuthylazine and terbutryn are highly effective herbicides [3], unfortunately also with strong adverse impacts on humans and the environment [4–9]. On the other hand, 1,3,5-triazines represent a remarkable platform for the design of potential drugs, especially with anti-infective (antiviral, antibacterial, antimycobacterial, antifungal, antiprotozoal, anthelmintic) and anticancer effects, but also with anti-inflammatory, antidiabetic, antioxidant, antiulcer, anticonvulsant and cardioprotective activities, depending on the specific substitution of the 1,3,5-triazine scaffold [1,10–14].

In addition to all these activities, triazines, both 1,2,4-isomers [15,16] and 1,3,5-isomers [17–19], were found to exhibit the ability to inhibit the beta-site amyloid precursor protein-cleaving enzyme 1 (BACE1), see Figure 1.

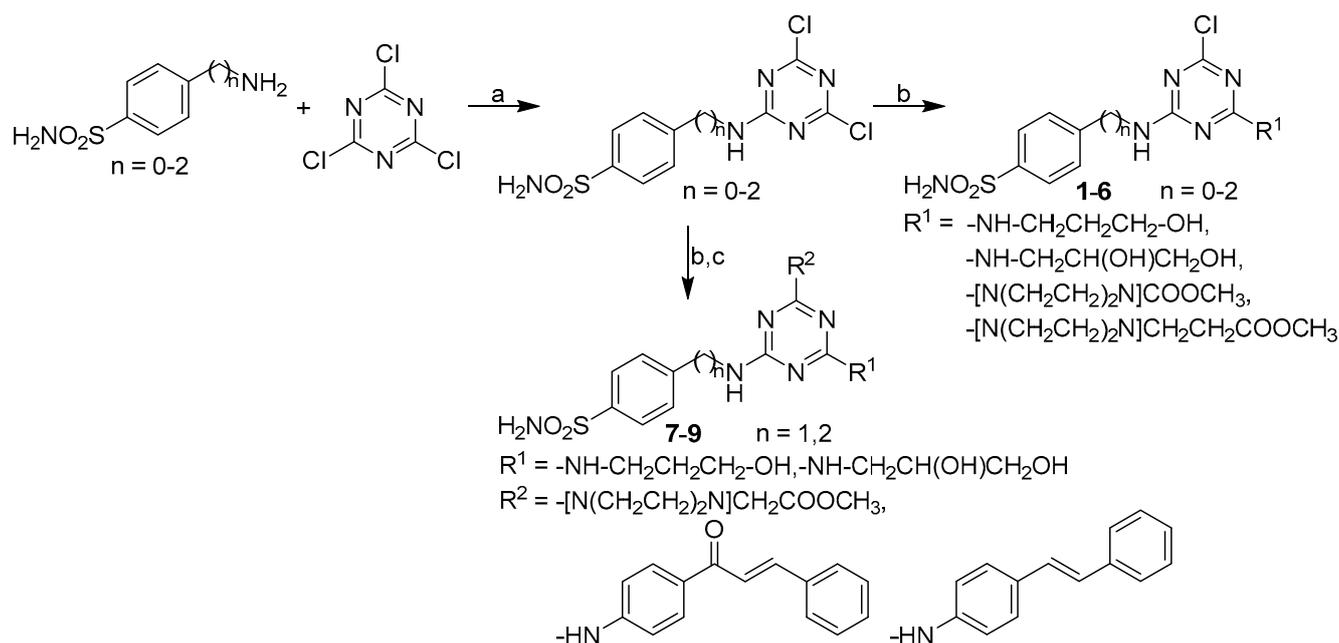
for the determination of the capacity factor k . The total flow of the column was 0.4 mL/min, injection 5 μ L, column temperature 30 $^{\circ}$ C and sample temperature 10 $^{\circ}$ C. A thiourea methanolic solution was used for the determination of the dead time (t_D). Retention times (t_R) were measured in minutes. The capacity factors k were calculated according to the formula $k = (t_R - t_D)/t_D$, where t_R is the retention time of the solute and t_D is the dead time obtained using an unretained analyte. Each experiment was repeated three times.

2.3. Determination of BACE1 Inhibitory Activity

The BACE inhibitory activity was determined via commercial assay according to manufacturer instructions (Merck Life Science, Bratislava, Slovakia) [40]. The principle of the assay is based on the fluorescence resonance energy transfer (FRET) method, in which the fluorescence signal enhancement is observed after substrate cleavage by BACE1, meaning that the lower the percentage of BACE activity, the more BACE1 is inhibited by the test compounds.

3. Results and Discussion

The synthesis of the discussed compounds is shown in Scheme 1. The starting 1,3,5-triazin-2-yl-aminoarylsulfonamides were synthesized according to Garaj et al. [39]. Subsequently, 4-[(4,6-dichloro-1,3,5-triazin-2-yl)amino]benzene-1-sulfonamide, 4-[(4,6-dichloro-1,3,5-triazin-2-yl)amino]methyl]benzene-1-sulfonamide and 4-[2-[(4,6-dichloro-1,3,5-triazin-2-yl)amino]ethyl]benzene-1-sulfonamide with appropriate reagents provided the final molecules as recently described [35–38]. The structures of the target compounds are shown in Table 1.

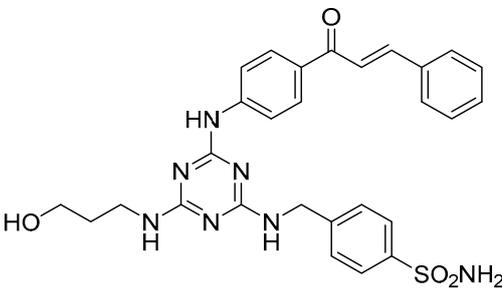
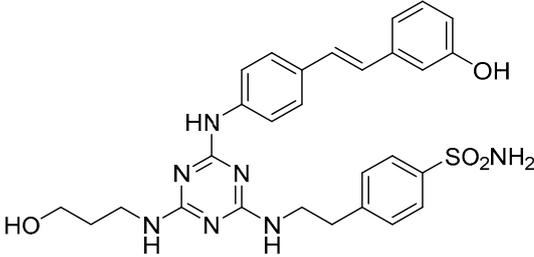


Scheme 1. Synthesis of target compounds 1–9. Reagents and conditions: (a) acetone, 0–5 $^{\circ}$ C [39]; (b) suitable amine derivative (1 eq.), DMF, K₂CO₃, 35 $^{\circ}$ C; (c) suitable amine derivative (1 eq.), DMF, K₂CO₃, 100 $^{\circ}$ C [35–38].

Table 1. Structures of discussed ring-substituted 1,3,5-triazine derivatives 1–9, experimentally determined lipophilicity ($\log k$), predicted topological polar surface area (tPSA) of investigated compounds and in vitro reduction of BACE1 activity (%).

No.		$\log k$	tPSA ¹ (Å ²)	Reduction of BACE1 Activity (%)
1 [35]		−0.3213	141.53	2.3
2 [36]		−0.3807	141.53	0
3 [36]		−0.2816	141.53	0
4 [37]		0.3490	142.05	0
5 [37]		−0.1155	142.05	0
6 [36]		−0.0666	142.05	0
7 [37]		0.4387	194.54	9.6

Table 1. Cont.

No.		log <i>k</i>	tPSA ¹ (Å ²)	Reduction of BACE1 Activity (%)
8 [38]		0.3597	170.63	0
9 [38]		0.3850	173.79	0

¹ Calculated using ChemBioDraw Ultra 13.0 (CambridgeSoft, PerkinElmer Inc., Cambridge, MA, USA).

Although the discussed compounds were previously structurally fully characterized [35–38], their lipo-hydrophilic properties have only now been determined via reversed-phase high-performance liquid chromatography (RP-HPLC) using an end-capped non-polar C18 stationary RP column and expressed as the logarithm of the capacity factor *k*. The retention times of the individual compounds were obtained under isocratic conditions with acetonitrile as an organic modifier in the mobile phase. The values of log *k* are given in Table 1.

Table 1 shows the structures of the tested compounds. The derivatives differ from each other in the length of the linker (*n* = 0–2) between the amino-triazine and benzenesulfonamide fragments. In addition, the compounds differ either by substitution of the second amino group of the triazine with 3-hydroxypropyl (compounds 1–3, 8, 9) or by the incorporation of the amino group into the substituted piperazine (compounds 4–7), and the last (third) substitution can be found on the triazine ring with either chlorine (compounds 1–6), 2,3-dihydroxypropylamino (compound 7), or a complex arylamine (compounds 8, 9).

Lipophilicity is one of the parameters that fundamentally influence not only the pharmacokinetics, but also the effect, of bioactive agents [41,42]. Even in this limited series of nine highly functionalized compounds, a wide range from –0.38 to 0.43 of log *k* values is evident, with 4-[(4-chloro-6-[(3-hydroxypropyl)amino]-1,3,5-triazin-2-yl)-amino)methyl]benzenesulfonamide (2) showing the lowest experimental log *k* value, while methyl (4-[4-[(2,3-dihydroxypropyl)amino]-6-[(4-sulfamoylbenzyl)amino]-1,3,5-triazin-2-yl]piperazin-1-yl)acetate (7) is the most lipophilic. However, in general, it can be stated that all derivatives are rather hydrophilic in nature, with five of them having a negative log *k* value, which indicates their problematic bioavailability due to limited transport through membranes.

In addition to lipophilicity, the topological polar surface area (tPSA), which is defined as the sum of the surfaces of the polar atoms (most often oxygens, nitrogens and attached hydrogens) in a molecule [43], has become a widely used molecular descriptor in the study of drug properties (to ensure so-called drug-likeness) [42]. This descriptor, showing a correlation with passive molecular transport through membranes, is also logically related to the magnitude of the drug–receptor interactions [44]. Therefore, the tPSA values for the individual studied compounds were calculated using the ChemBioDraw Ultra 13.0

program. The most common value is approx. 142 Å² (compounds 1–6) and ca. 172 Å² (compounds 8 and 9). The most lipophilic compound 7 also achieved the highest tPSA value, namely 194.54 Å², which, similarly to its lipophilicity, is largely different from the values of the other derivatives.

All the investigated compounds (see Table 1) were tested for their ability to inhibit BACE1 using a commercially available kit [40]. Performance of the test is described in Section 2.3. As the IC₅₀ values of known BACE1 inhibitors (including the kit reference standard [40]) are in the nanomolar range, all the evaluated compounds were tested at a concentration of 10 µM (see, e.g., [27]), as it is conceivable that if there is no activity at 10 µM, the compound most likely does not inhibit BACE1 [27]. As can be seen from the results in Table 1, the compounds showed no activity; only compound 7 demonstrated some ability to slightly inhibit the BACE1 enzyme (approx. 9% reduction in BACE1 activity). It should be noted that 4-({4-chloro-6-[(3-hydroxypropyl)amino]-1,3,5-triazin-2-yl)-amino)benzenesulfonamide (1) also showed a 2% reduction in BACE1 activity. It can only be speculated whether the “significant” inhibition of BACE1 by derivative 7, compared to the zero activity of the other tested derivatives, is related to its highest log *k* and tPSA values within the series of the investigated compounds. Nevertheless, from the obtained data, it can be concluded that the mentioned trisubstituted triazines do not have the ability to interact with BACE1.

However, it must be admitted that, based on the latest studies and the situation with the early termination of all clinical trials of the main BACE1 inhibitors such as verubecestat atabecestat, elenbecestat lanabecestat umibecestat, LY2886721, RO5508887 or PF-06751979 by pharmaceutical companies (e.g., Pfizer, AstraZeneca, Merck, Eli Lilly, Roche, Novartis, Janssen, Biogen, Amgen) due to serious side effects or actual ineffectiveness, it is possible to speculate to what extent the development of BACE inhibitors (the β-secretase inhibition hypothesis) represents a dead end, or how long the development of these inhibitors for AD treatment will continue to be preferred by pharmaceutical companies [27,45,46]. On the other hand, natural BACE1 inhibitors with different structures are still being discovered [47] and the role of the Aβ peptide is being investigated in depth [48,49].

4. Conclusions

Recently, a series of trisubstituted 1,3,5-triazine derivatives was published and significant inhibitions of therapeutically important carbonic anhydrases were found. The finding that triazines are able to inhibit BACE1 led to an assay of their ability to inhibit BACE1. However, none of the evaluated compounds demonstrated a significant decrease in BACE1 activity.

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