Contents lists available at ScienceDirect

Bioorganic Chemistry

journal homepage: www.elsevier.com/locate/bioorg

Selenides bearing benzenesulfonamide show potent inhibition activity against carbonic anhydrases from pathogenic bacteria Vibrio cholerae and Burkholderia pseudomallei



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ARTICLE INFO

Keywords: Carbonicanhvdrase Inhibitors Metalloenzymes Selenium Vibrio cholerae Burkholderia pseudomallei

ABSTRACT

A series of selenides bearing benzenesulfonamide moieties was evaluated as carbonic anhydrase (CA, EC 4.2.1.1) inhibitors against the pathogenic bacteria Vibrio cholerae (VchCAa and VchCAB) and Burkholderia pseudomallei (BpsCAB) enzymes. The molecules represent an interesting lead for antibacterial agents with a possibly new mechanism of action showing excellent inhibitory action and selectivity for inhibiting VchCAa and BpsCAB over the human (h) off-target isoforms hCA I and II. Identification of potent and possibly selective inhibitors of bacteria CAs over the human counterparts may lead to pharmacological tools useful for understanding the physiological role(s) of these under-investigated proteins.

1. Introduction

The carbonic anhydrases (CAs, EC 4.2.1.1) are a superfamily of metalloenzymes which catalyse the interconversion reaction of CO₂ and H_2O with HCO_3^- and H^+ to perform a pivotal role in pH regulation [1]. It is an essential enzyme in all life kingdoms, from Archaea to Bacteria and Eukaryotes [2]. To date, was discovered seven genetically distinct families, named α – (present in vertebrates, protozoa, algae, cytoplasm of green plants, and in many gram-negative Bacteria), β – (in both gram-negative and -positive Bacteria, mono and dicotyledons plants, fungi and Archaea), γ – (Bacteria, cyanobacteria and Archaea), δ -, ζ -, θ - (in marine diatoms) and η - CAs (protozoa belonging to the *Plasmo*dium spp) [1-3]. Recently, the inhibition of bacterial CAs was demonstrated to influence both growth and pathogenicity of microorganisms [4–6], thus, could be a new approach for obtaining anti-infective agents with a new mechanism of action compared to classical antibiotics [7]. Discover and develop new anti-infective agents, at present, is inevitable because infectious diseases are the second-leading cause of death in the world. In this particular contest, Burkholderia pseudomallei and Vibrio cholerae are Gram-negative bacteria that cause different endemic diseases. The first is a saprophytic bacterium responsible for melioidosis, an endemic disease present in tropical and sub-tropical areas of the world [8,9]. B. pseudomallei is highly virulent by inhalation and also acquired resistance to penicillin, ampicillin, first-generation and

aminoglycoside antibiotics [8,9]. The second Gram-negative bacterium, Vibrio cholerae, is causative of cholera, a human disease that is characterized by massive loss of water and electrolytes, which leads to severe dehydration and hypovolemic shock if the condition is not treated [10]. This study may be of interest for designing new organoselenium inhibitors of VchCAa, VchCAB and BpsCAB that may have clinical applications [11].

second-generation cephalosporins, macrolides, guinolones and most

2. Results and discussion

2.1. Chemistry

Commercially available sulfanilamide 1, has been used for preparing the corresponding selenocyanate derivative 2 and diselenide 3, as reported earlier [12] (Scheme 1).

Thereafter, 3 was reduced with NaBH₄ to the corresponding selenolate, which in turn was treated in situ with the appropriate halo-alkyl moieties 4a-j, affording the selenides 5a-j in good yield (Table 1) [13].

Finaly, the selenide 6 was obtained from reduction of diphenyldiselenide with NaBH₄ and in situ was added 4-(bromomethyl)benzenesulfonamide to afford in good yield as outlined in Scheme 2[13].

Our main interest was to investigate structure-activity relationship related to the inhibition of the two pathogenic bacteria from Vibrio

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https://doi.org/10.1016/j.bioorg.2018.05.015

Received 21 April 2018; Received in revised form 15 May 2018; Accepted 16 May 2018 Available online 18 May 2018

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Scheme 1. Synthesis of selenocyanate 2 and diselenide 3 derivative.

Table 1



^a Yields are referred to isolated products



Table 2

Inhibition data against human (h) isoforms hCA I, II and bacterial enzymes VchCA α , VchCA β and BpsCA β of derivates **2**, **3**, **5a-j**, **6** and acetazolamide (AAZ) by a stopped flow CO₂ hydrase assay [14].

Cmp	$K_{I} (nM)^{a}$				
	VhCAa	VhCAβ	BpsCAβ	hCAI	hCAII
2	4.6	2172.5	75.3	95.6	53.1
3	5.7	6804.3	83.6	1522.7	7.9
5a	5.9	7597.7	754.2	338.3	355.3
5b	4.1	5941.7	845.6	256.8	9.3
5c	6.1	8726.9	529.8	352.2	73.2
5d	6.5	5264.4	51.9	9.7	69.9
5e	28.6	7853.1	9.1	5.2	36.5
5f	6.7	2589.9	21.4	7.3	9.3
5g	6.0	6900.8	5.4	293.5	7.6
5h	4.4	5068.3	823.9	297.1	70.8
5i	4.3	5897.1	876.9	21.9	6.3
5j	7.5	1852.8	866.6	261.7	41.2
6	5.4	7697.1	650.4	226.1	53.0
AAZ	6.8	451	745	250	12.1

^a Mean from 3 different assays, by a stopped flow technique (errors were in the range of \pm 5–10% of the reported values).

cholerae (VchCAα, VchCAβ) and, Burkholderia pseudomallei (BpsCAβ).

2.2. Carbonic anhydrase inhibition

The inhibition studies of VchCA α , VchCA β and BpsCA β with compounds **2**, **3**, **5a-j** and **6** were performed in order to detect possible candidates for anti-infective studies. We tested *in vitro* these compounds for their inhibitory activity against the two off-targets physiologically relevant hCA isoforms I, II and the bacterial enzymes VchCA α , VchCA β and BpsCA β by means of the stopped-flow carbon dioxide hydration assay [14]. These activities were compared to those of the standard, clinically used CAI acetazolamide (AAZ) (Table 2).

The following structure–activity relationship (SAR) can be drawn from the data of this table:

(i) Selenides 2, 5a-j, 6 and diselenide 3 exhibited potent inhibitory activity towards VchCAα with inhibition constants in the low-nanomolar range, except for compound 5c, which was active in the medium nanomolar range (overall, the K₁s ranged between 4.3 and 28.6 nM). However, small structural differences in the tail moiety, such as the replacement of methyl moiety (5c) with different methylene chains (5a-b, 5d-j, 6), did not lead to particular effects on

Scheme 2. Synthesis of 4-((phenylselanyl)methyl) benzenesulfonamide **6**.

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