

ANTIMICROBIAL ACTIVITY OF *meta*-ALKOXYPHENYLCARBAMATES CONTAINING SUBSTITUTED *N*-PHENYLPIPERAZINE FRAGMENT

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ABSTRACT

In the present investigation, the basic esters of *meta*-alkoxyphenylcarbamic acid bearing variously substituted *N*-phenylpiperazine fragment were screened for their *in vitro* antimicrobial activity against *Staphylococcus aureus*, *Escherichia coli* and *Candida albicans*, respectively. The most effective against *Escherichia coli* was found the compound 6d (MIC=195,3 µg/mL) bearing simultaneously *para*-fluoro substituent at the 4-phenylpiperazin-1-yl core and *meta*-methoxy side chain in the lipophilic part of the molecule. From whole analyzed set of the molecules the substance 8e with propoxy side chain forming *meta*-alkoxyphenylcarbonyl fragment and lipophilic, sterically bulky *meta*-trifluoromethyl group attached at *N*-phenylpiperazine moiety was evaluated as the most active against *Candida albicans* (MIC=97,7 µg/mL). On the contrary, all investigated structures were practically inactive against *Staphylococcus aureus* (MIC>1000 µg/mL)

Key words: Phenylcarbamates; substituted *N*-phenylpiperazines; *Candida albicans*

INTRODUCTION

To survive in the environment, bacteria and yeasts must respond to several stress factors that lead to non-ideal growth conditions. The emergence of a pathogen community depends on its ability to survive in different environments and to interact successfully with the host (18). As an additional stress, they may be exposed to a wide range of antimicrobially active agents, such as antibiotics, that can act as a selective pressure for the development of microorganisms resistance. As a result of antibiotic use and misuse, the prevalence of antimicrobial resistance among bacterial pathogens has increased resulting in more complicated treatment. In general, bacteria have genetic

ability to transmit and acquire resistance to the drugs, which are utilized as the therapeutic agents. The problem of microbial resistance is growing and the outlook for the use of antimicrobial drugs in the future is still uncertain. Therefore, actions must be taken to reduce the resistance by controlling the use of antibiotics, by development of the research for a better understanding the genetic mechanisms of resistance. Projection and synthesis of new antimicrobial agents active against emerging resistant pathogens is thus an essential process (15, 16, 17).

The objective of present study is to investigate *in vitro* susceptibility of selected clinically significant microbial strains, the opportunist pathogens, to novel lipophilic basic

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esters of *meta*-alkoxyphenylcarbamic acid containing variously substituted 4-phenylpiperazin-1-yl fragment in the structure.

MATERIALS AND METHODS

Chemistry

The preparation of evaluated compounds, labelled as 6d–6g and 8c–8e (Table 1), their spectral characteristics (^1H NMR, ^{13}C NMR, IR, MS, UV/VIS) as well as the elemental analyses data were published previously (9, 10). The procedures of physicochemical parameters determination (surface activity γ , dissociation constant $\text{p}K_{\text{a}}$, lipophilicity descriptors – $\log P_{\text{exp}}$ estimated by the shake-flask method in the octan-1-ol/ buffer medium with $\text{pH} = 7,3$, $\log k'$ from RP-HPLC, R_{M} from RP-TLC) and the corresponding readouts and conclusions were published in paper of Malik *et al.* (11).

In vitro antimicrobial activity assay

Microorganisms: The antimicrobial activity of concerned compounds was investigated against Gram-positive bacteria *Staphylococcus aureus* ATCC 6538 (*Micrococcaceae*), Gram-negative bacteria *Escherichia coli* CNCTC 377/79 (*Enterobacteriaceae*) and yeast *Candida albicans* CCM 8186 as well. The tested bacterial strains were purchased from American Type Culture Collection and Czech Collection of Type Cultures (National Institute of Public Health, Prague, Czech Republic), yeast was obtained from Czech Collection of Microorganisms (Brno, Czech Republic).

Culture media: The blood agar, Endo agar and Sabouraud's agar were used for cultivation of microorganisms (Imuna, Šarišské Michaľany, Slovak Republic). The blood agar was prepared by adding of 10% of defibrine sheep's blood to the melted and cooled (50°C) competent components.

Determination of minimum inhibitory concentration (MIC): The values of MIC of studied structures were estimated according to approach which was detailed in the paper of Mlynarčík *et al.* (14).

The tested substances were solubilised in dimethyl sulfoxide medium due to their very limited solubility in water. The standard suspension of microorganisms was prepared from 24h bacterial cultures cultivated at a blood agar (Gram-positive bacteria) and Endo agar (Gram-negative bacteria) as well as from the 48h cultures cultivated at the Sabouraud's agar for yeasts. The suspension exhibited the concentration of 5×10^7 colony forming unit (cfu)/mL of bacteria and 5×10^5 cfu/mL of *Candida*, respectively. The UV/VIS spectrophotometry was used for the determination of the microorganisms concentration, all the prepared suspensions were adjusted to the absorbance value of 0,35 at the wavelength of 540 nm.

The microorganism suspension prepared in such a way was then particularly added in amount of 5 μL into the solutions consisting of evaluated substances (100 μL) and to the double concentrated peptone broth medium (8%) assigned to bacteria or the Sabouraud's medium (12%) related to *Candida*, respectively (14). For the prepared stock solutions of all evaluated substances there was a concentration of 25000 $\mu\text{g}/\text{mL}$.

The solutions were then serially diluted by a half (the final concentrations were 12500; 6250; 3125; 1562,5; 781,3; 390,6; 195,3 and 97,7 $\mu\text{g}/\text{mL}$, respectively). The quantitative screening was performed using 96-well microtiter plates, microorganisms were incubated there at 37°C for 24h. After that, from particular well the amount of 5 μL of evaluated suspension was taken and cultured on a blood agar (bacteria) or on Sabouraud's agar (yeasts). Then were the Petri dishes incubated for 24h at 37°C .

The values of MIC were read as the lowest concentration of antimicrobially active compound which inhibited the visible microbial growth (14).

RESULTS AND DISCUSSION

The investigated substances, chemically 1-[3-(3-alkoxyphenylcarbamoxyloxy)-2-hydroxypropyl]-4-(4-fluoro-/3-

trifluoromethylphenyl)piperazinium chlorides (alkoxy = methoxy to butoxy group; labelled as 6d–6g and 8c–8e, respectively), were previously *in vitro* tested against *Mycobacterium tuberculosis* CNCTC My 331/88 (19). The estimated values of MIC for 6d–6g were in the range of 55–241 µg/mL (related to the interval of 125–500 µmol/L), in the series of 8c–8e the evaluated compounds exhibited the MICs in the area of 7,8–8,3 µg/mL which corresponded to 16 µmol/L in all cases. However, they were less active than isoniazide (INH) for which the value of 0,07 µg/mL (0,5 µmol/L) as MIC was assigned. On the other hand, considered molecules performed relatively higher *in vitro* efficiency against potential pathogenic non-tuberculous *Mycobacterium kansasii* My 235/80 comparing the standard INH (20). The structures 6d–6g had the MICs against listed mycobacterial strain in the range of 27,5–129,5 µg/mL (62,5–250 µmol/L), the molecules 8c–8e were actually more effective, their MICs were in the area of 8,3–15,7 µg/mL (16–32 µmol/L). Simultaneously estimated MIC related to INH was more than 34 µg/mL (250 µmol/L).

Consequently, there was continuous interest to extend the knowledge about the spectrum of antimicrobial activities of the target structures from both sets 6 and 8 which were then tested against *Staphylococcus aureus* ATCC 6538 (Gram-positive bacteria, *Micrococcaceae*), *Escherichia coli* CNCTC 377/79 (Gram-negative bacteria, *Enterobacteriaceae*) and *Candida albicans* CCM 8186 (yeast) as well. The idea for a selection of such structures was based on the knowledge that *meta*- as well as *para*-alkoxy substituted phenylcarbamic acid derivatives were previously more antimicrobially active comparing the *ortho*-substituted ones (13).

The level of a concrete antimicrobial activity of tested structures from both sets 6 and 8 was dependent on the *meta*-alkoxysubstituent length and type of the substituent attached at the phenyl ring in the basic part of the molecule.

The experimental investigations revealed that all evaluated substances were practically inactive against *Staphylococcus aureus* ATCC 6538, the most virulent

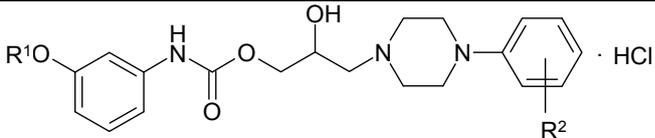
Staphylococcus species, with the MICs higher than 1000 µg/mL. Previously analyzing the influence of 2-piperidinoethyl-4-heptyloxyphenylcarbamate hydrochloride on some metabolic function of mentioned Gram-positive bacterial strain Mlynarčík *et al.* 1981 (14) suggested that the bacteriostasis could be equated to a loss of the cell's ability to synthesize ATP, which, in turn, may stem from an uncoupling of oxidative phosphorylation. Čižmárik *et al.* 1987 (5) comprehensively studied influence of piperidinoethyl esters of *ortho*-, *meta*- and *para*-alkoxyphenylcarbamic acid (alkoxy=metoxy to decyloxy) against given strain. They observed that *meta*- and *para*-substituted structures were antibacterially more active than *ortho*-substituted ones because of the steric effect. The proximity of *ortho*-alkoxy string to carbamate bond led to the twist of the benzene ring plain towards the carbamate group. Described process resulted in the planarity violation of molecule implying subsequent conjugation of π -bonds of benzene ring over NH fragment up to carbonyl group. The result was the different electronic density (charge) at carbonyl moiety which could be one of the possible binding sites to the reactive membrane locations. In the case of *meta*- and *para*-alkoxy derivatives described secondary steric effect did not manifest, even *para*-substituted derivatives were practically linear type of the molecule. In response to the paper (5), the lipophilicity of *meta*- and *para*-alkoxyphenylcarbamates was found as the factor conducted to the activity of such structures. More lipophilic structures displayed relatively higher efficiency. Carbonyl group also represents the centre at which the split occurred because of degradation (5). Čižmárik and Trupl (4) previously studied in very detail an antimicrobial effect of one structure from mentioned alkoxyphenylcarbamic acid esters set, heptacaine. They suggested that *Staphylococcus aureus* enzymically splitted heptacaine to *ortho*-heptyloxyaniline, which was then considered the actual antimicrobial agent. The observations from current evaluation of series 6 and 8 could confirmed a hypothesis postulated in a paper (4). There was an assumption

that the carbamate bond in evaluated compounds was splitted. After that, the activities of formed products, especially appropriate *meta*-alkoxyanilines, were much more lower than effectiveness of *ortho*-heptyloxyaniline due to a lower lipophilicity.

The investigated molecules in which structure was

incorporated only a single atom of fluorine, 6d–6g, were more active against *Escherichia coli* CNCTC 377/79 with MICs in the interval of 195,3–781,3 $\mu\text{g/mL}$ (444–1721 $\mu\text{mol/L}$) than the substances with trifluoromethyl substitution, 8c–8e, which were against listed bacterial strain inactive (Table 1).

Table 1. *In vitro* antimicrobial activity of structures 6d–8e against selected microbial strains (MICs expressed in $\mu\text{g/mL}$ and $\mu\text{mol/L}$ units, respectively).



Entry	R^1	R^2	<i>Staphylococcus aureus</i> ATCC 6538		<i>Escherichia coli</i> CNCTC 377/79		<i>Candida albicans</i> CCM 8186	
			$\mu\text{g/mL}$	$\mu\text{mol/L}$	$\mu\text{g/mL}$	$\mu\text{mol/L}$	$\mu\text{g/mL}$	$\mu\text{mol/L}$
6d	CH ₃	4'-F	12500	28414	195,3	444	1562,5	3552
6e	C ₂ H ₅	4'-F	12500	27537	781,3	1721	1562,5	3442
6f	C ₃ H ₇	4'-F	6250	13356	390,6	835	781,3	1670
6g	C ₄ H ₉	4'-F	6250	12967	390,6	810	781,3	1621
8c	CH ₃	3'-CF ₃	6250	12757	6250	12757	195,3	399
8d	C ₂ H ₅	3'-CF ₃	12500	24804	6250	12402	195,3	388
8e	C ₃ H ₇	3'-CF ₃	6250	12066	6250	12066	97,7	189

These preliminary findings indicated that for the activity of such tested phenylcarbamates against *Escherichia coli* CNCTC 377/79 was important the suitable length of alkoxy side string (methoxy substitution was preferred) as well as a presence of a moiety, attached at the *N*-phenylpiperazine fragment, with a comparable van der Waals radii towards atom of hydrogen but with different electronic configuration. The lowest MIC value against this Gram-negative bacterial strain was noticed for the compound 6d (195,3 $\mu\text{g/mL}$, 444 $\mu\text{mol/L}$).

The effect of fluorination of an aromatic system in the basic part of tested molecules could be documented by the hydrophobic Hansch-Leo π value ($\pi = \log P_{\text{C}_6\text{H}_5\text{X}} - \log P_{\text{C}_6\text{H}_6}$ for substituted benzenes) taken from the literature (1, 8). The π readout related to fluorine is 0,14; according to Bondi's van der Waals radii estimation, fluorine is moderately larger (1,47 Å) than hydrogen (1,20 Å). Its Hammett sigma constant for *para*-position (σ_p) is 0,06 and represents only slightly higher value

than is a readout for hydrogen ($\sigma_p=0,00$). For fluorine atom is also typical inductive electron-withdrawing as well as resonance electron-donating influencing (1, 8).

Presumably the electronic effects were primarily responsible for a better internalisation of the compounds tested (series 6) through the Gram-negative outer membrane. Present idea is supported by the research of Beveridge 1999 (2) or Hamadi *et al.* 2008 (7). One of an unusual features of the outer membrane of mentioned bacteria is its asymmetric distribution of lipids over inner and outer faces. The outer face contains (virtually) all of the lipopolysaccharides (LPSs), whereas the inner face has most of the phospholipids. LPSs contain more charge *per* unit of surface area than any phospholipids, and most of this charge is anionic at neutral pH because of exposed phosphoryl and carboxyl groups which can be readily ionized. The outer face of the outer membrane is highly charged. The resonance electron-donating effect of fluorine atom attached at

the *para*-position of aromatic ring performed there relatively electronic density enhancement. The consequent electronic interactions with different membrane components of *Escherichia coli* generally favoured molecules from the series 6 comparing the members from the set 8, in which was evident strongly deactivating influence of *meta*-trifluoromethyl fragment towards phenyl ring. Except of mentioned, for fluorine atom is typical relatively smaller van der Waals radii (and possible better internalisation into the membrane's structures) comparing trifluoromethyl group.

On the contrary, the significant enhancement of an activity against *Candida albicans* CCM 8186 was observed for the compounds 8c–8e, their estimated MICs were in the interval of 97,7–195,3 µg/mL (189–399 µmol/L). The substances 6d–6g were against this yeast practically inactive (781,3–1562,5 µg/mL, 1621–3552 µmol/L). In general, the evaluated series 8c–8e was even more effective against the yeast than previously tested dimethylaminoethyl, piperidinoethyl as well as perhydroazepinoethyl esters of *meta*-alkoxyphenylcarbamic acids with an identical *meta*-alkoxy substituent attached at the lipophilic aromatic ring (12, 13).

The experimental findings revealed that the level of an activity against *Candida albicans* of tested structures was dependent on the length and type of the substituent attached at the *N*-phenylpiperazine moiety (Table 1). From both analyzed sets 6 and 8 as the most active was considered compound 8e (with the lowest value of MIC) containing *meta*-propoxy side chain and lipophilic, sterically bulky trifluoromethyl fragment with its strong inductive electron-withdrawing effect. For an illustration, Hansch-Leo π value for trifluoromethyl group is 0,88; its electron-withdrawing influencing, expressed by the value of $\sigma_m=0,43$, is evident (1, 8). According to Tafts steric effects constants E_s , given substituent is also much larger in steric bulk than hydrogen or methyl moiety; for the comparison the corresponding E_s data are present: $E_{s_{CF_3}}=1,16$ (trifluoromethyl); $E_{s_{CH_3}}=0,00$ (methyl); $E_{s_H}=-1,24$ (hydrogen).

Following present experimental findings, the lipophilicity

of tested structures played a key role in their ability to act against *Candida albicans* CCM 8186. Approximately 80 to 90% of the cell wall of this yeast consists of the carbohydrate, three basic constituents represent the major polysaccharides of its cell wall (3) – β -glucans, chitin and glyco[manno]proteins. In addition, cell wall of mentioned yeast contains proteins (6 to 25%) and minor amounts of lipids (1 to 7%). The inner layer, enriched for chitin and polysaccharide matrix, is more electron translucent than outer layers which are enriched for glyco[manno]proteins. Relatively higher lipophilicity allowed easier internalisation of the compounds tested into the eukaryotic pathogens, as compared with the prokaryotic ones, causing a perturbation of the membranes especially by the substances from the series 8 resulting in the anticandidacidal effect at relatively lower MIC values. Present suggestion of a possible acting of evaluated structures is consistent with another published conclusions of Limban *et al.* 2011 (6).

In the future, we intend to extend our research within this class of alkoxyphenylcarbamic acid derivatives containing *N*-phenylpiperazine ring by the testing of new compounds with *meta*- and *para*-alkoxy substitution and with various substituents attached at the basic moiety for which are typical with electron-donating or electron-withdrawing effects.

CONCLUSION

In order to find new structures in the group of *meta*-alkoxyphenylcarbamic acid derivatives with improved antimicrobial activity, nine molecules were tested. Their chemical structures were previously confirmed by spectral (1H NMR, ^{13}C NMR, IR, MS, UV/VIS) data and by elemental analysis. The tested compounds exhibited a different level of antimicrobial activity against some selected Gram-positive (*Staphylococcus aureus* ATCC 6538), Gram-negative (*Escherichia coli* CNCTC 377/79) bacterial strains and against a yeast (*Candida albicans* CCM 8186). The activity was dependent on the length of *meta*-alkoxy side string co-creating

the lipophilic part, electronic and lipophilic properties of substituent attached at *N*-phenylpiperazine ring which formed a basic part of the molecule. The presence of relatively shorter alkoxy side chain and a single atom of fluorine in *para*-position at aromatic system favoured the antimicrobial activity against *Escherichia coli* CNCTC 377/79, while the highest inhibitory effect against *Candida albicans* CCM 8186 was exhibited by the most lipophilic structure from whole tested series containing propoxy fragment and *meta*-trifluoromethyl group attached at *N*-phenylpiperazine fragment. On the other hand, the substances tested were inactive against *Staphylococcus aureus* ATCC 6538.

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