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IN VITRO AND *IN SILICO* ASSESSMENT OF UREA/THIOUREA DERIVATIVES IN ANTIMICROBIAL THERAPY

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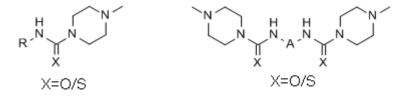
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The emergence and spread of antimicrobial resistance, together with the lack of newly developed antimicrobial drugs present serious public health issues worldwide. Apart from bacteria that do not respond to antibiotics, therapy-resistant mutants of viruses and multidrug resistant fungal strains introduce additional difficulties to effective therapy.^[1] Novel antimicrobial compounds containing chalcogenic elements (S, Se, Te) have attracted reasonable attention in the field of experimental chemotherapy.^[2]

The aim of our study was to evaluate the potency of novel urea and thiourea derivatives (1-10) against various microorganisms (aerobic and anaerobic bacteria, fungi, Herpes Simplex Virus 1) *in vitro*, as well as their conformity to the attributes of successful drug candidates, using computational *in silico* methods.

Figure 1.: General structure of the tested compounds



The antimicrobial activities of the tested compounds against reference strains of aerobic or anaerobic bacteria and yeasts were evaluated using disk diffusion tests and when warranted, broth microdilution method, according to EUCAST standards.^[3] A double-disk synergy test was used for the detection of synergism between the derivatives and antibiotics (ampicillin, cefuroxime, imipenem, vancomycin) against multiresistant bacterial strains.^[4] A MIC reduction assay was performed to quantify the effects of the (thio)ureas on the minimal inhibitory concentrations of reference drugs. The anti-HSV activity of the tested compounds was evaluated using MTT assay on Vero cells.^[5] The predicted physicochemical and pharmacokinetic properties of the tested compounds were determined *in silico*, using OSIRIS Molecular Property Explorer and PreADMET software.^[6,7]

Bacteria			Fungi
Aerobic	Anaerobic	Multiresistant	
Staphylococcus aureus	Clostridium perfringens	Staphylococcus aureus	Candida albicans
ATCC 25923	ATCC 13124	ATCC 43300 (MRSA)	ATCC 10231
Staphylococcus	Propionibacterium acnes	Enterococcus faecium	Candida krusei
epidermidis ATCC 12228	ATCC 11827	QC/2008/12/03 (VRE)	ATCC 14243
Enterococcus faecalis	Bacteroides fragilis	Acinetobacter baumanii	Candida glabrata
ATCC 29212	ATCC 25285	clin. isol. no.: 59738	ATCC 2001
Escherichia coli		Pseudomonas aeruginosa	Candida parapsilosis
ATCC 25922		ATCC 27863	ATCC 22019
Klebsiella pneumoniae			Candida tropicalis
ATCC 49619			ATCC 13803
Proteus mirabilis			Cryptococcus diffluens
PMI 60007			ATCC 32059
Salmonella Derby			
HWCMB 10032			
Pseudomonas aeruginosa			
PAE 170022			

Table 1.: Bacterial and fungal reference strains used in our experiments

The ureas/thioureas did not show antibacterial activity against the aerobic and anaerobic bacteria, nor did they inhibit the growth of the fungal strains included in our study (MICs of the tested compounds were >200 μ M; inhibition zone diameters <10 mm in all experiments). Some of the tested compounds presented modest adjuvant properties, reducing the inhibitory concentrations of antibiotics by 25-50% against susceptible bacteria, however they did not influence the effects of reference drugs against resistant strains. Based on the percentages of maximum cell recovery and selectivity, a 4-nitrophenyl-derivative and a 1,4-phenylene-substituted compound showed potent anti-HSV properties. All tested compounds complied with Lipinsky's Rule of Five and the predicted percentages of intestinal absorption are excellent (\geq 95%) for all the respective compounds.^[8,9]

The tested thio(ureas) lacked any antimicrobial activity against the bacterial and fungal strains, and their potential as antibiotic adjuvants is modest at best. On the other hand, their predicted ADME properties are of the highest standard and some of the compound showed promising antiviral activity. The synthesis and biological evaluation of novel derivatives in the future could introduce alternative antiviral drugs with selective activity.

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