

Synthesis and preliminary hepatotoxicity evaluation of new caffeine-8-(2-thio)-propanoic hydrazid-hydrazone derivatives

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Abstract

New series of caffeine-8-(2-thio)-propanoic hydrazid-hydrazone derivatives were designed and synthesized. The targeted compounds were obtained in yields of 51 to 96% and their structures were elucidated by FTIR, ¹H NMR, ¹³C NMR, MS and microanalyses. All of the compounds were found to be “drug-like” as they fulfill the criteria of drug-likeness, which includes the MDDR-like rule. The tested compounds were subjected to *in silico* prediction of substrate/metabolite specificity and Drug Induced Liver Injury (DILI). The prediction for indicated that the evaluated compounds would most probably act as CYP1A2 substrates. The performed *in vitro* studies didn't reveal statistically significant hepatotoxicity of the tested compounds, probably due to the pro-oxidant effects expressed on sub-cellular (isolated rat liver microsomes) level. The obtained experimental results confirmed the predicted low hepatotoxicity for the tested structures. Based on these results the compounds may be considered as promising structures for design of future molecules with low hepatotoxicity.

Keywords

caffeine, isolated rat liver microsomes, pro-oxidant effects

1. Introduction

Caffeine (1,3,7-trimethylxanthine) is a naturally occurring compound and one of the methylxanthines classified as a member of the alkaloid class of natural compounds. It is commonly distributed in plants throughout the world. Caffeine is also a key component of many beverages and food (coffee, tea, cocoa) and is widely consumed for its stimulating effect on the central nervous system. Nowadays, Caffeine is medically considered as the most popular, unregulated, and legal psychoactive drug in approximately all regions in the world (Alsabri et al. 2018). Caffeine and other

methylxanthines are used in clinical medicine as diuretics, analgesics, and muscle relaxants, and they can aid in the treatment of brain disorders such as headaches and Parkinson's disease (Al Deeb et al. 2002; Kolayli et al. 2004). Some of the effects of caffeine are due to caffeine-derived metabolites that have their own biological activities (Franco et al. 2013; Hetzler et al. 1990; Welsh et al. 2012). Caffeine and its derivatives in human liver are metabolized by the cytochrome P450 oxidase enzyme system (in particular by the CYP1A2 isoenzyme (Bloomer et al. 1995)). This metabolism

results in the formation of three major dimethylxanthine metabolites (paraxanthine, theobromine and theophylline) and one hydroxylated metabolite (1, 3, 7-trimethyluric acid) (Kot and Daniel 2008a), where paraxanthine is the predominant caffeine-derived metabolite in humans and 1, 3, 7-trimethyluric acid and theophylline are determined as the major metabolites in rodents and monkeys, respectively (Berthou et al. 1992; Kot and Daniel 2008b).

Oxidative processes are very important for living organisms. Under pathological or oxidative stress conditions reactive oxygen species (ROS) are overproduced which results in peroxidation of membrane lipids, leading to accumulation of lipid peroxides. On the other hand, oxidative stress is known to be involved in ageing and in various diseases, including diabetes mellitus, atherosclerosis, rheumatoid arthritis, Alzheimer's disease, Parkinson's disease and cancer (Metro et al. 2017). Some contradictory evidences are available for the influence of caffeine on oxidative stress processes (Acheson et al. 2004). It is known that some of the effects of caffeine lead to protection against cellular damage by producing beneficial antioxidant effects (Pasaoglu et al. 2011), since caffeine has been recognized as an efficient biological antioxidant (Shi et al. 1991). However, some data suggest that this property is observed only when caffeine acts as an effective inhibitor of lipid peroxidation, at millimolar concentrations, in rat liver microsomes (Devasagayam et al. 1996). On the other hand, caffeine shows *in vitro* prooxidant effect (Gulcin 2008) and is capable of inducing oxidative damage by fostering lipid peroxidation in rat liver (Dianzani et al. 1991). The role of caffeine in lipid peroxidation is varied and possibly depends on the dose of caffeine ingested.

Acyldiazones are another molecular scaffold, on the basis of which new biologically active compounds can be generated. They are compounds formed from the condensation of an acylhydrazine and a carbonyl reagent. Azomethine group (-NH-N=CH-) contained in their structure is connected with carbonyl group and is responsible for the different pharmacological applications of cited compounds. Hydrazid-hydrazones have, therefore, attracted considerable attention to their wide range of biological activities (Sarigöl et al. 2015), including antioxidant, anticancer and cytoprotective potency (Hristova-Avakumova et al. 2017; Nikolaevskii et al. 2012).

Microsomes are widely used test systems for investigation of the metabolic stability and metabolic profile of a large number of molecules during the drug discovery and development phases (Barcelos et al. 2014). Microsomal liver fractions combined with the possibility of automation of incubating process are elevated to high-tech applications. High storage stability of microsomes provides an opportunity to create a human bank for hepatic fractions and to study differences in enzyme activities in the population. Correlation analysis is applied to study metabolic pathways in the context of enzymatic topology (Barcelos et al. 2014), where malondialdehyde (MDA) is known to be the most frequently used biomarker of lipid peroxidation and oxidative stress evaluation.

In this study, based on the biological activity profiles of xanthines and acyl hydrazones, we hybridized these two

ring systems into one unit thus reporting the synthesis of series novel caffeine-8-(2-thio)-propanoic hydrazid-hydrazone derivatives. Based on some literary data, in addition, we set as purpose to evaluate the effects of the newly synthesized compounds on isolated rat liver microsomes as a model of hepatotoxicity on subcellular level, applying the level of generated malondialdehyde (MDA) in the rat liver microsomes as a measure of the hepatotoxicity.

2. Materials and methods

2.1. Materials

All chemicals and solvents were purchased from Merck AG (Merck, Darmstadt, Germany). The chromatographic system for TLC control and purity elucidation is based on an aluminium sheets Silica gel F254 (Merck, Darmstadt, Germany), using the following mobile phases: *Phase 1*: 25% NH₄OH/Acetone/CHCl₃/CH₃CH₂OH (1/3/3/4) and *Phase 2*: H₂O/n-butanol/CH₃COOH (5/4/1), with detection at UV 254 nm. Yields were calculated for purified products. A Buchi 535 capillary apparatus (Switzerland) was used to determine the melting points. The IR spectra 400 – 4000cm⁻¹ were recorded on a Nicolet iS10 FT-IR Spectrometer using ATR technique with Smart iTR adapter. UV-spectra were recorded on a Jenway 6715 UV/VIS Spectrophotometer, Japan. A Bruker Spectrospin WM250 spectrometer (Faerlands, Switzerland) was used to acquire ¹H-NMR and ¹³C-NMR spectra at 250 and 75 MHz, respectively. TMS as internal standard and DMSO-d₆ were used as internal standard and solvent, respectively, all OH and NH protons were D₂O exchangeable. The coupling constants (*J*) are expressed in Hertz (Hz). The mass spectra were recorded on a Dionex Ultimate 3000 RSLC Ultrasonic Liquid Chromatography System (Thermo Scientific) equipped of a six-channel SRD-3600 Degaser, a HPG-3400RS High Pressure Binary Gradient Pump, an Automatic Sample Injector WPS-3000TRS and TCC-3000RS column thermostat. The chromatographic system is connected to a Q-Exact Plus mass spectrometer (Thermo Scientific, Bremen, Germany) with Enhanced Resolution Mode enabled (up to 280,000 FWHM at m/z 200). Microanalyses, determined for C, N and H, were within ±0.4% of theoretical values and were performed on Euro EA 3000-Single, EUROVECTOR SpA analyser. All names were generated by using structure-to-name algorithm of ChemBioDraw Ultra software, Version 11.0, CambridgeSoft.

2.2. Chemistry

2.2.1. *Synthesis of initial intermediates 8-bromocaffeine (1), caffeine-8-(2-thio)-propanoic acid (3), methyl ester of caffeine-8-(2-thio)-propanoic acid (4) and caffeine-8-(2-thio)-propanehydrazide (5)*

The starting 8-bromocaffeine (1) was obtained using oxidative bromination of caffeine according to protocol, described by Mitkov et al (Mitkov et al. 2012). Intermediates 3, 4 and 5 are obtained in excellent yields and purity,

following the synthetic procedure published in our earlier paper (Mitkov et al. 2019).

2.2.2. General procedure for syntheses of substituted *N'*-substituted 2-(1,3,7-trimethyl-2,6-dioxo-2,3,6,7-tetrahydro-1H-purin-8-ylthio)propanehydrazides (6a-i)

In a round-bottomed flask equipped with a reflux condenser and an electromagnetic stirrer, the hydrazide **5** (0.0032 mol) and the corresponding ketone (0.004 mol) were mixed in 50 ml of ethanol. The reaction mixture is stirred under reflux. During the reaction, the precipitation of the product begins. Reaction time is determined by exhaustion of the starting reagents (TLC monitoring in phase 2) and varies from 40 minutes to 4 hours. The reaction mixture was cooled and filtered. If necessary, the filtrate is concentrated in vacuum to 10 ml and after several hours additional amount of product is separated. The product is recrystallized from ethanol.

***N'*-(1-phenylethylidene)-2-(1,3,7-trimethyl-2,6-dioxo-2,3,6,7-tetrahydro-1H-purin-8-ylthio)propanehydrazide (6a)**

FTIR (ATR), cm^{-1} : 3176, 1701, 1655, 1601, 1538, 1445, 1159; λ_{max} , nm: 217, 298; $^1\text{H-NMR}$ (DMSO- d_6), δ : 1.30 (3H, d, $J = 6.7$ Hz, -CH- CH_3), 2.16 (3H, s, =C- CH_3), 3.30 (s; 3H; N1- CH_3), 3.39 (s; 3H; N3- CH_3), 3.69 (s; 3H; N7- CH_3), 3.82 (1H, q, $J = 6.7$ Hz, S- CH), 7.32 (1H, t, $J = 7.4$, Ar-H), 7.43 (2H, m, Ar-H), 8.09 (2H, m, Ar-H); ^{13}C NMR (DMSO- d_6), δ : 13.1 (CH- CH_3), 16.1 (=C- CH_3), 27.9 (N¹- CH_3), 29.8 (N³- CH_3), 32.5 (N⁷- CH_3), 33.7 (S- CH), 138.3 (C_{Ar}-1), 126.1 (C_{Ar}-2 and C_{Ar}-6), 128.7 (C_{Ar}-3 and C_{Ar}-5), 128.9 (C_{Ar}-6), 107.1 (C-5), 145.4 (N=C), 147.0 (C-8), 148.8 (C-4), 151.4 (C-2), 154.7 (C-6), 167.3 (CO-NH); C₁₉H₂₂N₆O₂S (414.48); % calculated: C 55.06, H 5.35, N 20.28, S 7.73; and % found: C 54.98, H 5.25, N 20.47, S 7.70. LC-MS (70 eV) m/z (%): 415 (M+1), 416 (M+2).

***N'*-(1-*p*-tolylethylidene)-2-(1,3,7-trimethyl-2,6-dioxo-2,3,6,7-tetrahydro-1H-purin-8-ylthio)propanehydrazide (6b)**

FTIR (ATR), cm^{-1} : 3178, 1701, 1655, 1601, 1538, 1446, 1159; λ_{max} , nm: 216, 288; $^1\text{H-NMR}$ (DMSO- d_6), δ : 1.30 (3H, d, $J = 6.7$ Hz, -CH- CH_3), 2.12 (3H, s, =C- CH_3), 2.21 (3H, s, Ar- CH_3), 3.30 (s; 3H; N1- CH_3), 3.39 (s; 3H; N3- CH_3), 3.69 (s; 3H; N7- CH_3), 3.82 (1H, q, $J = 6.7$ Hz, S- CH), 7.13 (2H, m, Ar-H), 7.43 (2H, m, Ar-H); ^{13}C NMR (DMSO- d_6), δ : 13.1 (=C- CH_3), 16.1 (CH- CH_3), 21.3 (Ar- CH_3), 27.9 (N¹- CH_3), 29.8 (N³- CH_3), 32.5 (N⁷- CH_3), 33.7 (S- CH), 138.5 (C_{Ar}-1), 125.2 (C_{Ar}-2 and C_{Ar}-6), 129.3 (C_{Ar}-3 and C_{Ar}-5), 139.8 (C_{Ar}-6), 107.1 (C-5), 145.4 (N=C), 147.0 (C-8), 148.8 (C-4), 151.4 (C-2), 154.7 (C-6), 167.3 (CO-NH); C₂₀H₂₄N₆O₃S (428.51); % calculated: C 56.06, H 5.65, N 19.61, S 7.48; and % found: C 56.03, H 5.45, N 19.47, S 7.20. LC-MS (70 eV) m/z (%): 429 (M+1), 430 (M+2).

***N'*-(1-(4-methoxyphenyl)ethylidene)-2-(1,3,7-trimethyl-2,6-dioxo-2,3,6,7-tetrahydro-1H-purin-8-ylthio)propanehydrazide (6c)**

FTIR (ATR), cm^{-1} : 3171, 1701, 1652, 1603, 1538, 1446, 1159; λ_{max} , nm: 216, 294; $^1\text{H-NMR}$ (DMSO- d_6), δ : 1.30 (3H, d, $J = 6.7$ Hz, -CH- CH_3), 2.12 (3H, s, =C- CH_3), 3.30 (s; 3H;

N1- CH_3), 3.39 (s; 3H; N3- CH_3), 3.69 (s; 3H; N7- CH_3), 3.79 (3H, s, OCH₃), 3.82 (1H, q, $J = 6.7$ Hz, S- CH), 7.40 (2H, m, Ar-H), 7.28 (2H, m, Ar-H); ^{13}C NMR (DMSO- d_6), δ : 13.1 (=C- CH_3), 16.1 (CH- CH_3), 55.5 (OCH₃), 27.9 (N¹- CH_3), 29.8 (N³- CH_3), 32.5 (N⁷- CH_3), 33.7 (S- CH), 138.5 (C_{Ar}-1), 128.7 (C_{Ar}-2 and C_{Ar}-6), 114.1 (C_{Ar}-3 and C_{Ar}-5), 160.4 (C_{Ar}-6), 107.1 (C-5), 145.4 (N=C), 147.0 (C-8), 148.8 (C-4), 151.4 (C-2), 154.7 (C-6), 167.3 (CO-NH); C₂₀H₂₄N₆O₄S (444.51); % calculated: C 54.04, H 5.44, N 18.91, S 7.21; and % found: C 54.03, H 5.42, N 18.87, S 7.18. LC-MS (70 eV) m/z (%): 445 (M+1), 446 (M+2), 447 (M+3).

***N'*-(1-(2-hydroxyphenyl)ethylidene)-2-(1,3,7-trimethyl-2,6-dioxo-2,3,6,7-tetrahydro-1H-purin-8-ylthio)propanehydrazide (6d)**

FTIR (ATR), cm^{-1} : 3176, 1699, 1654, 1612, 1538, 1489, 1447, 1157; λ_{max} , nm: 216, 288; $^1\text{H-NMR}$ (DMSO- d_6), δ : 1.30 (3H, d, $J = 6.7$ Hz, -CH- CH_3), 2.19 (3H, s, =C- CH_3), 3.30 (s; 3H; N1- CH_3), 3.39 (s; 3H; N3- CH_3), 3.69 (s; 3H; N7- CH_3), 3.82 (1H, q, $J = 6.7$ Hz, S- CH), 7.53 (1H, d, $J = 7.3$ Hz, Ar-H), 7.23 (1H, d, $J = 8.8$ Hz, Ar-H), 7.16 (1H, s, Ar-H), 6.95 (2H, d, $J = 8.8$ Hz, Ar-H); ^{13}C NMR (DMSO- d_6), δ : 13.7 (=C- CH_3), 16.1 (CH- CH_3), 27.9 (N¹- CH_3), 29.8 (N³- CH_3), 32.5 (N⁷- CH_3), 33.7 (S- CH), 120.5 (C_{Ar}-1), 156.7 (C_{Ar}-2), 117.2 (C_{Ar}-3), 131.9 (C_{Ar}-4), 118.7 (C_{Ar}-5), 128.1 (C_{Ar}-6), 107.1 (C-5), 148.6 (N=C), 147.0 (C-8), 148.8 (C-4), 151.4 (C-2), 154.7 (C-6), 167.3 (CO-NH); C₁₉H₂₂N₆O₄S (430.48); % calculated: C 53.01, H 5.15, N 19.52, S 7.45; and % found: C 52.95, H 5.05, N 19.49, S 7.38. LC-MS (70 eV) m/z (%): 431 (M+1), 432 (M+2).

***N'*-(1-(4-hydroxyphenyl)propylidene)-2-(1,3,7-trimethyl-2,6-dioxo-2,3,6,7-tetrahydro-1H-purin-8-ylthio)propanehydrazide (6e)**

FTIR (ATR), cm^{-1} : 3274, 3173, 1694, 1652, 1607, 1538, 1514, 1447, 1157; λ_{max} , nm: 216, 308; $^1\text{H-NMR}$ (DMSO- d_6), δ : 1.30 (3H, d, $J = 6.7$ Hz, -CH- CH_3), 2.16 (3H, s, =C- CH_3), 3.30 (s; 3H; N1- CH_3), 3.39 (s; 3H; N3- CH_3), 3.69 (s; 3H; N7- CH_3), 3.82 (1H, q, $J = 6.7$ Hz, S- CH), 7.47 (2H, d, $J = 7.3$ Hz, Ar-H), 7.18 (2H, d, $J = 8.8$ Hz, Ar-H); ^{13}C NMR (DMSO- d_6), δ : 13.1 (=C- CH_3), 16.3 (CH- CH_3), 27.9 (N¹- CH_3), 29.8 (N³- CH_3), 32.5 (N⁷- CH_3), 33.7 (S- CH), 138.5 (C_{Ar}-1), 128.7 (C_{Ar}-2 and C_{Ar}-6), 116.9 (C_{Ar}-3 and C_{Ar}-5), 157.8 (C_{Ar}-4), 145.4 (N=C), 147.0 (C-8), 148.8 (C-4), 151.4 (C-2), 154.7 (C-6), 167.3 (CO-NH); C₂₀H₂₄N₆O₄S (444.51); % calculated: C 54.04, H 5.44, N 18.91, S 7.21; and % found: C 53.98, H 5.37, N 18.48, S 7.22. LC-MS (70 eV) m/z (%): 445 (M+1), 446 (M+2).

***N'*-(1-phenylpropylidene)-2-(1,3,7-trimethyl-2,6-dioxo-2,3,6,7-tetrahydro-1H-purin-8-ylthio)propanehydrazide (6f)**

FTIR (ATR), cm^{-1} : 3175, 1700, 1654, 1603, 1539, 1446, 1159; λ_{max} , nm: 216, 288; $^1\text{H-NMR}$ (DMSO- d_6), δ : 1.05 (3H, t, $J = 7.3$ Hz, -CH₂- CH_3), 1.30 (3H, d, $J = 6.7$ Hz, -CH- CH_3), 2.44 (2H, q, $J = 7.3$ Hz, =C- CH_2 -), 3.30 (s; 3H; N1- CH_3), 3.39 (s; 3H; N3- CH_3), 3.69 (s; 3H; N7- CH_3), 3.82 (1H, q, $J = 6.7$ Hz, S- CH), 7.27 (2H, m, Ar-H), 7.43 (2H, m, Ar-H), 8.09 (2H, m, Ar-H); ^{13}C NMR (DMSO- d_6), δ : 25.1 (=C- CH_2), 10.2 (CH₂- CH_3), 27.9 (N¹- CH_3),

29.8 (N³-CH₃), 32.5 (N⁷-CH₃), 33.7 (S-CH), 137.1 (C_{Ar}-1), 126.9 (C_{Ar}-2 and C_{Ar}-6), 128.7 (C_{Ar}-3 and C_{Ar}-5), 129.8 (C_{Ar}-6), 107.1 (C-5), 157.4 (N=C), 147.0 (C-8), 148.8 (C-4), 151.4 (C-2), 154.7 (C-6), 167.3 (CO-NH); C₂₀H₂₄N₆O₃S (428.51); % calculated: C 56.06, H 5.65, N 19.61, S 7.48; and % found: C 56.01, H 5.55, N 19.57, S 7.35. LC-MS (70 eV) *m/z* (%): 429 (M+1), 430 (M+2).

N'-(1-phenylbutylidene)-2-(1,3,7-trimethyl-2,6-dioxo-2,3,6,7-tetrahydro-1H-purin-8-ylthio)propanehydrazide (**6g**) FTIR (ATR), cm⁻¹: 3179, 1699, 1654, 1603, 1538, 1447, 1159; λ_{max}, nm: 216, 306; ¹H-NMR (DMSO-d₆), δ: 1.00 (3H, t, *J* = 6.7 Hz, -CH₂-CH₃), 1.33 (3H, d, *J* = 6.7 Hz, -CH-CH₃), 1.67 (2H, m, -CH₂-), 2.43 (2H, q, *J* = 7.3 Hz, -CH₂-CH₃), 3.30 (s; 3H; N1-CH₃), 3.39 (s; 3H; N3-CH₃), 3.69 (s; 3H; N7-CH₃), 3.88 (1H, q, *J* = 6.7 Hz, S-CH-), 7.27 (1H, m, Ar-H), 7.43 (2H, m, Ar-H), 8.22 (2H, m, Ar-H); ¹³C NMR (DMSO-d₆), δ: 28.4 (=C-CH₂), 13.8 (CH₂-CH₃), 21.1 (-CH₂-), 27.9 (N¹-CH₃), 29.8 (N³-CH₃), 32.5 (N⁷-CH₃), 33.7 (S-CH), 137.1 (C_{Ar}-1), 126.9 (C_{Ar}-2 and C_{Ar}-6), 128.7 (C_{Ar}-3 and C_{Ar}-5), 128.9 (C_{Ar}-6), 107.1 (C-5), 157.4 (N=C), 147.0 (C-8), 148.8 (C-4), 151.4 (C-2), 154.7 (C-6), 167.3 (CO-NH); C₂₁H₂₆N₆O₃S (442.54); % calculated: C 57.00, H 5.92, N 18.99, S 7.24; and % found: C 56.95, H 5.55, N 18.77, S 7.30. LC-MS (70 eV) *m/z* (%): 443 (M+1), 444 (M+2).

N'-(4-methoxyphenyl)(phenyl)methylene)-2-(1,3,7-trimethyl-2,6-dioxo-2,3,6,7-tetrahydro-1H-purin-8-ylthio)propanehydrazide (**6h**)

FTIR (ATR), cm⁻¹: 3272, 3175, 1699, 1652, 1603, 1538, 1445, 1157; λ_{max}, nm: 216, 288; ¹H-NMR (DMSO-d₆), δ: 3.82 (3H, s, -O-CH₃), 1.33 (3H, d, *J* = 6.7 Hz, -CH-CH₃), 3.30 (s; 3H; N1-CH₃), 3.39 (s; 3H; N3-CH₃), 3.69 (s; 3H; N7-CH₃), 3.88 (1H, q, *J* = 6.7 Hz, S-CH-), 7.32 (1H, m, Ar-H), 7.32 (2H, m, Ar-H), 7.43 (2H, m, Ar-H), 8.20 (2H, m, Ar-H), 8.22 (2H, m, Ar-H); ¹³C NMR (DMSO-d₆), δ: 55.5 (O-CH₃), 27.9 (N¹-CH₃), 29.8 (N³-CH₃), 32.5 (N⁷-CH₃), 33.7 (S-CH), 133.4 (C_{Ar}-1), 126.4 (C_{Ar}-2 and C_{Ar}-6), 128.7 (C_{Ar}-3 and C_{Ar}-5), 128.9 (C_{Ar}-6), 133.4 (C_{Ar}-1'), 131.2 (C_{Ar}-2' and C_{Ar}-6'), 114.1 (C_{Ar}-3' and C_{Ar}-5'), 160.4 (C_{Ar}-6'), 107.1 (C-5), 153.8 (N=C), 147.0 (C-8), 148.8 (C-4), 151.4 (C-2), 154.7 (C-6), 167.3 (CO-NH); C₂₅H₂₆N₆O₄S (506.58); % calculated: C 59.28, H 5.17, N 16.59, S 6.33; and % found: C 59.18, H 5.14, N 16.87, S 6.25. LC-MS (70 eV) *m/z* (%): 507 (M+1).

N'-(2-methyl-1-phenylpropylidene)-2-(1,3,7-trimethyl-2,6-dioxo-2,3,6,7-tetrahydro-1H-purin-8-ylthio)propanehydrazide (**6i**)

FTIR (ATR), cm⁻¹: 3218, 1694, 1655, 1607, 1522, 1465, 1158; λ_{max}, nm: 218, 294; ¹H-NMR (DMSO-d₆), δ: 1.29 (6H, d, *J* = 6.7 Hz, 2 x-CH-CH₃), 1.33 (3H, d, *J* = 6.7 Hz, -CH-CH₃), 2.45 (1H, m, =C-CH-), 3.30 (s; 3H; N1-CH₃), 3.39 (s; 3H; N3-CH₃), 3.69 (s; 3H; N7-CH₃), 3.88 (1H, q, *J* = 6.7 Hz, S-CH-), 7.37 (1H, m, Ar-H), 7.43 (2H, m, Ar-H), 8.35 (2H, m, Ar-H); ¹³C NMR (DMSO-d₆), δ: 30.3 (=C-CH), 21.0 (2 x CH-CH₃), 27.9 (N¹-CH₃), 29.8 (N³-CH₃), 32.5 (N⁷-CH₃), 33.7 (S-CH), 132.6 (C_{Ar}-1), 127.2 (C_{Ar}-2 and C_{Ar}-6), 128.7 (C_{Ar}-3 and C_{Ar}-5), 128.9 (C_{Ar}-6), 107.1 (C-5), 153.8 (N=C), 147.0 (C-8), 148.8 (C-4), 151.4 (C-2), 154.7 (C-6), 167.3

(CO-NH); C₂₁H₂₆N₆O₃S (442.54); % calculated: C 57.00, H 5.92, N 18.99, S 7.24; and % found: C 56.88, H 5.85, N 18.87, S 7.18. LC-MS (70 eV) *m/z* (%): 443 (M+1), 444 (M+2).

2.3. Biological evaluation

Animals

Male Wistar rats (body weight 200–250 g) were used. The rats were housed in plexiglass cages (3 per cage) in a 12/12 light/dark cycle, under standard laboratory conditions (ambient temperature 20 °C ± 2 °C and humidity 72 % ± 4 %) with free access to water and standard pelleted rat food 53-3, produced according to ISO 9001:2008.

Animals were purchased from the National Breeding Center, Sofia, Bulgaria. At least 7 days of acclimatization was allowed before the commencement of the study. The health was monitored regularly by a veterinary physician. The vivarium (certificate of registration of farm № 0072/01.08.2007) was inspected by the Bulgarian Drug Agency in order to check the husbandry conditions (№ A-11-1081/03.11.2011). All performed procedures were approved by the Institutional Animal Care Committee and made according Ordinance № 15/2006 for humane behavior to experimental animals.

Isolation and incubation of microsomes

The liver microsomes were isolated by ultracentrifugation, using Beckman L8-M centrifuge with a 70Ti rotor, for 1 hour, following Guengerich's methodology (Guengerich 1989). The protein measurement was carried out using method of Lowry et al (Lowry et al. 1951).

Measurement of malondialdehyde (MDA) in isolated rat microsomes

The concentration of MDA was determined spectrophotometrically at 535 nm by the Debyy & Gutier method (Deby and Goutier 1990).

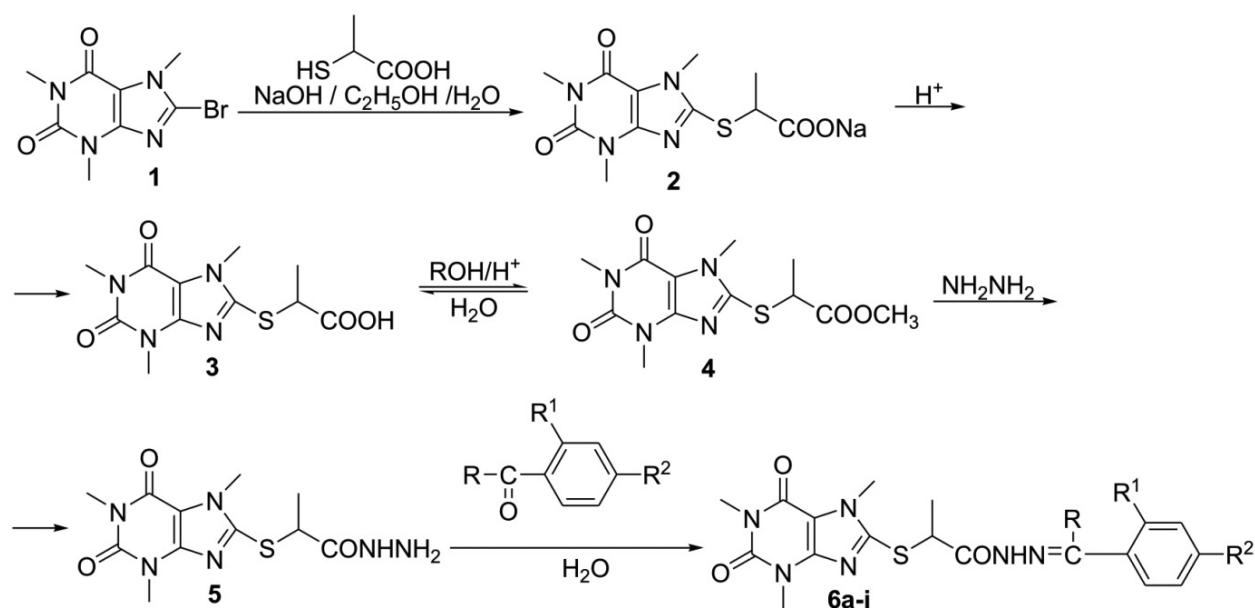
Statistical analysis

Statistical analysis was performed using statistical program "MEDCALC". Results are expressed as mean ± SEM for 7 experiments. The significance of the data was assessed using the nonparametric Mann-Whitney test. Values of *P* ≤ 0.05; *P* ≤ 0.01 and *P* ≤ 0.001 were considered statistically significant. Three parallel samples were used.

3. Results and discussion

3.1. Chemistry

The synthesis of described hydrazid-hydrazone derivatives was carried out following a procedure outlined on Scheme 1. The starting 8-bromocaffeine (**1**) was obtained using oxidative bromination of caffeine according to protocol, described



Scheme 1. General pathway for synthesis of a new caffeine-8-(2-thio)-propanoic hydrazid-hydrazone derivatives **6a-i**.

Table 1. ID, structures, m.p. and yields of newly synthesized hydrazones.

Compound	R	R ¹	R ²	M.p. (°C)	Yield (%)
6a	CH ₃	H	H	242–243	51
6b	CH ₃	H	CH ₃	248–250	69
6c	CH ₃	H	OCH ₃	24–249	73
6d	CH ₃	OH	H	239–243	68
6e	C ₂ H ₅	H	OH	228–230	80
6f	C ₂ H ₅	H	H	231–233	95
6g	C ₃ H ₇	H	H	242–246	96
6h	C ₆ H ₅	H	OCH ₃	219–221	62
6i	CH(CH ₃) ₂	H	H	243–245	82

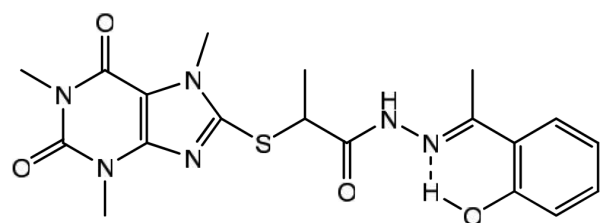


Figure 1. Formation of a 6-membered ring in **6d** due to an intramolecular hydrogen bond.

by Mitkov et al (Mitkov et al. 2012). Intermediates **3**, **4** and **5** are obtained in excellent yields and purity, following the synthetic procedure published in our earlier paper (Mitkov et al. 2019). The final products **6a-i** were prepared via condensation of the hydrazide **5** with the corresponding ketone in a molar ratio of 1: 1.25 in an ethanol medium. The compounds **6a-i** were obtained with purity (within \pm

0.4% of the calculated value), evaluated by microanalysis. The structures of the target compounds were confirmed on the basis of their FT-IR, ¹H-, ¹³C NMR and mass spectral data. The physical data of all synthesized compounds **6a-i** are given in Table 1.

In the performed spectral analysis a slight discrepancy in the expected and the measured IR spectra for compound **6d** was observed. This may be due to the intramolecular hydrogen bond which may be found in **6d** resulting in formation of a stable 6-membered ring as shown in structure below (Figure 1).

Thus no absorption bands due to the free O-H stretching vibration were found in the infrared spectra of any of the salicylidene anilines. Instead, a broad, weak band having some fine structure is found in the region from 2700 to 3100 cm⁻¹ as is the case with **6d**.

Also, a slight absorption band at 1157–1159 cm⁻¹, which is not observed in the spectra of starting xanthenes **1** and **3**, is clearly visible in the spectra of studied compounds **6a-i**. We consider this band to be due to N-N stretching vibration from the hydrazide residue. The identifications of this stretching vibration is a difficult task, but there is literary evidence to support our assumption (Bharanidharan et al. 2017; Gulaczyk et al. 2003).

Prediction of toxicity and drug-likeness

The toxicity and drug-likeness of compounds were predicted by using PROTOX II (“ProTox-II”) web server. The ProTox-II web server provides several advantages over existing computational models. It includes both chemical and molecular target knowledge. The prediction scheme is classified into different levels of toxicity such as oral toxicity as well as organ toxicity (hepatotoxicity). ProTox-II incorporates molecular similarity and pharmacophore based fragment propensities (Banerjee et al. 2018b).

Table 2. The values of the calculated molecular descriptors of tested compounds 6a–i.

Comp.	Molweight	Num. of hydrogen bond acceptors	Num. of hydrogen bond donors	Num. of atoms	Num. of bonds	Num. of rings	Num. of rotatable bonds	Total charge	Molecular Polar Surface Area
Cff	194.19	3	0	14	15	2	0	0	61.82
3	294.37	3	0	20	21	2	3	0	90.28
4	312.34	5	0	21	22	2	4	0	113.42
5	312.35	6	0	21	22	2	4	0	142.24
6a	414.48	6	0	29	31	3	6	0	128.58
6b	428.51	6	0	30	32	3	6	0	128.58
6c	444.51	7	0	31	33	3	7	0	137.81
6d	430.48	7	0	30	32	3	6	0	148.81
6e	428.51	6	0	30	32	3	7	0	128.58
6f	444.51	7	0	31	33	3	7	0	148.81
6g	442.53	6	0	31	33	3	8	0	128.58
6h	506.58	7	0	36	39	4	8	0	137.81
6i	442.53	6	0	31	33	3	7	0	128.58

Table 3. Results of toxicity prediction with ProTox-II webserver.

Comp.	Hepatotoxicity prediction	Probability	Predicted LD ₅₀ [mg/kg b.w.]	Predicted toxicity class
Cff	Inactive	0.97	127	3
3	Active	0.54	910	4
4	Inactive	0.65	196	3
5	Active	0.50	550	3
6a	Inactive	0.56	550	4
6b	Inactive	0.56	550	4
6c	Inactive	0.53	150	3
6d	Inactive	0.52	150	3
6e	Inactive	0.55	660	4
6f	Inactive	0.51	150	3
6g	Inactive	0.54	790	4
6h	Inactive	0.53	3000	5
6i	Inactive	0.56	660	4

An MDDRlike rule published by Oprea et al. (Oprea 2000; Oprea et al. 2000) was applied for drug-likeness evaluation of the newly synthesized compounds. The values of the calculated descriptors (Table 2) indicate that all compounds can be defined as “drug-like” as they fulfill the criteria of drug-likeness which includes the MDRP filter. MDDR-like rule (Oprea 2000; Oprea et al. 2000) is using as molecular descriptors in the means of number of rings, number of rigid bonds and number of rotatable bonds, while the probability of finding a ‘druglike’ compound being higher in the ranges of No. Rings ≥ 3 , No. Rigid bonds ≥ 18 , No. Rotatable bonds ≥ 6 , while the probability of finding a ‘non drug-like’ compound being higher in the ranges of No. Rings ≤ 2 , No. Rigid bonds ≤ 17 , No. Rotatable bonds ≤ 5 .

Drug-induced hepatotoxicity is a significant cause of acute liver failure and one of the major reasons for the withdrawal of drugs from the market. Drug-induced liver injury (DILI) is either a chronic process or a rare event. However, prediction of DILI is important and one of the safety concerns for the drug developers, regulators and clinicians (Liu et al. 2015). The data for the prediction of DILI are taken from DILIRank (Chen et al. 2016) and the NIH LiverTox database (Thakkar et al. 2018). The ProTox-II hepatotoxicity prediction model has a balanced accuracy of 82.00% on cross validation and 86.00% on exter-

nal validation. The AUC–ROC scores of cross-validation and external validation are 0.86 and 0.91 respectively. The kappa value is 0.69 for the model (Banerjee et al. 2018b). The model applied to predict hepatotoxicity is described in detail in the literature (Banerjee et al. 2018a).

The results of prediction are shown in Table 3. All the compounds appeared to be low toxic belonging to the category III and IV of GHS (Globally Harmonized System of Classification and Labeling of Chemicals) categories (“GHS-unece”).

Prediction of substrate/metabolite specificity

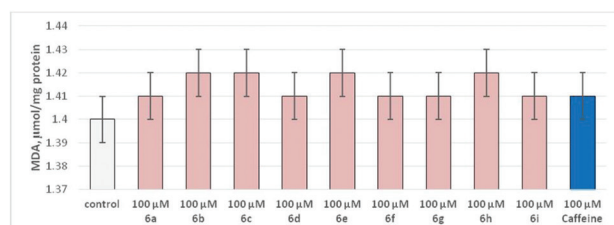
The evaluated set of compounds was subjected to a prediction of substrate/metabolite specificity, using an SMP web-service (Bezhtentsev et al. 2016) for *in silico* prediction of the substrate/metabolite specificity. This service provides the possible interaction of the tested group of compounds with 18 cytochrome P450 and UGT isoforms: CYP1A2, CYP2C9, CYP2C19, CYP2D6, CYP3A4, UGT1A10, UGT1A1, UGT2B7, UGT1A7, UGT2B15, UGT1A8, UGT1A4, UGT2B17, UGT2B10, UGT1A3, UGT1A9, UGT1A6, UGT2B4; defined as the most probable targets. The performed prediction is based on the PASS algorithm and is defined in the software Multilevel Neighborhoods of Atoms (MNA) descriptors. The obtained data

Table 4. “Probability to be Active” (Pa) values for the substrate based prediction result of **6a–i**, **3**, **4**, **5** and **Caffeine**.

Cmpnd	1A2	2A6	2E1	2C8
Cff	0.912	0.942	0.921	0.770
3	0.631	0.654	0.693	0.772
4	0.650	0.714	0.701	0.781
5				0.526
6a	0.672	0.697	0.687	0.594
6b	0.635	0.732	0.729	0.606
6c	0.679	0.705	0.695	0.629
6d	0.667		0.521	0.539
6e	0.754	0.794	0.788	0.673
6f	0.798	0.621	0.688	0.546
6g	0.822	0.858	0.839	0.623
6h	0.636	0.657	0.659	0.704
6i	0.721	0.764	0.765	0.712

Table 5. “Probability to be Active” (Pa) values for the metabolite based prediction result of **6a–i**, **3**, **4**, **5** and **Caffeine**.

Cmpnd	1A2	2E1	3A4
Cff	0.944	0.858	
3	0.677	0.677	
4	0.833		0.504
5	0.735	0.520	
6a	0.808	0.568	0.554
6b	0.818	0.557	0.544
6c	0.855		0.539
6d	0.667	0.583	
6e	0.859	0.686	
6f	0.781	0.742	
6g	0.903	0.762	0.515
6h	0.833		
6i	0.841	0.637	

**Figure 2.** Effect of the hydrazid-hydrazones **6a–i**, administered alone at concentration 100 µM, on isolated rat liver microsomes.

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for the substrate based predictions are presented in Table 4 for compounds **6a–i**, **3**, **4**, **5** and **Caffeine**. The results for the corresponding metabolite based predictions for the same set of structures are presented in Table 5. Based on the obtained data we consider the tested compounds to perform most probably CYP4501A2 substrate activity.

In vitro evaluation of effect on isolated rat liver microsomes

It is a well known fact that Caffeine is metabolized by CYP1A2. It is of interest to establish whether during their metabolism the newly synthesized caffeine derivatives **6a–i** would be biotransformed into reactive metabolites and by this to reveal possible pro-oxidative properties.

Thus we decided to evaluate their effect on isolated rat liver microsomes, as an *in vitro* system, which contains most of the isoforms of the cytochrome P450 system.

Administered alone, all of the tested compounds didn't reveal statistically significant pro-oxidant effects on isolated rat liver microsomes (Figure 2). Caffeine (as positive control) also didn't induce lipid peroxidation in statistical manner on isolated microsomes.

4. Conclusion

The present study describes the synthesis of series new caffeine-8-(2-thio)-propanoic hydrazid-hydrazone derivatives which are subjected to *in silico* prediction of substrate/metabolite specificity. Based on the obtained data we consider the tested compounds to perform most probably as CYP1A2 substrates. The newly synthesized hydrazid-hydrazones were also subjected to *in silico* prediction of the Drug Induced Liver Injury (DILI). The prediction for all compounds indicates that they would not show hepatotoxicity. The carried out *in vitro* studies confirmed that the tested compounds didn't reveal statistically significant hepatotoxicity, due to pro-oxidant effects, on sub-cellular (isolated rat liver microsomes) level. These results underlined the tested molecules as promising structures for design of future compounds with low hepatotoxicity.

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