



Research Article

A novel approach for apoptosis and caspase-3 inhibition using new candidates of 1,5-Diaryl triazole-3-carboxamides

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ABSTRACT

Decreasing or inhibiting apoptosis is a therapeutic strategy to reduce the pathological states of diseases like stroke, neurodegeneration, retinal cell death, myocardial and liver ischemia, and inflammatory diseases such as sepsis, osteoarthritis, rheumatoid arthritis, and asthma. This study aims to design and synthesize a new series of apoptosis inhibitors using a 1,5-diaryl-1,2,4-triazole-3-carboxamide scaffold. Then we conducted a comprehensive assessment of the effects of compounds **8a**, **8b**, **9a**, **10a**, and **10b** on two critical organs: the liver and ovary. *In vivo*, general biomarkers in the liver and ovary were measured and specific biomarkers were. Different stains such as Perl's iron stain, H & E stain and immune stain were utilized to evaluate caspase-3 inhibition. The findings are consistent with the docking study, i.e., compounds with two carbon linkers are better than those with longer or shorter ones as antiapoptotic. The most promising compounds **8a** and **10a** showed the highest potencies and selectivity against caspase-3 amongst other caspases like caspase -7, -8, -9, and 10 that could be considered for further clinical studies.

1. Introduction

Apoptosis is a form of cell death in which a programmed sequence of events leads to the elimination of cells [1]. Although apoptosis occurs typically during development and aging as a homeostatic mechanism to maintain cell populations in tissues, it is represented as a defense mechanism in immune reactions or when cells are damaged by diseases or noxious agents [2]. Too little or too much apoptosis due to imbalance probably occurs in some cases between apoptotic and antiapoptotic mediators leading to some diseases that are attributed to overexpression or inhibition of apoptotic mediators [2]. When apoptosis does not work correctly, cells that should be eliminated may persist and become immortal, for example, in cancer and autoimmunity [3, 4]. When

apoptosis works overly well, it kills too many cells and inflicts grave tissue damage; this is the case in strokes[5] and neurodegenerative disorders such as Alzheimer's [6], retinal cell death [7], myocardial and liver ischemia [8, 9], inflammatory diseases such as sepsis [10], osteoarthritis [11], rheumatoid arthritis [12], and asthma [13]. The Caspase family is a hot target for drug design because dysregulation of caspase expression leads to severe diseases. To date, 14 mammalian caspases have been identified, among them, caspases -2, -3, -6, -7, -8, -9, and -10 which were described as apoptotic caspases [14]. The nucleophilic cysteine in the caspase active site is involved in breaking down aspartic acid peptide bonds within proteins [3]. Caspase-3 is generally more promiscuous than caspase-7 and is revealed as the major

executioner of space during the demolition phase of apoptosis [15, 16]. Additionally, caspase-3 plays a key executioner role, and its inhibition can drastically prevent apoptosis *in vitro* and *in vivo* [14].

Liver ischemia is a life-threatening disease that may occur due to abnormal heart rhythms, heart failure, dehydration, sepsis, severe bleeding, blood clotting in the hepatic artery during liver transplantation, burns, and blood vessel swelling [17]. Ovary ischemia is one of the leading causes of polycystic ovary syndrome, the most common cause of ovulatory infertility in about 10% of women of childbearing age [18-20]. Ischemia/reperfusion (I/R) injury is the restoration of blood supply to organs after an ischemic period leading to parenchymal damage due to exaggerated apoptosis [21]. I/R injury elevates mortality rates to 60–80%. Because of the emergence of acute hepatic and ovarian ischemia and its ongoing impact on the global health system, there has been a growing interest in developing novel antiapoptotic agents to combat this disorder. As a result, we here reported our medicinal chemistry efforts, starting from the core scaffold of caspase inhibitors to the design of simpler and more efficient antiapoptotic compounds simulating the general structure of caspase inhibitors as illustrated in **Figure 1**. Most caspases recognize tetrapeptide sequences from the N-terminus to the C-terminus, positions of each residue on the existing substrate that could be named P4, P3, P2, and P1. The corresponding binding sites on the protein are named S4-S1 [22, 23]. The enzyme can hydrolyze the peptide bond after the P1 Asp [24]. Throughout the designed structure, **P1** was designed to be a straight-chain aliphatic carboxylic acid with different lengths. This provides more flexibility for the COOH group, which acts as a cysteine trap, to be more available for interaction with the suitable amino acid residues in the binding site of caspase. While **P2** was designed to be 1,2,4-triazole; triazole incorporation in some compounds increases the caspase inhibition activity as in compound **I** with $IC_{50} = 84.9$ nM when H1,4-disubstituted-1,2,3-triazole replaces H atom gave compound **II**, the inhibitory activity improved to be with $IC_{50} = 9$ nM for caspase-3 inhibition and as in case of Ald-SC-2, compound **III**, with $IC_{50} = 4.67$ μ M [25]. Another objective of this study was to identify the effect of **P3** and **P4** on inhibitory activity by trying *para*-substitution on ring **A** and the addition of ring **B** to increase the inhibitory activity of the designed compounds (**Figure S2**).

2. Results and discussion

2.1. Docking study

The target compounds, 1,2,4-triazole-3-carboxamide derivatives of different substituents on ring **A** with different lengths ($X=1-5$) (**Figure 2**), NAC as a reference

drug and the caspase 3 co-crystallized ligand NA3 ((3S)-3-(((5-bromopyridin-3-yl)carbonyl)amino)-4-oxobutanoic acid) were docked on caspase-3 protein (PDB: 1re1) [26]. The more the binding, the more caspase inhibitory activity and the more antiapoptotic activity along with the affinity and the value of the root-mean-square deviation of atomic positions (rmsd), which indicates the better fit and more stability of the compound conformation into the pocket [27, 28]. Based on the validation process, the obtained (rmsd) is 1.3341 Å. It is less than 2 Å indicating that the docking protocol with the receptor (PDB code: 1RE1) is valid and feasible for the advanced docking process. Docking results revealed that compounds of the linker 2 carbons i.e., carboxamides of 3-amino propionic acid (β alanine), are the best in affinity to the binding site of caspase 3. In an attempt to explain the highest potency of **10a** which is lower in affinity than **8a**, **9a** and **10d**. **10a** is a prodrug as it is an ester. Its carboxylate form was docked and the results were impressive high-affinity $E_{conf} - 69.9612$ kcal/mole while the ester form E_{conf} is 8.8229 kcal/mole. In addition to S-the final score, rmsd_refine between the pose before and after refinement, E_{conf} -The energy of the conformer, The interaction with the catalytic site of caspase 3, Cys 285 and His 237 was highly considered (**Figures 1** and **S3**). Compounds **8a**, **9a**, **10a(a)** and **10a** showed binding with the caspase-3 catalytic binding site (Cys 285 and Hist 237) and NAC and NA3. The docking results revealed that all compounds **8a**, **9a**, **10a**, NA3 and NAC showed similar bondings. O- of carboxylate formed ionic bonds with Arg(A) 179, and Arg(B) 341. C=O, the carboxylic group, showed interaction with Cys(A)285. The amidic carbonyl played as a hydrogen bond acceptor with His(A) 237. Triazole ring showed pi-cation interaction with His(A) 237. Compounds **8b**, **10b**, **9b** and **10b** did not show good binding with the catalytic site as they did not form binding with His(A)237.

2.2. Chemistry

Based on the docking study, compounds **8a**, **9a**, and **10a** may have expected good antiapoptotic effects, while **8b**, **9b**, and **10b** with low or no activity. At the same time, compounds of linkers of the length ($x=1, 3, 4$ or 5) may be of partial activity. So, compounds of the expected high activity like **8a**, **9a**, **10a** and the lowest activity **8b** and **10b** were selected to save time and money. The target 1,2,4-triazole-3-carboxamide derivatives **8a**, **8b**, **9a**, **10a** and **10b**) and their intermediates were prepared as outlined in **Scheme 1**. According to Schotten-Baumann, a nucleophilic substitution reaction was carried out throughout the acylation of glycine with benzoyl chloride, producing one equivalent 2-(benzamido)acetic acid **5**. Erlenmeyer azlactone synthesis of oxazolone intermediate

6 was performed through intramolecular condensation in a dehydrating agent such as acetic anhydride.[29, 30] Treatment of appropriate aniline derivatives with the nitrosonium ion generated *in situ* upon the reaction of sodium nitrite with HCl at 0 °C afforded the corresponding diazonium salt **2a**, **2b** and **2c** [31]. According to the Kuskov-like reaction, coupling of the diazonium salts **2a**, **2b**, and **2c** with the active methylene carbon of lactone **6** in the presence of sodium acetate afforded hydrazone derivatives **7a,7b** and **7c** in 75-85% yield. Based on Sawdey rearrangement, refluxing the mixture of oxazolones and the appropriate amino acid derivatives or heating it in the microwave for 5 min produced the target derivatives **8a**, **8b**, **9a**, **10a** and **10b** in a good yield. ¹HNMR and ¹³CNMR identified the newly synthesized compounds, while the purity of the newly prepared compounds was checked by elemental analysis.

2.3. Biological Study

2.3.1. The study depended on causing ischemia followed by reperfusion (I/R) in the liver and ovary in rats and studying the protective effect of the test compound in comparison with NAC. I/R results in a series of pathophysiological events, including elevation in mitochondrial permeability, the release of cytochrome c, caspase-3 activation[32], the release of inflammatory mediators such as TNF α , IL- β , etc., and the increase of oxidative stress that leads to apoptosis [33-35]. The decrease in tissue damage induced by I/R in a dose equivalent to 30 mg of NAC as an antiapoptotic reference [36-38]. The antiapoptotic mechanism was studied by biochemical, histological, immunohistochemical, and morphometric. Furthermore, serum biomarker concentrations were measured using suitable methods to detect liver and ovary injury. The biomarkers measured were either specific for each organ, liver and ovary or general for both as shown in Table S1. *In vivo* antiapoptotic study in liver

2.3.1.1. Assay of general biomarkers; (TNF α & MDA) inhibition and (SOD, GSH and NOx) activation in liver

2.3.1.2. Hepatic *in vivo* antiapoptotic study of compounds **8a**, **8b**, **9a**, **10a** and **10b** based on the conc. And the percentage of inhibition for the liver general biomarkers TNF α and MDA compared to sham, model, and NAC. Complying with docking results, compounds **8a**, and **10a** (with linker n=2 and *p*-methoxy or *p*-ethyl ester in ring A, respectively) displayed an excellent inhibitory effect on TNF α with 90.6 % and 90.47 %; respectively in comparison with NAC (45.2 %) and on MDA with 90.77% and 88.1 % better than NAC (39%). Although compound **9a** (with linker n=2 and *p*- F in ring A) showed a moderate TNF α inhibitory effect (50.14%), it had better inhibition than NAC on MDA. Compounds **8b** and **9a** (with linker n = 5) also possessed a weak to no inhibition. Interestingly,

compounds **8a**, and **10a** showed excellent activation of SOD (98.57 % and 79.12%, respectively), GSH (101.33% and 98.5% respectively) and NOx (105.48 % and 96.46 %, respectively) greater than NAC (52.4%, 77% and 67.71%, respectively) as listed in Table 1. In contrast, compounds **9a**, **8b**, and **10b** had a weak activation. These findings further support our hypothesis that compounds with short-chain linkers (n=2) are the best antiapoptotic compounds.

2.3.2.1 Assay of specific liver biomarkers (ALT and AST inhibition)

The liver-specific biomarkers (ALT and AST) were found in the case of apoptosis induction due to I/R. Remarkably, compounds **8a** and **10a** demonstrated robust inhibition of ALT activity (93.62% and 99.17%, respectively) and AST activity (101.79% and 99.35%, respectively) when compared to NAC (42% and 27.5%, respectively) (Table 2).

2.3.2.2 Assay of specific ovarian biomarker (AMH)

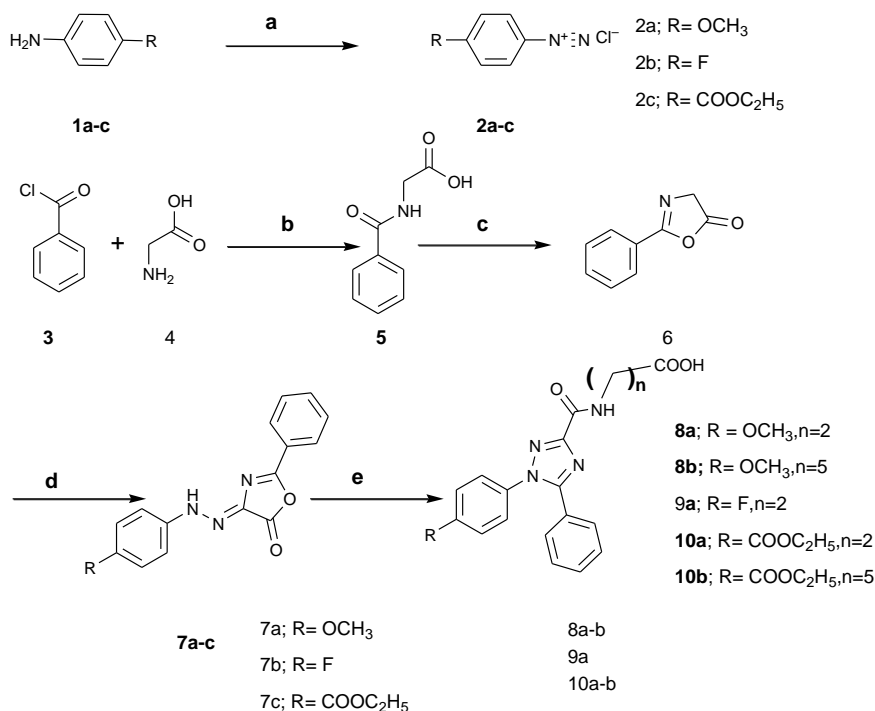
Based on the results of general biomarkers estimations, it is predictable good antiapoptotic evidence on ovarian biomarkers. **8a**, **10a** and **9a** revealed outstanding AMH activation (88.23% 110.6% and 109.41%, respectively) as listed in Table 4. Additionally, others with five carbon linker lengths were worth more than NAC. These experiments prove that some of the newly docked compounds will be good candidates as antiapoptotic agents. The above biochemical results for the ovary and liver match those observed before docking. Whereas **8a** and **10a** showed good binding with Cys285, which means more caspase-3 inhibition and less apoptosis, compounds with no interaction with Cys285 as in compounds **8a** and **10b** result in low caspase inhibition and more apoptosis. Moreover, **9a** showed excellent amelioration in the case of the ovary, which may be attributed to the fluorine group that probably enhances the lipophilicity of the drug, increasing the drug cell penetration.

Table 4: AMH concentrations in the ovary of I/R rats treated with 0 or 1.83 mM of **8a**, **8b**, **9a**, **10a**, **10b** and NAC.

Group	Serum AMH (ng/mol)	Activation %
Sham	2.18	100
Model	0.48	0
NAC	1.211	43
8a	1.98	88.23
8b	0.62	8.24
9a	2.34	109.41
10a	2.36	110.6
10b	0.70	12.94

2.3.3 *In vivo* Antiapoptotic study in the ovary:

2.3.3.1 Assay of general biomarkers; (SOD, GSH and NOx) activation and (MDA) inhibition in the ovary



Reagents and conditions: (a) NaNO₂, HCl, stirring; (b) NaOH, HCl, stirring; (c) Ac₂O/ 60 °C; (d) compound (5a-e) NaOAc, 2-8 °C, 4h, stirring; (e) H₂N(CH₂)_nCOOH (n = 1,2,3,4), AcOH, NaOAc, reflux three h (55-60% yield) or microwave 3 min (80-85% yield)

Scheme 1: Synthesis of the target compounds **8a-b**, **9a**, **10a-b**

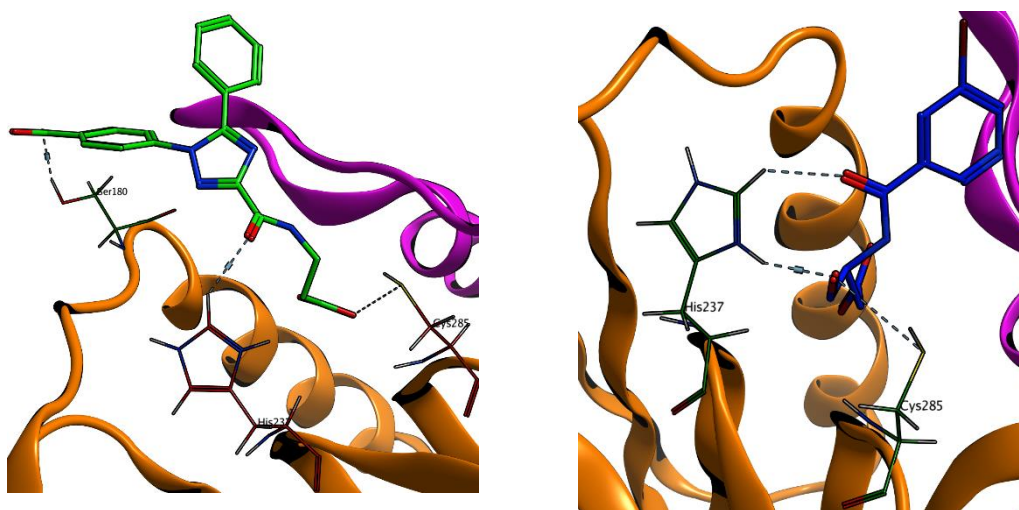


Figure 1: A 3D representation of the docking poses of compound **10a(a)** (green) and the co-crystallized ligand **NA3** (blue) in the caspase-3 catalytic site

Table 1: TNF α , MDA inhibition %, and SOD, GSH, and NOx activation in the liver of I/R rats treated with 0 or 1.83 mM of **8a**, **8b**, **9a**, **10a**, **10b** and NAC.

Biomarker		Sham ^a	Model ^b	NAC	8a	8b	9a	10a	10b
TNF α	Conc (pg/mL)	9.86	31.48	19.63	11.89	30.74	20.64	11.92	32.14

	% inhibition	100	0	45.2	90.6	3.4	50.14	90.47	-3.05
MDA	Conc (nmol/g tissue)	27.08	61.43	40.47	30.25	64.25	65.32	31.17	66.78
	% inhibition	100	0	39	90.77	-8.2	-11.32	88.1	-15.57
SOD	Conc (nmol/mg protein)	40.01	10.74	26.1	39.59	13.29	11.15	33.90	11.87
	% Activation	100	0	52.4	98.57	8.71	1.40	79.12	3.86
GSH	Conc (µmol / mg protein)	83.24	56.11	77.19	83.60	50.11	53.82	82.83	50.79
	% Activation	100	0	77	101.33	-22.12	-8.44	98.5	-19.61
NOx	Conc (nmol/ gm tissue)	141.79	83.6	123	145.19	80.57	81.68	139.73	79.60
	% Activation	100	0	67.71	105.48	-5.21	-3.3	96.46	-6.87

Sham^a = No drug, without I/R Model^b = No drug, with I/R

Table 2: The inhibition percent of ALT and AST in the liver of I/R rats treated with 0 or 1.83 mM of **8a**, **8b**, **9a**, **10a**, **10b** and NAC.

Group		Sham ^a	Model ^b	NAC	11b	11d	13b	14b	14d
ALT	Conc. (U/I)	28.55	69.6	45.75	31.17	72.26	70.64	29.89	70.46
	% inhibition	100	0	42	93.62	-6.48	-2.1	99.17	-13.86
AST	Conc. (U/I)	37.75	80.65	49.33	36.98	75.99	81.1	37.47	76.45
	% inhibition	100	0	27.5	101.79	10.87	9.8	99.35	-1.05

Sham^a = No drug, without I/R Model^b = No drug, with I/R

Table 3: SOD, GSH, and NOx concentrations in the ovary of I/R rats treated with 0 or 1.83 mM of **8b**, **8d**, **10b**, **11b**, **11d** and NAC.

Group	Test	Sham ^a	Model ^b	NAC	8a	8b	9a	10a	10b
SOD	Conc (nmol/mg protein)	142.01	65.93	109.45	133.11	73.22	136.29	140.97	69.22
	% Activation	100	0	57.21	88.30	9.6	92.5	98.63	4.32
GSH	Conc (µmol / mg protein)	44.95	25.51	41.84	46.35	22.28	45.87	47.03	23.77
	% Activation	100	0	84	107.20	-16.62	104.37	110.7	-8.95
NOx	Conc (nmol/ gm tissue)	171.61	105.92	145.33	168.42	107.45	170.24	165.34	100.51
	% Activation	100	0	60	95.14	2.33	97.91	90.45	-8.24
MDA	Conc (nmol/g tissue)	39.24	72.71	59.60	38.05	73.06	41.53	37.42	69.65
	Inhibition %	100	0	28	103.56	-1.05	93.16	94.56	9.14

Sham^a = No drug, without I/R Model^b = No drug, with I/R

The ovary was previously subjected to I/R to induce apoptosis. **Table 3** listed the results of **8a**, **10a** and **9a** (with linker n= 2) to exhibit again relatively significant activation for SOD (88.30 % 98.63% and 92.5 %, respectively), GSH (107.20% and 110.7%, 104.37 %, respectively) and NOx (95.14%, 90.45%, 97.91%, respectively). Moreover, they exhibited MDA inhibitory effects (103.56%, 94.56% and 93.16 %, respectively). On the other hand, **8b** and **10b** (with linker n= 5) possessed very weak activity. These findings provide further evidence for the suggested aspect; derivatives of short linkers showed the greatest antiapoptotic effect.

2.3.3.2. Assay of specific ovarian biomarker (AMH)

Based on the results of general biomarkers estimations, it is predictable good antiapoptotic evidence on ovarian biomarkers. **8a**, **10a** and **9a** revealed outstanding AMH activation (88.23% 110.6% and 109.41%, respectively) as listed in **Table 4**. Additionally, others with five carbon linker lengths were worth more than NAC. These experiments prove that some of the newly docked compounds will be good candidates as antiapoptotic agents. The above biochemical results for the ovary and liver match those observed before docking. Whereas **8a** and **10a** showed good binding with Cys285, which means more caspase-3 inhibition and less apoptosis, compounds with no interaction with Cys285 as in compounds **8a** and **10b** result in low caspase inhibition and more apoptosis.

Moreover, **9a** showed excellent amelioration in the case of the ovary, which may be attributed to the fluorine group that probably enhances the lipophilicity of the drug, increasing the drug cell penetration.

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10a	2.36	110.6
10b	0.70	12.94

2.4 Histopathological studies of liver and ovarian tissues

Hematoxylin and eosin (H& E) were used for histological examination while Perl's stain was used to detect iron [39, 40]. Activated caspase-3 immunohistochemistry was used as a marker for detecting apoptosis in tissue sections [41]. Numerous apoptotic hepatocytic cells with intense reactions were detected by immunohistochemical study. This is in good harmony with previous researchers, who stated an increase in caspase-3 immunoreaction in the myocardium after six hours of ischemia [40]. The apoptotic histological changes observed in the present work were in accordance with previous studies that confirmed early damage of cells after reperfusion mainly in the form of apoptosis. They stated that numerous amounts of both cytochrome c and other apoptosis-enhancing molecules are released in ischemia-reperfusion, triggering the cells' apoptosis [42]. While other researchers described apoptosis as a specific histological feature of cell death, characterized by shrunken cells, chromatin condensation, and nuclear fragmentation [43]. Some previous workers reported that increased lipid peroxidation that occurred during I/R, led to the synthesis of both proteolytic lysosomal and mitochondrial matrix enzymes into the cytoplasm, leading to intracellular proteolysis and tissue damage [44]. Ovarian tissue that was taken from the sham group displayed a thin germinal layer covered the ovaries. It showed normal morphology with multiple follicles in various stages of development, ranging from primordial follicles to mature Graafian follicles. The growing follicles showed oocytes surrounded by distinct zona pellucida surrounding the oocytes. The Parenchymal ovarian cells surrounding the growing follicle are organized into concentric sheaths named theca folliculi. The secondary follicles were formed when little fluid-filled spaces

between the granulosa cells appeared. When these spaces were enlarged and fused, they formed prominent mature Graafian follicles. The oocyte appeared peripherally in the follicle enclosed in the cumulus oophorus and surrounded by granulosa cells. Corpora lutea was also observed in lightly stained lutein cells between minimal connective tissue stroma (**Figure S8**).

In the current study, in the I/R group (model), the ovaries showed secondary follicles with intra-antral desquamated follicular cells, multiple atretic follicles without oocytes, and multiple dark nuclei. The corpora lutea appeared with hemorrhage and massive hemorrhage (**Figure S9**). These findings were under previous histological studies done on ischemic ovaries. These results are referred to the action of reactive oxygen species (ROS) formed during I/R. The I/R group exhibited massive interstitial hemorrhage and congested vessels in this work. It was in line with those who stated that torsion of the ovarian blood supply led to severe vascular congestion, hemorrhage, and eventually multiple necroses of ovarian tissue [45].

Using Perl's iron staining, it was found that intracellular cytoplasmic ferritin increased in the I/R group, and it was in line found the same finding in the case of compounds **8a** and **10a**; this ferritin staining appeared more extensive if compared with the sham group and compounds **9a**, **8b** and **10b**. Also, it was seen more than the I/R group. It could be explained that ferritin H could suppress apoptotic cell death, with subsequently increasing tubular epithelial viability and improving function by limiting oxidative stress after I/R injury. In the present study, an immunohistochemical study for caspase-3 showed more markedly evident positive brownish staining for apoptosis, especially in follicular cells and interstitial cells in the I/R group if compared with the other groups. Previous researchers demonstrated that caspase-3 immunoreactivity was observed in the ovarian tissues in the I/R group and other groups. It was explained that apoptosis was performed by activating proteases called caspases, resulting in DNA fragmentation. The apoptotic process was detected by immune detecting active caspase-3. Furthermore, Caspase-3 enhances endogenous endonucleases that lead to DNA fragmentation.

2.4.1 H and E results for liver tissue (better to be before immune effect)

Sham sections of liver tissue stained with H&E revealed normal lobular architecture. The hepatocytes radiated as cords from the central veins. They had acidophilic cytoplasm and vesicular nuclei. Some hepatocytes appeared with two nucleoli. The blood sinusoids lay between the hepatic cells. The ischemic reperfusion (I/R) group showed that the hepatocytes had a variable amount of severe cytoplasmic vacuolations. Apoptotic cells were noticed in H&E staining (**Figure 2**).

2.4.1 Ovary

It is worth mentioning that both compounds **8a** and **10a** caused amelioration of all previous ovarian injuries caused by ischemic reperfusion injury if given before ischemia. The more or less standard lobular architecture was noticed. At the same time, the still variable amount of atretic follicles with the absence of its ova was seen. Furthermore, the corpora lutea appeared with minor hemorrhage (**Figure 3**).

H& E ovarian sections (**Figure S4 and S5**): Compound **9a** group showed a thick epithelial germinal layer covering ovarian tissue. Disturbed lobular architecture with congested blood vessels was also noticed. The focal ovarian stromal loss was seen. Stromal cells are arranged in clusters and show pale cytoplasm. Some ovarian follicles in this group displayed severe follicular cell loss without their ova. Multiple atretic follicles with separation of their follicular cells were detected. The corpora lutea showed degenerative features and appeared with congested blood vessels. H& E ovarian sections for compound **10b** showed disturbed ovarian lobular architecture. Stroma showed severe atretic follicles, dilated congested blood vessels, and interstitial hemorrhage. The proliferation of stromal cells was also noticed. Homogenous acidophilic material appeared in the ovarian stroma. Corpora lutea with severe vacuolations were frequently seen. Sever demarcations were seen between aggregated Corpora lutea (**Figure S6**).

2.4.2 Perl's staining results for liver and ovarian tissues.

By using Perl's iron staining (**Figure 4**), it was observed that intracellular cytoplasmic ferritin granules increased in the I/ R group, and it was per another researcher noticed the same finding. Using Perl's staining for compounds **8a** and **10a** groups, this ferritin staining was more intense and diffuse if compared with sham and I/R groups. It could be explained by Hatcher et al., who reported that ferritin H could decrease apoptotic cell death, with subsequent enhanced tubular epithelial viability and ameliorate renal function by limiting oxidative stress after I/R injury.

In the ovary, the sham group showed faint blue cytoplasmic staining, while the I/R group exhibited more blue cytoplasmic coloration compared with the sham group. Both compounds **8a** and **10a** group showed more cytoplasmic staining than a sham and previous I/R group. Compounds **8b,9a** and **10b** groups showed cytoplasmic staining less than **8a** and **10a** groups but still more than the sham group (**Figure S7**).

2.4.3 Immunohistochemical study for caspase-3 in the liver and ovary tissues.

An immunohistochemical study for caspase in the Sham group showed negative expression, while in the ischemic group, intense and massive hepatic cytoplasmic immunoreaction in liver cells was noticed. (**Figure S8**) In the ovary, an immunohistochemical study for caspase in the Sham group showed negative expression, while in the ischemic group, intense and massive ovarian cytoplasmic immunoreaction in ovarian cells was noticed. Regarding **9a, 8b**, and **10b** groups; the ovarian tissue showed more intense cytoplasmic and nuclear reactions than observed in the sham group. It was noticed that the **10a** and **8a** groups showed little cytoplasmic expression for caspase if compared with ischemic reperfusion, **9a, 8b**, and **10b** (**Figure S9**).

2.4.4 Morphometric studies for liver and ovary tissues.

For the morphometrical study; ten non-overlapping fields from three sections of each rat of all groups were used [46]. It was found that there was a high significance increase in liver cells immunostained for caspase-3 in the model group if compared with the sham group ($P1 < 0.001$). Additionally, it was a significant increase in liver cells immunoreactivity in **9a, 8b** and **10b** ($p < 0.001$, $P < 0.001$, $P5 < 0.00$) consequently if compared with the sham group. At the same time, there was a significant decrease in this expression in the case of **8a** and **10a** if compared with **9a, 8b** and **10b**, as shown in **Table S1**.

For ovary tissues, it was found that there was a high significance increase in ovarian cells immuno-stained for caspase-3 in the model group if compared with the sham group ($P1 < 0.001$). Additionally, it was a significant increase in ovarian cells immunoreactivity in both **9a, 8b** and **10b** ($p < 0.001$, $P < 0.001$, $P5 < 0.00$) consequently if compared with the sham group. At the same time, there was a significant decrease in this expression in the case of **8a** and **10a** if compared with **9a, 8b** and **10b**, as shown in **Table S1**.

2.5 Caspase-3 enzyme inhibition and selectivity assay

Caspase-3 enzyme inhibition IC_{50} assay was done in Vacsera using the ELISA method for the most active compounds **8a** and **10a**, which had 85137.62 ng/mL and 2432.82 ng/mL. So, the selectivity assay was done for **10a** toward other apoptotic caspases such as caspase 7, 8, 9, 10, and the IC_{50} equaled 6150.643 ng/mL, 884.7528 ng/mL, 35737 ng/mL, 1161.181 ng/mL.

3. Conclusion

A novel class of 1,5-diaryl triazole-3-carboxamide hybrid scaffolds was designed, synthesized, and characterized by

different spectroscopic techniques. The prepared compounds retain good-to-moderate caspase-3 inhibitory activity and possess substantial antiapoptotic activity. Our investigation unveiled that, among the synthesized target compounds, **8a** and **10a** both possessing the shortest chain linker (n=2)- emerged as the most potent antiapoptotic candidates when compared to NAC. Furthermore, these compounds offered valuable insights into their robust caspase-3 inhibitory activity.. A detailed mechanistic study of molecular docking and biochemical and histopathological studies in the liver and the ovary prove this unexpectedly potent antiapoptotic activity that is expected to be a priority of future work.

4. Experimental

4.1 Chemistry

Chemicals were purchased from Aldrich, Alfa Aesar, Across Organics and El-Nasr Pharmaceutical Chemical Companies and used without further purification. The reaction progress was monitored using precoated TLC plates (Kiesslgel 60, 254 Merck); the spots were detected by exposure to a UV lamp at 254 nm. Analytical grade chemicals and solvents were used, and the progress of the reactions was monitored by thin-layer chromatography pre-coated Merck silica gel 60 F254 aluminum sheets. Melting points were determined on Stuart's electro-thermal melting point apparatus in degrees Celsius (°C) and were uncorrected. NMR spectra were recorded using Bruker AG (500 MHz ¹H NMR, 125 MHz ¹³C NMR), Switzerland, Faculty of Pharmaceutical Sciences, Umm Al-Qura University, Mecca, Saudi Arabia and on Bruker Advance III (400 MHz ¹H NMR, 100 MHz ¹³C NMR), Columbia, Vancouver, Canada; using TMS as an internal reference. Chemical shifts (δ) were expressed in ppm relative to TMS (δ=0 ppm) as the internal standard and CDCl₃ or DMSO-*d*₆ as a solvent. Chemical shifts (δ) were expressed in parts per million (ppm), and coupling constants (*J*) were expressed in Hertz. The signals were designated as follows: s, singlet; d, doublet; t, triplet; m, multiplet. Elemental analyses were recorded on Shimadzu GC/MS-QP5050A, The Regional Center for Mycology and Biotechnology, Al-Azhar University, Egypt. The microwave device (household oven) was operated at maximum power.

4.1.1 General procedure for the synthesis of hippuric acid (**5**)

Benzoyl chloride (3.96 g, 0.028 mol) was added in three portions to a cold stirred solution of glycine (2.25g, 0.028 mol) in an aqueous sodium hydroxide (6 M, 15 mL). After each addition of benzoyl chloride, sodium hydroxide (6 M) was added to keep the pH between 7.4 and 8.0. The stirring was continued for about 2 h until a clear yellowish solution formed and then diluted hydrochloric acid was

added in a portion-wise manner. The formed precipitate was filtered off, washed with water, air dried, and recrystallized in distilled water to afford white crystals of compound **5** (m.p: 187 – 189 °C) (reported: 187-188 °C) in 90% yield

General procedure for the syntheses of (Z)-2-Phenyl-4-(2-arylhydrazono)oxazol-5(4H)-one derivatives (**7a,b,c**) [48, 49]

Hippuric acid (0.013 mol, 3.50 g) in acetic anhydride (7 mL) was heated until a clear yellow solution was obtained; this solution was cooled to room temperature (solution A). To a cold solution of the *p*-substituted aniline derivatives (0.013 mol) in 5 N HCl (3 mL) in an ice-salt bath at 0-5°C, a solution of sodium nitrite (0.013 mol, 0.897 g) in the least amount of water was added in a dropwise manner. The reaction mixture was left for 10 min. (solution B). Solution A was added to solution B in the presence of anhydrous sodium acetate (0.018 mol, 1.5 g). The reaction mixture was stirred at 0-10°C for 2 h; the formed precipitate was filtered off, air-dried, and recrystallized from ethyl acetate affording the appropriate oxazolone derivatives in a good yield.

4-(2-(4-Methoxyphenyl)hydrazono)-2-phenyloxazol-5(4H)-one (**7a**)

Orange-red powder (yield: 85%, m.p: 205-206°C, reported m.p: 201-203°C)[50]

4-(2-(4-Fluorophenyl)hydrazono)-2-phenyloxazol-5(4H)-one (**7b**)

Deep orange powder (yield: 87%, m.p: 220-222°C); ¹H NMR (500 MHz, DMSO-*d*₆) δ(ppm): 11.84 (s, 1H, NH), 8.12-8.09 (m, 2H, ArH), 7.73-7.70 (m, 1H, ArH), 7.65-7.60 (m, 2H, ArH), 7.50-7.45 (d, 2H, ArH), 7.24-7.21 (m, 2H, ArH); ¹³C NMR (125MHz, DMSO-*d*₆) δ(ppm): 161.72, 159.87, 157.96, 139.83, 134.06, 132.16, 129.83, 128.28, 125.83, 117.09, 116.53; DEPT¹³C NMR (125MHz, DMSO-*d*₆) δ(ppm): 134.06, 129.83, 128.28, 117.09, 116.53.

Ethyl-4-(2-(5-oxo-2-phenyloxazol-4(5H)-ylidene)hydrazinyl) benzoate (**7c**)

Orange powder (yield 86%, m.p: 209-210 °C);¹H NMR (500 MHz, DMSO-*d*₆) δ(ppm): 12.66 (bs, 1H, OH), 11.98 (s, 1H, NH), 8.86 (s, 1H, NH), 8.02 – 7.87 (m, 3H, ArH), 7.83 – 7.72 (m, 2H, ArH), 7.59 – 7.47 (m, 4H, ArH), 4.39-4.34 (m, 2H, CH₂), 1.35-1.29 (m, 3H, CH₃);¹³C NMR (125 MHz, DMSO-*d*₆) δ(ppm): 166.93, 163.15, 147.19, 134.28, 131.91, 131.33, 129.90, 129.05, 128.84, 128.56, 127.70, 114.95, 60.87, 14.73; DEPT¹³C NMR (125 MHz, DMSO-*d*₆) δ(ppm): 131.91, 131.33, 128.56, 127.70, 114.95, 60.87, 14.73

General procedure of *p*-substituted (1,5-diaryl-1H-1,2,4-triazole-3-carboxamido) acid derivatives (**8a,8b**), **9a**, (**10a**, **10b**)[51, 52]

A mixture of oxazolone derivatives (0.01 mol), the appropriate amino acid analogs (0.012 mol), and

anhydrous sodium acetate (0.50g) are dissolved in acetic acid glacial (15 mL). The reaction mixture was heated under reflux for 3 h or under the microwave; the reaction ran for 5 minutes (divided into 30 sec for each run to prevent explosive of solvent) and was left to cool in the air. The reaction mixture was then diluted with ice water (4 mL for each 1 mL acetic acid). The precipitate was filtered off, washed with water, and re-crystallized from toluene to obtain white crystals.

3-(1-(4-Methoxyphenyl)-5-phenyl-1H-1,2,4-triazole-3-carboxamido)propanoic acid (8a)

White crystals; (r = 53% yield, MW = 76%) ; mp: 160°C; ¹H NMR (500 MHz, CDCl₃) δ (ppm): 8.09 (bs, 1H, NH), 7.50 (d, J = 7.5 Hz, 2H, ArH), 7.45 (t, J = 7.5 Hz, 1H, ArH), 7.40 – 7.35 (m, 2H, ArH), 7.31–7.27 (m, 2H, ArH), 6.93 (d, J = 7.5 Hz, 2H, ArH), 3.86 (s, 3H, OCH₃), 3.82–3.78 (m, 2H, CH₂NH), 2.76–2.73 (m, 2H, CH₂C=O); ¹³C NMR (125 MHz, CDCl₃) δ (ppm): 176.13, 160.24, 159.19, 155.71, 154.84, 130.69, 130.33, 129.06, 128.73, 126.90, 126.47, 114.57, 55.60, 34.98, 33.78; Anal. Calcd for C₁₉H₁₈N₄O₄ (366.38): C, 62.29; H, 4.95; N, 15.29; Found; C, 62.05; H, 5.13; N, 15.47.

6-(1-(4-Methoxyphenyl)-5-phenyl-1H-1,2,4-triazole-3-carboxamido)hexanoic acid (8b)

White crystals; (r = 72% yield, MW = 87%, mp: 95°C); ¹H NMR (400 MHz, CDCl₃) δ (ppm): 9.41 (s, 1H, OH), 7.51 (d, J = 7.5 Hz, 2H, ArH), 7.47 (t, j = 6.8 Hz, 1H, NH), 7.43 – 7.37 (m, 2H, ArH), 7.36 – 7.28 (m, 3H, ArH), 6.92 (d, J = 7.5 Hz, 2H, ArH), 3.84 (s, 3H, OCH₃), 3.50 (q, j = 6.8 Hz, 2H, CH₂NH), 2.36 (t, J = 7.2 Hz, 2H, CH₂C=O), 1.74 – 1.52 (m, 4H, 2CH₂), 1.56 – 1.19 (m, 2H, CH₂CH₂); ¹³C NMR (100 MHz, CDCl₃) δ (ppm): 178.63, 160.21, 159.05, 155.99, 154.67, 130.58, 130.47, 128.98, 128.69, 126.90, 126.70, 114.56, 55.60, 39.28, 33.87, 29.19, 26.32, 24.36; Anal. Calcd for C₂₂H₂₄N₄O₄ (408.46): C, 64.69; H, 5.92; N, 13.72; Found; C, 64.97; H, 6.09; N, 14.01.

3-(1-(4-Fluorophenyl)-5-phenyl-1H-1,2,4-triazole-3-carboxamido)propanoic acid (9a)

White crystals; (r = 62% yield, MW = 88%, m.p = 153°C); ¹H NMR (400 MHz, DMSO-*d*₆) δ (ppm) : 12.33 (bs, 1H, COOH), 8.68 (t, J = 5.6 Hz, 1H, NHCH₂), 7.59–7.54 (m, 2H, ArH), 7.51–7.47 (m, 3H, ArH), 7.47–7.43 (m, 2H, ArH), 7.4–7.37 (m, 2H, ArH), 3.50 (q, J = 6.8 Hz, 2H, CH₂ NH), 2.56–2.51 (m, 2H, CH₂C=O); ¹³C NMR (125 MHz, DMSO-*d*₆) δ (ppm): 173.45, 163.59, 161.78, 158.98, 156.66, 155.17, 130.95, 129.30, 129.16, 128.99, 127.34, 117.11, 35.48, 34.22; Anal. Calcd for C₁₈H₁₅FN₄O₃ (354.34): C, 61.01; H, 4.27; N, 15.81; Found C, 61.26; H, 4.45; N, 16.07.

3-(1-(4-(ethoxycarbonyl)phenyl)-5-phenyl-1H-1,2,4-triazole-3-carboxamido)propanoic acid (10a)

White crystals; (r = 74% yield, MW = 89%, m.p = 164°C); ¹H NMR (400 MHz, CDCl₃) δ (ppm): 8.09 (d, J = 6.8 Hz, 2H, ArH), 8.01 (t, J = 6.0 Hz, 1H, NHCH₂), 7.53–7.46 (m, 2H, ArH), 7.46–7.41 (m, 3H, ArH), 7.39 (d, J = 6.8 Hz, 2H, Ar.

H), 4.41 (q, J = 7.2 Hz, 2H, OCH₂CH₃), 3.84–3.76 (m, 2H, CH₂CH₂NH), 2.74 (t, J = 6.0 Hz, 2H, CH₂C=O), 1.41 (t, J = 7.2 Hz, 3H, CH₃CH₂); ¹³C NMR (100 MHz, CDCl₃) δ (ppm): 176.29, 165.34, 159.03, 156.40, 155.23, 140.74, 131.24, 131.00, 130.76, 129.11, 128.91, 126.37, 125.08, 61.56, 35.00, 33.75, 14.27; Anal. Calcd for C₂₁H₂₀N₄O₅ (408.41): C, 61.76; H, 4.94; N, 13.72; Found C, 61.54; H, 5.03; N, 14.01.

6-(1-(4-(Ethoxycarbonyl)phenyl)-5-phenyl-1H-1,2,4-triazole-3-carboxamido)hexanoic acid (10b)

White crystals; (r = 53% yield, MW = 67%, m.p = 145°C); ¹H NMR (500 MHz, DMSO-*d*₆) δ (ppm): 11.98 (s, 1H, COOH), 8.67 (bs, 1H, NH), 8.05 (d, J = 7.0 Hz, 2H, ArH), 7.59 (d, J = 7.0 Hz, 2H, ArH), 7.50–7.48 (m, 3H, ArH), 7.46–7.44 (m, 2H, ArH), 3.32–3.25 (m, 4H, OCH₂ and CH₂NH), 2.24–2.20 (m, 2H, CH₂C=O), 1.56–1.50 (m, 6H, 3CH₂), 1.34–1.28 (m, 3H, CH₂CH₃); ¹³C NMR (125 MHz, DMSO-*d*₆) δ (ppm): 174.96, 166.85, 158.92, 157.28, 155.14, 141.17, 132.07, 131.07, 131.00, 129.42, 129.22, 127.38, 126.28, 61.54, 39.06, 34.08, 29.30, 26.42, 24.71, 14.56; Anal. Calcd for C₂₄H₂₆N₄O₅ (450.50): C, 63.99; H, 5.82; N, 12.44; Found C, 64.15; H, 5.97; N, 12.70.

4.2 Docking study

The 3D of caspase-3 was obtained from Protein Data Bank (<https://www.rcsb.org/>; PDB ID code: 1re1). Protein contains a reference molecule called nicotinic acid succinic semialdehyde derivative (NA3) as a co-crystallized ligand. The protein binding sites were managed using modules of Ligand Atoms Selection of MOE, and the NA3 binding sites were used in the crystal structure as a reference. The sequence order of the binding sites was set using the MOE program. Hydrogen was added to complete the protein structure using selected Protonate 3D of the MOE program; only ligand and protein chains A and B of the crystal structure were used. The protein was then aligned using Align module of MOE. For the docking simulation process, the placement was managed on a triangular matcher, rescoring was managed on the London dG parameter, and the conformation number was managed in 5 poses with 30 refinements; conformation was selected as the forcefield of MOE and was used to generate five poses. All the docking conformations were analyzed, and the best value with the particular pose was selected for further interactional study.

4.3 Biology

Sera were separated and stored in aliquots at 80 °C till used for estimation of liver transaminases, including alanine transaminase (ALT) and aspartate transaminase (AST) by enzymatic colorimetric methods using commercial kits (spectrum diagnostic, Cairo, Egypt). MDA, an index of lipid peroxidation, was determined by using 1,1,3 tetra methoxy propane as standard [53]. A

tumor necrosis factor (TNF α) ELISA kit (IDlabsT- Minc. Biotechnology, Canada) was used. Anti-Mullerian Hormone (AMH) ELISA kit (Abcam, USA) was utilized. Total NO $_x$, the stable oxidation end product of nitric oxide, served as an index of nitric oxide level and was measured by reduction of nitrate into nitrite using activated cadmium granules, followed by color development with Griess reagent in an acidic medium [54].

The activity of SOD was measured by the method with a slight modification. This method depends on the inhibition of the pyrogallol autooxidation by SOD. Calculating the percentage of inhibition for the samples was done by running a control with no sample under the same conditions. [55] SOD activity was expressed in units; where the unit equals the amount of the enzyme that inhibited 50% of pyrogallol autooxidation.

The percentage of activation or inhibition is calculated as follows:[56]

Calculation % of activation = [(X-min)/(max-min)] x 100

Calculation % of inhibition = 1-[(X-min)/(max-min)] x 100

X = specific conc. (U/I), **min** = least conc., **max** = last conc

4.4 Caspase inhibition assay method

Using abcam kits for caspases

4.5 Histology

4.5.1 Material and Methods

Ovarian specimens were obtained from each group and fixed in 10% formol saline for 24 h. Paraffin blocks were carried on, and 5- μ m thick sections were subjected to H&E, Perls' stain. Other sections were further used for immunohistochemical study.

4.5.2 Immunohistochemical study

The caspase-3 antibody is a rabbit polyclonal for staining formalin-fixed paraffin-embedded tissues (Biovision, Milpitas, California, USA). The primary antibody (diluted at 1:20) was applied to the sections for 60 min. Then, the sections were incubated with the secondary antibody (biotinylated rabbit polyclonal antibody). The reaction was visualized with diaminobenzidine tetrahydrochloride and was noticed to be cytoplasmic. Sections were then counterstained with Mayer's hematoxylin.

4.5.3 Image capture

Histological sections were captured using Olympus digital camera mounted on a BX51 microscope (Olympus, Japan) connected to a computer programmed with LC micro application software in the light microscopic unit of the Histology Department, Faculty of Medicine, Minia University. Images were saved as jpg. Adobe Photoshop 7 standardizes brightness, contrast, and background color.

4.5.4 Morphometric analysis

Quantitative data were collected to measure the mean number of immune-stained cells for caspase. Measurements were performed in 8 randomly selected non-overlapping fields from the rat of each group under magnification X 400.

4.5.5 Data handling and Statistical analysis

Quantitative data were analyzed using SPSS (IBM Corp. Released 2010. Windows, Version 19.0). The mean and standard error (SE) was calculated for the parameters of each group. Values were expressed as means \pm SE. One-way analysis of variance (ANOVA) test was carried out to estimate significant differences between groups, followed by Tukey-Kramer as a post hoc test. The results were considered statistically significant if the p-values were <0.05.

Conflicts of Interest

The authors declare no conflict of interest

Supplementary File

https://odr.journals.ekb.eg/article_357547.html

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