Design, Synthesis, Docking and 2D **QSAR** studies of novel 3,5-diaryl Pyrazole Derivatives and their evaluation as Antioxidants and as Immunomodulators, inhibitors of TNF- α , IL-2, IL-6

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Introduction

- Orally active small molecules that modify the proinflammatory cytokine release associated with
 many auto-immune disorders such as rheumatoid
 arthritis (RA) have generated considerable interest
 in the pharmaceutical industry. They offer a costeffective and convenient alternative to biologics
 such as Enbrel, Remicade, Humira and Kineret
- These agents are expensive, parenterally administered.
 They are also under review for increased risk of cancer, infection, multiple sclerosis, and for the potential to induce neutralizing antibodies over the long term

- Tumor necrosis factor-α (TNF-α), one of the major pro-inflammatory cytokines, has been proven to be a potential target for these agents. TNF-α has been called a sentinel cytokine or "the body's fire alarm"
- The overexpression of TNF-α has been implicated in a number of serious inflammatory disorders such as rheumatoid arthritis, multiple sclerosis, inflammatory bowel disease, graft-versus-host disease, and adult respiratory distress syndrome.
- TNF-α is a strong inducer of other proinflammatory cytokines such as interleukins IL-1, IL-6 and IL-8

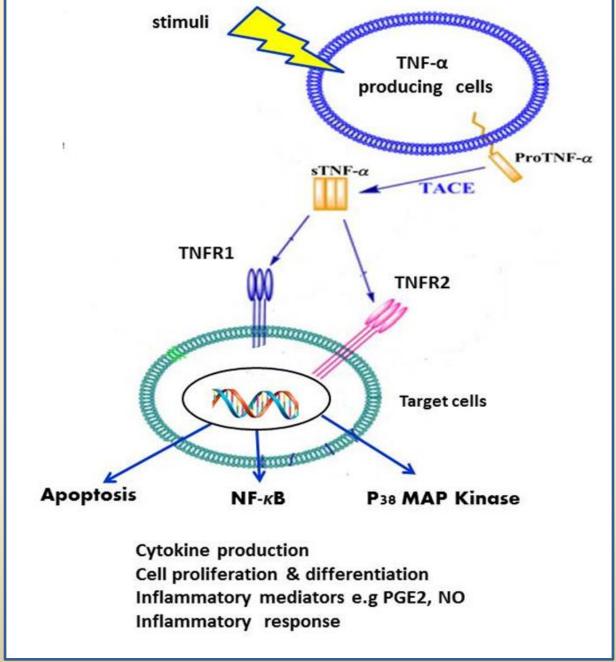


Figure (1): Receptor binding and biological actions of TNF-α (Chem Biol Drug Des 2010)

- **IL-6** is a potent pro-inflammatory agent that plays a crucial role in the pathogenesis of **systemic** inflammatory disease. Targeting this pathway in rheumatoid arthritis (RA) seems an attractive route as IL-6 is important for both **joint destruction** and **systemic manifestations**.
- It promotes inflammatory events through the expansion and activation of T cells and the differentiation of B cells.
- IL-6 blockade is a major advancement in the treatment of RA as it targets a unique molecule.

- IL-2 proved to play a pivotal role in regulating immune response, its suppression has been widely used to prevent allograft rejection in organ transplantion.
- IL-2 inducible T-cell kinase (ITK) has been found to play an important role in T-cell activation and proliferation, where it is primarily expressed.
- Therefore, ITK represents a novel potential target for anti-inflammatory therapy in a variety of indications such as psoriasis and allergic asthma.

- **P38-** also known as cytokine-suppressive antiinflammatory drug binding protein (CSBP), is a member of the **mitogen activated protein** (MAP) kinase family that is involved in stress and inflammatory response signal transduction pathways.
 - It is critical for the production and activity of multiple pro-inflammatory cytokines, including TNF-α, IL-1, IL-6, and IL-8, in cells such as macrophages, monocytes, synovial cells, and endothelial cells.

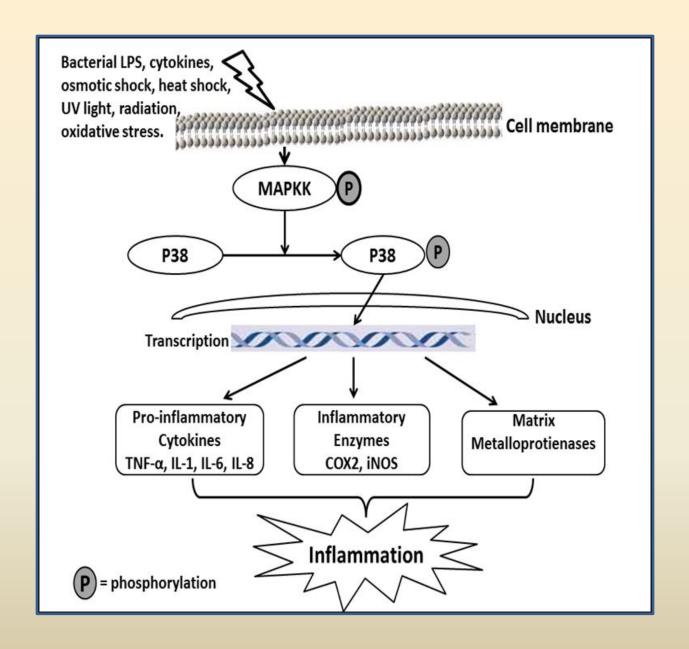


Figure (2): P38 MAPK regulation of inflammation (*Pharmacol Ther 1999*)

Therefore, inhibition of these targets has become a major focus of current drug discovery and development in treatment of severe inflammatory disorders.

Examples of pyrazole based scaffolds as immunomodulators

Clinical candidate SD0006

Burnette *et al.*, Pharmacology (2009) TNF- α = 0.016 µmol/L

Das *et al.*, Bioorg Med Chem (2010) $p38\alpha$ IC50 = 2 nM

TNF α = 75 %

Pfizer (SC 806) 2003 p38 α IC50 = 50 Nm

TNF α = 98 %, at 5mg/kg

B. P. Bandgar et al. Bioorg. Med. Chem. (2010)

IL-6 = 47%

TNF- α = 24% inhibition at 10 μ M

BTP-3 = 314 nM

Wu Chen et al., Cellular Immunology (2002)

Celecoxib

Liu Y et al., Cancer Prev Res (Phil 2011) Wang et al., Oncol Rep (2014)

Scuto et al., Leukemia (2011)

Aim of The Work

Fig (3): Some representative examples of pyrazole-based cytokine inhibitors and the novel compounds

Synthesis

Scheme 1: Synthesis of N-(4-acetyl phenyl) benzamide derivatives

5b R = 4-F 5c R = 2-OH 5d R = 4-OCH₃ 5e R = 4-CH₃ 5f R = 3,4,5-tri-OCH₃ 5g R = 2-OH-3-OCH₃ 5h R = Furyl 5i R = 4-Cl 5j R = 2,4-Cl 5k R = 2-Cl 5l R = 2-CH₂CH₃

Scheme 2: Synthesis of N-(4-3-(pheny)-1-prop-2-en-1-one phenyl)benzamide derivatives

c R = 2-OH d R = 4-OCH₃ e R = 4-CH₃ f R = 3,4,5-tri-OCH₃ g R = 2-OH-3-OCH₃ h R = Furyl i R = 4-Cl j R = 2,4-Cl k R = 2-Cl l R = 2-CH₂CH₃

a R = H **b** R = 4-F

Scheme 3: Synthesis of 4-chloro-N-(subs. phenyl-1H-pyrazol-3-)benzamide derivatives

BIOLOGICAL EVALUATION

In vivo TNF-α, IL-2, IL-6 Assay In Rat (Acute LPS Model)

- The novel compounds were evaluated for their ability to inhibit LPS-induced production of TNF-α, IL-2 and IL-6 in rat at 30 mg/kg p.o.
- Enzyme-linked immunosorbent assay kit, life science inc. (E90133Ra), (E90073Ra), (E90079Ra).
- Dexamethasone was used as a reference drug.

$$H_3CO$$
 OCH_3
 $OCH_$

OCH₃

Ю

6f = 58 %.

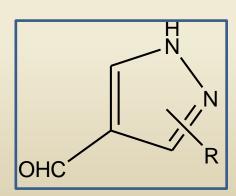
6g = 62 % Dexamethasone = 66%

6f = 42 %

Dexamethasone = 57 %

Antioxidant Activities

 Increased generation of reactive oxygen species (ROS) has been observed in degenerative diseases. It has been reported that the pyrazole core possesses radical-scavenging ability and even its modulation in inflammatory response was sometimes related to its considerable antioxidant activity.



- Moreover, it has been reported that certain antioxidants reduce LPS-induced inflammation and fever.
- Glutathione peroxidase is an important enzyme in cellular antioxidant defense systems, detoxifying peroxides and hydroperoxides.
- If GPX activity is decreased, more hydrogen peroxide is present, which leads to direct tissue damage and activation of nuclear factor-κB–related inflammatory pathways.

- <u>Superoxide Dismutase (SOD)</u> is one of the most important antioxidative enzymes.
- It catalyzes the dismutation of the superoxide (O_2^-) radical into either ordinary molecular oxygen (O_2) or hydrogen peroxide (H_2O_2) .

Antioxidant Activities

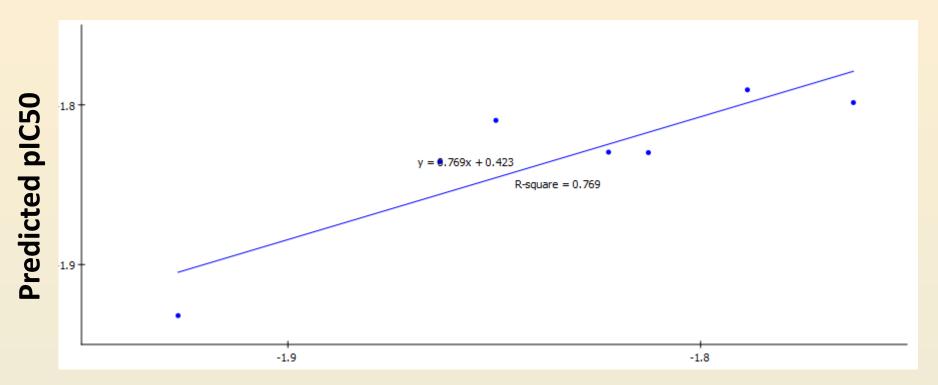
- Glutathione Peroxidase Cellular activity Assay Kit was used to measure GPX.
- LPS reduced it by 52%
- Dexamethasone 51 %
- All the tested compounds reduced GPX by 45-47 %.
- Superoxide Dismutase Activity Assay KIT
- LPS reduced it by 64%
- Dexamethasone 62 %
- All the tested compounds reduced SOD in the range 57-52 %
- Finally, Compound 6i reduced the enzyme by 57%.

2D QSAR Studies

- Development of QSAR Models
- QSAR analyses for inhibitory activities of the synthesized pyrazole derivatives against TNF-α, IL-2 and IL-6 were performed in order to determine the crucial factors governing this activity. The analysis was run by means of the DS 2.5 software (Discovery Studio 2.5, Accelrys, Co., Ltd., San Diego, CA, USA).
- Training set was prepared from the synthesized compounds with their measured pIC50s
- "Calculate Molecular Properties" module was used for calculating different molecular properties for the training set compounds

- Genetic function approximation (GFA) was utilized to search for the best possible QSAR regression equation capable of correlating the variations in the biological activities of the training compounds with variations in the generated descriptors
- multiple linear regression modeling (MLR)
- QSAR model was validated employing leave one-out cross-validation, r₂ (squared correlation coefficient value) and r₂ prediction (predictive squared correlation coefficient value), residuals between the predicted and experimental activity of the test set and training set

TNF-α

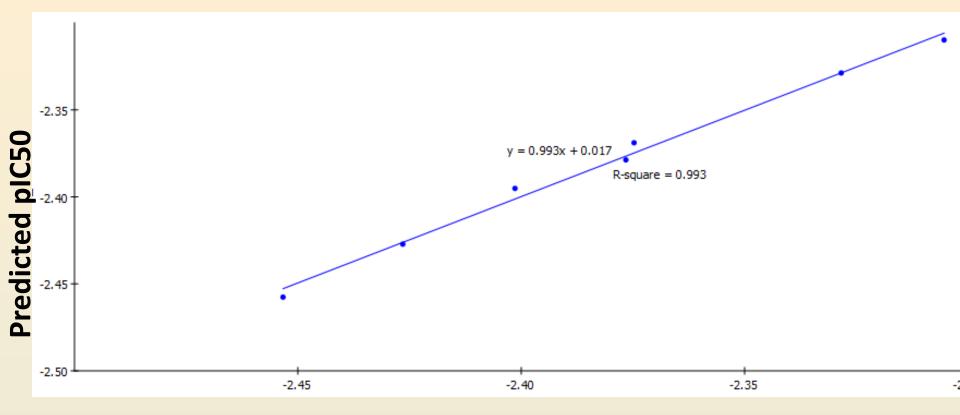


Experimental pIC50

Fig (4) Predicted versus experimental PIC50 of the tested compounds against TNF- α according to equation 1 r^2 = 0.769, r^2 (prediction) = 0.654, Least square error = 0.000572

Equation (1) representing the best performing QSAR model for the activity against TNF- α :

-logIC50 = -1.42953 - 0.16598 **SC_3_C** - 0.0033134 **Jurs_WNSA_2**.



Experimental pIC50

Fig (5): Predicted versus experimental pIC50 of the tested compounds against IL-2 according to equation $2 r^2 = 0.993$, r^2 (prediction) = 0.892, Least square error = 0.0054

Equation (2) = $-2.2525 + 0.00049194 \text{ PMI_Y} - 0.0039522 \text{ Molecular_Volume}$

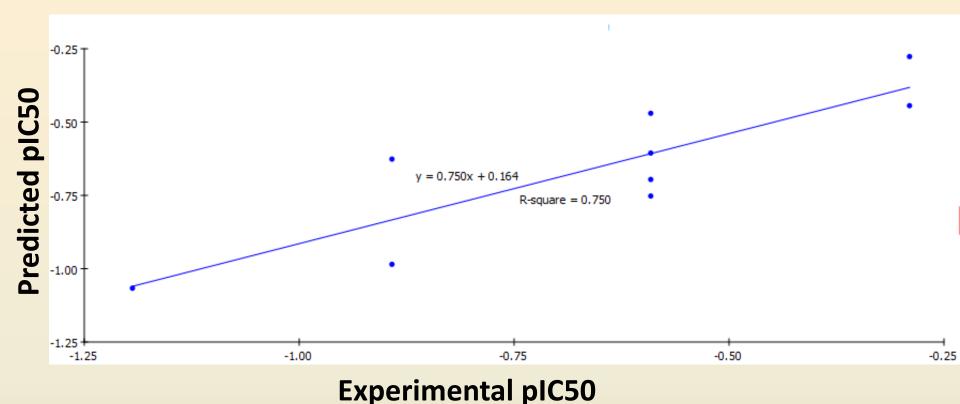


Fig (6) Predicted versus experimental pIC50 of the tested compounds against IL-6 according to equation $3 \, r^2 = 0.750, \, r^2$ (prediction) = 60, Least square error = 0.019

Equation (3) = -2.0117 - 0.36899 Kappa_3_AM + 0.00066893 PMI_Y

Docking Studies

IL-2

Schematic representation of important interactions between SB 203580 and ITK.

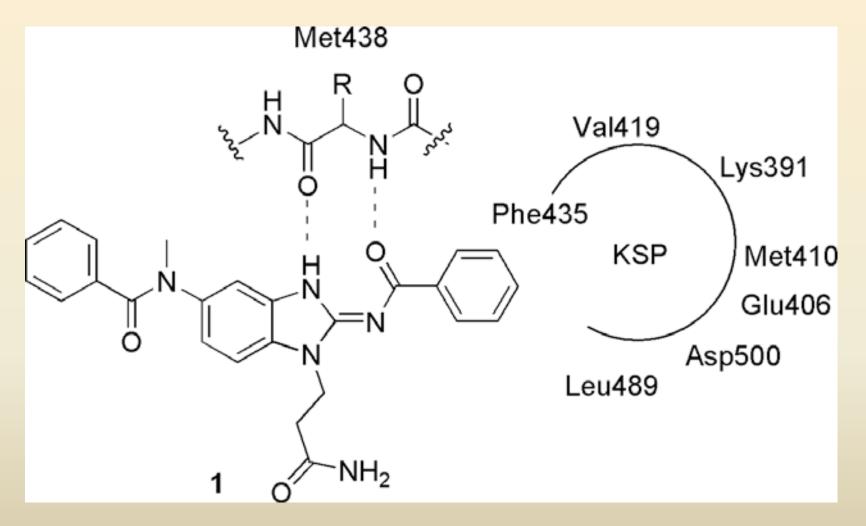


Fig (7). The structure and binding mode into the active site of ITK

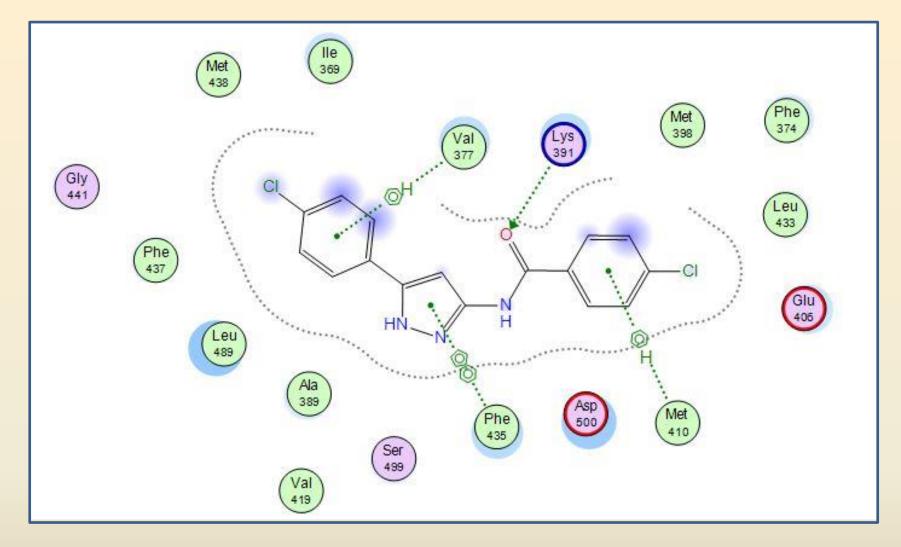


Fig (8): Binding interactions of 6i into the active site of ITK (PDB ID: 1SM2). The important amino acid residues are shown together with their respective number.

S = -11.721, E - Conf = -6.627 Kcal/mol, RMSD = 1.014

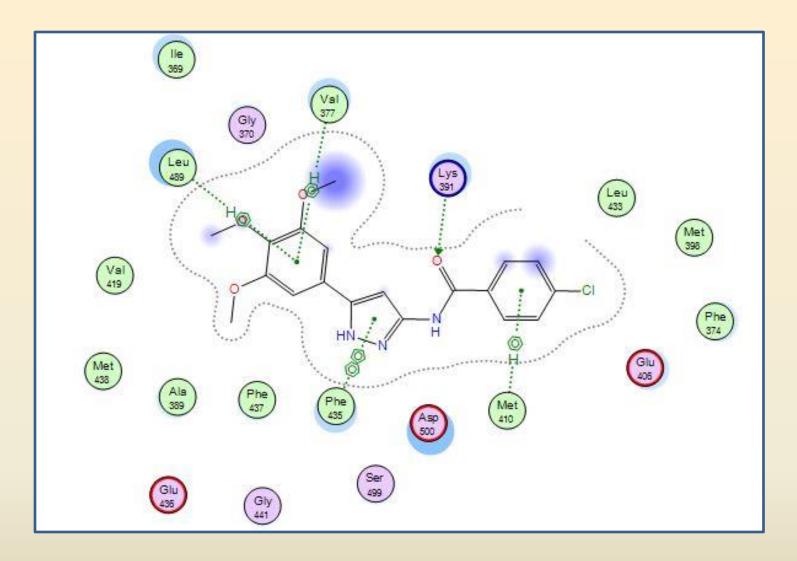


Fig (9). Binding mode of 6f into the active site of ITK (PDB ID: 1SM2). The important amino acid residues are shown together with their respective number.

S = -13.221, E-Conf = -3.626 Kcal/mol, RMSD = 1.26

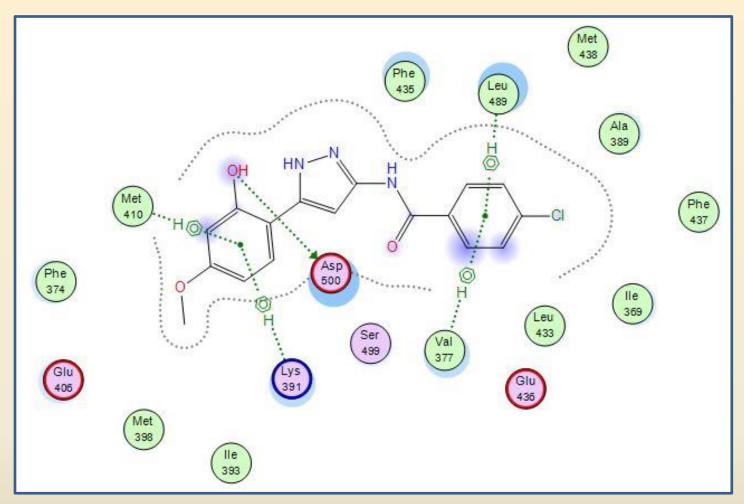
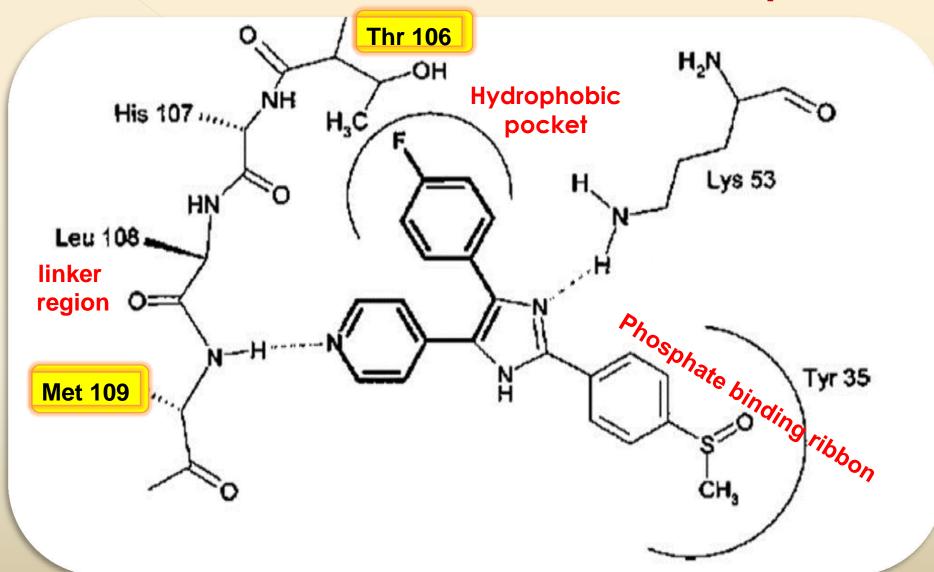


Fig (10): Binding mode of 6f into the active site of ITK (PDB ID: 1SM2). The important amino acid residues are shown together with their respective number.

S = -12.991, E-Conf = -5.442 Kcal/mol, RMSD = 1.454

TNF-α

Schematic representation of important interactions between SB 203580 and p38a.



Med Res Rev. 2006, 26, 1-62.

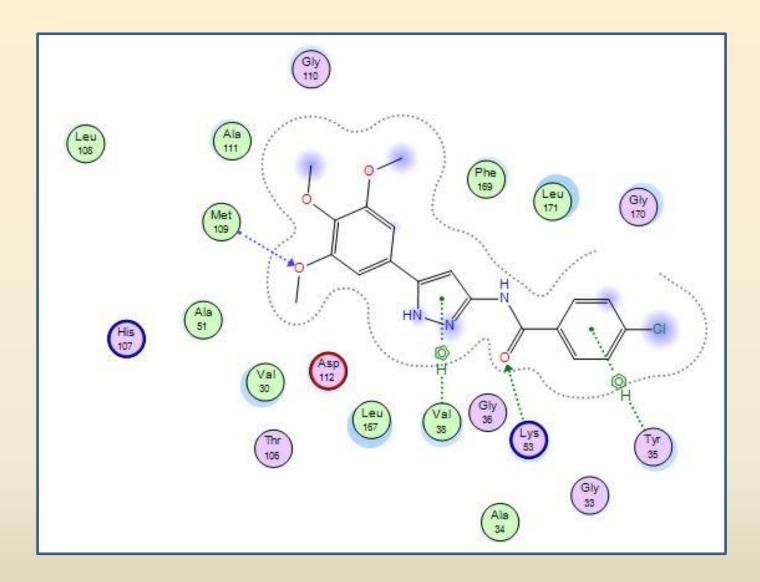


Fig (11): Binding interactions of **6f** into the active site of p38α (PDB ID: 1GM2). The important amino acid residues are shown together with their respective number.

S = -13.59, E-Conf = -4.842 Kcal/mol, RMSD = 0.843

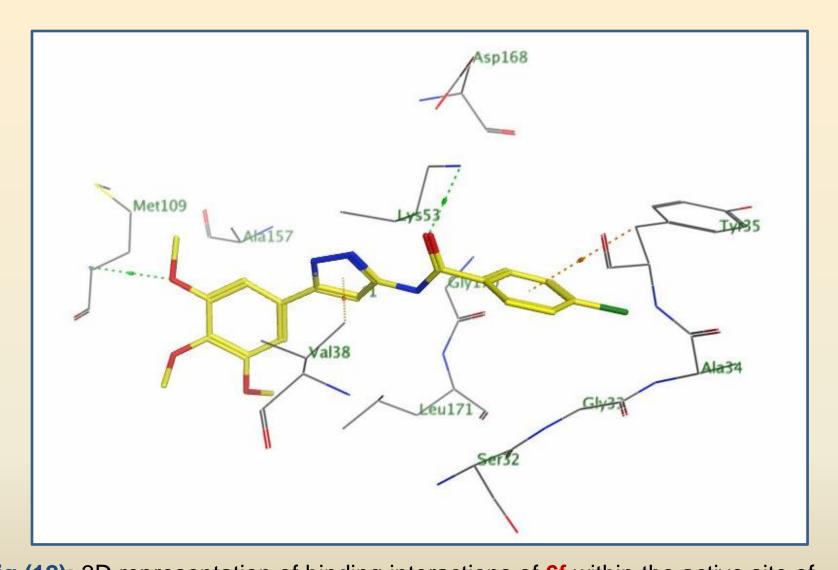


Fig (12): 3D representation of binding interactions of 6f within the active site of p38α (PDB ID: 1GM2). The important amino acid residues are shown together with their respective number.

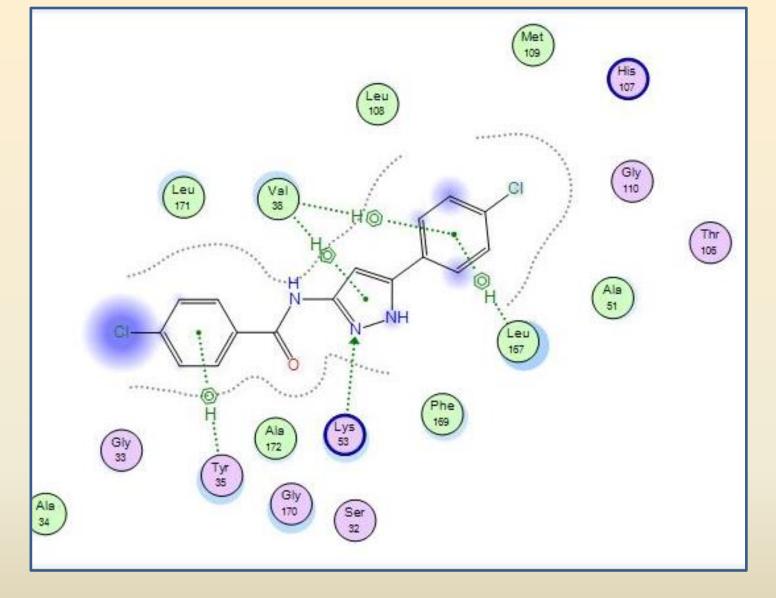


Fig (13): Binding interactions of **6i** into the active site of p38α (PDB ID: 1GM2). The important amino acid residues are shown together with their respective number.

S = -11.650, E-Conf = -3.842 Kcal/mol, RMSD = 2.23

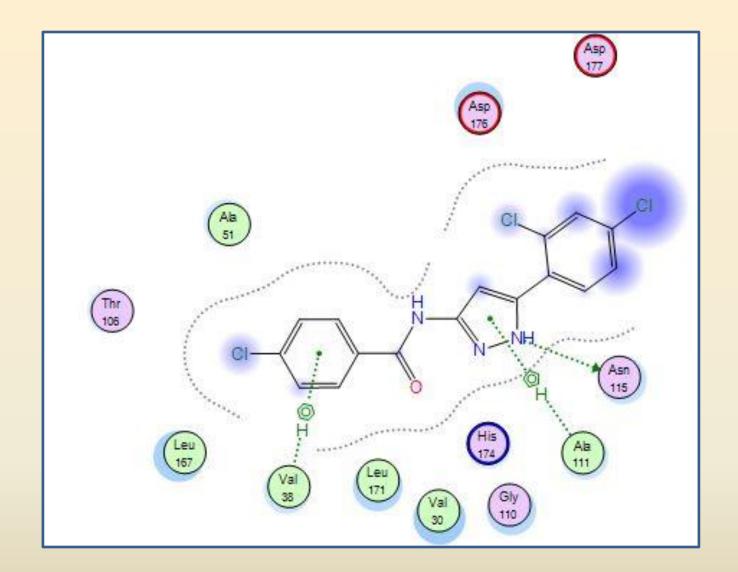


Fig (14): Binding interactions of **6j** into the active site of p38α (PDB ID: 1GM2). The important amino acid residues are shown together with their respective number.

S = -11.897, E-Conf = 1.032, Kcal/mol, RMSD = 1.762

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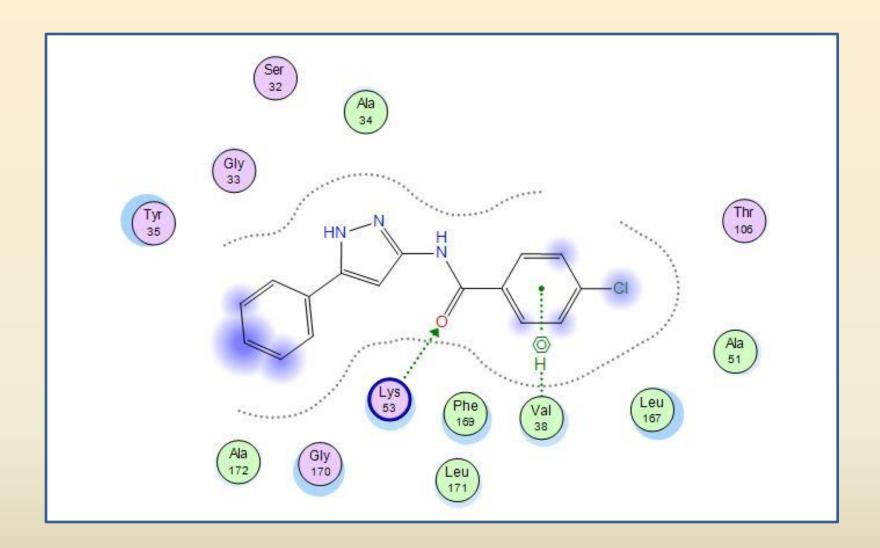


Fig (15): Binding interactions of **6a** into the active site of p38α (PDB ID: 1GM2). The important amino acid residues are shown together with their respective number.

$$S = -13.11$$
, E -Conf = 1.184, K cal/ m ol, $RMSD = 1.184$

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Conclusions

- In an attempt to generate 3,5-diaryl pyrazoles as immunomodularors through inhibition of multiple pro-inflammatory cytokines such as TNF-α, IL-2, and IL-6 nove derivatives were synthesized and evaluated against these cytokines.
- Compounds 6i and 6f demonstrated significant inhibitory activities against the three cytokines compared to the reference dexamethasone.
- A 2D QSAR model was generated for each of these activities where the models were characterized by high correlation coefficient values and good predictive ability.
- The biological results were in agreement with docking scores and binding interactions into the active sites of the enzymes.

Synthesis: Al-Azhar University, Pharmaceutical Chemistry Department.

Biological evaluation: Biochemistry lab at Animal Reproduction Institue.

In Silico studies: were performed on MOE.10 and DS 2.5 softwares.

Thank you