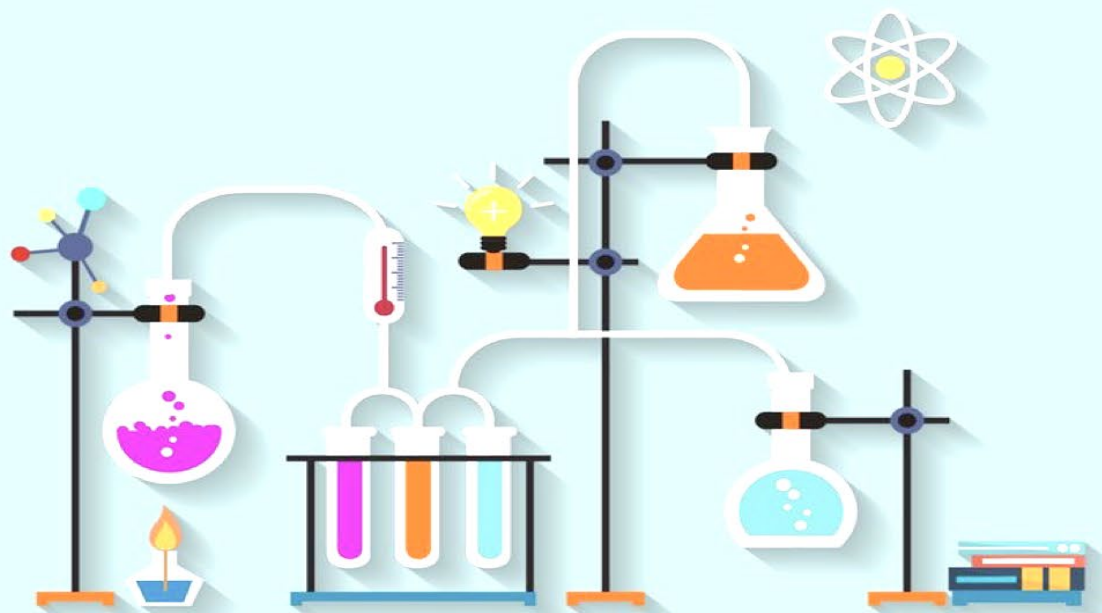


4'-fluoro-5,7-dihydroxyflavone – Piperazine Hybrids as VEGFR-2 Inhibitors: Design, In-silico Study, Synthesis, And Anticancer Activity

**DEPARTMENT OF PHARMACEUTICAL
CHEMISTRY**

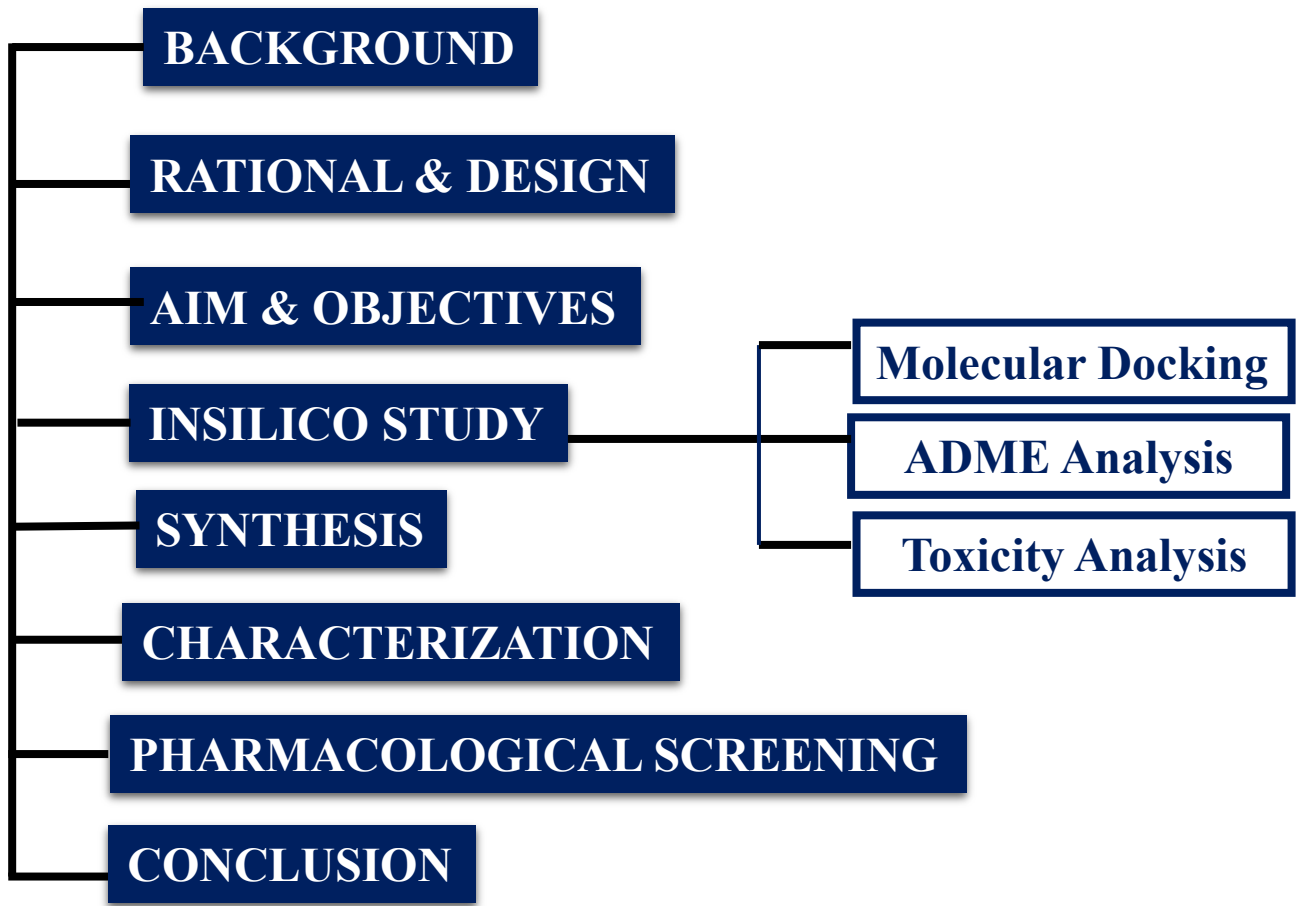


Presented by
Ms. Kalyani R. Thombre
Ph.D. Research Scholar

Guided by
Dr. Krishna R. Gupta
M. Pharm, PhD, DBM, PGDRA

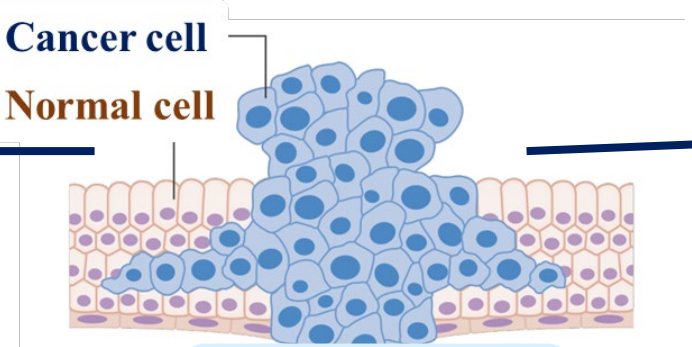
Smt. Kishoritai Bhoyar College of Pharmacy, Kamptee, Nagpur, Maharashtra, India

CONTENT



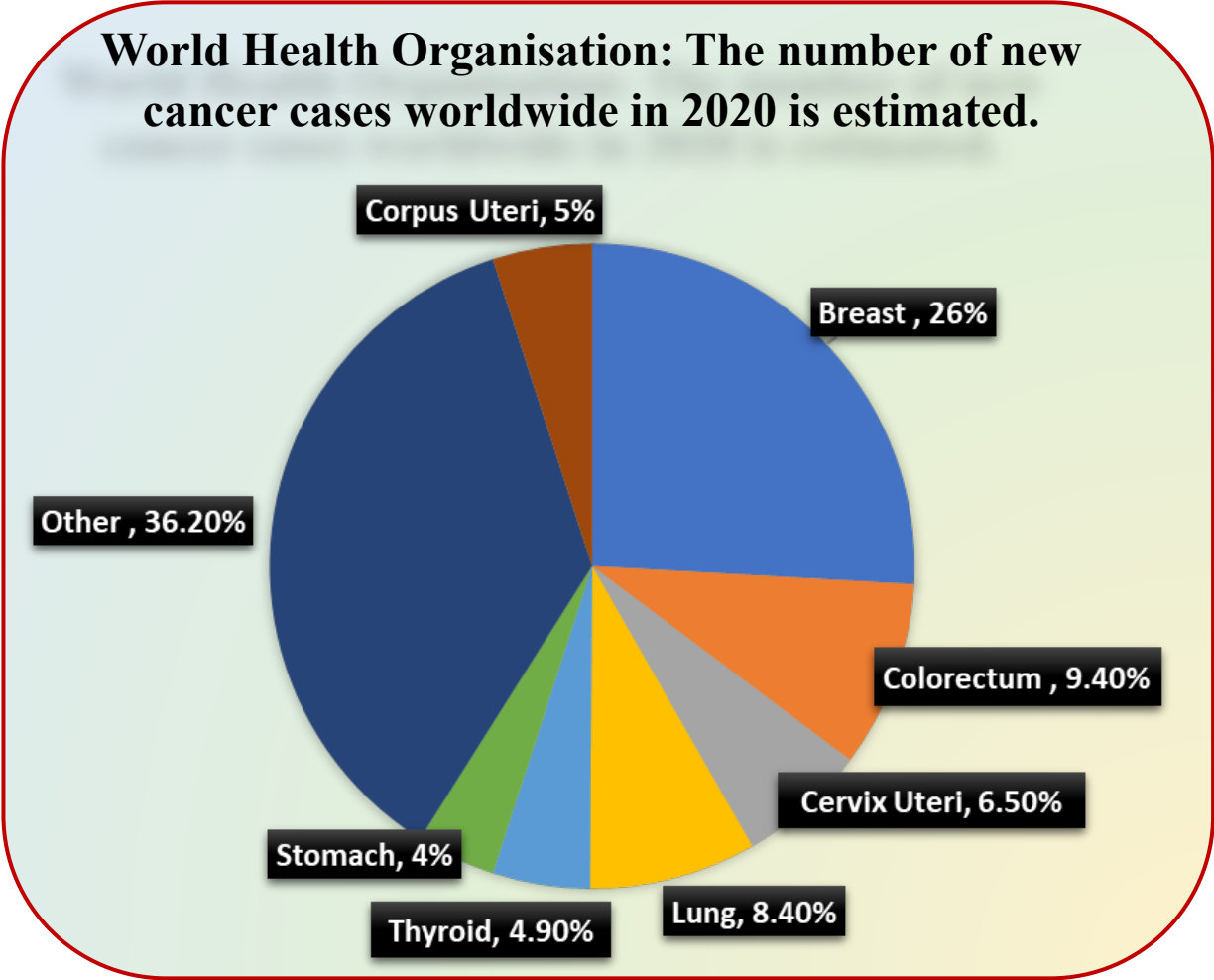
BACKGROUND

Cancer



Uncontrolled cell division

Prevalent cause of mortality in world at present



Despite advancements in diagnosis, treatment, and prevention

Global burden of cancer death

drastically ↑↑↑↑

Development of new anticancer agents

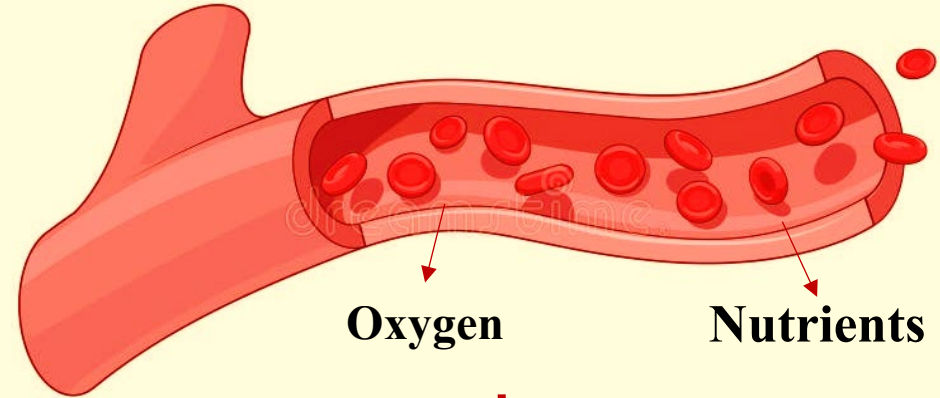


**Maximum Effectiveness
&
Minimal Toxicity**



**Vital trend in anticancer
Drug Development and
Research**

ANGIOGENESIS

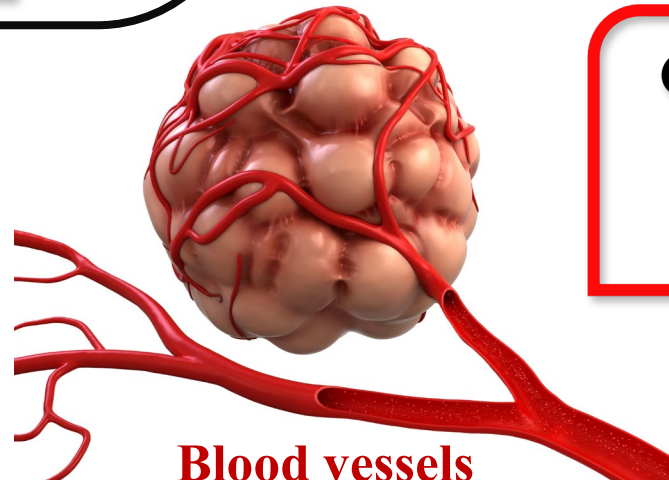


Essential Factor



**Cancer Growth
&
Proliferation**

Tumour



Blood vessels

Vascular Endothelial Growth Factor Receptor 2 (VEGFR-2)

VEGF – exert angiogenesis effect via binding with **VEGFR-2**

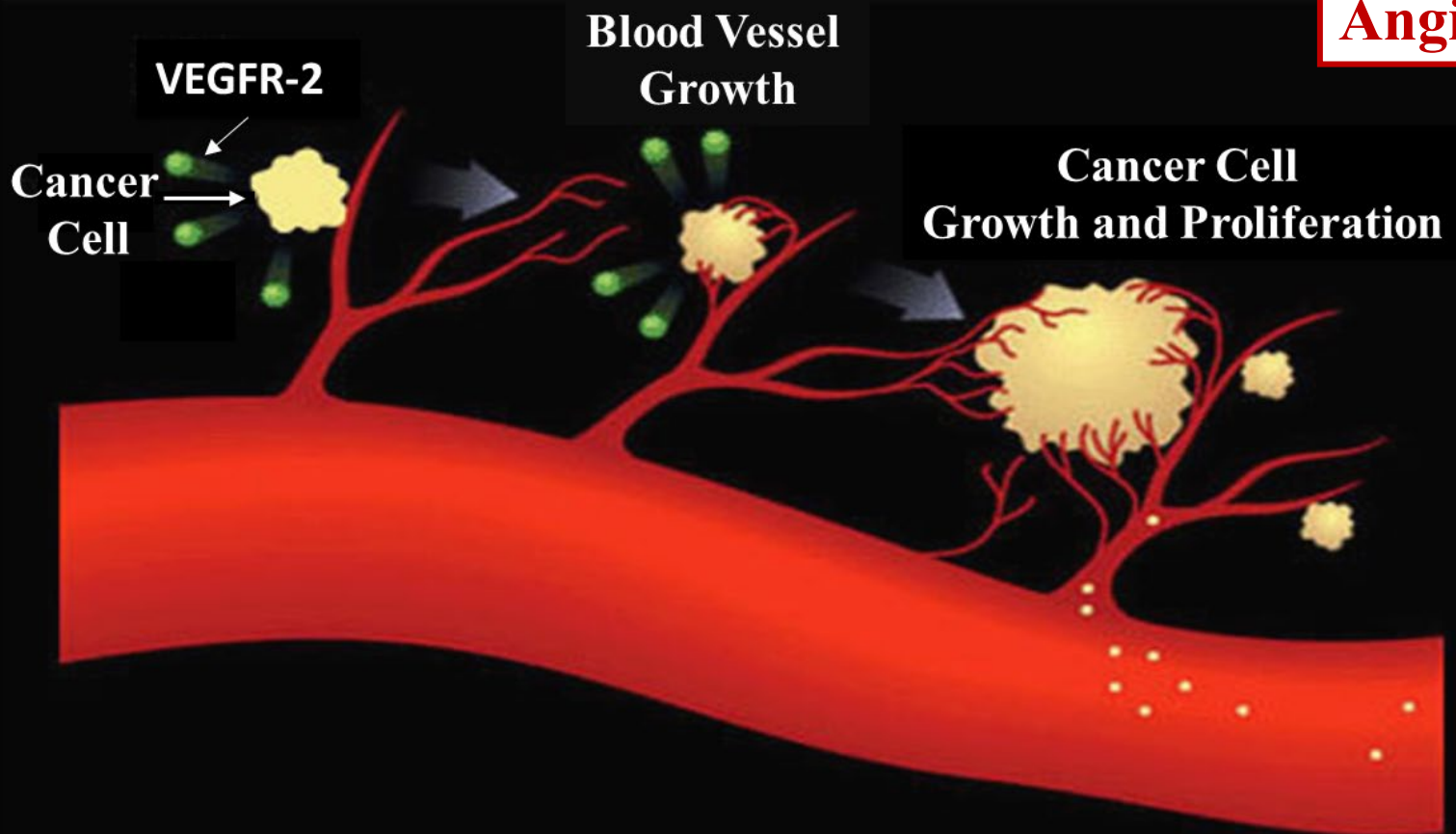
↓
Play major role
↓

Angiogenesis

VEGFR-2 pathway

↓
Competent Target

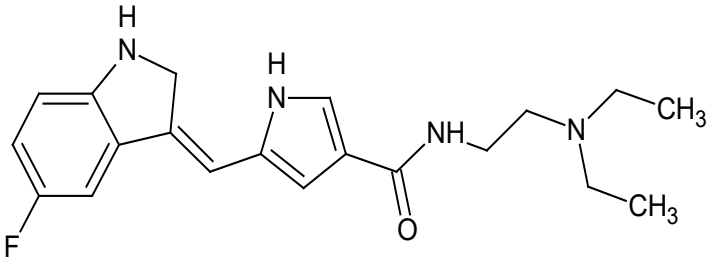
↓
Vital Selectivity for Cancer Cells



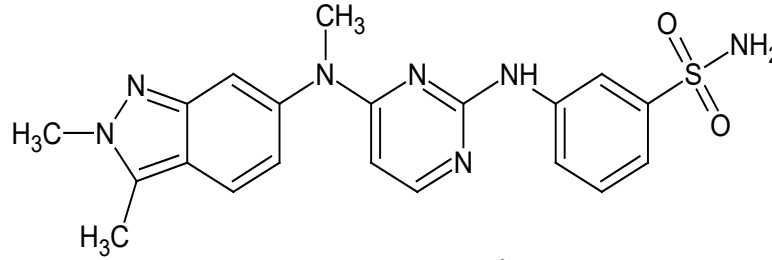
By inhibiting VEGFR-2

↓
Discovery of New Anticancer Drug

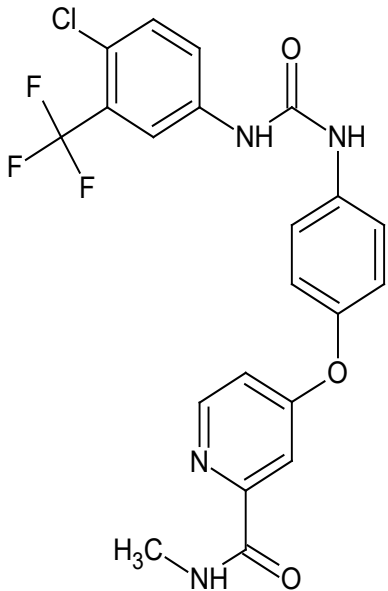
WHY NEW VEGFR-2 INHIBITORS?



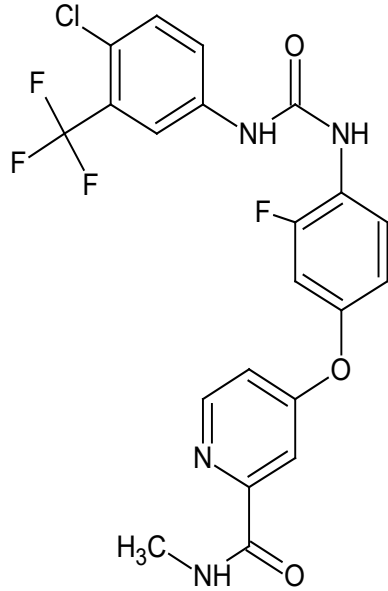
Sunitinib



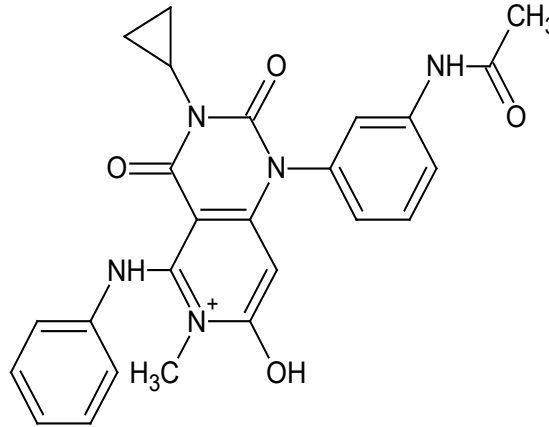
Pazopanib



Sorafenib



Regorafenib



Trametinib



Drug resistance

**Unwanted side effects
(Hypertension)**

Decreasing efficacy



DRAWBACKS

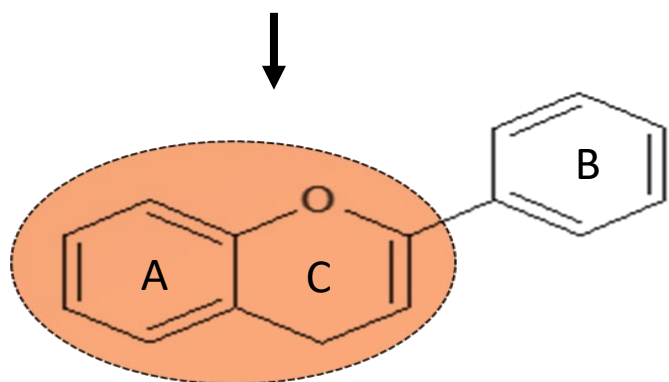
FDA APPROVED VEGFR-2 Inhibitors

FLAVONOIDS

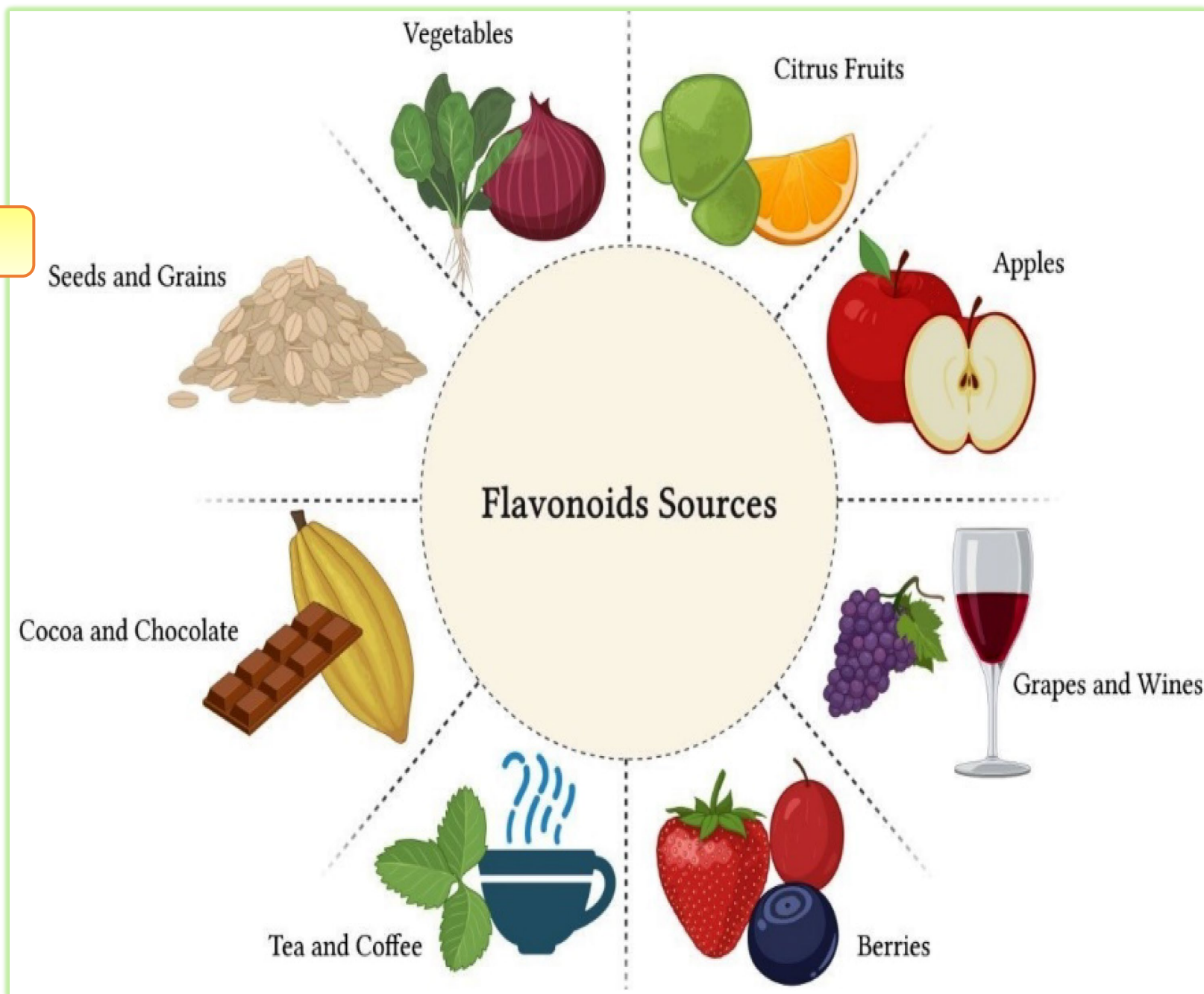
Natural Bioactive Compound

Secondary Metabolite

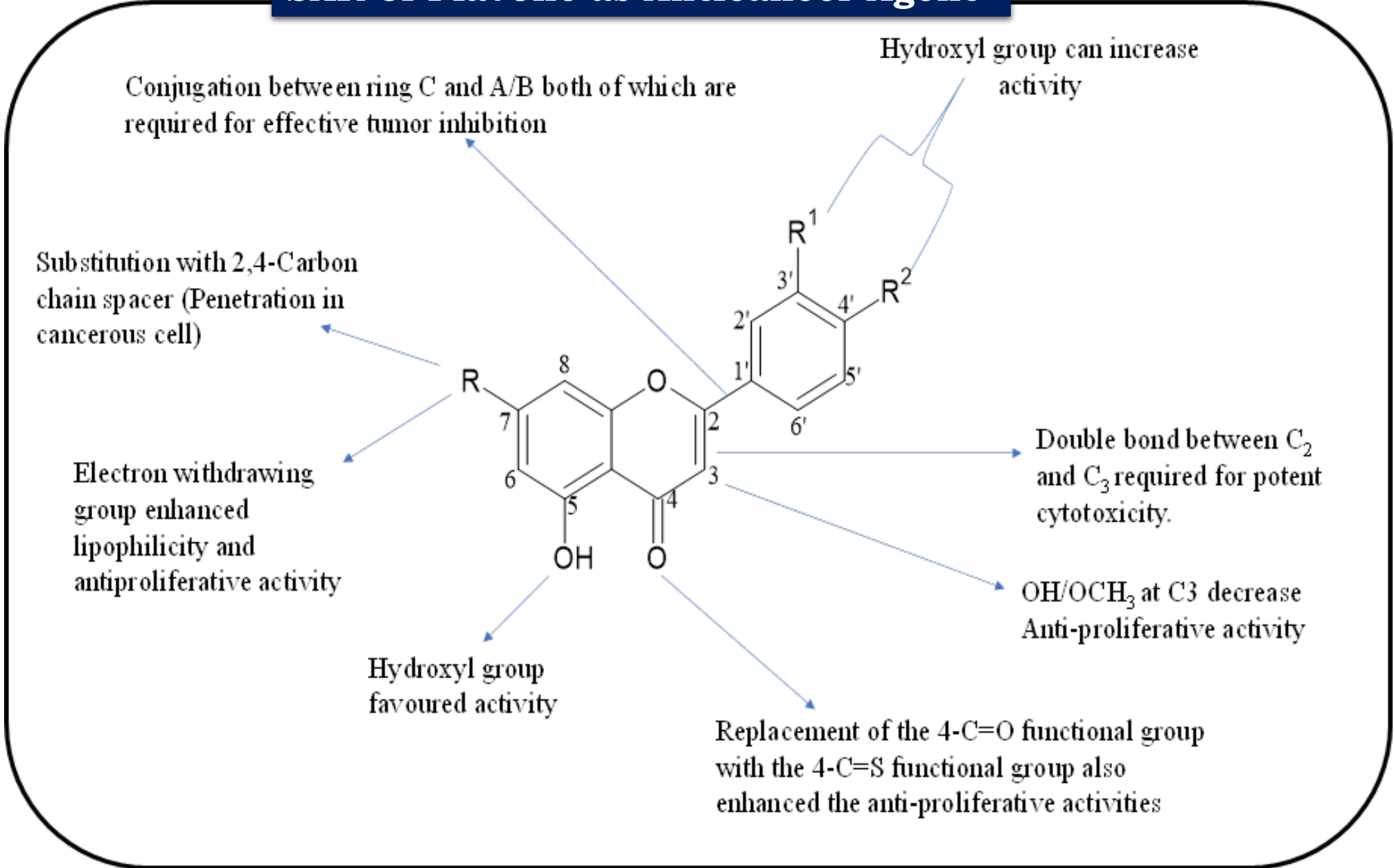
Consists of **benzopyrane** ring bearing **Phenolic** or **Polyphenolic** groups



Benzopyran



SAR of Flavone as Anticancer Agent



SAR analysis of VEGFR-2 inhibitors

VEGFR-2 Inhibitors

FOUR basic characteristics - common with their core structures

Hetero aromatic ring, or head, occupies the catalytic ATP binding domain and engages in hydrogen bonding interactions with the hinge region's Cys919 amino acid residue.

A segment known as a "linker" sits on the gatekeeper residues and joins the ATP-binding and DFG domains.

Amino acids (Glu885 and Asp1046) engage with a hydrogen-bonding moiety (pharmacophore) through H-bonds in the protein's DFG motif.

Terminal hydrophobic (tail) molecule binds to it via various hydrophobic interactions.

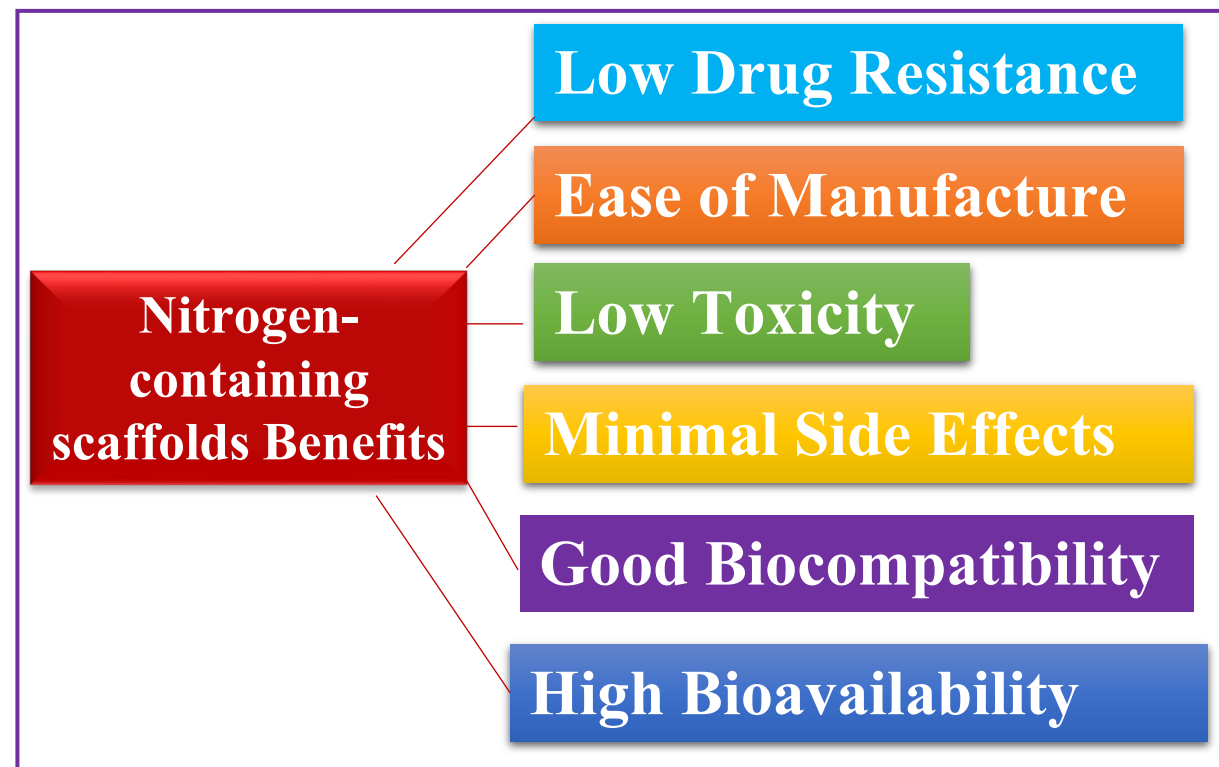
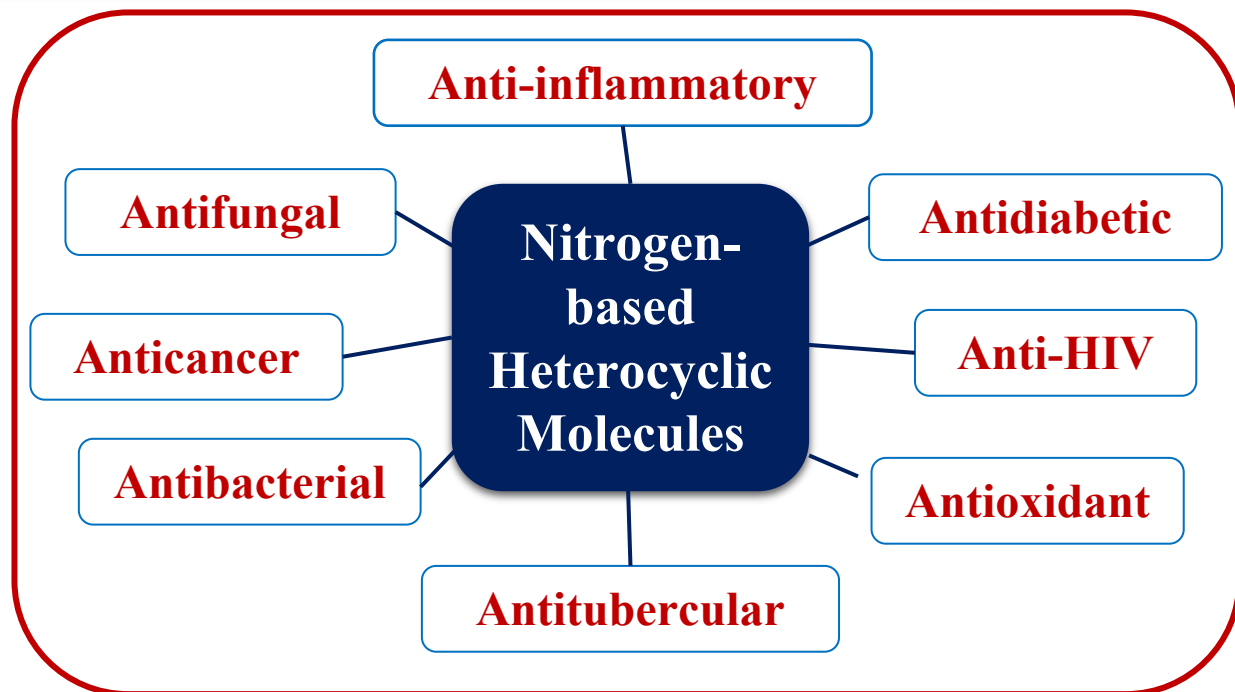
Head Hing Bonding

2-5 Carbon Linker

Hydrogen Binding Moiety

Tail Hydrophobic moiety

Importance of N-Heterocycles



Several nitrogen-containing moieties (Ex. Benzotriazole, Pyrazole, Morpholine, Benzothiazole, Pyrimidine, Nicotinamide, Piperazine derivatives, etc) with different scaffolds have shown strong anti-cancer and anti-angiogenesis properties, as they **target several receptors such as fibroblast and vascular endothelial growth factor (VEGF), tumor growth factor (TGF), and other kinases required for cancer growth and progression**

AIM & OBJECTIVES

AIM:

Design and Synthesis of New, Potent, and Safe 4'-fluoro-5,7-dihydroxyflavone – Piperazine Hybrids as VEGFR-2 Inhibitor.

OBJECTIVES:

- 1. To conduct Insilco Drug Design and docking study of 4'-fluoro-5,7-dihydroxyflavone – Piperazine Hybrids on relevant receptors.**
- 2. To Synthesis of these derivatives.**
- 3. Characterization of synthesized Derivatives.**
- 4. Pharmacological Evaluation of synthesized Derivatives as effective Anti-cancer agents.**

EXPERIMENTAL PROTOCOL

1. INSILICO STUDY

1. Molecular Docking

By utilizing Molsoft ICM pro X64 software

To determine docking score of designed derivatives with biological target VEGFR-2 using PDB ID: 4ASD



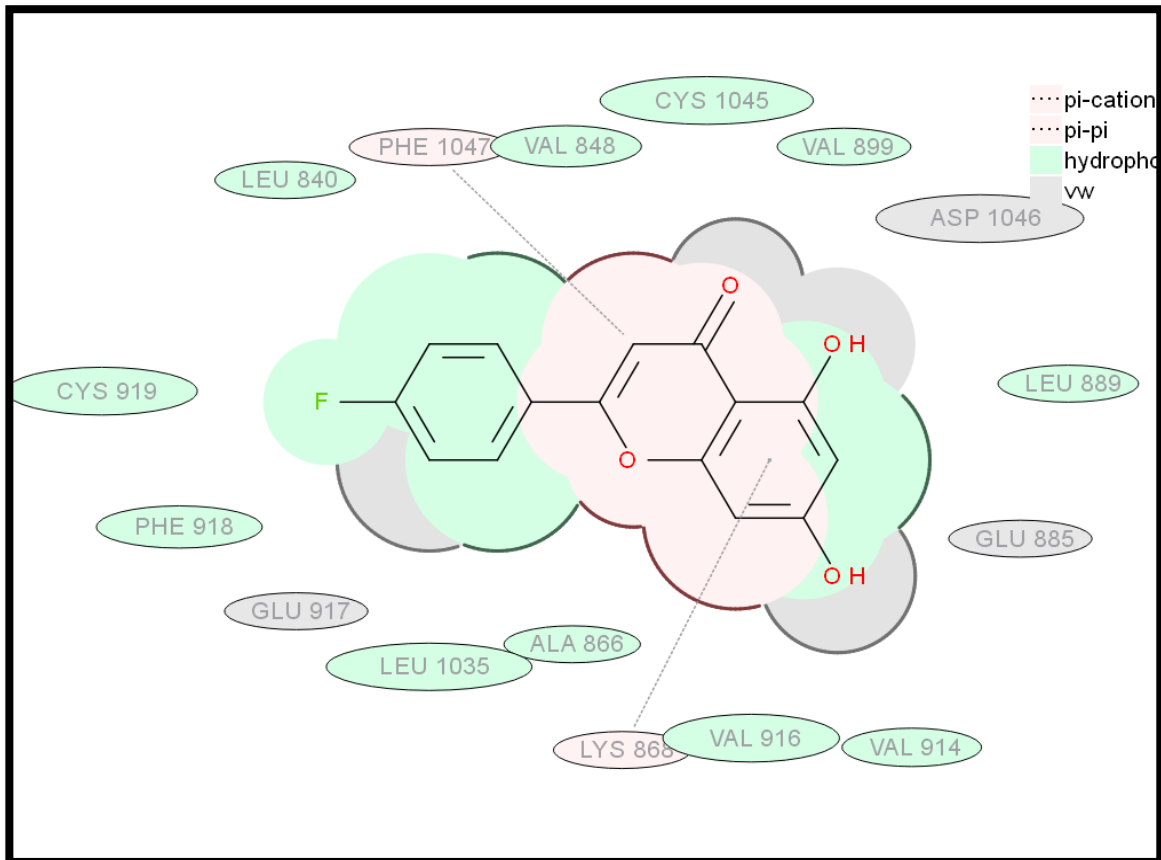
4ASD - CRYSTAL STRUCTURE OF VEGFR2
(JUXTA MEMBRANE AND KINASE DOMAINS) IN
COMPLEX WITH SORAFENIB (BAY 43-9006)



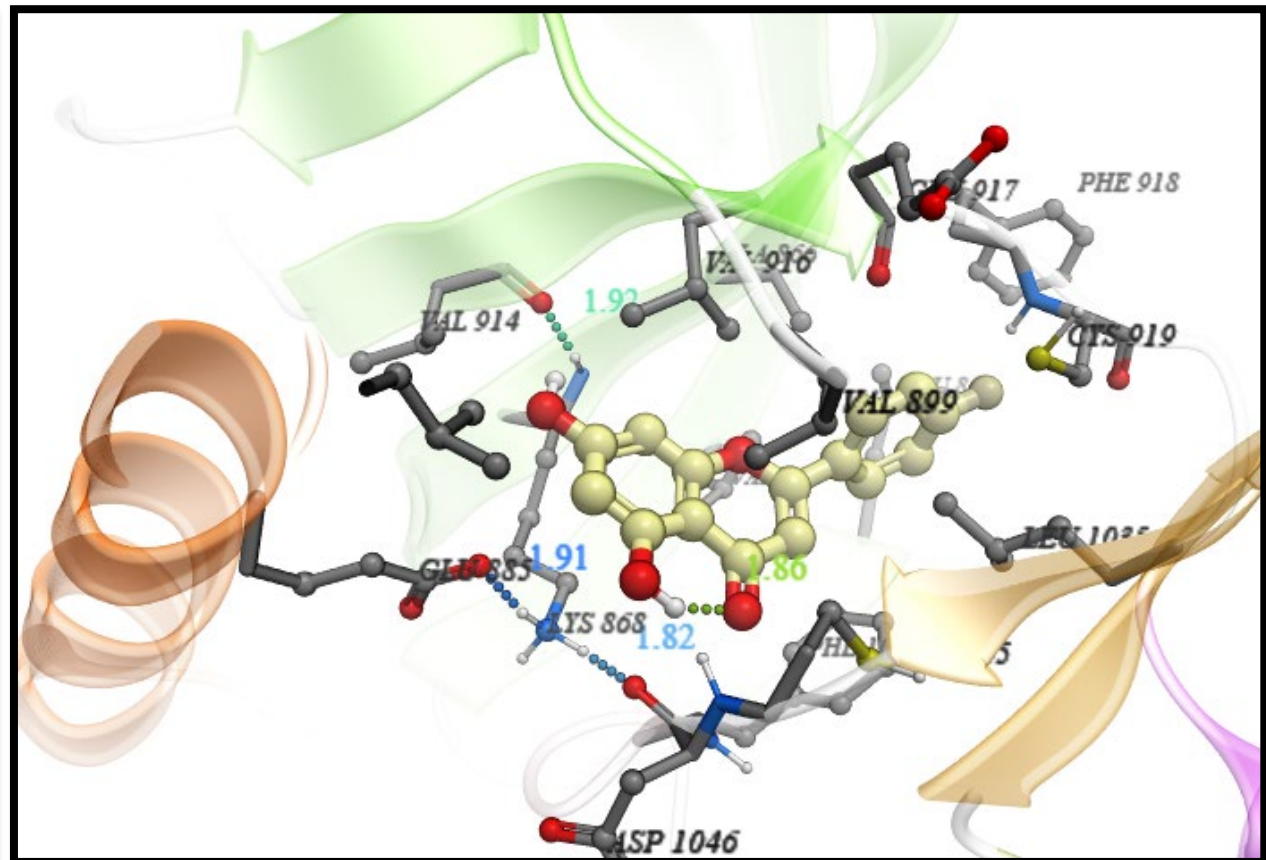
RMSD: 0.91

Compounds	Docking Score
S	-16.4
S1	-23.1
S2	-22.7
S3	-27.5
S4	-26.0
S5	-25.8
S6	-23.8
S7	-29.4
S8	-21.2
S9	-24.4
S10	-29.2

Docking Score: -16.4

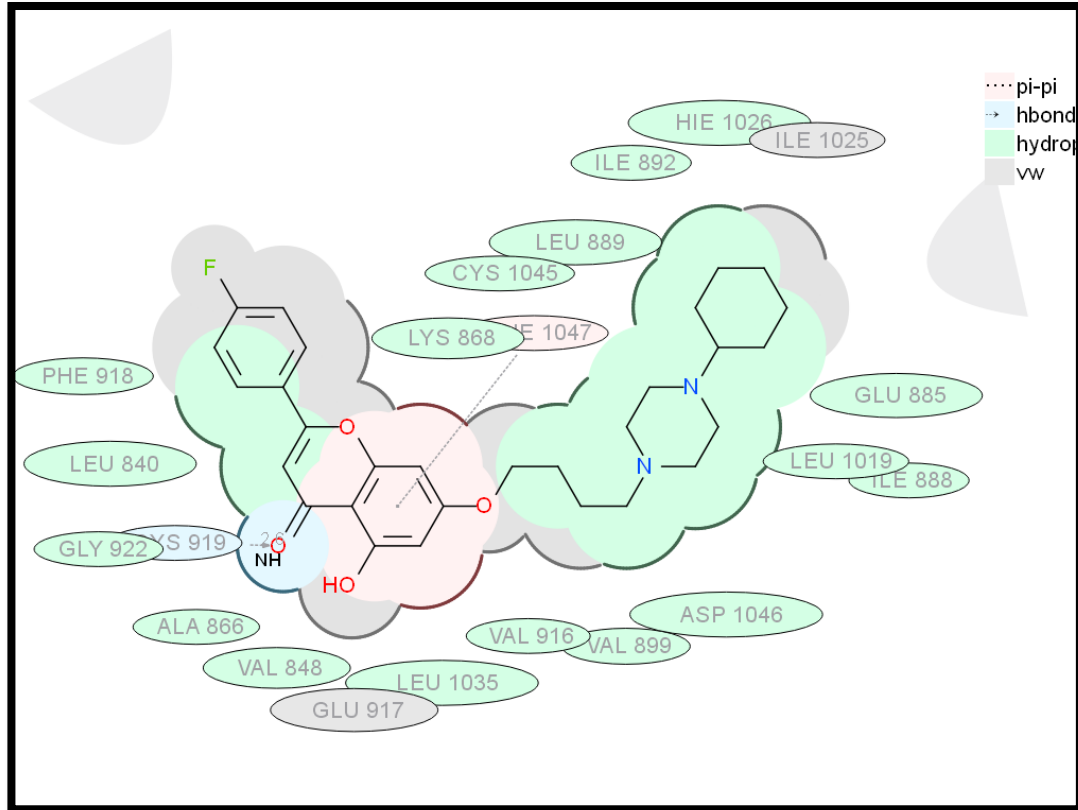


2D

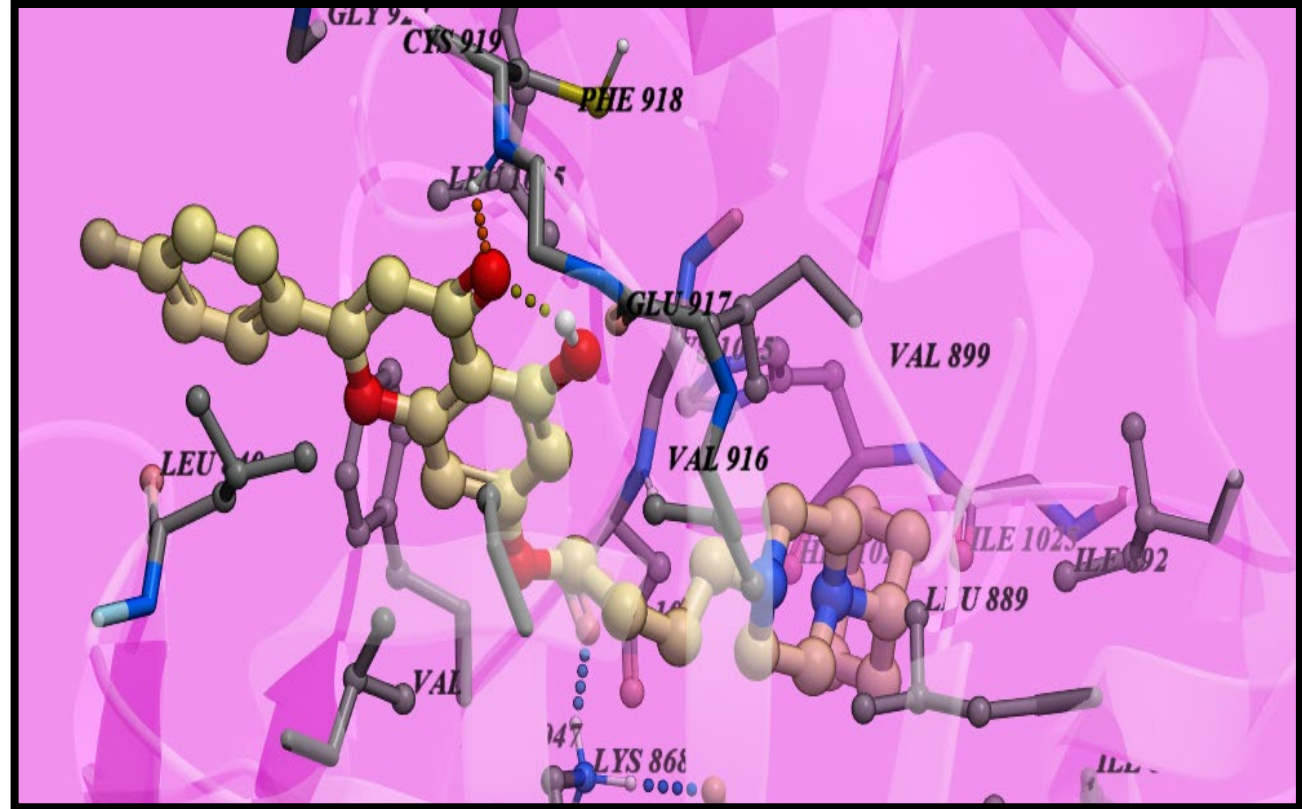


3D

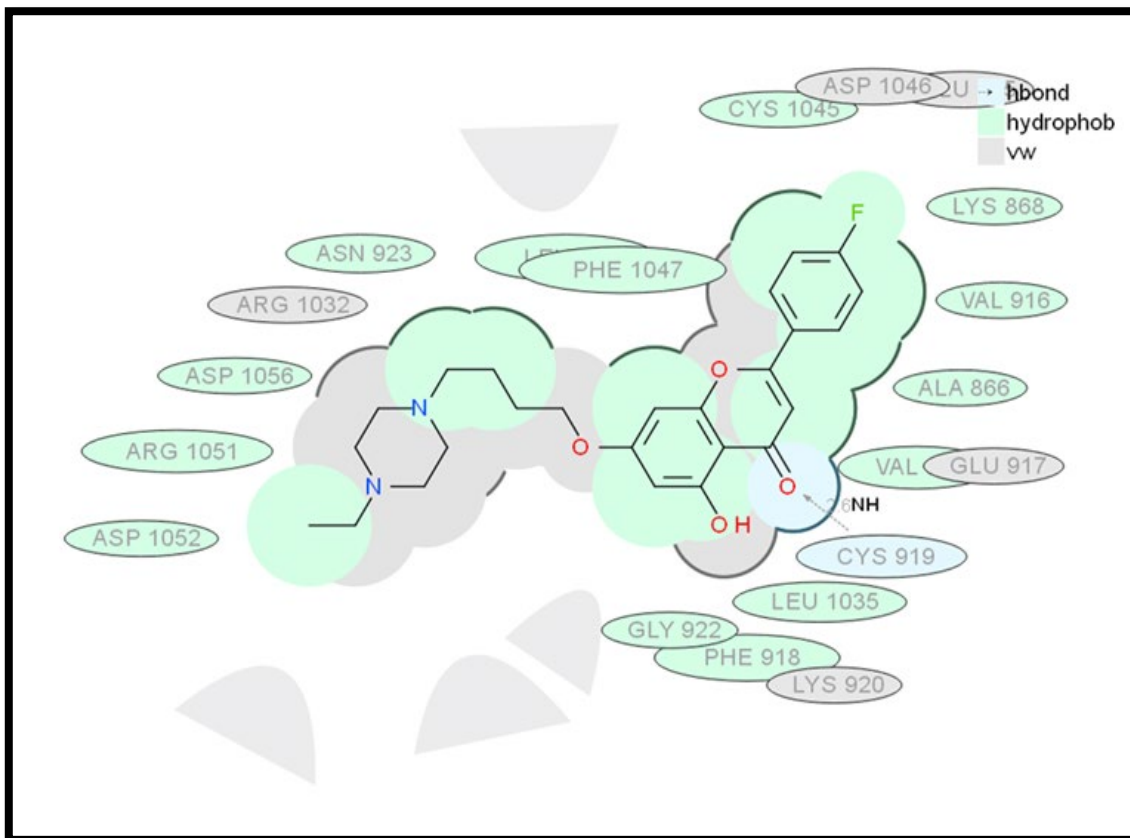
Docking Score: -24.4



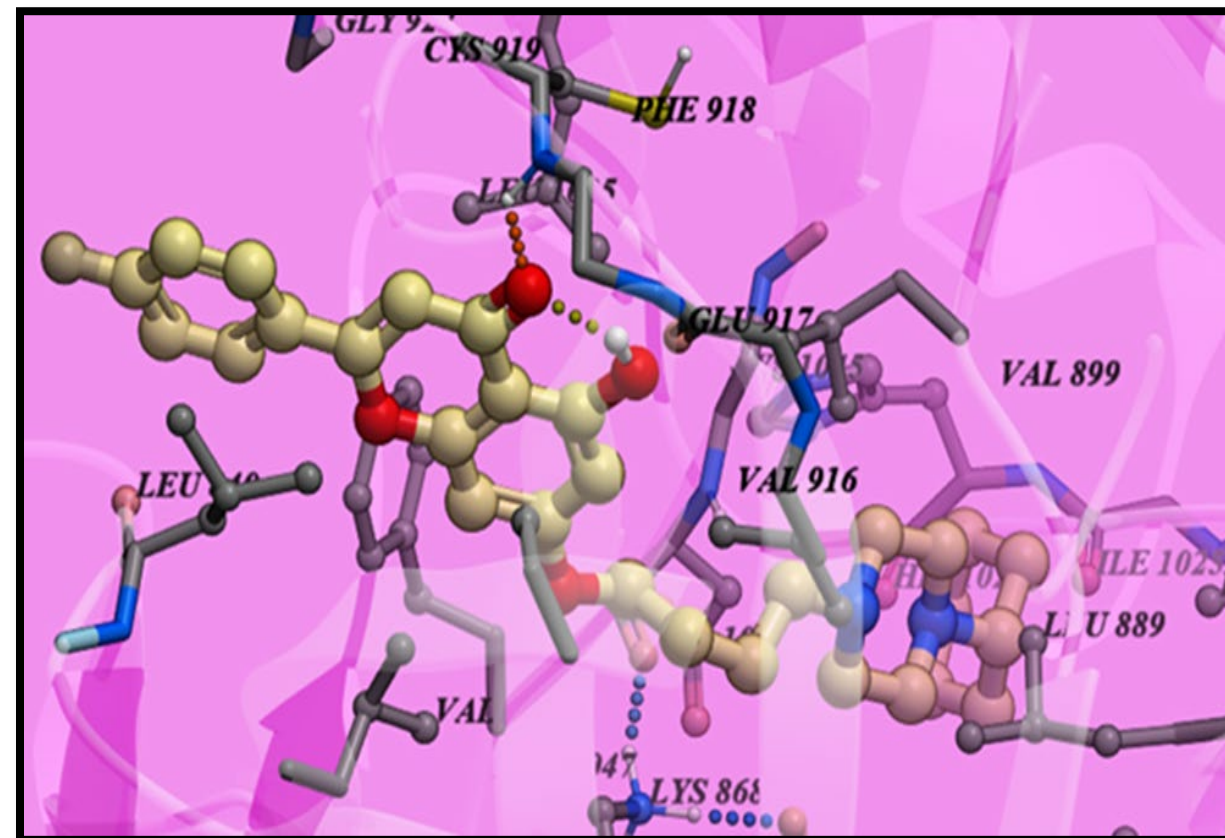
2D



3D

Docking Score: -29.2

2D



3D

2. ADME Analysis

The pharmacokinetic characteristics of the designed compounds were analyzed by using Molsoft ICM Pro X64.

Compounds	Lipinski Rule of 5				
	Log P	Mol. Wt. (g/mol)	HBD	HBA	Lipinski's Rule Violation
S	2.38	272.23	2	5	0
S1	4.13	410.38	2	6	0
S2	2.83	485.38	2	7	0
S3	3.86	410.48	1	6	0
S4	4.52	440.51	1	7	0
S5	5.05	454.53	1	7	0
S6	1.05	616.36	1	7	1
S7	4.6	489.62	1	7	0
S8	4.47	456.51	2	8	0
S9	4.51	455.52	2	8	0
S10	4.89	488.55	2	7	0

All designed derivatives follows Lipinski Rule of Five except S6

3. Toxicity Analysis

Toxicity profile of the designed compounds was estimated using Protox II software.

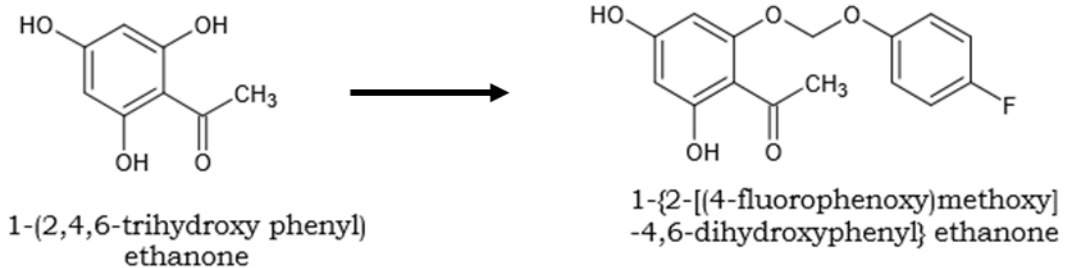
Compounds	Toxicity Class
S	CLASS IV
S1	CLASS IV
S2	CLASS IV
S3	CLASS IV
S4	CLASS IV
S5	CLASS IV
S6	CLASS III
S7	CLASS IV
S8	CLASS IV
S9	CLASS IV
S10	CLASS IV

- **Class I:** fatal if swallowed ($LD50 \leq 5$)
- **Class II:** fatal if swallowed ($5 < LD50 \leq 50$)
- **Class III:** toxic if swallowed ($50 < LD50 \leq 300$)
- **Class IV:** harmful if swallowed ($300 < LD50 \leq 2000$)
- **Class V:** may be harmful if swallowed ($2000 < LD50 \leq 5000$)
- **Class VI:** non-toxic ($LD50 > 5000$)

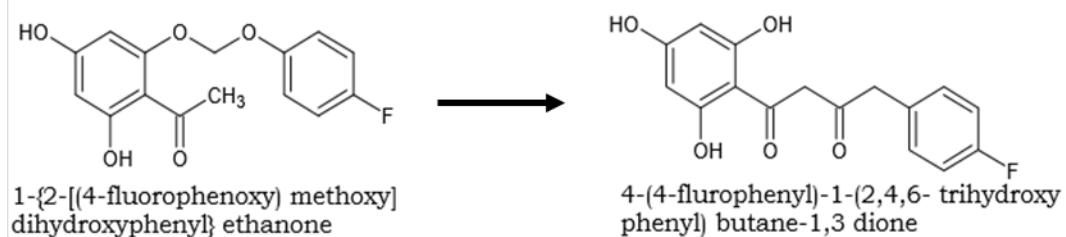
2. SYNTHESIS

Scheme

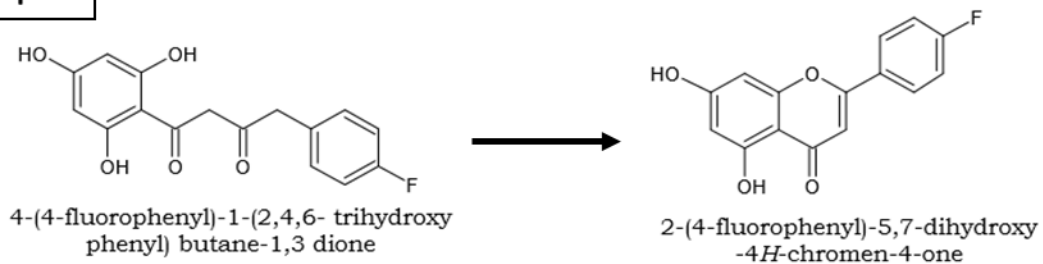
Step 1



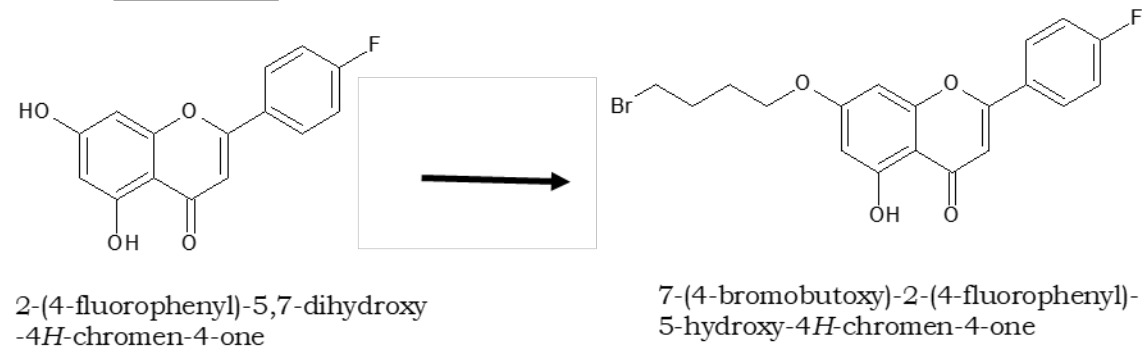
Step 2



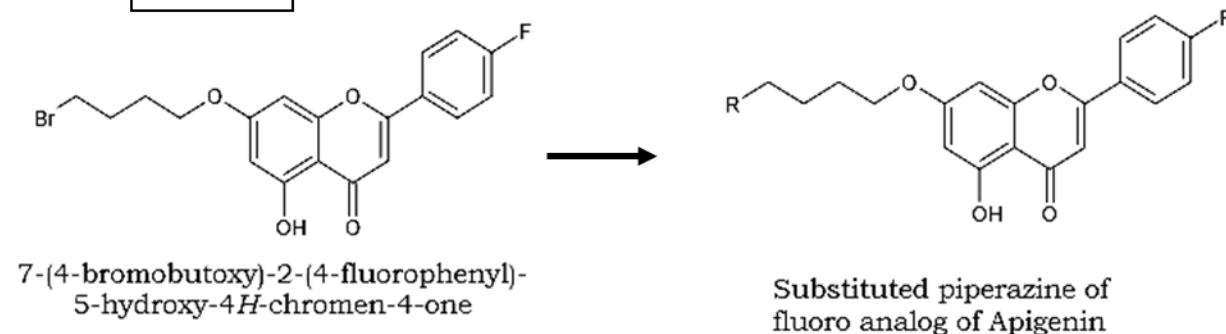
Step 3



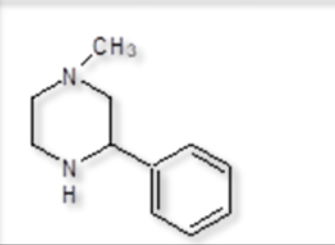
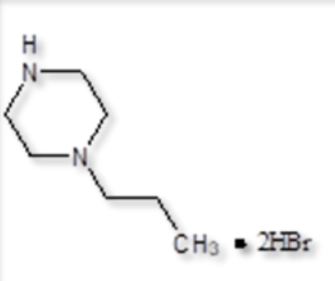
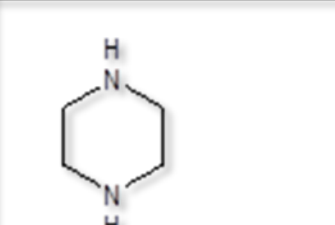
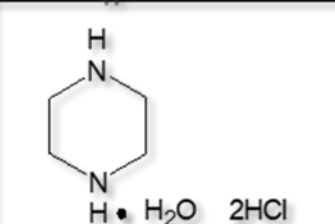
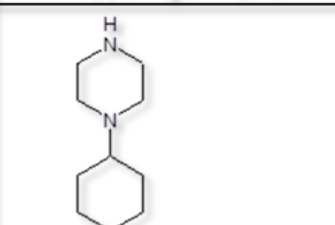
Step 4

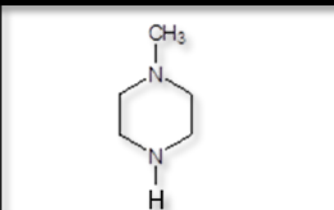
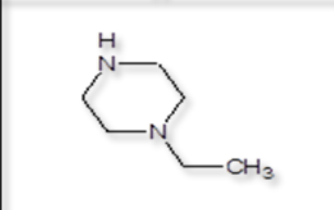
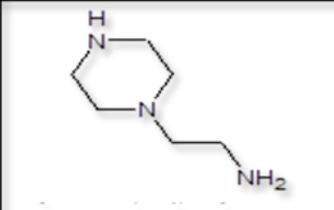
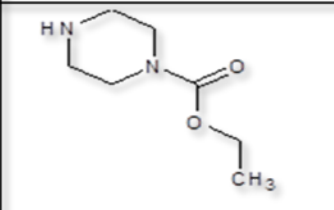
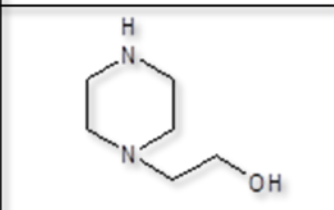


Step 5



Where R- Piperazine Derivatives

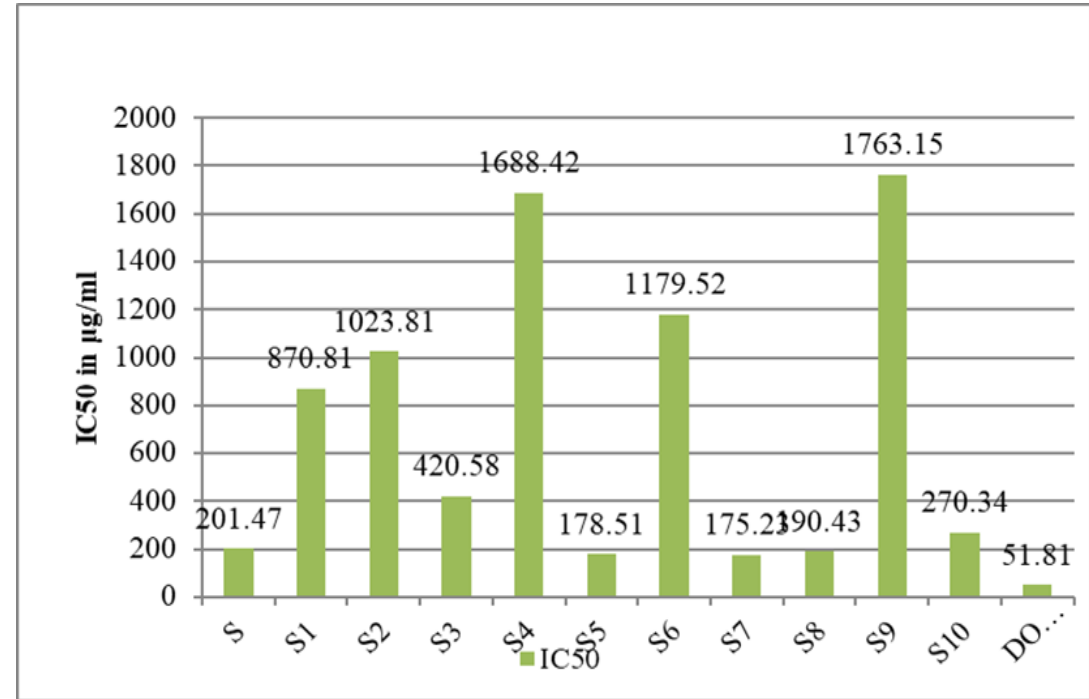
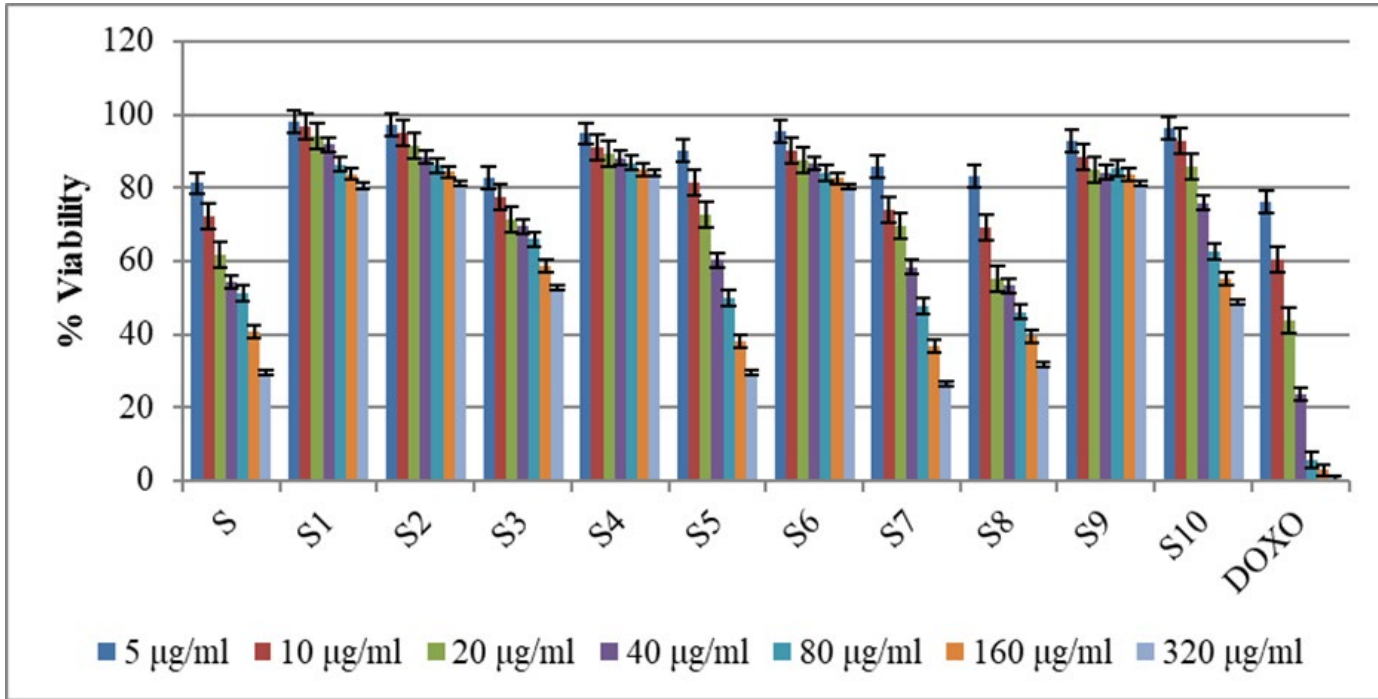
Code	R group	Rf value
S1		0.91
S2		0.68
S3		0.71
S4		0.60
S5		0.72

S6		0.61
S7		0.65
S8		0.75
S9		0.80
S10		0.75

3. PHARMACOLOGICAL SCREENING

In Vitro Anti-Proliferative Activities

Performed by using MCF-7 Cell line followed by MTT assay



Potency: S7 > S5 > S8 > S > S10

CONCLUSION

- **Substituted Piperazine Derivatives of 4'-fluro-5,7-dihydroxyflavone was synthesized as a VEGFR-2 inhibitor to improve Therapeutic Activity.**
- **These findings have encouraged us to continue the synthesis and testing of Substituted Piperazine Derivatives of 4'-fluro-5,7-dihydroxyflavone as VEGFR-2 inhibitor**

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3. Mohamed AE, Arif MA, Sevil K, and T. Dardeer, "Evaluation of Vascular Endothelial Growth Factor (VEGF) levels and Survival among Triple-Negative Breast Cancer (TNBC) and non-TNBC Cases", International Journal of Advanced Scientific and Technical Research. 2013;6(3);22-36.
4. Takahashi H, Shibuya M. The vascular endothelial growth factor (VEGF)/VEGF receptor system and its role under physiological and pathological conditions. Clinical Science. 2005 Sep 1;109(3):227–41.
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6. Dudley AC, Griffioen AW. Pathological angiogenesis: mechanisms and therapeutic strategies. Angiogenesis. 2023 Aug;26(3):313–47.
7. Frelin C, Ladoux A, d'angelo G. Vascular endothelial growth factors and angiogenesis. Annales d'endocrinologie. 2000 Mar 1;61:70–4.
8. Wang X, Bove AM, Simone G, Ma B. Molecular Bases of VEGFR-2-Mediated Physiological Function and Pathological Role. Front Cell Dev Biol. 2020 Nov 16;8:599281.
9. Shibuya M, Claesson-Welsh L. Signal transduction by VEGF receptors in regulation of angiogenesis and lymphangiogenesis. Experimental Cell Research. 2006 Mar;312(5):549–60.
10. Terman BI, Dougher-Vermazen M, Carrion ME, Dimitrov D, Armellino DC, Gospodarowicz D, et al. Identification of the KDR tyrosine kinase as a receptor for vascular endothelial cell growth factor. Biochemical and Biophysical Research Communications. 1992 Sep;187(3):1579–86.

THANK YOU