

Original Article

Novel 2,4-Thiazolidinedione Derivatives as Potential Therapeutics for NAFLD: Molecular Design, Synthesis, and Biological Validation Targeting FFAR4

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ABSTRACT

Aim: To design, synthesize, and biologically validate novel 2,4-thiazolidinedione derivatives as potential FFAR4 modulators for the treatment of non-alcoholic fatty liver disease (NAFLD).

Objective: This study aims to investigate the interactions between synthesized thiazolidinedione derivatives and the FFAR4 binding site, assess their lipid-lowering efficacy, and evaluate their safety profiles in vitro.

Method: In silico molecular docking studies were conducted to identify potential interactions between the derivatives and the FFAR4 binding site, followed by the synthesis of selected compounds using conventional methods. Comprehensive physicochemical characterization was performed using UV, FTIR, NMR, and mass spectrometry. In vitro evaluations included lipid accumulation assays using hepatocyte cell lines and cytotoxicity assessments via MTT assays.

Result: The docking studies revealed favourable binding affinities with key amino acid residues in FFAR4, guiding the rational design of lead compounds. The synthesized derivatives demonstrated significant lipid-lowering activity in hepatocyte cell lines, with several compounds exhibiting promising safety profiles in cytotoxicity tests.

Conclusion: The results support the hypothesis that 2,4-thiazolidinedione derivatives can effectively act as FFAR4 agonists, contributing to reduced lipid accumulation in hepatocytes. This study underscores the potential of these derivatives as therapeutic agents for NAFLD, highlighting the importance of further development and clinical evaluation in addressing this critical health concern. The integrated approach utilized here provides a valuable foundation for advancing the discovery of novel FFAR4-targeted agents in the fight against metabolic liver diseases.

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Introduction

Disease Target

The term non-alcoholic fatty liver disease (NAFLD) was first introduced by Schaffner in 1986. NAFLD is now recognized as the leading cause of chronic liver disease globally, constituting a significant and often overlooked public health crisis [1]. Current estimates indicate that approximately 25% of the general population suffers from NAFLD, with 3-5% affected by non-alcoholic steatohepatitis (NASH) [2].

NAFLD is characterized by excessive fat accumulation in the liver among individuals who consume little to no alcohol, ranging from simple fatty liver (steatosis) to more severe forms like NASH, which can lead to liver

fibrosis, cirrhosis, and hepatocellular carcinoma [3]. This condition is closely associated with metabolic disorders, including obesity, type 2 diabetes mellitus (T2DM), dyslipidemia, and hypertension [3-5]. Symptoms are frequently absent or nonspecific, such as fatigue or abdominal discomfort, which complicates early diagnosis [6].

At present, there is no approved pharmacological treatment specifically for NAFLD. However, potential therapeutic options include insulin sensitizers, weight loss medications (such as rimonabant, a cannabinoid [CB1] receptor inhibitor), lipid-lowering agents (including statins and fibrates), and hepatoprotective antioxidants (like vitamin E, ursodeoxycholic acid, betaine, and lipoic acid) [6-8].

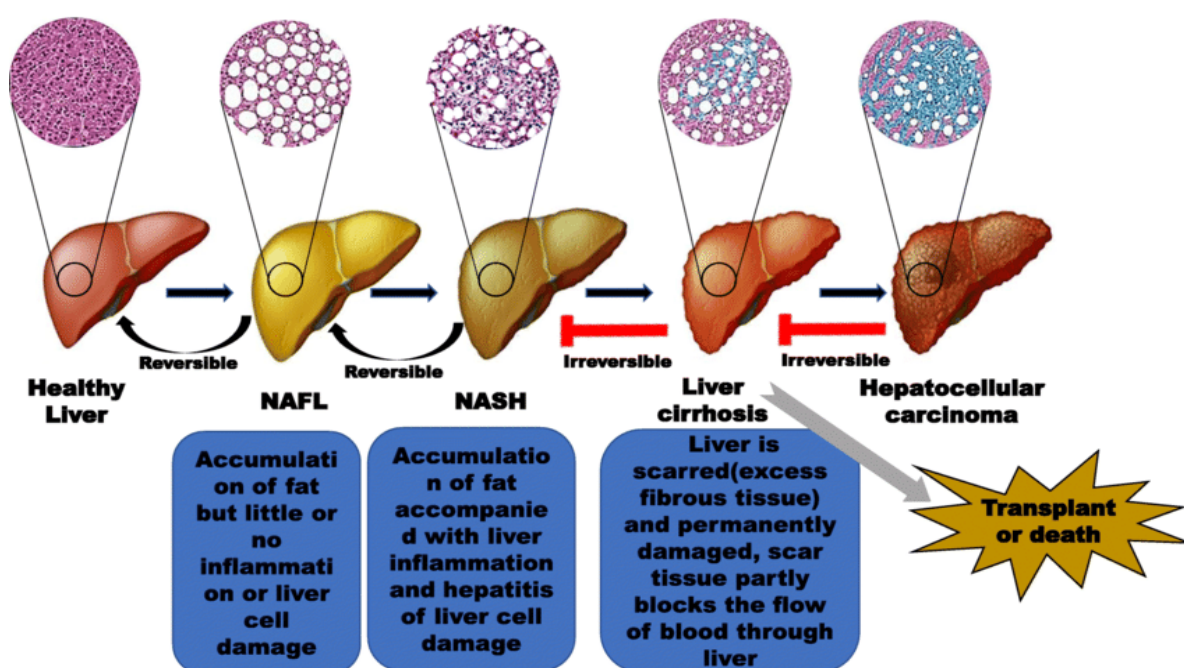


Figure 1: Clinical progression of NAFLD/NASH.

Biological Target

The Free Fatty Acid Receptor 4 (FFAR4), also referred to as GPR120, is pivotal in regulating inflammation by modulating pro-inflammatory cytokine release and enhancing insulin sensitivity [9]. FFAR4 is activated by long-chain fatty acids, initiating powerful anti-inflammatory signaling pathways. Upon activation, FFAR4 effectively inhibits the Toll-like receptor (TLR)-mediated activation of NF- κ B and significantly reduces the production of pro-inflammatory cytokines such as TNF- α and IL-6. This anti-inflammatory action is mediated via β -arrestin-2-dependent signaling, which also enhances insulin sensitivity [10].

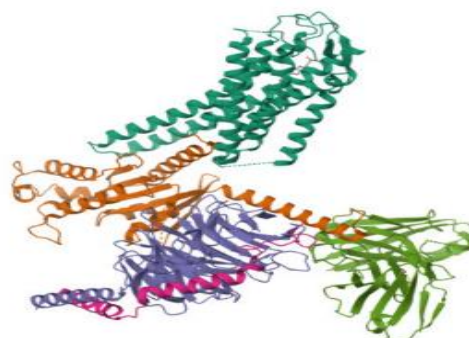


Figure 2: 3D Structure of FFAR4 (8ID6).

Drug Design

Drug design, often called rational drug design, is an innovative process for developing new drug molecules based on known interactions with biological targets [11].

Computational methods ideally predict the binding affinity of compounds prior to synthesis, allowing researchers to focus on the most promising candidates, thereby saving significant time and resources. These computational techniques have transformed the discovery process by reducing the number of required iterations and consistently yielding novel structures [12,13].

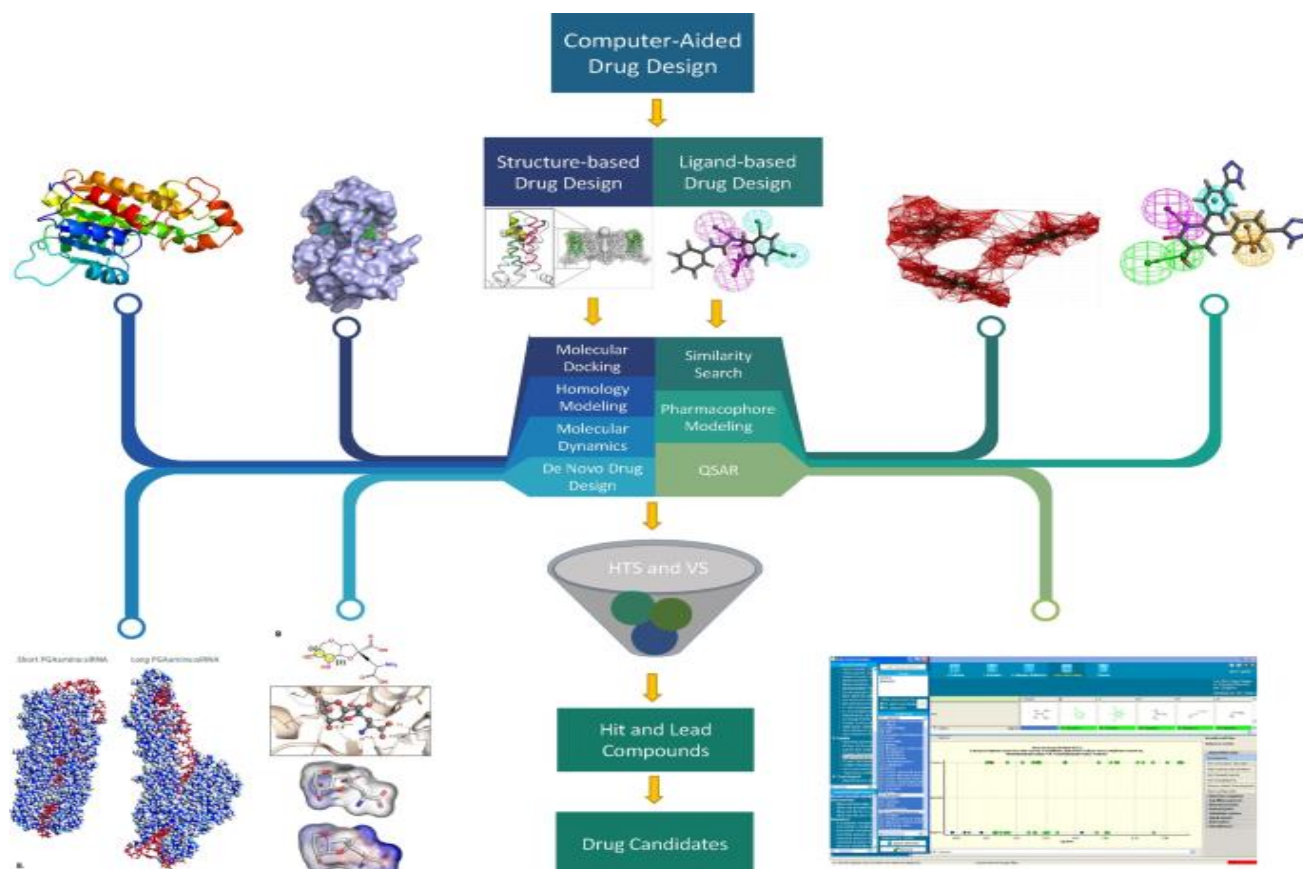


Figure 3: Computer-Aided Drug Design.

Chemistry of Scaffold

2,4-Thiazolidinedione (TZD) is a distinctive five-membered heteroaryl ring structure, featuring nitrogen and sulfur atoms alongside two adjacent carbonyl groups. It stands out as a highly privileged scaffold for developing pharmaceutically active compounds [14-16].

2,4-TZD derivatives are well-established for their anti-inflammatory and insulin-sensitizing properties. The administration of TZD influences the production of adipokines and lowers pro-inflammatory cytokines, particularly TNF- α , which correlates positively with the degree of steatosis and fibrosis [17-19].

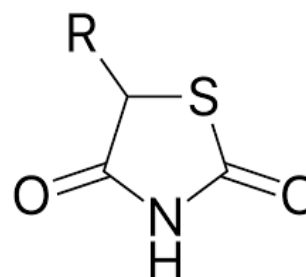


Figure 4: Structure of 2,4 Thiazolidinedione.

Materials and Methods

Target Selection

Based on the literature review, FFAR4 (PDB ID - 8ID6) was chosen as the target.

Scaffold Selection

2,4-thiazolidinedione was chosen as the basic nucleus. Structural modification at 3 and 5 positions was done to get novel compounds.

Designing Of Novel Compounds

About 162 ligands were designed using Chems sketch software.

Novelty Assessment

Novelty of those ligands were assessed using Pubchem and Zinc15 database.

In-Silico Drug-Likeness & Toxicity

Drug-likeness property and Toxicity profile of those ligands were evaluated using Molinspiration, a free web tool and Osiris Property explorer respectively.

Molecular Docking

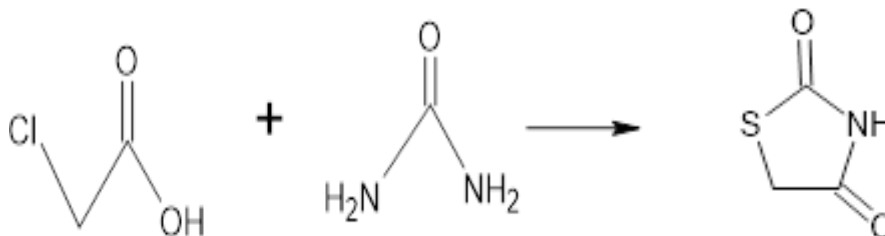
Energy minimization of the ligands was performed using Chem3D software. The energy minimized ligands were docked against the target using Autodock tools1.5.6. The interaction between the molecule and the target was viewed through Molegro Molecular viewer and Biovia Discovery Studio visualizer.

Synthesis

3 molecules (SN1C, SN15E, SN33C) were selected for further synthesis based on drug-likeness property, toxicity profile, optimum docking score and synthetic feasibility.

Synthetic Scheme [22-23]:

Step 1: Nucleus synthesis

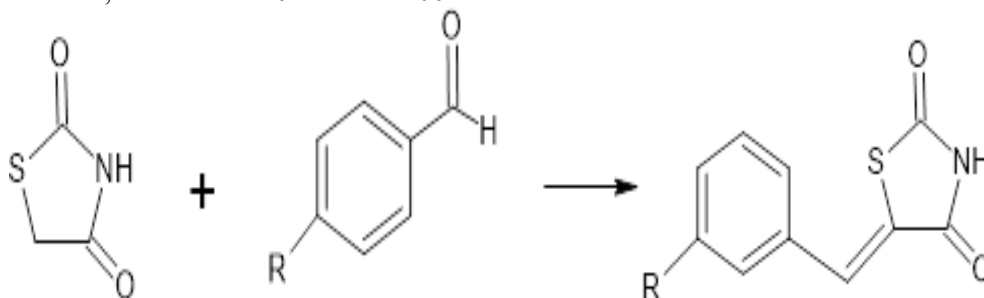


Procedure

Chloroacetic acid (5.64g, 0.06 mol) in 6ml of H₂O + Thiourea (4.56g, 0.06 mol) in 6ml of H₂O - stirred for 15 mins - white solid precipitates - now slowly added 6ml of conc. HCl - stirred, refluxed for 10-12 hrs at 100-

110°C - on cooling - mass of clusters of white needles formed. The product is filtered, washed, dried, purified and recrystallized.

Step 2: Knoevenagel Condensation

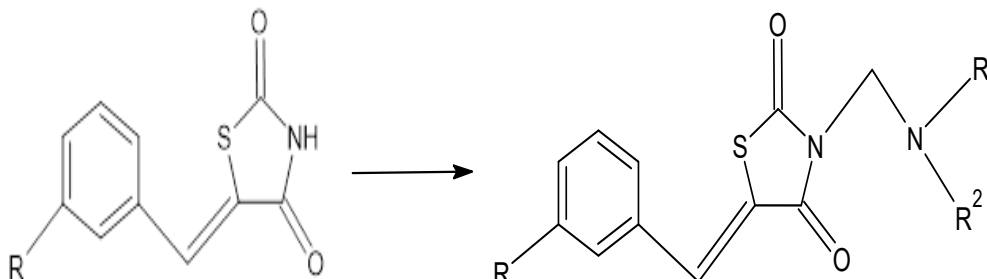


Procedure

Thiazolidinedione (0.1mol) + Substituted benzaldehyde (0.1mol) + Toluene (10 ml) + catalyst - [piperidine (2-3 drops)] - stirred, refluxed at 110 °C for 8-10 hrs - on

cooling, corresponding TZD derivatives precipitates. The product is filtered, washed, dried, purified and recrystallized.

Step 3: Mannich Reaction



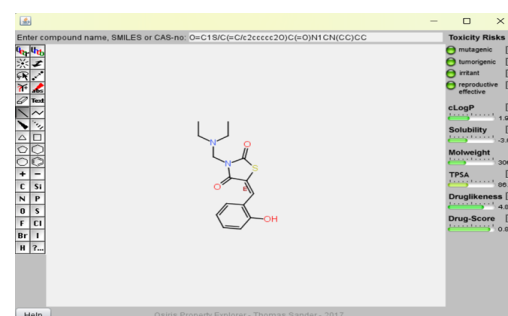
SN33C	Novel	 <p>The screenshot shows the Molinspiration software interface. The central window displays the chemical structure of SN33C, which is a 2,4-thiazolidinedione derivative. The right-hand panel lists various drug-likeness and toxicity metrics, including cLogP, Solubility, Molecular Weight, TPSA, Drug-likeness, and Drug Score. The bottom status bar indicates the software version as 2017.</p>
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Table 2: Drug-likeness property of synthesized compounds.

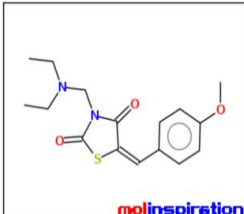

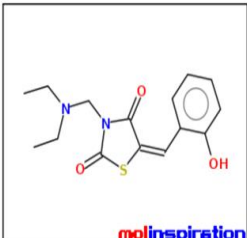
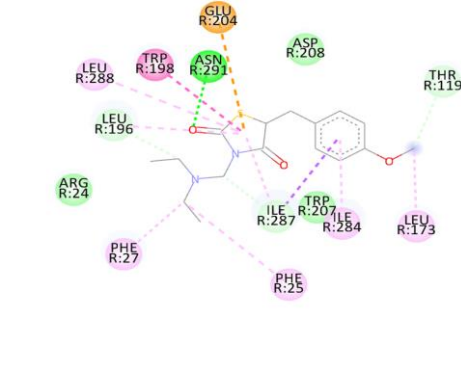
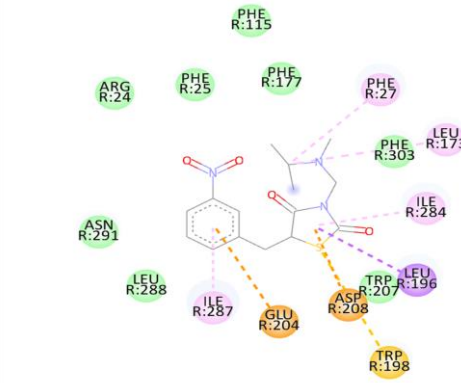
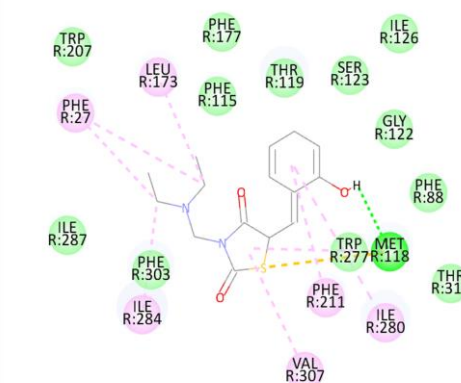
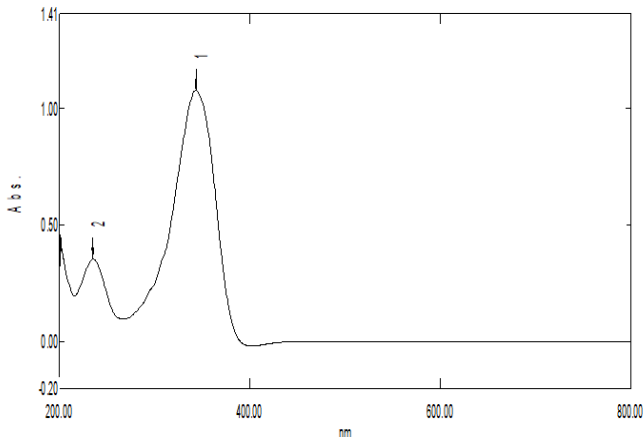
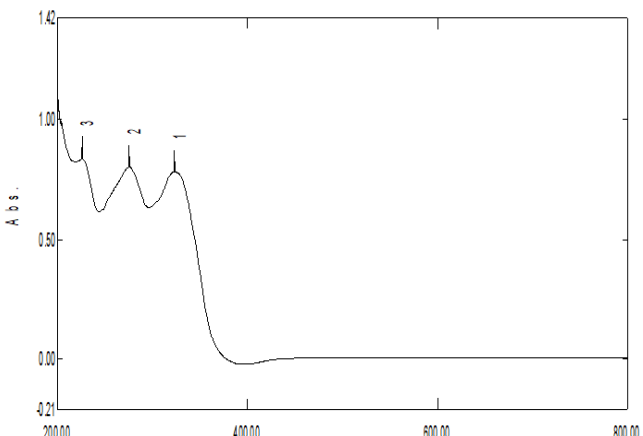
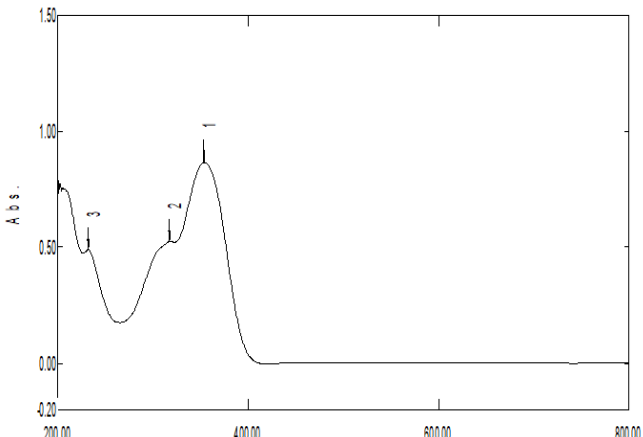
Compound ID	Drug-likeness property																		
SN1C	<p>miSMILES: <chem>CCN(CC)Cn2c(=O)sc(=Cc1ccc(OC)cc1)c2=O</chem></p>  <p>Molinspiration property engine v2022.08</p> <table> <tr><td>miLogP</td><td>2.38</td></tr> <tr><td>TPSA</td><td>51.55</td></tr> <tr><td>atoms</td><td>22</td></tr> <tr><td>MW</td><td>320.41</td></tr> <tr><td>nON</td><td>5</td></tr> <tr><td>nOHNH</td><td>0</td></tr> <tr><td>nviolations</td><td>0</td></tr> <tr><td>nrotb</td><td>6</td></tr> <tr><td>volume</td><td>291.36</td></tr> </table>	miLogP	2.38	TPSA	51.55	atoms	22	MW	320.41	nON	5	nOHNH	0	nviolations	0	nrotb	6	volume	291.36
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nOHNH	0																		
nviolations	0																		
nrotb	6																		
volume	291.36																		
SN15E	<p>miSMILES: <chem>[H]C(c1cccc(N(=O)=O)c1)=c2sc(=O)n(CN(C)C(C)C)c2=O</chem></p>  <p>Molinspiration property engine v2022.08</p> <table> <tr><td>miLogP</td><td>2.17</td></tr> <tr><td>TPSA</td><td>88.14</td></tr> <tr><td>atoms</td><td>23</td></tr> <tr><td>MW</td><td>335.38</td></tr> <tr><td>nON</td><td>7</td></tr> <tr><td>nOHNH</td><td>0</td></tr> <tr><td>nviolations</td><td>0</td></tr> <tr><td>nrotb</td><td>5</td></tr> <tr><td>volume</td><td>288.93</td></tr> </table>	miLogP	2.17	TPSA	88.14	atoms	23	MW	335.38	nON	7	nOHNH	0	nviolations	0	nrotb	5	volume	288.93
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nviolations	0																		
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SN33C	<p>miSMILES: <chem>[H]C(c1cccc1O)=c2sc(=O)n(CN(CC)CC)c2=O</chem></p>  <p>Molinspiration property engine v2022.08</p> <table> <tr><td>miLogP</td><td>2.26</td></tr> <tr><td>TPSA</td><td>62.54</td></tr> <tr><td>atoms</td><td>21</td></tr> <tr><td>MW</td><td>306.39</td></tr> <tr><td>nON</td><td>5</td></tr> <tr><td>nOHNH</td><td>1</td></tr> <tr><td>nviolations</td><td>0</td></tr> <tr><td>nrotb</td><td>5</td></tr> <tr><td>volume</td><td>273.83</td></tr> </table>	miLogP	2.26	TPSA	62.54	atoms	21	MW	306.39	nON	5	nOHNH	1	nviolations	0	nrotb	5	volume	273.83
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nON	5																		
nOHNH	1																		
nviolations	0																		
nrotb	5																		
volume	273.83																		

Table 3: Docking score and Docking interaction of synthesized compounds.

Compound ID	Docking score against biological target (Kcal/mol) FFAR4 (PDB ID:8ID6)	Docking interaction
SN1C	-9.48	
SN15E	-8.9	
SN33C	-8.32	

Characterization Studies
UV Spectroscopy

Table 4: UV spectrum of synthesized compounds.

Compound ID	UV Spectrum	λ_{max}
SN1C		343.5 nm
SN15E		323.5 nm
SN33C		353.5 nm

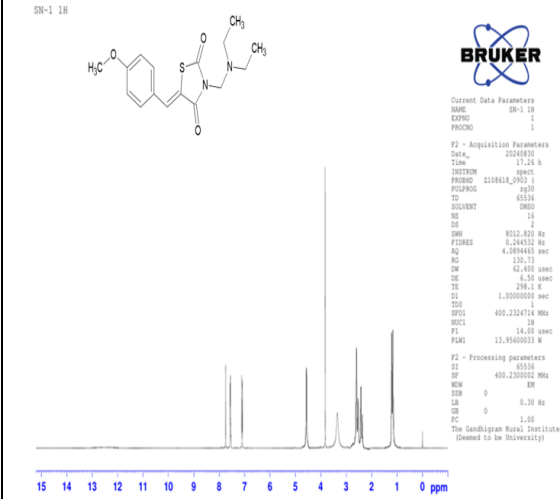
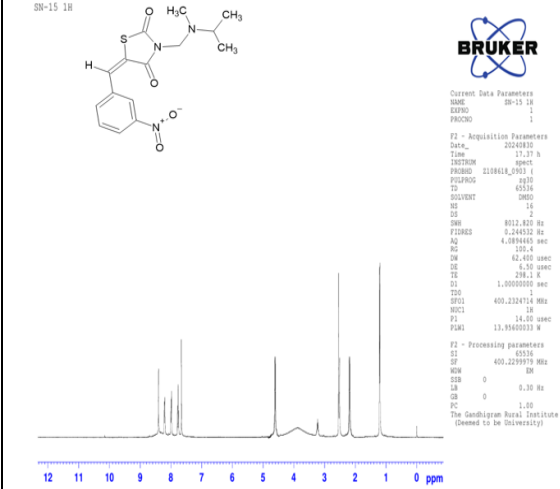
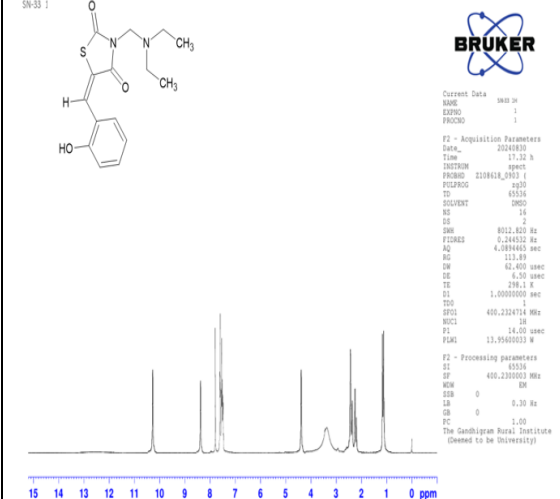
IR Spectroscopy

Table 5: IR spectrum of synthesized compounds.

Compound ID	IR Spectrum	Interpretation														
SN1C		<table><tr><th>FUNCTIONAL GROUP</th><th>STRETCHING FREQUENCY (cm⁻¹)</th></tr><tr><td>C-H (Aromatic)</td><td>3094</td></tr><tr><td>C-H (Aliphatic)</td><td>2962</td></tr><tr><td>C=C</td><td>1589</td></tr><tr><td>C=O</td><td>1697</td></tr><tr><td>C-N</td><td>1180</td></tr><tr><td>C-O</td><td>1011</td></tr></table>	FUNCTIONAL GROUP	STRETCHING FREQUENCY (cm ⁻¹)	C-H (Aromatic)	3094	C-H (Aliphatic)	2962	C=C	1589	C=O	1697	C-N	1180	C-O	1011
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C-O	1011															
SN15E		<table><tr><th>FUNCTIONAL GROUP</th><th>STRETCHING FREQUENCY (cm⁻¹)</th></tr><tr><td>C-H (Aromatic)</td><td>3032</td></tr><tr><td>C-H (Aliphatic)</td><td>2993</td></tr><tr><td>C=C</td><td>1620</td></tr><tr><td>C=O</td><td>1690</td></tr><tr><td>C-N</td><td>1157</td></tr><tr><td>NO₂</td><td>1566</td></tr></table>	FUNCTIONAL GROUP	STRETCHING FREQUENCY (cm ⁻¹)	C-H (Aromatic)	3032	C-H (Aliphatic)	2993	C=C	1620	C=O	1690	C-N	1157	NO ₂	1566
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NO ₂	1566															
SN33C		<table><tr><th>FUNCTIONAL GROUP</th><th>STRETCHING FREQUENCY (cm⁻¹)</th></tr><tr><td>C-H (Aromatic)</td><td>3024</td></tr><tr><td>C-H (Aliphatic)</td><td>2924</td></tr><tr><td>C=C</td><td>1589</td></tr><tr><td>C=O</td><td>1682</td></tr><tr><td>C-N</td><td>1157</td></tr><tr><td>O-H</td><td>3425</td></tr></table>	FUNCTIONAL GROUP	STRETCHING FREQUENCY (cm ⁻¹)	C-H (Aromatic)	3024	C-H (Aliphatic)	2924	C=C	1589	C=O	1682	C-N	1157	O-H	3425
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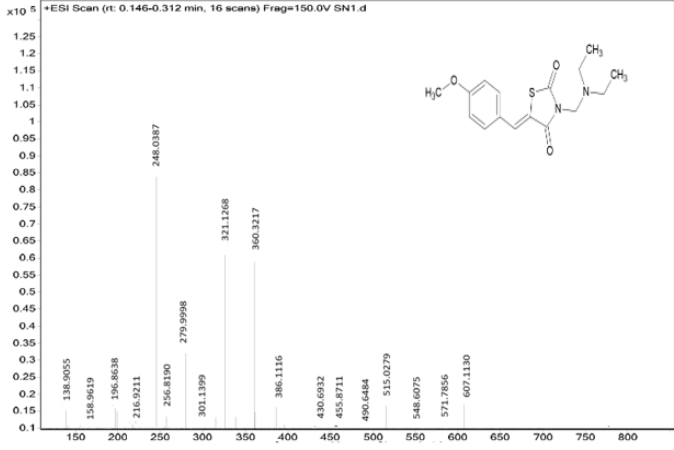
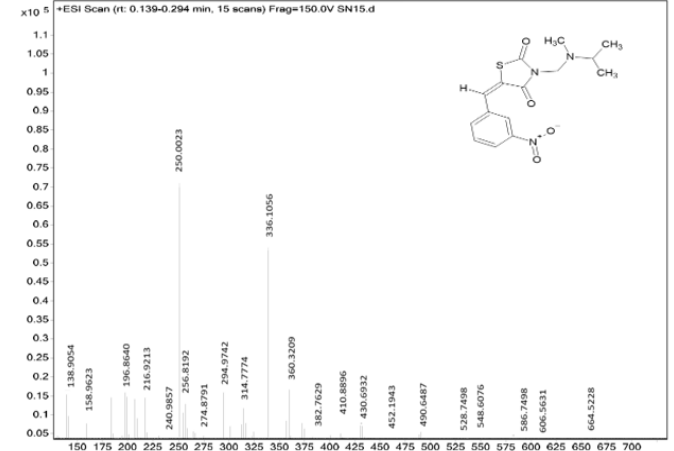
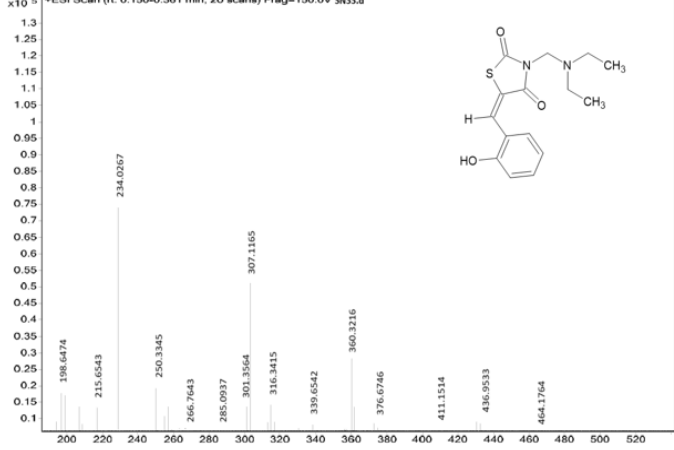
NMR Spectroscopy

Table 6: NMR spectrum of synthesized compounds.

Compound ID	NMR Spectrum	Interpretation			
SN1C		δ VALUE (PPM)	NATURE OF PROTONS	NATURE OF PEAKS	NUMBER OF PROTONS
		7.85	1-ethylene -H	Singlet	1
		7.63, 7.63	Aromatic C-H	Doublet	4
		7.14, 7.14		Doublet	
		4.55	Methylene -CH ₂	Singlet	6
		2.64, 2.64		Quartet	
		3.81		Singlet	9
		1.02, 1.02	Methyl -CH ₃	Triplet	
SN15E		δ VALUE (PPM)	NATURE OF PROTONS	NATURE OF PEAKS	NUMBER OF PROTONS
		8.49	1-ethylene -H	Singlet	1
		7.87 – 8.27	Aromatic C-H	Multiplet	4
		4.55	Methylene -CH ₂	Singlet	2
		2.69	Methine -CH	Multiplet	1
		2.26	Methyl -CH ₃	Singlet	9
		1.00, 1.00		Doublet	
SN33C		δ VALUE (PPM)	NATURE OF PROTONS	NATURE OF PEAKS	NUMBER OF PROTONS
		10.27	Alcohol -OH	Singlet	1
		8.34	1-ethylene -H	Singlet	1
		6.72 – 7.49	Aromatic C-H	Multiplet	4
		4.55	Methylene -CH ₂	Singlet	6
		2.64, 2.64		Quartet	
		1.02, 1.02		Triplet	6

Mass Spectroscopy

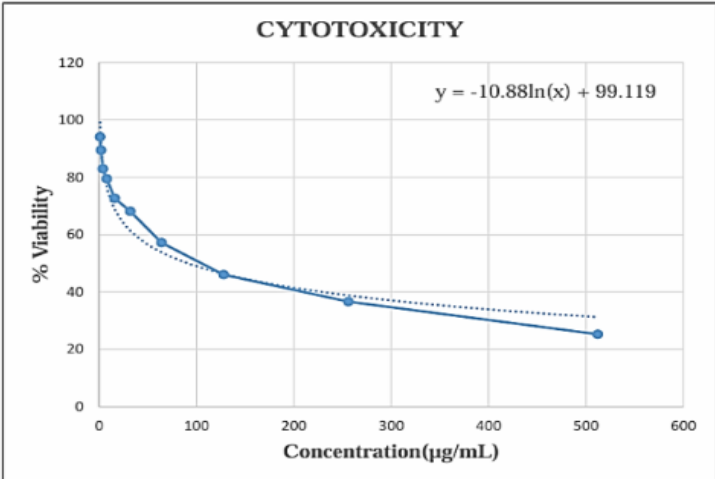
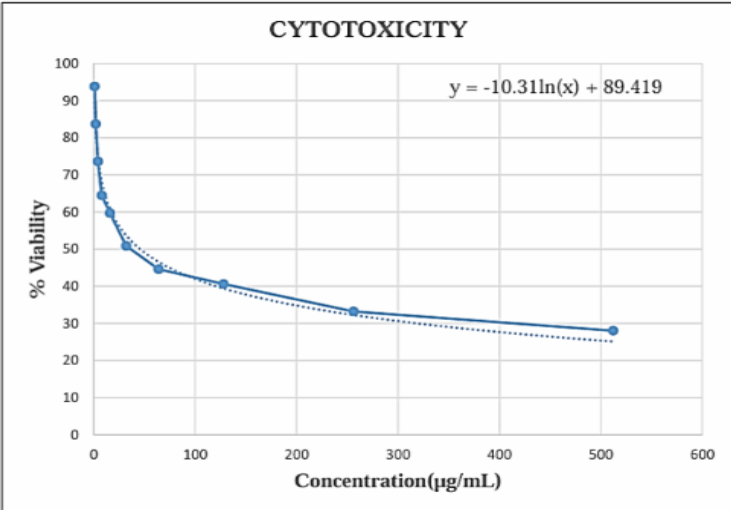
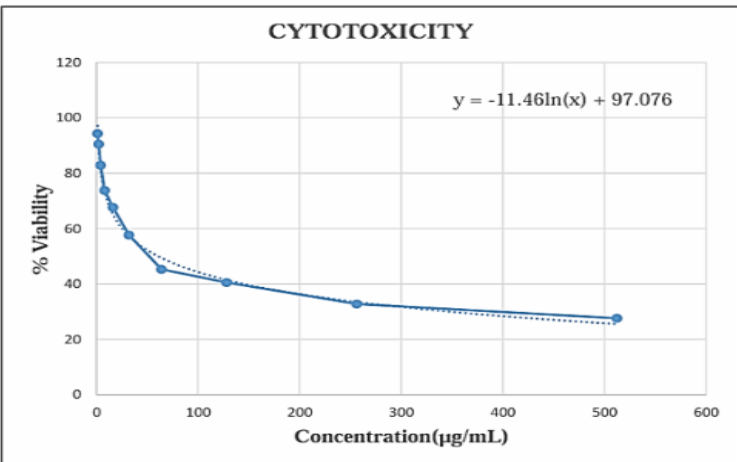
Table 7: Mass spectrum of synthesized compounds.

Compound ID	Mass Spectrum	Molecular weight
SN1C	<p>Sample Name: SN1, Inj Vol: 1, Data Filename: SN1.d, Position: InjPosition, ACQ Method: union.m, P1-F2, Instrument Name: Instrument 1, SampleType: Sample, User Name: User Name, IRM Calibration Status: Success, Acquired Time: 10/21/2024 2:23:59 PM</p> <p>*ESI Scan (rt: 0.146-0.312 min, 16 scans) Frag=150.0V SN1.d</p>  <p>Chemical structure of SN1C: <chem>CCN(CC)C1=CC(=O)N(C1=CC=C2C(=O)N(C2)C3=CC=C(C=C3)OC)C4=CC=CC=C4</chem></p>	321.12 g/mol
SN15E	<p>Sample Name: SN15, Inj Vol: 1, Data Filename: SN15.d, Position: InjPosition, ACQ Method: union.m, P1-F4, Instrument Name: Instrument 1, SampleType: Sample, User Name: User Name, IRM Calibration Status: Success, Acquired Time: 10/21/2024 2:27:54 PM</p> <p>*ESI Scan (rt: 0.130-0.294 min, 15 scans) Frag=150.0V SN15.d</p>  <p>Chemical structure of SN15E: <chem>CCN(CC)C1=CC(=O)N(C1=CC=C2C(=O)N(C2)C3=CC=C(C=C3)[O-])C4=CC=CC=C4</chem></p>	336.10 g/mol
SN33C	<p>Sample Name: SN33, Inj Vol: 1, Data Filename: SN33.d, Position: InjPosition, ACQ Method: union.m, P1-F6, Instrument Name: Instrument 1, SampleType: Sample, User Name: User Name, IRM Calibration Status: Success, Acquired Time: 10/21/2024 2:31:55 PM</p> <p>*ESI Scan (rt: 0.150-0.361 min, 20 scans) Frag=150.0V SN33.d</p>  <p>Chemical structure of SN33C: <chem>CCN(CC)C1=CC(=O)N(C1=CC=C2C(=O)N(C2)C3=CC=C(C=C3)O)C4=CC=CC=C4</chem></p>	307.11 g/mol

In-Vitro Studies

MTT Cytotoxicity Assay (Using HepG2 cell line)

Table 8: IC₅₀ value and Concentration Response Curve of synthesized compounds

Compound ID	Concentration (µg/mL) IC ₅₀ value	Concentration Response Curve
SN1C	91.342	 <p>CYTOTOXICITY</p> <p>$y = -10.88\ln(x) + 99.119$</p>
SN15E	45.758	 <p>CYTOTOXICITY</p> <p>$y = -10.31\ln(x) + 89.419$</p>
SN33C	60.816	 <p>CYTOTOXICITY</p> <p>$y = -11.46\ln(x) + 97.076$</p>

Free Fatty Acid Induction & treatment - Oil Red O Staining

Based on the desired balance between potency and safety, sample SN1C was chosen for further *in-vitro*

studies such as Oil Red O Staining and Gene Expression Study.

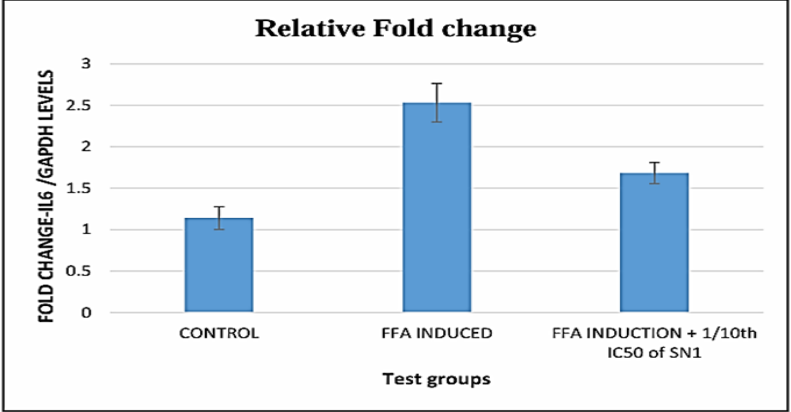
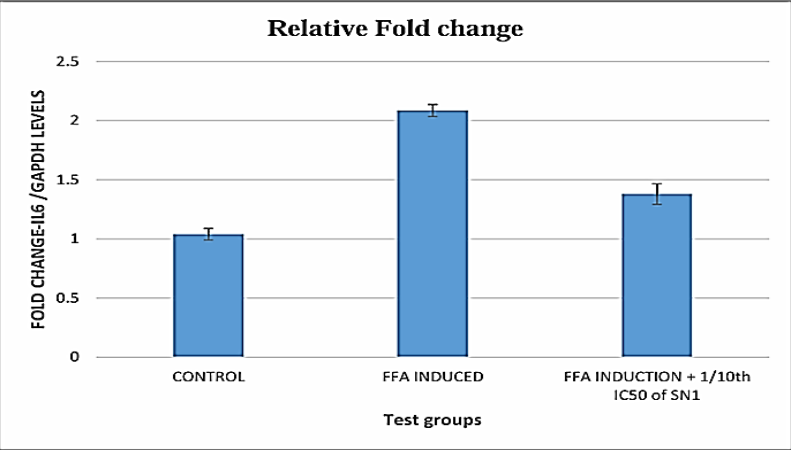


Figure 5: Oil Red O Staining of Compound SN1C.

Gene Expression Study

Gene expression study was carried out using sample SN1C for TNF- α and IL-6 gene (β -actin - reference gene).

Table 9: Gene Expression Study of Compound SN1C.

Compound ID	SN1C
Gene expression of TNF α	<div><p>Relative Fold change</p></div>
Gene expression of IL6	<div><p>Relative Fold change</p></div>

Conclusion

The findings of this research not only highlight the effectiveness of 2,4-thiazolidinedione derivatives in targeting FFAR4 but also pave the way for the development of new pharmacological treatments for NAFLD.

The *in-vitro* evaluation of synthesized compounds demonstrated that:

- SN1C is a promising candidate for NAFLD treatment due to its low toxicity, ability to reduce lipid accumulation (Oil Red O staining), and anti-inflammatory effects (gene expression study), while maintaining cell viability.
- By selecting SN1C based on its safety profile and moderate potency, the study highlights the importance of balancing efficacy and toxicity in developing therapeutics.

Through a comprehensive approach that included *in silico* molecular docking, synthetic chemistry, and *in vitro* biological assays, we successfully identified promising compounds that exhibited significant lipid-lowering activity and favourable safety profiles, underscoring their potential as FFAR4 agonists for NAFLD management.

Our integrated approach—merging computational modelling with synthetic and biological evaluations—provides a robust framework for future drug development in this therapeutic area. Given the increasing prevalence of NAFLD globally, continued exploration of these compounds could play a vital role in addressing this major public health challenge.

Author Contributions

All authors contributed equally to this research. All authors read and approved the final manuscript.

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Conflict of Interest

The authors declare no conflict of interest.

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Ethical Approvals

This study does not involve experiments on animals or human subjects.

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