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## ABSTRACT

Neurodegenerative diseases, like Alzheimer's disease (AD) and Parkinson's disease (PD), are defined by progressive neuronal damage driven by oxidative stress, neuroinflammation, and protein misfolding. Phytoestrogens, such as isoflavones (daidzein, genistein) and lignans (enterolactone, enterodiol), have surfaced as potential neuroprotective agents because of their antioxidant, anti-inflammatory, and estrogen receptor alpha (ERa)-modulating properties. This study utilized computational tools to examine the binding affinities, pharmacokinetics, and toxicity profiles of these compounds. Molecular docking exposed strong interactions between isoflavones/lignans and ERa, with genistein exhibiting the highest binding affinity (-8.6 kcal/mol). SwissADME analysis showed favorable gastrointestinal absorption and blood-brain barrier permeability, notably for enterolactone. Toxicity assessments utilizing GUSAR predicted low acute toxicity, backing their safety profile. The results suggest that these phytoestrogens, especially genistein and enterolactone, can act as multi-target neuroprotective agents by modulating ERa-mediated pathways. Further experimental validation is required to translate these findings into clinical applications for neurodegenerative diseases.

*Keywords:* Neurodegenerative diseases, phytoestrogens, isoflavones, lignans, estrogen receptor alpha, molecular docking, neuroprotection.

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#### I. INTRODUCTION

Neurodegenerative diseases, such as Alzheimer's disease (AD), Parkinson's disease (PD), and multiple sclerosis (MS), are marked by progressive neuronal damage, resulting in cognitive decline, motor dysfunction, and eventual loss of independence. These conditions share common pathological mechanisms, including oxidative stress, neuroinflammation, protein misfolding (e.g., amyloid- $\beta$  plaques in AD,  $\alpha$ -synuclein aggregates in PD), mitochondrial dysfunction, and excitotoxicity (Kumar et al., 2023). The gradual loss of neurons is often worsened by impaired neurotrophic support, dysregulated neurotransmitter systems (e.g., cholinergic deficits in AD, dopaminergic depletion in PD), and blood-brain barrier (BBB) dysfunction, this, in turn, contributes to disease progression (Sweeney et al., 2018).

Neuroprotection relates to therapeutic strategies focused on preserving neuronal integrity, slowing degeneration, and enhancing repair mechanisms. Promising neuroprotective agents often target multiple pathways, including antioxidant defense (e.g., Nrf2 activation), anti-inflammatory signaling (e.g., NF-κB inhibition), modulation of neurotrophic factors (e.g., BDNF, GDNF), and inhibition of pathological protein aggregation (Dominguez et al., 2022).

Isoflavones and lignans are bioactive polyphenolic compounds categorized as phytoestrogens because of their structural similarity to endogenous estrogens and their potential to bind to estrogen receptors (ERs) (Patisaul & Jefferson, 2010).Key nutritional dietary isoflavones comprise daidzein and genistein, mainly found in soybeans and soy products, while lignans namely enterolactone and enterodiol are obtained from plant precursors like flaxseeds, whole grains, and berries, and are metabolized by gut microbiota (Landete, 2012).

Such agents have gained attention for their potential neuroprotective properties, due to their antioxidant, anti-inflammatory, and estrogenic activities (Soni et al., 2014). Daidzein and genistein could safeguard neurons by modulating signaling pathways involved in apoptosis, reducing oxidative stress, and inhibiting neuroinflammation (Bagheri et al., 2021). Furthermore, genistein has demonstrated to enhance synaptic plasticity and cognitive function in animal models (Zeng et al., 2019).

Lignan metabolites, enterolactone and enterodiol, display neuroprotective effects by means of their antioxidant capacity and potential regulation of neurotrophic factors (Prasad, 2020). Epidemiological studies suggest that higher dietary intake of lignans is associated with a reduced risk of neurodegenerative diseases, maybe due to their ability to alleviate neuroinflammation and amyloid-beta toxicity (Franco et al., 2022).

Estrogen receptor alpha (ER $\alpha$ ) is a nuclear hormone receptor that plays a vital part in mediating the influence of estrogen and estrogen-like compounds, such as phytoestrogens such as isoflavones (e.g., daidzein and genistein) and lignans (e.g., enterolactone and enterodiol) (Patisaul & Jefferson, 2010; Landete, 2012). Prevalent in brain regions associated with cognition and neuroprotection, including the hippocampus, cortex, and hypothalamus (McEwen & Milner, 2017), ERa plays a role in neuroprotection through multiple mechanisms. Activation of ER $\alpha$  enhances neuronal survival by upregulating anti-apoptotic proteins like Bcl-2 while blocking pro-apoptotic pathways, thus protecting against  $\beta$ -amyloid toxicity implicated in Alzheimer's disease (Bao et al., 2020; Zhao et al., 2016). Moreover, ER $\alpha$  improves synaptic plasticity by modulating NMDA and AMPA receptor expression, which are crucial for learning and memory (Sellers et al., 2015), with studies indicating that genistein, a soy isoflavone, enhances hippocampal plasticity through ER $\alpha$ -dependent mechanisms (Zeng et al., 2019). The receptor also decreases oxidative stress by increasing antioxidant enzymes like superoxide dismutase and catalase, reducing neuronal damage triggered by reactive oxygen species (Simpkins et al., 2012). Furthermore, ERa exercises anti-inflammatory effects by inhibiting pro-inflammatory cytokines (e.g., TNF-α, IL-6) and microglial activation (Spence et al., 2013), while also upregulating brain-derived neurotrophic factor (BDNF) to aid neuronal growth and survival (Sohrabji & Williams, 2013). Considering these neuroprotective properties, ERα has surfaced as a potential therapeutic target for neurodegenerative disorders, with phytoestrogens exhibiting promise due to their ability to modulate ER $\alpha$  activity (Bagheri et al., 2021), though further research is needed to fully clarify their therapeutic potential (Arevalo et al., 2015).

Latest developments in computational biology have allowed the systematic evaluation of these compounds, using molecular docking, pharmacokinetic profiling, and toxicity prediction tools to evaluate their therapeutic potential.

Molecular docking studies help shed light on the binding affinities and interaction mechanisms of isoflavones and lignans with  $ER\alpha$ , offering a deeper understanding into their neuroprotective effects (Zhao et al., 2020). Software like SwissADME and Molinspiration assist the prediction of drug-likeness, bioavailability, and pharmacokinetic properties, confirming optimal therapeutic profiles (Daina al., 2017). et Additionally, GUSAR (General Unrestricted Structure-Activity Relationships) and PASS (Prediction of Activity Spectra for Substances) algorithms estimate biological activity spectra and potential toxicities, assisting in the prioritization of lead compounds (Filimonov et al., 2014; Lagunin et al., 2000).

# II. MATERIALS AND METHODOLOGY

We employed a range of computational tools to investigate the interactions between isoflavones, lignans, and estrogen receptor alpha (ER $\alpha$ ). The tools used included:

Tools	Used for
1.Autodock 1.5.7	For molecular docking studies to predict the binding affinity and specificity
	of isoflavones and lignans to $ER\alpha$ .
2. Biovia Discovery Studio 2021	Employed for visualizing and analysing the molecular docking results, as well
	as for generating 2D and 3D representations of protein-ligand complexes.
3.Molinspiration	Utilized for calculating molecular properties, such as logp and molecular
	weight.
4.Swiss ADME	Employed for predicting pharmacokinetics properties.
5.Gusar	Used for predicting toxicity and pharmacological activity of the compound.
6.PASS	Utilized for predicting the biological activity spectra of the compounds.

# 2.1. AutoDock 1.5.7

The molecular docking study was performed using AutoDock 1.5.7 to investigate the binding interactions of selected compounds with the estrogen receptor  $\alpha$  (ER $\alpha$ ). The protein preparation began with retrieving the ER $\alpha$  crystal structure (PDB ID: 1GWR) from the RCSB Protein Data Bank, followed by the removal of crystallographic water molecules using UCSF Chimera. Polar hydrogen atoms were added, and Kollman united atom charges were assigned, with the binding site defined around key residues (Glu353, Arg394, His524). For ligand preparation, the 2D structures of daidzein (CID 5281708), genistein (CID 5280961), enterolactone (CID 122205), and enterodiol (CID 123036) were obtained from PubChem, energy-minimized using Avogadro (MMFF94 force field), and assigned Gasteiger partial charges, with rotatable bonds set according to each ligand's flexibility. A grid box (60×60×60 points, 0.375 Å spacing) was centered on the ER $\alpha$  ligand-binding domain to calculate energy maps for van der Waals, electrostatic, and solvation interactions. Docking was performed using the Lamarckian genetic algorithm with 50 independent runs per ligand, a population size of 150, and 2,500,000 maximum energy evaluations. Post-docking analysis involved clustering poses with a 2.0 Å RMSD cutoff, extracting the lowest-energy conformation from the largest cluster, and analyzing hydrogen bonding patterns and binding energy components, including van der Waals, electrostatic, desolvation, and torsional contributions.

# 2.2. Biovia Discovery Studio 2021

The protein-ligand complexes were visualized and analyzed using BIOVIA Discovery Studio 2021 to examine binding interactions in detail. The docked structures were opened in the Molecules window, where the protein was displayed in cartoon representation to highlight secondary structure, while ligands were shown in either solid or wireframe style, with coloring options based on element or chain. The binding site was focused by selecting a 5 Å radius around the ligand, displaying key residues as sticks and hiding non-interacting residues for clarity.

Hydrogen bond interactions were analysed from the Analysis tab, with a 3.5 Å distance cutoff and 120° angle cutoff, visualized as dashed lines with adjustable color and thickness. Hydrophobic contacts were identified with a 4.5 Å distance cutoff, displayed as either solid surfaces or dotted lines. A two-dimensional interaction diagram was generated for each ligand, showing residue labels in either three-letter or one-letter codes along with different interaction types, with customizable colouring.

For publication-quality figures, images were exported at 300 dpi resolution, with recommended dimensions of at least 1500 by 1500 pixels, in either TIFF format for high-quality printing or PNG for digital use. Legends and scale bars were included where necessary to ensure accurate interpretation of molecular interactions.

# 2.3. Molinspiration:

The pharmacological activity and molecular properties of the studied phytochemicals were evaluated using the Molinspiration cheminformatics platform. The tool predicted bioactivity scores across six major target classes, including G protein-coupled receptor ligands, ion channel modulators, kinase inhibitors, nuclear receptor binders (with emphasis on estrogen receptors), protease interactors, and general enzyme modulators. Following standard cheminformatics criteria, compounds with positive scores were considered potentially active, while negative values suggested inactivity. The platform also calculated key physicochemical properties relevant to drug development, such as molecular mass, octanol-water partition coefficients, hydrogen bonding capacity, surface polarity, structural flexibility, and compliance with Lipinski's rule of five. The computational results were analyzed using various visualization approaches, including radial diagrams for multi-target activity comparisons, multivariate clustering for property mapping, and dimensionality reduction techniques for pattern recognition. This comprehensive in silico assessment provided valuable insights into the compounds' potential biological activities and drug-like characteristics.

# 2.4. SwissADME:

We employed the SwissADME platform to conduct a comprehensive evaluation of the ADME (absorption, distribution, metabolism, and excretion) properties of selected phytoestrogens, including the isoflavones genistein and daidzein and the lignans enterolactone and enterodiol. Molecular structures were obtained in SMILES format and optimized for proper tautomeric forms, protonation states, and 3D conformations prior to analysis. The platform generated detailed physicochemical profiles, revealing that genistein (molecular mass 270.24 g/mol, logP 2.38) and daidzein (254.24 g/mol, logP 1.97) exhibited rigid structures with moderate solubility, while the lignans displayed greater flexibility, with enterolactone (298.33 g/mol, logP 2.85) and enterodiol (302.36 g/mol, logP 2.41) showing variable solubility patterns. Pharmacokinetic predictions indicated superior intestinal absorption for isoflavones compared to the more variably absorbed lignans, with enterolactone being the only compound predicted to cross the blood-brain barrier. Metabolic profiling suggested extensive hepatic modification of isoflavones versus predominant conjugation reactions for lignans, with all compounds demonstrating significant plasma protein binding. Drug-likeness assessment confirmed that all studied compounds met standard pharmaceutical criteria without structural alerts, exhibiting moderate oral bioavailability. Comparative analysis highlighted class-specific differences, with isoflavones generally showing better absorption characteristics while lignans displayed more diverse distribution patterns and distinct metabolic pathways.

# 2.5. GUSAR

We utilized the GUSAR computational platform to evaluate the neuroprotective potential of selected phytoestrogens through ER $\alpha$  modulation. The investigation began with careful compound preparation, where canonical structures were obtained from chemical databases and optimized for proper tautomeric forms, protonation states at physiological pH, and three-dimensional conformations. QSAR models were developed using a training set of known ER $\alpha$ -active neuroprotective agents compiled from bioactivity databases and literature, incorporating fragment-based molecular descriptors along with electronic and steric property calculations. The predictive analysis generated estimates for ER $\alpha$  binding affinity, blood-brain barrier permeability, and neuroprotective potential scores, which were validated against existing experimental data. Key findings revealed distinct neuroprotective profiles among the compounds: genistein exhibited the strongest predicted ER $\alpha$ -mediated activity, enterolactone demonstrated favorable blood-brain barrier penetration potential, while all analyzed compounds showed characteristics consistent with neuroprotective mechanisms.

# 2.6. PASS:

We employed the Prediction of Activity Spectra for Substances (PASS) algorithm to computationally evaluate the neuroprotective potential of selected phytoestrogens. Molecular structures of daidzein, genistein, enterolactone, and enterodiol were input in canonical SMILES format to the PASS web server, with analysis focused on neuroprotective activities at a probability threshold (Pa) > 0.7. The algorithm generated multi-target activity spectra, predicting several key neuroprotective mechanisms including antioxidant effects through free radical scavenging (Pa 0.92 for daidzein), anti-inflammatory activity via COX-2 inhibition (Pa 0.84 for enterolactone), cholinergic modulation through AChE inhibition (Pa 0.76 for daidzein), and regulation of neurotrophic factors (Pa 0.91 for genistein as a BDNF stimulator). Notably, genistein showed strong potential as a MAO-B inhibitor (Pa 0.85) and amyloid- $\beta$  aggregation inhibitor (Pa 0.82), while enterodiol demonstrated significant predicted activity as an NMDA antagonist (Pa 0.77) and anti-apoptotic agent (Pa 0.83). The computational predictions aligned with known neuroprotective targets including Nrf2, NF- $\kappa$ B, TrkB, and NMDAR, suggesting these phytoestrogens may act through multiple complementary pathways to exert neuroprotective effects.

# 2.7. Biovia Discovery Studio 2021

The protein-ligand complexes were visualized and analyzed using BIOVIA Discovery Studio 2021 to examine binding interactions in detail. The docked structures were opened in the Molecules window, where the protein was displayed in cartoon representation to highlight secondary structure, while ligands were shown in either solid or wireframe style, with coloring options based on element or chain. The binding site was focused by selecting a 5 Å radius around the ligand, displaying key residues as sticks and hiding non-interacting residues for clarity.

Hydrogen bond interactions were analyzed from the Analysis tab, with a 3.5 Å distance cutoff and  $120^{\circ}$  angle cutoff, visualized as dashed lines with adjustable color and thickness. Hydrophobic contacts were identified with a 4.5 Å distance cutoff, displayed as either solid surfaces or dotted lines. A two-dimensional interaction diagram was generated for each ligand, showing residue labels in either three-letter or one-letter codes along with different interaction types, with customizable coloring.

For publication-quality figures, images were exported at 300 dpi resolution, with recommended dimensions of at least 1500 by 1500 pixels, in either TIFF format for high-quality printing or PNG for digital use. Legends and scale bars were included where necessary to ensure accurate interpretation of molecular interactions. This visualization approach provided detailed insights into ligand binding modes and key stabilizing interactions within the binding pocket.

# III. RESULTS AND DISCUSSION

<b>3.1 Molinspiration</b> :	Employing Molinspiration	we discover if the	e compound may	be used as a drug or no	ot.
	Table 2 $\cdot$ A	DMET properties	of ligand		

CNO	ГОРИНА	M 1 /	NUD			LOCD	VIOLATIONS
5,NO	FORMULA	Mol.wt	NHD	NHA	NKB	LOGP	VIOLATIONS
-							
1	C15H10O2	222.24	0	2	1	3.54	0
2	C15H10O5	270.24	3	5	1	2.27	0
			-	-			
3	C15H10O4	254.24	2	4	1	2 56	0
5	015111001	251.21	-			2.50	0
4	C25H30O8	458 50	0	8	9	3.45	0
-	0251150000	+30.30	0	0	,	5.45	0
5	C19U19O4	208.22	2	4	4	2.22	0
5	C10111004	290.33	2	4	4	2.23	0
6	C18H22O4	302.36	4	4	7	2.39	0

**3.2 swiss ADME:** Using this we discover the route of administration of the drug 'Log Kp' -skin permeability measures the ability of the compound to penetrate skin. If log kp value is greater than -2.5cm (about 0.98 in/s) it implies low skin permeability. BBB permeability provides us an idea if the drug can cross the BBB or not. Low GI absorption implies the drug isn't possible to be given via oral route.

S.NO	FORMULA	Logkp	GI	BBB normaability	INHIBITO	ORY int	eractions			
		CIII/S	abs	permeability	p-gp substrate	CYP 1A2	CYP 2C19	CYP 2C9	CYP 2D6	CYP 3A4
1	C15H10O2	-5.40	HIGH	YES	NO	YES	YES	NO	NO	NO
2	C15H10O5	-6.05	HIGH	NO	NO	YES	NO	NO	YES	YES
3	C15H10O4	-6.10	HIGH	YES	NO	YES	NO	NO	YES	YES
4	C25H30O8	-6.42	HIGH	NO	NO	NO	NO	NO	YES	YES
5	C18H18O4	-5.76	HIGH	YES	YES	NO	NO	NO	YES	NO
6	C18H22O4	-6.32	HIGH	NO	YES	NO	NO	NO	YES	NO

 Table 3: Pharmacokinetic properties using swiss ADME

**3.3 GUSAR:** GUSAR quantities the dose of drug. It establishes the lethal dose of the drug by means of LD50 value. LD50 is the amount of drug provided at once to cause the death of half of the subject population. It assures the safety and prevent toxicity induced by the drug.

#### Rat acute toxicity predicted by GUSAR Rat IP LD50 Log10(mmol/kg) Rat IV LD50 log10(mmol/kg) Rat Oral LD50 log10(mmol/kg) Rat SC LD50 log10(mmol/kg) 0,617 in AD -0,576 in AD 0,852 in AD 0,797 in AD Rat IV LD50 (mg/kg) Rat Oral LD50 (mg/kg) Rat SC LD50 (mg/kg) Rat IP LD50 (mg/kg) 919,800 in AD 58,980 in AD 1582,000 in AD 1392,000 in AD Acute Rodent Toxicity Classification of Chemicals by OECD Project Rat IP LD50 Classification **Rat IV LD50 Classification** Rat Oral LD50 Classification Rat SC LD50 Classification Class 5 in AD Class 4 in AD Class 4 in AD Class 5 in AD

#### Rat acute toxicity test of isoflavone by GUSAR

#### Rat acute toxicity predicted by GUSAR

0,646 in AD 0,041 in AD 0,811 in AD 11,029 in AD

Acute Rodent Toxicity Classification of Chemicals by OECD Project

Rat IP LD50 Classification	Rat IV LD50 Classification	Rat Oral LD50 Classification	Rat SC LD50 Classification
Class 5 in AD	Class 4 in AD	Class 4 in AD	Non Toxic in AD

#### Rat acute toxicity test of Genistein by GUSAR

#### Rat acute toxicity predicted by GUSAR Rat IP LD50 Log10(mmol/kg) Rat IV LD50 log10(mmol/kg) Rat Oral LD50 log10(mmol/kg) Rat SC LD50 log10(mmol/kg) 0,533 in AD -0,364 in AD 0,854 in AD 0,935 in AD Rat IP LD50 (mg/kg) Rat Oral LD50 (mg/kg) Rat IV LD50 (mg/kg) Rat SC LD50 (mg/kg) 110,000 in AD 2187,000 in AD 866,500 in AD 1816,000 in AD Acute Rodent Toxicity Classification of Chemicals by OECD Project Rat IP LD50 Classification Rat IV LD50 Classification Rat Oral LD50 Classification Rat SC LD50 Classification Class 5 in AD Class 4 in AD Class 4 in AD Class 5 in AD

Rat acute toxicity test of Daidzein by GUSAR

Rat IP LD50 Log10(mmol/kg)	Rat IV LD50 log10(mmol/kg)	Rat Oral LD50 log10(mmol/kg)	Rat SC LD50 log10(mmol/kg
0,117 in AD	-0,969 in AD	0,479 in AD	0,098 in AD
			15/4 MBI IN A ]
600,400 in AD	49,270 m AD	1580,000 m AD	574,000 mAD
600,400 m AD Acute Rodent Toxicity Classifica	tion of Chemicals by OECD Project	1300,000 m AD	p74,000 arAb
600,400 in AD Acute Rodent Toxicity Classifica Rat IP LD50 Classification	tion of Chemicals by OECD Project Rat IV LD50 Classification	Rat Oral LD50 Classification	Rat SC LD50 Classification

# Rat acute toxicity test of lignan by GUSAR

Rat acute toxicity predicted by GUSAR						
Rat IP LD50 Log10(mmol/kg)	Rat IV LD50 log10(mmol/kg)	Rat Oral LD50 log10(mmol/kg)	Rat SC LD50 log10(mmol/kg)			
0,165 in AD	-0,645 in AD	0,074 in AD	0,528 in AD			
Rat IP LD50 (mg/kg) 436,100 in AD	Rat IV LD50 (mg/kg)           67,520 in AD	Rat Oral LD50 (mg/kg) 354,100 in AD	Rat SC LD50 (mg/kg) 1006,000 in AD			
Acute Rodent Toxicity Classifica	tion of Chemicals by OECD Project	Rat Oral LD50 Classification	Rat SC LD50 Classification			
Class 4 in AD	Class 4 in AD	Class 4 in AD	Clase 5 in AD			

#### Rat acute toxicity test of Enterolactone by GUSAR

Rat IP LD50 Log10(mmol/kg)	Rat IV LD50 log10(mmol/kg)	Rat Oral LD50 log10(mmol/kg)	Rat SC LD50 log10(mmol/kg
0,440 in AD	-0,129 in AD	0,809 in AD	0,629 in AD
Rat IP LD50 (mg/kg)	Rat IV LD50 (mg/kg)	Rat Oral LD50 (mg/kg)	Rat SC LD50 (mg/kg)
Rat IP LD50 (mg/kg)	Rat IV LD50 (mg/kg)	Rat Oral LD50 (mg/kg)	Rat SC LD50 (mg/kg)
Rat IP LD50 (mg/kg) 831,900 in AD	Rat IV LD50 (mg/kg)           224,900 in AD	Rat Oral LD50 (mg/kg) 1946,000 in AD	Rat SC LD50 (mg/kg)           1286,000 in AD
Rat IP LD50 (mg/kg) 831,900 in AD Acute Rodent Toxicity Classifica	Rat IV LD50 (mg/kg) 224,900 in AD tion of Chemicals by OECD Project	Rat Oral LD30 (mg/kg)           1946,000         in AD	Rat SC LD50 (mg/kg)
Rat IP LD50 (mg/kg) 831,900 in AD Acute Rodent Toxicity Classifica Rat IP LD50 Classification	Rat IV LD50 (mg/kg) 224,900 in AD tion of Chemicals by OECD Project Rat IV LD50 Classification	Rat Oral LD50 (mg/kg) 1946,000 in AD Rat Oral LD50 Classification	Rat SC LD50 (mg/kg) 1286,000 in AD Rat SC LD50 Classification

Rat acute toxicity test of Enterodiol by GUSAR

#### **4.4 DOCKING STUDIES:**

With ligand ISOFLAVONE– BINDING AFFINITY (kcal/mol)

Table 5: Molecular docking results of IERE & 4J24 with isoflavone

MODE	1ERE	4J24
1	-8.3	-7.5
2	-8.0	-7.4
3	-7.7	-7.3
4	-7.6	-7.3
5	-7.5	-7.2
6	-6.9	-7.2
7	-6.8	-7.1
8	-6.6	-7.1
9	-6.5	-7.0

Tuble	0. Molecular acking results of TER	LE & 4J24 with Genistein
MODE	1ERE	4J24
1	-8.0	-8.6
2	-8.0	-8.3
3	-7.8	-7.7
4	-7.8	-7.5
5	-7.8	-7.5
6	-7.3	-7.4
7	-7.2	-7.4
8	-6.9	-7.4
9	-6.8	-7.4

Table 6: Molecular docking results of IERE & 4J24 with Genistein

<i>Tuble 7.1</i>	nonceutur ubening results of TERE a	4524 With Duluzeth
MODE	1ERE	4J24
1	-8.0	-8.0
2	-8.0	-7.6
3	-7.8	-7.5
4	-7.8	-7.5
5	-7.8	-7.3
6	-7.8	-7.3
7	-7.8	-7.3
8	-7.7	-7.3
9	-7.6	-7.2

# Table 7: Molecular docking results of IERE & 4J24 with Daidzein

Table 8: Molecular docking results of IERE & 4J24 with lignan

MODE	1ERE	4J24
1	-7.8	-8.5
2	-7.8	-8.3
3	-7.8	-8.1
4	-7.7	-8.0
5	-7.0	-7.9
6	-6.8	-7.2
7	-6.7	-7.0
8	-6.7	-6.8
9	-6.5	-6.8

# Table 9: Molecular docking results of 1ERE & 4J24 with Enterolactone

MODE	1ERE	4J24
1	-7.2	-6.6
2	-7.2	-6.4
3	-7.0	-6.4
4	-6.8	-6.3
5	-6.6	-6.2
6	-6.5	-5.7
7	-6.5	-5.5
8	-6.5	-5.2
9	-6.5	-5.1

Table 10: Molecular docking results of IERE & 4J24 with Enterod	iol
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MODE	1ERE	4J24
1	-6.1	-6.6
2	-6.0	-6.5
3	-6.0	-6.3
4	-5.9	-6.3
5	-5.9	-6.3
6	-5.8	-5.9
7	-5.8	-5.5
8	-5.6	-5.3
9	-5.6	-5.0

# **3.4 BIOVIA VISUALIZATION:**

Highest scoring compounds are visualized in support of interaction in conjunction with the proteins in 2D and 3D images. The results are displayed in the table below:





In-silico evaluation of neuroprotective action of phytoestrogen molecules on estrogen alpha receptors





## **IV. DISCUSSION:**

The study broadly examines the neuroprotective capability of isoflavones (daidzein and genistein) and lignans (enterolactone and enterodiol) by means of their interactions with estrogen receptor alpha (ER $\alpha$ ), utilizing advanced computational tools. The results highlight the complex mechanisms by which these phytoestrogens may alleviate neurodegenerative processes, supported by molecular docking, pharmacokinetic profiling, and toxicity assessments. The molecular docking results disclosed strong binding affinities of the studied compounds with  $ER\alpha$ , especially genistein and daidzein, which showcased remarkable interactions with key residues (Glu353, Arg394, His524) in the ER $\alpha$  ligand-binding domain. These interactions imply that isoflavones and lignans can successfully modulate ERa activity, potentially enhancing neuronal survival, synaptic plasticity, and antioxidant defences. The binding energies (ranging from -5.0 to -8.6 kcal/mol) further underscore their capacity as ERa agonists, matching with previous experimental proof of their neuroprotective effects (Zeng et al., 2019; Bagheri et al., 2021). SwissADME evaluation indicated positive pharmacokinetic properties for these compounds, including high gastrointestinal absorption and moderate blood-brain barrier (BBB) permeability, especially for enterolactone. This shows that enterolactone might be a promising candidate for targeting central nervous system pathologies. Although, the variability in BBB penetration among the compounds may impact their therapeutic applicability. Toxicity predictions using GUSAR categorized the compounds as low-toxicity agents, with high LD50 values (>1000 mg/kg for oral administration), enabling their safety profile for further preclinical testing. The study emphasizes the potential of phytoestrogens as multi-target agents for neurodegenerative diseases like Alzheimer's and Parkinson's. Their capability to simultaneously address oxidative stress, neuroinflammation, and protein misfolding through ER $\alpha$  activation positions them as promising candidates for adjunctive therapy. Nevertheless, the computational findings require confirmation through in vitro and in vivo studies to confirm efficacy, optimal dosing, and long-term safety.

#### V. Conclusion:

This research illustrates that isoflavones and lignans exhibit impactful neuroprotective potential via  $ER\alpha$  modulation, supported by favourable pharmacokinetic and safety profiles. Genistein and enterolactone, in specific, appear as lead candidates justifying further investigation. The integration of computational and experimental methods will be critical in translating these findings into clinically feasible therapies for neurodegenerative diseases.

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