

# Molecular Docking Studies on Isolated and Characterized Compounds of Marine and Plant Origin with Antidiabetic Activity

## Abstract

Introduction: Diabetes Mellitus is a chronic disease characterized by hyperglycemia and disturbance in protein and fat metabolism. This study was carried out to determine the antidiabetic activities and molecular docking interactions of some marine and plant-derived compounds earlier reported.

Methods: The antidiabetic activities were determined by using Virtual screening and rigid docking protocol in identifying the compounds' binding affinity with the target protein while toxicity, pharmacokinetics and physicochemical profiles were established using online web servers: Protox 11 and SwissAdme respectively to determine their drug-likeness. The target receptor (PDB Code: 3VI8) which was used for the molecular docking studies was retrieved from the Protein Data Bank (PDB), (<http://www.pdb.org>) database while the structure of the ligand, rosiglitazone was retrieved from PubChem database (<https://pubchem.ncbi.nlm.nih.gov>). MMFF94 force field was used for energy minimization of the ligand molecule. The prepared compounds were then subjected to interact with the receptor through molecular docking. The protocol facilitates flexible compound docking for various compound conformers within the rigid receptor. The Best conformation for each compound was chosen and the interaction was visualized in Discovery studio).

Results: Compounds; 6, 7, 12, 32, 66, 80, 89, 121, 138 and, 139 showed greater binding affinity with PPAR $\gamma$  target protein (3vi8) with free binding energy of (-6.2 to 8.1 kcal/mol) comparable to the standard drug with 5.2 kcal/mol. Similarly, the selected compounds possess acceptable physicochemical and toxicity profiles which promise good oral bioavailability.

Conclusion: Ten (10) compounds extracted from seven (7) naturally occurring (marine) species were found as potential peroxisome proliferators-activated receptor gamma (PPAR $\gamma$ ) agonist with promising antidiabetic activity. The *insilico* studies showed that the compounds had strong binding interactions with the drug receptors.

**Keywords:** Antidiabetic Activity, Marine and plant-derived, Molecular Docking, PPAR $\gamma$ , physicochemical.

## 1.0 Introduction

Diabetes Mellitus (DM) is a chronic disease condition which is majorly characterized by hyperglycemia and disturbance of protein, fat and carbohydrate metabolism. There are basically two types of DM, namely, the types 1 (T1DM) and 2 (T2DM), otherwise known as insulin-dependent and none insulin dependent, respectively. Antidiabetic drugs or hypoglycemic agents are medications that work to lower the blood glucose concentrations (i.e. the amount of sugar in the blood). There are different classes of antidiabetic drugs currently used in clinical practice and their selection depends on several patients' factors including; the nature of the diabetes, age and underlying diseases etc. Antidiabetic drugs exert their useful effects through different mechanisms of action such as (a) increasing insulin levels in the body, (b) increasing the body's sensitivity (or decreasing its resistance) to insulin, or (c) decreasing glucose absorption in the intestines.<sup>1,5</sup>

, However, due to actual adverse side effects of some of these drugs, their use in treatments are somewhat considered to be unsatisfactory in terms of the prevention of complications and preservation of quality of life. For example, the  $\alpha$ -glucosidase inhibitors, such as acarbose and miglitol, while effective at decreasing the absorption of glucose by interfering with the action of  $\alpha$ -glucosidases present in the small intestinal brush border, are often associated with abdominal bloating, diarrhea and flatulence. Conventional insulin secretagogues, such as sulfonylureas and the class of meglitinides, both result in the induction of hypoglycemia. While metformin is the only therapeutic agent that has been demonstrated to reduce macrovascular events in T2DM, its use may be restricted to certain conditions. For example, Metformin is not recommended in decreased renal or hepatic function.

<sup>14</sup> Metformin is the first-line drug of choice for the treatment of T2DM, particularly in overweight and obese patients and those with normal kidney function.<sup>2</sup> Agonists of the peroxisome proliferator-activated nuclear receptor (PPAR), thiazolidinediones, are able to reduce insulin resistance but are under intense scrutiny because of concerns with their safety. In fact, the use of rosiglitazone has now been severely restricted in the US and has been completely suspended in Europe as a result of concerns regarding its cardiovascular safety.<sup>3</sup> Notably, insulin, which is used to treat T1DM patients (for whom the hormone is no longer produced internally), is also occasionally

used for patients with T2DM when other medications fail to adequately control blood glucose levels. However, hypoglycemia and weight gain are common side effects. Thus, new approaches are needed to treat T2DM. One of the desirable approaches to achieve this goal would be to identify agents that promote/enhance glucose (nutrient)-dependent insulin secretion.<sup>5</sup>

Extensive research has been conducted on the molecular targets for T2DM, including PPAR $\gamma$ , protein tyrosine phosphatase-1B (PTP1B), DPP-IV, glycogen synthase kinase-3 (GSK-3), pyruvate dehydrogenase kinase (PDHK), cannabinoid receptors, fructose-bisphosphatases, and  $\beta$ 3-adrenoceptor ( $\beta$ 3-AR), in an attempt to develop newer antidiabetic agents.<sup>6-7,12-13</sup> These therapeutic targets are important, and most of them are suitable for the *in silico* analysis.<sup>8</sup>

Peroxisome proliferator-activated receptors (PPAR) are fatty acid-activated transcription factors that belong to the nuclear hormone receptor family<sup>1,2</sup>. Three PPAR isotypes, PPAR $\alpha$ , PPAR $\beta/\delta$  and PPAR $\gamma$ , have previously been identified. Each of these subtypes appears to be differentiated in a tissue-specific manner and plays a pivotal role in glucose and lipid homeostasis.<sup>3,4</sup> PPAR $\gamma$  constitutes a primary target for the development of drug candidates for the treatment of type II diabetes. Thiazolidinediones (TZDs) represent the first known PPAR $\gamma$  agonists used as oral antidiabetic agents.<sup>4,5</sup> In addition, several studies have suggested that oral PPAR $\gamma$  full agonists not only exert an antidiabetic effect but also may act as a promising therapeutic target for a broad variety of skin disorders, including inflammatory skin diseases, such as psoriasis and atopic dermatitis, melanoma and other skin malignancies<sup>6-9</sup>. Furthermore, PPAR $\gamma$  full agonists may even induce cell growth arrest, apoptosis and terminal differentiation in various human malignant tumors.<sup>7,11</sup>

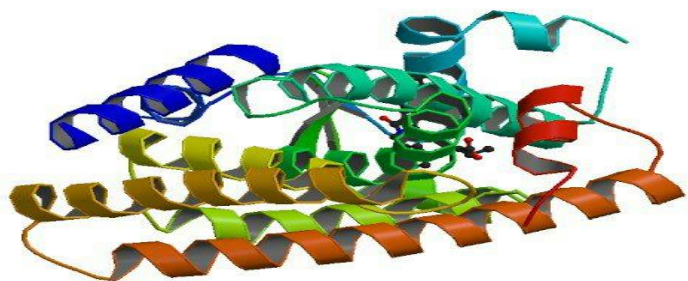


Fig 1: PPAR $\gamma$  receptor protein target; PDB code: 3VI8

## 1.2 Computer-aided drug design

Great advances have been made on Computer-aided drug design (CADD) methodologies and these have contributed significantly to the discovery and/or optimization of many clinically used drugs in recent years<sup>15</sup>. Drug discovery and development is a time-consuming and expensive process. On average, it takes 10–15 years and \$500–800 million to introduce a drug into the market<sup>16,17</sup>. Accordingly, CADD approaches have been widely used in the pharmaceutical industry to accelerate the process of new drug development<sup>18,19</sup>. CADD helps scientists focus on the most promising compounds so that they can minimize the synthetic and biological testing efforts. In practice, the choice of employing CADD approaches is usually determined by the availability of experimentally determined 3D structures of the target proteins. Thus, there are two major types of drug design: ligand-based drug design and structure-based drug design. If protein structures are unknown, various methods of ligand-based drug design can be employed, such as quantitative structure activity relationship (QSAR) and pharmacophore analysis. If the target structures are known, structure-based approaches can be used, such as molecular docking, which employs the 3D structures of the targets to design novel active compounds with improved potency. As more structures are becoming available, the prediction accuracy will likely improve<sup>18</sup>.

### The advantages of CADD over the traditional /conventional drug screening

CADD is capable of increasing the 'hit' rate of novel drug compounds because it uses a much more targeted search than the traditional high throughput screening and combinatorial chemistry. It not only aims to explain the molecular basis of therapeutics activity but also predicts possible derivatives that would improve activity. In the overall drug discovery campaign, CADD can be used for three(3) major purposes: (i) filter large compound libraries into smaller sets of predicted active compounds that can be tested experimentally; (ii) guide the optimization of 'lead' candidates, whether to increase their affinity or optimize their drug metabolism, other pharmacokinetic properties and, the potential for toxicity; (iii) design novel compound, either by 'group' starting molecules; one functional group at a time or by piecing together fragments into novel chemotype<sup>127</sup>.

**TABLE 1: List of some clinically approved drug discovered through CADD approaches**

<b>Drug</b>	<b>Year of approval</b>	<b>Therapeutic action</b>
Captopril	1981	Antihypertensive
Saquinavir	1995	Human immunodeficiency Virus (HIV) inhibitor
Dorzolamide	1995	Carbonic anhydrase inhibitor
Indinavir	1996	Human immunodeficiency Virus (HIV) inhibitor
Ritonavir	1996	Human immunodeficiency Virus (HIV) inhibitor
Trofiban	1998	Fibrinogen antagonist
Zanamivir	1999	Neuraminidase inhibitor
Oseltamivir	1999	Active against influenza A and B viruses.
Raltegravir	2007	Human immunodeficiency Virus (HIV) inhibitor
Aliskiren	2007	Human renin inhibitor
TMI-005	Phase II clinical trials	In Rheumatoid arthritis
LY-517717	Phase II clinical trials	Serine protease Inhibitor
Boceprevir	Phase III clinical trials	Hepatitis C virus (HCV) inhibitor
Nolatrexed	Phase III clinical trials	In Liver cancer
NVP- AUY922	Phase I clinical trials	Inhibitor for HSP90

SOURCE: (Neeema, B and Singh, B.K, 2017)

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## 2.1 METHODOLOGY

## 2.2 MATERIALS

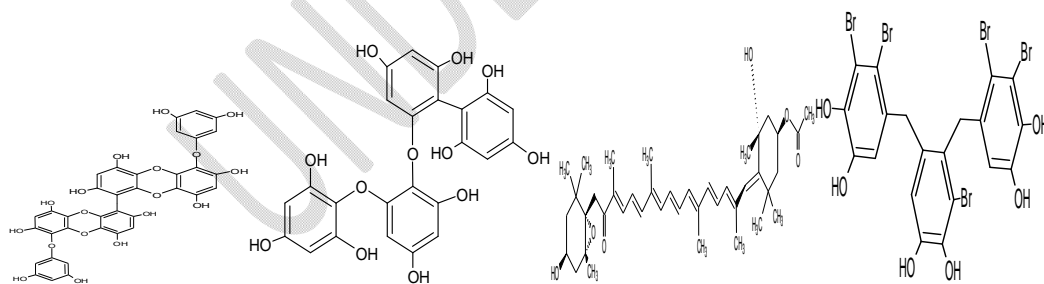
- Laptop
- Chems sketch
- Molecular Operating Environment (MOE)2014
- Discovery studio 2017
- Protox 11 (online software)
- SwissAdme (online software)
- Protein target; PPAR $\gamma$ , PDB ID:3VI8 (<http://www.rcsb.org/>)

## 2.3 Development of dataset

The 141 compounds used in the study structures were originally sourced from existing literatures wherein isolated compounds from marine and plants organisms with antidiabetic activity is established. Initial dataset was built using the chemsketch.

## 2.4 Generation of molecular structures and SMILES of compounds

The 2D structures of the 141 compounds and their SMILES notations were established using the chemsketch. The molecules were converted to pdb forms and saved after energy minimization and subsequent conversation to mdb formats and then stored to be used in the docking process.



Compound 6 (6,6-Bieckol)

compound 7 (Fucodiphloroethol-G)

compound 12 (Thunberol)

plant-derived c(OC)cc3-c(n4)sc(C)c4C

### 2.3 The molecular docking studies

The default parameters of MOE program were used for the molecular docking of the compounds. The target receptor (PDBCode: **3VI8**) used for the molecular docking studies was retrieved from the Protein Data Bank (PDB), (<https://www.wwpdb.org/>) database while the structure of the ligand, rosiglitazone was retrieved from PubChem database (<https://pubchem.ncbi.nlm.nih.gov>). MMFF94 force field was used for energy minimization of the ligand molecule. The prepared compounds were then subjected to interact with the receptor through molecular docking. The protocol facilitates flexible compound docking for various compound conformers within the rigid receptor. The Best conformation for each compound was chosen and the interaction was visualized in Discovery studio.

### 2.4 Study of physicochemical properties

The SwissAdme is an automated online tool used in the analysis of the physicochemical properties of chemical compounds. The SMILES of the entire compounds were uploaded into the software and run simultaneously. Results were extrapolated and analysed carefully and used in the prediction of the drug likeness of the selected compounds in tandem with other established parameters.

### 2.5 Toxicity studies using the protox11 software

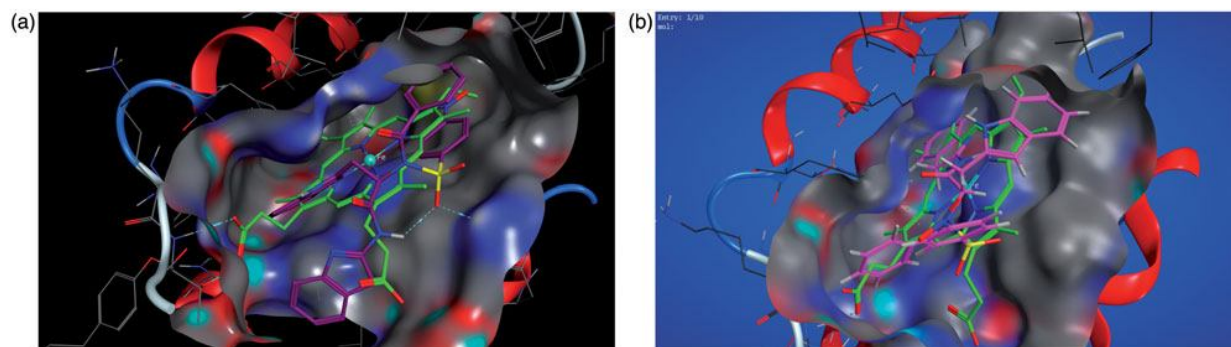
The toxicity profile of the compounds was evaluated using the protox11 (online software). The SMILES notation of each compound was copied into the software to run the procedure. The results of all the compounds were extrapolated, collated, tabulated and analysed.

## 4.0 RESULTS

**Table 2: virtual screening of the 141 compounds using PPAR $\gamma$  receptor target (3VI8)**

(Compound code)	E-score1 ( $\Delta G$ )	(Compound code)	E-score1 ( $\Delta G$ )
1	-11.5175	71	-13.6111
2	-18.7947	72	-15.0005
3	-15.3669	73	-12.8732
4	-16.9527	74	-13.7031
5	-17.9708	75	-14.5676
6	-16.8209	76	-13.2044
7	-19.0949	77	-14.3451
8	-9.81253	78	-12.5634
9	-11.5295	79	-12.7336
10	-12.7492	80	-14.4447
11	-11.4747	81	-13.4159
12	-18.1481	82	-13.8018
13	-9.86034	83	-13.1822
14	-8.10951	84	-15.8345
15	-10.2388	85	-12.56
16	-10.2502	86	-13.1504
17	-10.2502	87	-16.4061
18	-10.5263	88	-13.0626
19	-12.7871	89	-13.752
20	-13.5988	90	-12.6264
21	-10.8322	91	-14.9525
22	-9.55665	92	-14.0475
23	-13.7957	93	-11.6781
24	-14.5627	94	-14.7416
25	-12.461	95	-11.5576
26	-8.71318	96	-12.0841
27	-10.568	97	-12.1926
28	-10.6624	98	-14.6106
29	-16.3715	99	-12.3314
30	-15.939	100	-11.9029
31	-14.9434	101	-11.8649
32	-16.3715	102	-12.625
33	-14.5275	103	-12.625
34	-11.3469	104	-12.7995
35	-10.1483	105	-13.6341
36	-11.0534	106	-12.146
37	-11.2178	107	-13.2795
38	-11.0757	108	-12.4861
39	-16.0646	109	-13.5597
40	-14.9123	110	-12.8685
41	-9.22106	111	-12.9133
42	-11.0535	112	-13.6615
43	-11.9413	113	-12.6672
44	-13.5444	114	-12.7232
45	-13.5931	115	-13.0617
46	-9.40316	116	-12.8236
47	-10.5005	117	-12.863
48	-10.5448	118	-13.6777
49	-10.8395	119	-13.3214
50	-10.0082	120	-13.1005
51	-10.9029	121	-14.336
52	-11.0124	122	-14.5541
53	-12.6026	123	-13.3594
54	-12.0355	124	-13.9398
55	-12.812	125	-12.9934
56	-10.2006	126	-12.0891
57	-13.8667	127	-12.1504
58	-10.9227	128	-12.9668
59	-11.3084	129	-13.6258
60	-12.6284	130	-13.5446
61	-10.1857	131	-12.3831
62	-13.388	132	-13.4622
63	-12.1286	133	-12.9097
64	-12.0492	134	-11.7715
65	-14.1232	135	-13.0972
66	-16.8298	136	-13.243
67	-14.7508	137	-13.7865
68	-15.7552	138	-13.0769
69	-13.9737	139	-13.0291
70	-13.959	140	-12.8035
71	-13.6111	141	-19.5003
<b>Rosiglitazone</b>	-11.59		

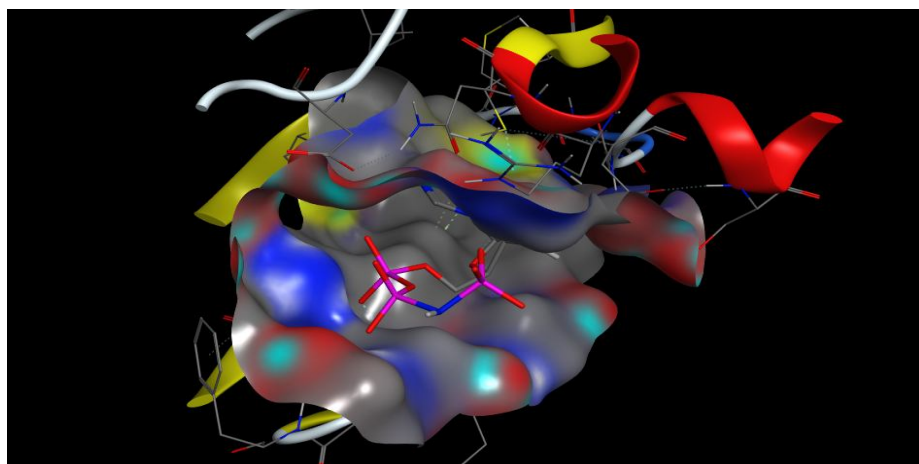
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**Figure 2.**(a, b) virtual screening protocol with protein target, 3VI8. The compound in green colour is the co-crystallized ligand of the target, the purple colour is the compounds fitting into the binding cavity of the target as the native ligand.

**Table 3: Rigid docking of 10 compounds with best predicted binding affinity with our target protein, 3VI8 PPAR $\gamma$  (lowest free binding energy, E scores).**

Compound code	E_score1 ( $\Delta G$ )	E_score2 ( $\Delta G$ )
6	-17.4864	-8.11895
7	-17.3851	-6.37149
12	-11.5589	-6.06639
32	-15.9291	-6.27781
66	-12.7987	-7.24062
80	-13.9822	-7.46954
89	-15.9846	-7.634
121	-15.5212	-7.17692
138	-13.5608	-7.04565
139	-13.165	-7.35982
rosiglitazone	-13.2378	-5.27409



**Fig3: rigid docking, ligand-receptor interaction model**

**Table 4: Toxicity profile of the 141 compounds**

z	LD50 (mg/kg) (predacc%)	Hepa (prob)	Carci (prob)	Imm (prob)	Muta (prob)	Cyto (prob)	AhR (prob)	AR (prob)	AR_LB D (prob)	aromas e (prob)	ER (prob)	ER_LB D (prob)	PPAR_Gamm a (prob)	nrf2/AR E (prob)	HSE (prob)	MMP (prob)	p53 (prob)	ATAD5 (prob)
1	5000 (70.97)	Inactiv e (0.88)	Inactiv e (0.61)	Active (0.89)	Inactiv e (0.93)	Inactiv e (0.95)	Inactiv e (1.0)	Inactiv e (0.93)	Inactiv e (0.95)	Inactive (0.99)	Inactiv e (0.92)	Inactiv e (0.95)	Inactive (1.0)	Inactive (0.57)	Inactiv e (0.57)	Inactiv e (0.56)	Inactiv e (0.97)	Inactiv e (0.99)
2	866 (69.26)	Inactiv e (0.78)	Inactiv e (0.64)	Inactiv e (0.94)	Inactiv e (0.55)	Inactiv e (0.91)	Active (0.76)	Inactiv e (0.97)	Inactiv e (0.90)	Inactive (0.67)	Inactiv e (0.50)	Inactiv e (0.62)	Inactive (0.96)	Inactive (0.86)	Inactiv e (0.86)	Active (0.64)	Inactiv e (0.84)	Inactiv e (0.85)
3	866 (69.26)	Inactiv e (0.78)	Inactiv e (0.64)	Inactiv e (0.93)	Inactiv e (0.55)	Inactiv e (0.91)	Active (0.76)	Inactiv e (0.97)	Inactiv e (0.90)	Inactive (0.67)	Inactiv e (0.50)	Inactiv e (0.62)	Inactive (0.96)	Inactive (0.86)	Inactiv e (0.86)	Active (0.64)	Inactiv e (0.84)	Inactiv e (0.85)
4	866 (69.26)	Inactiv e (0.78)	Inactiv e (0.64)	Inactiv e (0.93)	Inactiv e (0.55)	Inactiv e (0.91)	Active (0.76)	Inactiv e (0.97)	Inactiv e (0.90)	Inactive (0.67)	Inactiv e (0.50)	Inactiv e (0.62)	Inactive (0.96)	Inactive (0.86)	Active (0.86)	Inactiv e (0.64)	Inactiv e (0.84)	Inactiv e (0.85)
5	280	Inactiv	Inactiv	Active	Inactiv	Inactiv	Active	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Active	Inactiv	Inactiv	Inactiv

	(54.26)	e	e	(0.54)	e	e	(0.77)	e	e	(0.78)	e	e	(0.95)	(0.84)	(0.84)	e	e	e
		(0.75)	(0.50)	(0.53)	(0.91)	(0.77)	(0.96)	(0.92)	(0.78)	(0.50)	(0.77)	(0.95)	(0.84)	(0.84)	(0.66)	(0.77)	(0.78)	
6	10000	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Active	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Active	Inactiv	Inactiv	Inactiv
	(54.26)	e	e	e	e	e	(0.63)	e	e	(0.79)	e	e	(0.94)	(0.87)	(0.87)	e	e	e
		(0.82)	(0.70)	(0.97)	(0.53)	(0.82)	(0.94)	(0.92)	(0.57)	(0.70)	(0.94)	(0.92)	(0.93)	(0.64)	(0.85)	(0.58)	(0.78)	(0.92)
7	3600	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Active	Inactiv	Inactiv	Inactive	Active	Active	Inactive	Inactive	Inactiv	Active	Inactiv	Inactiv
	(67.38)	e	e	e	e	e	(0.68)	e	e	(0.66)	(0.77)	(0.70)	(0.93)	(0.64)	(0.85)	(0.58)	(0.78)	(0.92)
		(0.64)	(0.69)	(0.97)	(0.76)	(0.95)	(0.81)	(0.99)	(0.81)	(0.99)	(0.81)	(0.99)	(0.93)	(0.64)	(0.85)	(0.55)	(0.67)	
8	3474	Inactiv	Active	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Inactiv	Inactiv	Inactiv
	(100)	e	(0.67)	e	e	e	e	e	e	(0.99)	e	e	(0.99)	(0.95)	e	e	e	e
		(0.67)	(0.88)	(0.89)	(0.82)	(0.99)	(0.99)	(1.00)	(0.75)	(0.97)	(0.75)	(0.97)	(0.99)	(0.95)	(0.95)	(0.99)	(0.99)	(0.99)
9	1640	Inactiv	Inactiv	Active	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Inactiv	Inactiv	Inactiv
	(70.97)	e	e	(0.78)	e	e	e	e	e	(0.99)	e	e	(0.98)	(0.89)	e	e	e	e
		(0.77)	(0.65)	(0.84)	(0.85)	(0.91)	(0.99)	(0.99)	(0.88)	(0.94)	(0.88)	(0.94)	(0.98)	(0.89)	(0.95)	(0.95)	(0.99)	(0.99)
10	1680	Inactiv	Inactiv	Active	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Inactiv	Inactiv	Inactiv
	(68.07)	e	e	(0.55)	e	e	e	e	e	(0.97)	e	e	(0.91)	(0.82)	e	e	e	e
		(0.64)	(0.65)	(0.74)	(0.91)	(0.82)	(0.99)	(0.99)	(0.80)	(0.91)	(0.80)	(0.91)	(0.91)	(.82)	(0.81)	(0.90)	(0.98)	
11	219	Inactiv	Active	Active	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Inactiv	Inactiv	Inactiv
	(69.26)	e	(0.50)	(0.99)	e	e	e	e	e	(0.78)	e	e	(0.95)	(0.74)	e	e	e	e
		(0.81)	(0.50)	(0.70)	(0.98)	(0.91)	(0.92)	(0.84)	(0.95)	(0.84)	(0.95)	(0.95)	(0.74)	(0.64)	(0.72)	(0.91)		
12	1860	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Inactiv	Inactiv	Inactiv
	(70.97)	e	e	e	e	e	e	e	e	(0.89)	e	e	(0.97)	(0.68)	e	e	e	e
		(0.75)	(0.62)	(0.69)	(0.86)	(0.85)	(0.92)	(0.83)	(0.78)	(0.68)	(0.80)	(0.80)	(0.97)	(0.68)	(0.63)	(0.95)	(0.96)	
13	866	Inactiv	Active	Inactiv	Inactiv	Inactiv	Active	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Active	Inactiv	Inactiv
	(69.26)	e	(0.54)	e	e	e	(0.66)	e	e	(0.90)	e	e	(0.97)	(0.85)	e	(0.59)	e	e
		(0.74)	(0.59)	(0.55)	(0.88)	(0.98)	(0.86)	(0.59)	(0.63)	(0.85)	(0.84)	(0.89)	(0.85)	(0.84)	(0.89)			
14	5530	Inactiv	Inactiv	Active	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Inactiv	Inactiv	Inactiv
	(54.26)	e	e	(0.99)	e	e	e	e	e	(0.81)	e	e	(0.87)	(0.84)	e	e	e	e
		(0.66)	(0.58)	(0.62)	(0.78)	(0.79)	(0.95)	(0.93)	(0.77)	(0.88)	(0.77)	(0.88)	(0.87)	(0.84)	(0.84)	(0.50)	(0.76)	(0.91)
15	50	Inactiv	Active	Inactiv	Inactiv	Inactiv	Active	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Active	Inactiv	Inactiv
	(100)	e	(0.69)	e	e	e	(0.87)	e	e	(0.90)	e	e	(0.98)	(0.92)	e	(1.00)	e	e
		(0.68)	(0.94)	(0.99)	(0.68)	(0.99)	(0.99)	(0.87)	(0.97)	(0.87)	(0.97)	(0.97)	(0.92)	(0.92)	(0.92)	(0.51)	(1.0)	
16	50	Inactiv	Active	Inactiv	Inactiv	Inactiv	Active	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Active	Inactiv	Inactiv
	(100)	e	(0.69)	e	e	e	(0.87)	e	e	(0.90)	e	e	(0.98)	(0.92)	e	(1.00)	e	e
		(0.68)	(0.96)	(0.99)	(0.68)	(0.99)	(0.99)	(0.99)	(0.99)	(0.87)	(0.97)	(0.97)	(0.92)	(0.92)	(0.92)	(0.51)	(1.00)	
17	5000	Inactiv	Active	Inactiv	Inactiv	Inactiv	Active	Inactiv	Inactiv	Inactive	Active	Inactiv	Inactive	Inactive	Inactiv	Active	Inactiv	Inactiv
	(69.26)	e	(0.71)	e	e	e	(0.51)	e	e	(0.92)	(0.54)	e	(0.91)	(0.92)	e	(0.78)	e	e
		(0.60)	(0.97)	(0.90)	(0.76)	(0.99)	(0.99)	(0.99)	(0.92)	(0.54)	(0.96)	(0.96)	(0.91)	(0.92)	(0.92)	(0.96)	(0.98)	
18	1436	Inactiv	Active	Active	Active	Inactiv	Active	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Active	Inactiv	Inactiv
	(54.26)	e	(0.62)	(0.62)	(0.67)	e	(0.77)	e	e	(0.82)	e	e	(0.88)	(0.83)	e	(0.55)	e	e
		(0.51)	(0.66)	(0.93)	(0.95)	(0.57)	(0.82)	(0.57)	(0.82)	(0.88)	(0.83)	(0.83)	(0.83)	(0.83)	(0.79)	(0.90)		
19	1436	Inactiv	Active	Active	Active	Inactiv	Active	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Active	Inactiv	Inactiv
	(54.26)	e	(0.62)	(0.62)	(0.67)	e	(0.77)	e	e	(0.82)	e	e	(0.88)	(0.83)	e	(0.55)	e	e
		(0.51)	(0.66)	(0.93)	(0.95)	(0.57)	(0.82)	(0.57)	(0.82)	(0.88)	(0.83)	(0.83)	(0.83)	(0.83)	(0.79)	(0.90)		
20	1436	Inactiv	Active	Active	Active	Inactiv	Active	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Active	Inactiv	Inactiv
	(54.26)	e	(0.62)	(0.62)	(0.67)	e	(0.77)	e	e	(0.82)	e	e	(0.88)	(0.83)	e	(0.55)	e	e
		(0.51)	(0.66)	(0.93)	(0.95)	(0.57)	(0.82)	(0.57)	(0.82)	(0.88)	(0.83)	(0.83)	(0.83)	(0.83)	(0.79)	(0.90)		
21	5000	Inactiv	Active	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Active	Inactiv	Inactiv

	(70.97)	e (0.54)	(0.51) e (0.66)	e (0.76)	e (0.77)	e (0.56)	e (0.98)	e (0.98)	(0.96)	e (0.78)	e (0.79)	(0.94)	(0.86)	e (0.79)	(0.71) e (0.89)	e (0.94)		
22	220 (67.38)	Inactiv e (0.77)	Inactiv e (0.51)	Inactiv e (0.80)	Inactiv e (0.79)	Inactiv e (0.69)	Inactiv e (0.76)	Inactiv e (0.97)	Inactiv e (0.90)	Inactive (0.72)	Inactiv e (0.75)	Inactiv e (0.84)	Inactive (0.72)	Inactive (0.56)	Inactiv e (0.56)	Inactiv e (0.51)	Inactiv e (0.59)	Inactiv e (0.95)
23	1100 (68.67)	Inactiv e (0.54)	Inactiv e (0.50)	Inactiv e (0.82)	Inactiv e (0.77)	Inactiv e (0.66)	Inactiv e (0.72)	Inactiv e (0.97)	Inactiv e (0.94)	Inactive (0.88)	Inactiv e (0.84)	Inactiv e (0.83)	Inactive (0.94)	Inactive (0.90)	Inactiv e (0.90)	Inactiv e (0.71)	Inactiv e (0.85)	Inactiv e (0.98)
24	500 (67.38)	Inactiv e (0.64)	Inactiv e (0.52)	Active (0.78)	Inactiv e (0.69)	Inactiv e (0.74)	Inactiv e (0.89)	Inactiv e (0.96)	Inactiv e (0.96)	Inactive (0.94)	Inactiv e (0.94)	Inactiv e (0.98)	Inactive (0.98)	Inactive (0.98)	Inactiv e (0.98)	Inactiv e (0.88)	Inactiv e (0.97)	Inactiv e (0.96)
25	2000 23	Inactiv e 0.	Inactiv e 0.	Inactiv e 0.	Inactiv e 0.	Inactiv e 0.	Inactiv e 0.	Inactiv e 0.	Inactiv e 0.	Inactive 0.	Inactiv e 0.	Inactiv e 0.	Inactive 0.	Inactive 0.	Inactiv e 0.	Inactiv e 0.	Inactiv e 0.	Inactiv e 0.
26	500 23	Inactiv e 0.	Inactiv e 0.	Inactiv e 0.	Inactiv e 0.	Inactiv e 0.	Inactiv e 0.	Inactiv e 0.	Inactiv e 0.	Inactive 0.	Inactiv e 0.	Inactiv e 0.	Inactive 0.	Inactive 0.	Inactiv e 0.	Inactiv e 0.	Inactiv e 0.	Inactiv e 0.
27	500 23	Inactiv e 0.	Inactiv e 0.	Inactiv e 0.	Inactiv e 0.	Inactiv e 0.	Inactiv e 0.	Inactiv e 0.	Inactiv e 0.	Inactive 0.	Inactiv e 0.	Inactiv e 0.	Inactive 0.	Inactive 0.	Inactiv e 0.	Inactiv e 0.	Inactiv e 0.	Inactiv e 0.
28	500 (67.38)	Inactiv e (0.64)	Inactiv e (0.52)	Inactiv e (0.57)	Inactiv e (0.69)	Inactiv e (0.74)	Inactiv e (0.89)	Inactiv e (0.96)	Inactiv e (0.96)	Inactive (0.94)	Inactiv e (0.94)	Inactiv e (0.98)	Inactive (0.98)	Inactive (0.98)	Inactiv e (0.98)	Inactiv e (0.88)	Inactiv e (0.97)	Inactiv e (0.96)
29	1100 (68.07)	Inactiv e (0.84)	Active (0.50)	Inactiv e (0.96)	Inactiv e (0.77)	Inactiv e (0.66)	Inactiv e (0.72)	Inactiv e (0.97)	Inactiv e (0.94)	Inactive (0.88)	Inactiv e (0.84)	Inactiv e (0.83)	Inactive (0.94)	Inactive (0.90)	Inactiv e (0.90)	Inactiv e (0.71)	Inactiv e (0.85)	Inactiv e (0.98)
30	1700 (67.38)	Inactiv e (0.75)	Inactiv e (0.59)	Inactiv e (0.77)	Inactiv e (0.78)	Inactiv e (0.66)	Inactiv e (0.71)	Inactiv e (0.97)	Inactiv e (0.96)	Inactive (0.92)	Inactiv e (0.91)	Inactiv e (0.81)	Inactive (0.82)	Inactive (0.81)	Inactiv e (0.81)	Inactiv e (0.81)	Inactiv e (0.91)	Inactiv e (0.89)
31	2000 (68.07)	Inactiv e (0.71)	Active (0.54)	Inactiv e (0.92)	Inactiv e (0.78)	Inactiv e (0.64)	Inactiv e (0.70)	Inactiv e (0.97)	Inactiv e (0.79)	Inactive (0.58)	Inactiv e (0.68)	Inactiv e (0.58)	Active (0.57)	Active (0.64)	Active (0.64)	Active (0.84)	Active (0.63)	Inactiv e (0.96)
32	2000 (68.07)	Inactiv e (0.74)	Active (0.56)	Inactiv e (0.94)	Inactiv e (0.78)	Inactiv e (0.64)	Inactiv e (0.76)	Inactiv e (0.97)	Inactiv e (0.82)	Inactive (0.59)	Inactiv e (0.69)	Inactiv e (0.58)	Active (0.52)	Active (0.66)	Active (0.66)	Active (0.84)	Active (0.59)	Inactiv e (0.97)
33	1200 (67.38)	Inactiv e (0.77)	Inactiv e (0.60)	Inactiv e (0.85)	Inactiv e (0.82)	Inactiv e (0.670)	Inactiv e (0.85)	Inactiv e (0.97)	Inactiv e (0.96)	Inactive (0.70)	Inactiv e (0.83)	Inactiv e (0.75)	Inactive (0.64)	Active (0.52)	Active (0.52)	Active (0.61)	Inactiv e (0.62)	Inactiv e (0.97)
34	2000 (68.07)	Inactiv e (0.71)	Active (0.54)	Inactiv e (0.92)	Inactiv e (0.78)	Inactiv e (0.64)	Inactiv e (0.70)	Inactiv e (0.97)	Inactiv e (0.79)	Inactive (0.58)	Inactiv e (0.68)	Inactiv e (0.58)	Active (0.57)	Active (0.64)	Active (0.64)	Active (0.84)	Active (0.63)	Inactiv e (0.96)
35	1200 (67.38)	Inactiv e (0.78)	Inactiv e (0.55)	Inactiv e (0.86)	Inactiv e (0.78)	Inactiv e (0.70)	Inactiv e (0.84)	Inactiv e (0.97)	Inactiv e (0.94)	Inactive (0.77)	Inactiv e (0.84)	Inactiv e (0.85)	Inactive (0.73)	Inactive (0.59)	Inactiv e (0.59)	Inactiv e (0.55)	Inactiv e (0.69)	Inactiv e (0.97)
36	1380 (67.38)	Inactiv e (0.75)	Inactiv e (0.59)	Inactiv e (0.71)	Inactiv e (0.78)	Inactiv e (0.66)	Inactiv e (0.71)	Inactiv e (0.97)	Inactiv e (0.96)	Inactive (0.92)	Inactiv e (0.91)	Inactiv e (0.81)	Inactive (0.82)	Inactive (0.81)	Inactiv e (0.81)	Inactiv e (0.81)	Inactiv e (0.91)	Inactiv e (0.89)
37	2000 (68.07)	Active (0.55)	Inactiv e	Inactiv e	Inactiv e	Inactiv e	Inactiv e	Inactiv e	Inactiv e	Inactive (0.94)	Inactiv e	Inactiv e	Inactive (0.87)	Inactive (0.98)	Inactiv e	Inactiv e	Inactiv e	Inactiv e

38	860 (68.07)	Inactive (0.73)	Active (0.63)	Inactive (0.98)	Inactive (0.92)	Inactive (0.63)	Active (0.67)	Inactive (0.99)	Inactive (0.92)	Inactive (0.89)	Inactive (0.80)	Active (0.50)	Inactive (0.83)	Inactive (0.70)	Inactive (0.70)	Active (0.58)	Inactive (0.82)	Inactive (0.95)	Inactive (0.98)
39	2195 (67.38)	Inactive (0.58)	Active (0.65)	Inactive (0.98)	Inactive (0.85)	Inactive (0.63)	Inactive (0.62)	Inactive (0.95)	Inactive (0.95)	Inactive (0.88)	Inactive (0.63)	Inactive (0.71)	Inactive (0.83)	Inactive (0.82)	Inactive (0.82)	Inactive (0.67)	Inactive (0.85)	Inactive (0.97)	Inactive (0.95)
40	5000 (68.07)	Inactive (0.68)	Active (0.54)	Inactive (0.85)	Inactive (0.75)	Inactive (0.67)	Active (0.65)	Inactive (0.99)	Inactive (0.94)	Inactive (0.95)	Inactive (0.90)	Active (0.50)	Active (0.50)	Active (0.62)	Active (0.62)	Inactive (0.81)	Inactive (0.90)	Active (0.52)	Inactive (0.98)
41	2000 (68.07)	Inactive (0.71)	Active (0.54)	Inactive (0.99)	Inactive (0.78)	Inactive (0.64)	Inactive (0.70)	Inactive (0.97)	Inactive (0.79)	Inactive (0.58)	Inactive (0.68)	Inactive (0.58)	Active (0.57)	Active (0.64)	Active (0.64)	Active (0.84)	Active (0.63)	Active (0.96)	Inactive (0.98)
42	1380 (67.38)	Inactive -	Active -	Inactive -	Inactive -	Inactive -	Inactive -	Inactive -	Inactive -	Inactive -	Inactive -	Inactive -	Inactive -	Active -	Active -	Active -	Active -	Active -	Inactive -
42	1380 (67.38)	Inactive (0.71)	Active (0.53)	Inactive (0.99)	Inactive (0.77)	Inactive (0.66)	Inactive (0.73)	Inactive (0.97)	Inactive (0.82)	Inactive (0.57)	Inactive (0.66)	Inactive (0.59)	Inactive (0.55)	Active (0.57)	Active (0.57)	Active (0.82)	Active (0.50)	Inactive (0.93)	Inactive (0.98)
43	6430 (68.07)	Inactive (0.80)	Inactive (0.62)	Inactive (0.86)	Inactive (0.83)	Inactive (0.82)	Inactive (0.88)	Inactive (0.98)	Inactive (0.99)	Inactive (0.95)	Inactive (0.78)	Inactive (0.97)	Inactive (0.98)	Inactive (0.80)	Inactive (0.80)	Inactive (0.86)	Inactive (0.94)	Inactive (0.99)	Inactive (0.98)
44	860 (67.38)	Inactive (0.84)	Active (0.50)	Inactive (0.99)	Inactive (0.77)	Inactive (0.66)	Inactive (0.72)	Inactive (0.97)	Inactive (0.94)	Inactive (0.88)	Inactive (0.84)	Inactive (0.83)	Inactive (0.94)	Inactive (0.90)	Inactive (0.90)	Inactive (0.71)	Inactive (0.85)	Inactive (0.98)	Inactive (0.98)
45	1000 (68.07)	Inactive (0.74)	Inactive (0.57)	Active (0.97)	Inactive (0.55)	Inactive (0.52)	Inactive (0.59)	Inactive (0.95)	Inactive (0.98)	Inactive (0.83)	Inactive (0.89)	Inactive (0.97)	Inactive (0.99)	Inactive (0.97)	Inactive (0.97)	Inactive (0.73)	Inactive (0.90)	Inactive (0.94)	Inactive (0.98)
46	159 (67.38)	Inactive (0.68)	Inactive (0.64)	Active (0.99)	Inactive (0.60)	Inactive (0.82)	Inactive (0.64)	Inactive (0.95)	Inactive (0.98)	Inactive (0.62)	Active (0.53)	Inactive (0.68)	Inactive (0.81)	Inactive (0.78)	Inactive (0.78)	Active (0.81)	Inactive (0.60)	Inactive (0.72)	Inactive (0.98)
47	100 (54.26)	Inactive (0.71)	Inactive (0.62)	Inactive (0.87)	Inactive (0.71)	Inactive (0.77)	Inactive (0.82)	Inactive (0.96)	Inactive (0.99)	Inactive (0.77)	Inactive (0.74)	Inactive (0.88)	Inactive (0.90)	Inactive (0.76)	Inactive (0.76)	Active (0.57)	Inactive (0.75)	Inactive (0.94)	Inactive (0.98)
48	500 (69.26)	Inactive (0.65)	Active (0.51)	Active (0.93)	Inactive (0.54)	Inactive (0.88)	Inactive (0.67)	Inactive (0.99)	Inactive (0.98)	Inactive (0.68)	Inactive (0.70)	Inactive (0.90)	Inactive (0.98)	Inactive (0.87)	Inactive (0.87)	Inactive (0.57)	Inactive (0.74)	Inactive (0.70)	Inactive (0.98)
49	500 (69.26)	Inactive (0.70)	Inactive (0.62)	Active (0.99)	Inactive (0.55)	Inactive (0.82)	Inactive (0.70)	Inactive (0.98)	Inactive (0.95)	Inactive (0.66)	Inactive (0.81)	Inactive (0.91)	Inactive (0.95)	Inactive (0.88)	Inactive (0.88)	Inactive (0.57)	Inactive (0.81)	Inactive (0.83)	Inactive (0.98)
50	1000 (68.07)	Inactive (0.54)	Inactive (0.50)	Active (0.93)	Inactive (0.81)	Inactive (0.64)	Inactive (0.60)	Inactive (0.98)	Inactive (0.97)	Inactive (0.83)	Inactive (0.71)	Inactive (0.83)	Inactive (0.95)	Inactive (0.82)	Inactive (0.82)	Active (0.78)	Active (0.54)	Inactive (0.82)	Inactive (0.98)
51	570 (69.26)	Inactive (0.69)	Inactive (0.63)	Inactive (0.87)	Inactive (0.73)	Inactive (0.74)	Inactive (0.53)	Inactive (0.96)	Inactive (0.96)	Inactive (0.90)	Inactive (0.77)	Inactive (0.86)	Inactive (0.95)	Inactive (0.93)	Inactive (0.93)	Active (0.65)	Inactive (0.74)	Inactive (0.87)	Inactive (0.98)
52	2000 (68.07)	Inactive (0.67)	Inactive (0.56)	Active (0.96)	Inactive (0.55)	Inactive (0.84)	Inactive (0.58)	Inactive (0.96)	Inactive (0.95)	Inactive (0.89)	Inactive (0.63)	Inactive (0.84)	Inactive (0.95)	Inactive (0.84)	Inactive (0.84)	Inactive (0.52)	Inactive (0.82)	Inactive (0.77)	Inactive (0.98)

53	4000 (68.07)	Inactiv e (0.68)	Active (0.54)	Inactiv e (0.79)	Active (0.88)	Inactiv e (0.73)	Active (0.67)	Inactiv e (0.98)	Inactiv e (0.98)	Inactive (0.83)	Inactiv e (0.72)	Inactiv e (0.92)	Inactive (0.97)	Inactive (0.94)	Inactiv e (0.94)	Inactiv e (0.73)	Inactiv e (0.90)	Inactiv e (0.86)
54	500 (69.26)	Inactiv e (0.72)	Inactiv e (0.63)	Active (0.98)	Inactiv e (0.57)	Inactiv e (0.82)	Inactiv e (0.77)	Inactiv e (0.96)	Inactiv e (0.95)	Inactive (0.65)	Inactiv e (0.69)	Inactiv e (0.87)	Inactive (0.90)	Inactive (0.81)	Inactiv e (0.81)	Inactiv e (0.50)	Inactiv e (0.77)	Inactiv e (0.83)
55	5000 (70.97)	Inactiv e (0.69)	Inactiv e (0.57)	Active (0.65)	Inactiv e (0.70)	Inactiv e (0.99)	Inactiv e (0.86)	Inactiv e (0.99)	Inactiv e (0.97)	Inactive (0.65)	Active (0.58)	Active (0.54)	Inactive (0.97)	Inactive (0.93)	Inactiv e (0.93)	Active (0.81)	Inactiv e (0.82)	Active (0.69)
56	5000 (70.97)	Inactiv e (0.71)	Inactiv e (0.69)	Active (0.73)	Inactiv e (0.91)	Inactiv e (0.90)	Active (0.92)	Inactiv e (1.00)	Inactiv e (0.99)	Inactive (0.50)	Active (0.84)	Active (0.77)	Inactive (0.94)	Inactive (0.94)	Inactiv e (0.94)	Active (0.88)	Inactiv e (0.79)	Active (0.64)
57	1000 (69.26)	Inactiv e (0.66)	Inactiv e (0.60)	Active (0.99)	Inactiv e (0.74)	Inactiv e (0.89)	Inactiv e (0.87)	Inactiv e (0.97)	Inactiv e (0.98)	Inactive (0.84)	Inactiv e (0.55)	Inactiv e (0.65)	Inactive (0.82)	Inactive (0.55)	Inactiv e (0.55)	Active (0.68)	Inactiv e (0.63)	Inactiv e (0.93)
58	3000 (68.07)	Inactiv e (0.70)	Inactiv e (0.55)	Active (0.62)	Inactiv e (0.75)	Inactiv e (0.72)	Active (0.65)	Inactiv e (0.98)	Inactiv e (1.0)	Inactive (0.84)	Active (0.58)	Inactiv e (0.76)	Inactive (0.97)	Inactive (0.71)	Inactiv e (0.71)	Active (0.70)	Inactiv e (0.88)	Inactiv e (0.78)
59	1000 (68.07)	Inactiv e (0.67)	Inactiv e (0.62)	Active (0.92)	Inactiv e (0.65)	Inactiv e (0.76)	Inactiv e (0.76)	Inactiv e (0.97)	Inactiv e (0.94)	Inactive (0.82)	Inactiv e (0.63)	Inactiv e (0.83)	Inactive (0.89)	Inactive (0.68)	Inactiv e (0.68)	Active (0.83)	Active (0.53)	Inactiv e (0.92)
60	2000 (70.97)	Inactiv e (0.70)	Inactiv e (0.56)	Active (0.92)	Inactiv e (0.82)	Inactiv e (0.73)	Inactiv e (0.70)	Inactiv e (0.91)	Inactiv e (0.88)	Inactive (0.82)	Inactiv e (0.70)	Inactiv e (0.77)	Inactive (0.93)	Inactive (0.83)	Inactiv e (0.83)	Active (0.65)	Inactiv e (0.69)	Inactiv e (0.89)
61	2000 (70.97)	Inactiv e (0.57)	Inactiv e (0.55)	Active (0.76)	Inactiv e (0.74)	Inactiv e (0.68)	Inactiv e (0.79)	Inactiv e (0.95)	Inactiv e (0.91)	Inactive (0.80)	Inactiv e (0.74)	Inactiv e (0.86)	Inactive (0.93)	Inactive 0.80	Inactiv e (0.80)	Active (0.70)	Inactiv e (0.64)	Inactiv e (0.84)
62	3919 (70.97)	Inactiv e (0.70)	Inactiv e (0.69)	Active (0.94)	Inactiv e (0.82)	Inactiv e (0.75)	Active (0.84)	Inactiv e (0.99)	Inactiv e (0.99)	Inactive (0.54)	Active (0.69)	Active (0.60)	Inactive (0.96)	Inactive (0.93)	Inactiv e (0.93)	Active (0.86)	Inactiv e (0.73)	Active (0.62)
63	100 (68.07)	Inactiv e (0.74)	Active (0.50)	Active (0.51)	Inactiv e (0.75)	Inactiv e (0.83)	Inactiv e (0.94)	Active (0.55)	Active (0.58)	Inactive (0.81)	Active (0.50)	Inactiv e (0.64)	Inactive (0.99)	Inactive (0.59)	Inactiv e (0.59)	Inactiv e (0.61)	Inactiv e (0.91)	Inactiv e (0.98)
64	5000 (67.38)	Inactiv e (0.68)	Inactiv e (0.61)	Inactiv e (0.91)	Inactiv e (0.64)	Inactiv e (0.64)	Inactiv e (0.78)	Inactiv e (0.96)	Inactiv e (0.91)	Inactive (0.90)	Inactiv e (0.73)	Inactiv e (0.87)	Inactive (0.85)	Inactive (0.79)	Inactiv e (0.79)	Inactiv e (0.67)	Inactiv e (0.74)	Inactiv e (0.93)
65	552 (67.38)	Inactiv e (0.66)	Inactiv e (0.64)	Inactiv e (0.98)	Inactiv e (0.67)	Inactiv e (0.70)	Inactiv e (0.79)	Inactiv e (0.96)	Inactiv e (0.94)	Inactive (0.91)	Inactiv e (0.83)	Inactiv e (0.96)	Inactive (0.94)	Inactive (0.88)	Inactiv e (0.88)	Inactiv e (0.76)	Inactiv e (0.84)	Inactiv e (0.92)
66	1700 (67.38)	Inactiv e (0.70)	Inactiv e (0.64)	Inactiv e (0.99)	Inactiv e (0.62)	Inactiv e (0.63)	Inactiv e (0.79)	Inactiv e (0.96)	Inactiv e (0.93)	Inactive (0.88)	Inactiv e (0.77)	Inactiv e (0.90)	Inactive (0.88)	Inactive (0.79)	Inactiv e (0.79)	Inactiv e (0.69)	Inactiv e (0.71)	Inactiv e (0.93)
67	1000 (67.38)	Inactiv e (0.50)	Inactiv e (0.58)	Inactiv e (0.67)	Inactiv e (0.65)	Inactiv e (0.64)	Inactiv e (0.68)	Inactiv e (0.88)	Inactiv e (0.97)	Inactive (0.86)	Inactiv e (0.80)	Inactiv e (0.91)	Active (0.50)	Inactive (0.93)	Inactiv e (0.93)	Inactiv e (0.62)	Inactiv e (0.85)	Inactiv e (0.83)
68	800 (54.26)	Inactiv e (0.60)	Inactiv e (0.57)	Inactiv e (0.80)	Inactiv e (0.65)	Inactiv e (0.68)	Inactiv e (0.77)	Inactiv e (0.94)	Inactiv e (0.99)	Inactive (0.87)	Inactiv e (0.83)	Inactiv e (0.94)	Inactive (0.60)	Inactive (0.93)	Inactiv e (0.93)	Inactiv e (0.73)	Inactiv e (0.85)	Inactiv e (0.87)
69	5000 (69.26)	Inactiv e	Inactiv e	Inactiv e	Inactiv e	Inactiv e	Inactiv e	Inactiv e	Inactiv e	Inactive (0.90)	Inactiv e	Inactiv e	Inactive (0.90)	Inactive (0.93)	Inactiv e	Inactiv e	Inactiv e	Inactiv e

70	800 (54.26)	(0.72) Inactiv e	(0.66) Inactiv e	(0.99) Inactiv e	(0.65) Inactiv e	(0.74) Inactiv e	(0.69) Inactiv e	(0.98) Inactiv e	(0.99) Inactiv e	Inactive (0.88)	(0.65) Inactiv e	(0.91) Inactiv e	Inactive (0.90)	Inactive (0.92)	(0.93) Inactiv e	(0.70) Inactiv e	(0.86) Inactiv e	(0.82) Inactiv e
71	1500 (54.26)	(0.72) Inactiv e	(0.53) Inactiv e	(0.96) Active (0.52)	(0.67) Inactiv e	(0.61) Inactiv e	(0.79) Inactiv e	(0.97) Inactiv e	(0.99) Inactiv e	Inactive (0.86)	(0.90) Inactiv e	(0.90) Inactiv e	Inactive (0.72)	Inactive (0.93)	(0.92) Inactiv e	(0.77) Inactiv e	(0.92) Inactiv e	(0.96) Inactiv e
72	1500 (54.26)	(0.61) Inactiv e	(0.56) Inactiv e	(0.52) Active (0.52)	(0.68) Inactiv e	(0.63) Inactiv e	(0.76) Inactiv e	(0.96) Inactiv e	(0.98) Inactiv e	Inactive (0.86)	(0.88) Inactiv e	(0.95) Inactiv e	Inactive (0.72)	Inactive (0.93)	(0.93) Inactiv e	(0.78) Inactiv e	(0.87) Inactiv e	(0.89) Inactiv e
73	1000 (54.26)	(0.61) Inactiv e	(0.56) Active (0.50)	(0.52) Inactiv e	(0.68) Inactiv e	(0.63) Inactiv e	(0.76) Inactiv e	(0.96) Inactiv e	(0.98) Inactiv e	Inactive (0.88)	(0.88) Inactiv e	(0.95) Inactiv e	Inactive (0.74)	Inactive (0.94)	(0.93) Inactiv e	(0.78) Inactiv e	(0.87) Inactiv e	(0.89) Inactiv e
74	1000 (67.38)	(0.51) Inactiv e	(0.85) Inactiv e	(0.54) Inactiv e	(0.65) Inactiv e	(0.77) Inactiv e	(0.94) Inactiv e	(0.98) Inactiv e	(0.98) Inactiv e	Inactive (0.91)	(0.87) Inactiv e	(0.94) Inactiv e	Inactive (0.82)	Inactive (0.94)	(0.94) Inactiv e	(0.72) Inactiv e	(0.91) Inactiv e	(0.95) Inactiv e
75	1000 (67.38)	(0.61) Active (0.51)	(0.57) Inactiv e	(0.98) Inactiv e	(0.65) Inactiv e	(0.62) Inactiv e	(0.53) Inactiv e	(0.86) Inactiv e	(0.99) Inactiv e	Inactive (0.87)	(0.67) Inactiv e	(0.90) Inactiv e	Inactive (0.68)	Inactive (0.94)	(0.94) Inactiv e	(0.60) Inactiv e	(0.90) Inactiv e	(0.88) Inactiv e
76	1000 (54.26)	(0.53) Inactiv e	(0.52) Active (0.50)	(0.81) Inactiv e	(0.58) Inactiv e	(0.68) Inactiv e	(0.75) Inactiv e	(0.94) Inactiv e	(0.98) Inactiv e	Inactive (0.89)	(0.87) Inactiv e	(0.95) Inactiv e	Inactive (0.80)	Inactive (0.94)	(0.94) Inactiv e	(0.68) Inactiv e	(0.88) Inactiv e	(0.92) Inactiv e
77	3700 (54.26)	(0.53) Active (0.56)	(0.74) Inactiv e	(0.56) Inactiv e	(0.73) Inactiv e	(0.77) Inactiv e	(0.96) Inactiv e	(0.99) Inactiv e	(0.99) Inactiv e	Inactive (0.86)	(0.87) Inactiv e	(0.95) Inactiv e	Inactive (0.58)	Inactive (0.87)	(0.94) Inactiv e	(0.74) Inactiv e	(0.91) Inactiv e	(0.96) Inactiv e
78	3700 (54.26)	(0.53) Inactiv e	(0.55) Inactiv e	(0.87) Inactiv e	(0.69) Inactiv e	(0.68) Inactiv e	(0.79) Inactiv e	(0.96) Inactiv e	(0.98) Inactiv e	Inactive (0.89)	(0.83) Inactiv e	(0.95) Inactiv e	Inactive (0.54)	Inactive (0.88)	(0.87) Inactiv e	(0.71) Inactiv e	(0.88) Inactiv e	(0.92) Inactiv e
79	3700 (54.26)	(0.53) Active (0.50)	(0.54) Inactiv e	(0.86) Inactiv e	(0.63) Inactiv e	(0.66) Inactiv e	(0.79) Inactiv e	(0.94) Inactiv e	(0.98) Inactiv e	Inactive (0.89)	(0.88) Inactiv e	(0.97) Inactiv e	Active (0.52)	Inactive (0.91)	(0.88) Inactiv e	(0.72) Inactiv e	(0.84) Inactiv e	(0.88) Inactiv e
80	6000 (54.26)	(0.59) Active (0.50)	(0.89) Inactiv e	(0.61) Inactiv e	(0.64) Inactiv e	(0.77) Inactiv e	(0.92) Inactiv e	(0.98) Inactiv e	(0.98) Inactiv e	Inactive (0.89)	(0.84) Inactiv e	(0.96) Inactiv e	Active (0.52)	Inactive (0.91)	(0.91) Inactiv e	(0.67) Inactiv e	(0.84) Inactiv e	(0.87) Inactiv e
81	475 (67.38)	(0.72) Inactiv e	(0.55) Inactiv e	(0.58) Inactiv e	(0.69) Inactiv e	(0.70) Inactiv e	(0.86) Inactiv e	(0.90) Inactiv e	(0.95) Inactiv e	Inactive (0.89)	(0.75) Inactiv e	(0.86) Inactiv e	Inactive (0.88)	Inactive (0.92)	(0.92) Inactiv e	(0.62) Inactiv e	(0.91) Inactiv e	(0.90) Inactiv e
82	100 (54.26)	(0.52) Active (0.52)	(0.52) Inactiv e	(0.52) Inactiv e	(0.60) Active (0.51)	(0.69) Inactiv e	(0.69) Inactiv e	(0.90) Inactiv e	(0.97) Inactiv e	Inactive (0.88)	(0.80) Inactiv e	(0.91) Inactiv e	Inactive (0.72)	Inactive (0.91)	(0.91) Inactiv e	(0.62) Inactiv e	(0.83) Inactiv e	(0.86) Inactiv e
83	2000 (54.26)	(0.62) Inactiv e	(0.55) Inactiv e	(0.68) Inactiv e	(0.65) Inactiv e	(0.59) Inactiv e	(0.51) Inactiv e	(0.88) Inactiv e	(0.96) Inactiv e	Inactive (0.86)	(0.86) Inactiv e	(0.94) Inactiv e	Inactive (0.86)	Inactive (0.90)	(0.90) Inactiv e	(0.64) Inactiv e	(0.82) Inactiv e	(0.88) Inactiv e
84	3700 (54.26)	(0.51) Inactiv e	(0.52) Inactiv e	(0.94) Inactiv e	(0.65) Inactiv e	(0.69) Inactiv e	(0.80) Inactiv e	(0.94) Inactiv e	(0.98) Inactiv e	Inactive (0.90)	(0.88) Inactiv e	(0.97) Inactiv e	Active (0.56)	Inactive (0.88)	(0.88) Inactiv e	(0.75) Inactiv e	(0.85) Inactiv e	(0.89) Inactiv e
85	100 (54.26)	(0.51) Inactiv e	(0.57) Inactiv e	(0.64) Inactiv e	(0.64) Inactiv e	(0.63) Inactiv e	(0.69) Inactiv e	(0.89) Inactiv e	(0.97) Inactiv e	Inactive (0.87)	(0.83) Inactiv e	(0.91) Inactiv e	Inactive (0.50)	Inactive (0.92)	(0.92) Inactiv e	(0.64) Inactiv e	(0.85) Inactiv e	(0.85) Inactiv e
86	756	(0.51) Inactiv	(0.57) Active	(0.64) Active	(0.64) Inactiv	(0.63) Inactiv	(0.69) Inactiv	(0.89) Inactiv	(0.97) Inactiv	Inactive	(0.83) Inactiv	(0.91) Inactiv	Inactive	Inactive	(0.92) Inactiv	(0.64) Inactiv	(0.85) Inactiv	(0.85) Inactiv

	(67.38)	e	(0.52)	(0.72)	e	e	e	e	e	(0.91)	e	e	(0.86)	(0.96)	e	e	e	e
		(0.73)			(0.74)	(0.62)	(0.78)	(0.90)	(0.98)		(0.92)	(0.96)			(0.96)	(0.72)	(0.91)	(0.96)
87	250	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Inactiv	Inactiv	Inactiv
	(54.26)	e	e	e	e	e	e	e	e	(0.89)	e	e	(0.76)	(0.92)	e	e	e	e
		(0.60)	(0.56)	(0.56)	(0.65)	(0.62)	(0.72)	(0.92)	(0.96)		(0.79)	(0.91)			(0.92)	(0.67)	(0.86)	(0.88)
88	250	Inactiv	Active	Active	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Inactiv	Inactiv	Inactiv
	(54.26)	e	(0.50)	(0.93)	e	e	e	e	e	(0.90)	e	e	(0.85)	(0.94)	e	e	e	e
		(0.71)			(0.60)	(0.61)	(0.79)	(0.94)	(0.95)		(0.82)	(0.91)			(0.94)	(0.73)	(0.87)	(0.96)
89	3700	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Active	Inactive	Inactiv	Inactiv	Inactiv	Inactiv
	(54.26)	e	e	e	e	e	e	e	e	(0.88)	e	e	(0.62)	(0.89)	e	e	e	e
		(0.51)	(0.54)	(0.95)	(0.66)	(0.69)	(0.80)	(0.95)	(0.98)		(0.87)	(0.97)			(0.89)	(0.73)	(0.85)	(0.88)
90	250	Inactiv	Inactiv	Active	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Inactiv	Inactiv	Inactiv
	(54.26)	e	e	(0.87)	e	e	e	e	e	(0.89)	e	e	(0.79)	(0.92)	e	e	e	e
		(0.66)	(0.55)		(0.61)	(0.63)	(0.73)	(0.91)	(0.95)		(0.84)	(0.92)			(0.92)	(0.71)	(0.87)	(0.90)
91	800	Inactiv	Inactiv	Active	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Inactiv	Inactiv	Inactiv
	(54.26)	e	e	(0.69)	e	e	e	e	e	(0.86)	e	e	(0.88)	(0.91)	e	e	e	e
		(0.60)	(0.54)		(0.67)	(0.68)	(0.77)	(0.97)	(0.98)		(0.87)	(0.93)			(0.91)	(0.76)	(0.91)	(0.96)
92	250	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Inactiv	Inactiv	Inactiv
	(54.26)	e	e	e	e	e	e	e	e	(0.84)	e	e	(0.73)	(0.90)	e	e	e	e
		(0.65)	(0.54)	(0.90)	(0.62)	(0.61)	(0.77)	(0.91)	(0.94)		(0.83)	(0.87)			(0.90)	(0.63)	(0.88)	(0.96)
93	250	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Inactiv	Inactiv	Inactiv
	(67.38)	e	e	e	e	e	e	e	e	(0.85)	e	e	(0.82)	(0.93)	e	e	e	e
		(0.72)	(0.53)	(0.91)	(0.59)	(0.60)	(0.82)	(0.95)	(0.93)		(0.83)	(0.87)			(0.93)	(0.75)	(0.86)	(0.96)
94	250	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Inactiv	Inactiv	Inactiv
	(67.38)	e	e	e	e	e	e	e	e	(0.87)	e	e	(0.80)	(0.90)	e	e	e	e
		(0.72)	(0.54)	(0.50)	(0.59)	(0.62)	(0.83)	(0.95)	(0.93)		(0.83)	(0.86)			(0.90)	(0.73)	(0.87)	(0.95)
95	2000	Active	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Active	Inactive	Inactiv	Inactiv	Inactiv	Inactiv
	54.26	0.51	e	e	e	e	e	e	e	0.83	e	e	0.63	0.96	e	e	e	e
			0.56	0.99	0.65	0.68	0.82	0.95	0.99		0.84	0.97			0.96	0.57	0.83	0.91
96	1000	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Inactiv	Inactiv	Inactiv
	54.26	e	e	e	e	e	e	e	e	0.91	e	e	0.80	0.94	e	e	e	e
		0.67	0.59	0.91	0.59	0.67	0.80	0.97	0.97		0.85	0.96			0.94	0.73	0.82	0.94
97	2500	Inactiv	Inactiv	Active	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Inactiv	Inactiv	Inactiv
	68.07	e	e	0.71	e	e	e	e	e	0.94	e	e	0.88	0.73	e	e	e	e
		0.77	0.51		0.67	0.87	0.71	0.90	0.90		0.60	0.91			0.73	0.69	0.83	0.94
98	2500	Inactiv	Inactiv	Active	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Inactiv	Inactiv	Inactiv
	68.07	e	e	0.78	e	e	e	e	e	0.94	e	e	0.86	0.82	e	e	e	e
		0.76	0.54		0.68	0.87	0.72	0.89	0.91		0.61	0.89			0.82	0.72	0.85	0.94
99	2500	Inactiv	Inactiv	Active	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Inactiv	Inactiv	Inactiv
	68.07	e	e	0.85	e	e	e	e	e	0.87	e	e	0.90	0.82	e	e	e	e
		0.74	0.53		0.56	0.79	0.76	0.93	0.92		0.72	0.91			0.82	0.71	0.83	0.95
10	2500	Inactiv	Inactiv	Active	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Inactiv	Inactiv	Inactiv
	67.38	e	e	0.88	e	e	e	e	e	0.87	e	e	0.90	0.82	e	e	e	e
		0.74	0.53		0.56	0.79	0.76	0.93	0.92		0.72	0.91			0.82	0.71	0.83	0.95
10	2500	Inactiv	Inactiv	Active	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Inactiv	Inactiv	Inactiv
	68.07	e	e	0.91	e	e	e	e	e	0.94	e	e	0.89	0.81	e	e	e	e
		0.77	0.50		0.63	0.87	0.72	0.91	0.92		0.62	0.91			0.81	0.74	0.86	0.93
10	2500	Inactiv	Inactiv	Active	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Inactiv	Inactiv	Inactiv
	68.07	e	e	0.80	e	e	e	e	e	0.94	e	e	0.86	0.82	e	e	e	e



9		e	e	0.99	e	e	e	e	e	0.89	e	e	0.88	0.86	e	e	e	e
		0.74	0.55		0.68	0.77	0.76	0.93	0.97		0.84	0.95			0.86	0.75	0.88	0.92
12	2500	Inactiv	Inactiv	Active	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Inactiv	Inactiv	Inactiv
0	67.38	e	e	0.63	e	e	e	e	e	0.87	e	e	0.89	0.81	e	e	e	e
		0.71	0.50		0.69	0.74	0.74	0.93	0.96		0.86	0.95			0.81	0.75	0.87	0.93
12	5000	Inactiv	Inactiv	Active	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Inactiv	Inactiv	Inactiv
1	54.26	e	e	0.99	e	e	e	e	e	0.91	e	e	0.91	0.94	e	e	e	e
		0.64	0.55		0.61	0.79	0.81	0.89	0.96		0.92	0.97			0.94	0.71	0.87	0.92
12	200	Inactiv	Inactiv	Active	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Inactiv	Inactiv	Inactiv
2	67.38	e	e	0.80	e	e	e	e	e	0.88	e	e	0.84	0.83	e	e	e	e
		0.73	0.53		0.69	0.76	0.73	0.92	0.96		0.81	0.94			0.83	0.76	0.89	0.93
12	200	Inactiv	Inactiv	Active	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Inactiv	Inactiv	Inactiv
3	67.38	e	e	0.98	e	e	e	e	e	0.77	e	e	0.86	0.86	e	e	e	e
		0.70	0.54		0.64	0.70	0.75	0.96	0.97		0.83	0.94			0.86	0.74	0.85	0.93
12	1000	Inactiv	Inactiv	Active	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Inactiv	Inactiv	Inactiv
4	54.26	e	e	0.65	e	e	e	e	e	0.80	e	e	0.87	0.87	e	e	e	e
		0.60	0.96		0.75	0.68	0.92	0.95	0.80		0.94	0.69			0.	0.54	0.81	0.90
12	8000	Inactiv	Inactiv	Active	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Inactiv	Inactiv	Inactiv
5	67.38	e	e	0.77	e	e	e	e	e	0.84	e	e	0.74	0.91	e	e	e	e
		0.57	0.59		0.62	0.73	0.72	0.89	0.95		0.82	0.94			0.91	0.55	0.82	0.89
12	6200	Inactiv	Inactiv	Active	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Inactiv	Inactiv	Inactiv
6	67.38	e	e	0.76	e	e	e	e	e	0.86	e	e	0.75	0.89	e	e	e	e
		0.58	0.58		0.59	0.64	0.68	0.93	0.95		0.86	0.95			0.89	0.52	0.80	0.89
12	1000	Inactiv	Inactiv	Active	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Inactiv	Inactiv	Inactiv
7	54.26	e	e	0.75	e	e	e	e	e	0.86	e	e	0.77	0.87	e	e	e	e
		0.54	0.56		0.63	0.75	0.69	0.95	0.96		0.76	0.95			0.87	0.56	0.82	0.90
12	1000	Inactiv	Inactiv	Active	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Inactiv	Inactiv	Inactiv
8	54.26	e	e	0.96	e	e	e	e	e	0.86	e	e	0.69	0.90	e	e	e	e
		0.55	0.58		0.61	0.70	0.69	0.95	0.96		0.82	0.96			0.90	0.53	0.77	0.89
12	1000	Inactiv	Inactiv	Active	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Inactiv	Inactiv	Inactiv
9	54.26	e	e	0.87	e	e	e	e	e	0.84	e	e	0.74	0.88	e	e	e	e
		0.54	0.58		0.65	0.76	0.68	0.95	0.96		0.75	0.95			0.88	0.58	0.82	0.91
13	1000	Inactiv	Inactiv	Active	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Inactiv	Inactiv	Inactiv
0	54.26	e	e	0.99	e	e	e	e	e	0.83	e	e	0.67	0.91	e	e	e	e
		0.54	0.60		0.63	0.75	0.71	0.91	0.95		0.78	0.95			0.91	0.53	0.79	0.90
13	1000	Inactiv	Inactiv	Active	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Inactiv	Inactiv	Inactiv
1	54.26	e	e	0.99	e	e	e	e	e	0.81	e	e	0.66	0.89	e	e	e	e
		0.53	0.60		0.64	0.74	0.73	0.91	0.96		0.80	0.94			0.89	0.54	0.80	0.89
13	1000	Inactiv	Inactiv	Active	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Inactiv	Inactiv	Inactiv
2	54.26	e	e	0.99	e	e	e	e	e	0.83	e	e	0.68	0.89	e	e	e	e
		0.54	0.59		0.64	0.67	0.72	0.95	0.96		0.83	0.95			0.89	0.53	0.78	0.88
13	1000	Inactiv	Inactiv	Active	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Inactiv	Inactiv	Inactiv
3	54.26	e	e	0.95	e	e	e	e	e	0.84	e	e	0.73	0.86	e	e	e	e
		0.58	0.58		0.63	0.72	0.64	0.96	0.96		0.84	0.95			0.86	0.51	0.79	0.90
13	750	Inactiv	Inactiv	Active	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Inactiv	Inactiv	Inactiv
4	67.38	e	e	0.82	e	e	e	e	e	(0.84)	e	e	(0.70)	(0.86)	e	e	e	e
		0.59	0.59		0.63	0.72	0.64	0.95	(0.95)		(0.84)	(0.95)			(0.86)	(0.50)	(0.78)	(0.88)
13	1000	Inactiv	Inactiv	Active	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactive	Inactiv	Inactiv	Inactive	Inactive	Inactiv	Inactiv	Inactiv	Inactiv
5	(67.38)	e	e	(0.83)	e	e	e	e	e	(0.81)	e	e	(0.75)	(0.86)	e	e	e	e
		(0.57)	(0.58)		(0.66)	(0.70)	(0.67)	(0.95)	(0.95)		(0.80)	(0.94)			(0.86)	(0.60)	(0.82)	(0.92)

13	1000	Inactiv	Inactiv	Active	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	
6	(67.38)	e	e	(0.98)	e	e	e	e	e	e	(0.78)	e	e	(0.74)	(0.89)	e	e	e	
		(0.61)	(0.61)		(0.65)	(0.76)	(0.71)	(0.88)	(0.90)		(0.79)	(0.92)			(0.89)	(0.59)	(0.81)	(0.90)	
13	1000	Inactiv	Inactiv	Active	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	
7	(54.26)	e	e	(0.99)	e	e	e	e	e	e	(0.71)	e	e	(0.69)	(0.91)	e	e	e	
		(0.55)	(0.60)		(0.65)	(0.69)	(0.71)	(0.91)	(0.94)		(0.81)	(0.94)			(0.91)	(0.56)	(0.80)	(0.92)	
13	6200	Inactiv	Inactiv	Active	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Active	Inactiv	
8	(67.38)	e	e	(0.56)	e	e	e	e	e	e	(0.79)	e	e	(0.64)	(0.84)	e	(0.53)	e	
		(0.57)	(0.60)		(0.65)	(0.74)	(0.67)	(0.94)	(0.95)		(0.80)	(0.94)			(0.84)		(0.77)	(0.89)	
13	6200	Inactiv	Inactiv	Active	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	
9	(67.38)	e	e	(0.96)	e	e	e	e	e	e	(0.84)	e	e	(0.70)	(0.87)	e	e	e	
		(0.59)	(0.59)		(0.64)	(0.71)	(0.65)	(0.96)	(0.95)		(0.86)	(0.96)			(0.87)	(0.51)	(0.79)	(0.88)	
14	75	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	
0	(67.38)	e	e	e	e	e	e	e	e	e	(0.80)	e	e	(0.75)	(0.78)	e	e	e	
		(0.57)	(0.58)	(0.60)	(0.68)	(0.70)	(0.67)	(0.96)	(0.96)		(0.80)	(0.95)			(0.78)	(0.59)	(0.81)	(0.92)	
14	75	Inactiv	Inactiv	Active	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	Inactiv	
1	(67.38)	e	e	(0.97)	e	e	e	e	e	e	(0.80)	e	e	(0.70)	(0.87)	e	e	e	
		(0.57)	(0.59)		(0.66)	(0.70)	(0.69)	(0.93)	(0.96)		(0.81)	(0.93)			(0.87)	(0.87)	(0.57)	(0.80)	(0.91)

KEY: Z = parameter, Prob = probability, Hepa = hepatotoxicity, Carci = carcinogenicity, Imm = immunotoxicity, Muta = mutagenicity, Cyto = cytotoxicity, AhR = Aryl hydrocarbon Receptor, AR = Androgen Receptor, AR-LBD = Androgen Receptor Ligand, Binding Domain, ER = Estrogen Receptor Alpha, ER-LBD = Estrogen Receptor Ligand Binding Domain, PPAR-Gamma = Peroxisome Proliferator Activated Receptor Gamma, nrf2/ARE = Nuclear factor (erythroid-derived 2)-like 2/antioxidant responsive element (nrf2/ARE), HSE = Heat shock factor response element, MMP = Mitochondrial Membrane Potential, p53 = Phosphoprotein (Tumor Suppressor) p53, ATAD5 = ATPase family AAA domain-containing protein 5

**Table 5: Evaluation of pharmacokinetic and physicochemical properties (determination of Lipinski's rule of 5).**

Compound Code	MW	#Rotatable bonds	#H-bond acceptors	#H-bond donors	MR	TPSA	iLOGP	ESOL Class	Ali Class	Lipinski #violations
1	414.71	5	1	1	132.97	20.23	5.15	Poorly soluble	Insoluble	1
2	742.55	6	18	11	181.42	287.14	2.66	Poorly soluble	Insoluble	3
3	372.28	2	9	6	91.69	149.07	1.78	Moderately soluble	Moderately soluble	1
4	496.38	4	12	8	122.25	198.76	2.31	Moderately soluble	Poorly soluble	2
5	586.46	4	13	8	149.53	211.9	2.81	Poorly soluble	Poorly soluble	3

6	756.58	6	18	12	186.38	298.14	2.56	Poorly soluble	Insoluble	3
7	498.39	5	12	10	125.14	220.76	1.39	Moderately soluble	Poorly soluble	2
8	278.34	9	4	0	77.84	52.6	2.81	Moderately soluble	Moderately soluble	0
9	410.59	12	3	0	128.02	51.21	4.13	Moderately soluble	Poorly soluble	0
10	412.6	12	3	2	129.47	57.53	4.45	Moderately soluble	Poorly soluble	0
11	646.9	12	6	2	193.55	96.36	6.9	Poorly soluble	Insoluble	2
12	438.69	5	2	1	137.44	37.3	4.78	Poorly soluble	Insoluble	1
13	512.38	4	13	9	124.27	218.99	1.98	Moderately soluble	Poorly soluble	3
14	1000.82	14	24	17	244.06	408.52	2.68	Poorly soluble	Insoluble	3
15	330.8	0	1	1	51.57	20.23	2.41	Moderately soluble	Moderately soluble	0
16	251.9	0	1	1	43.87	20.23	2.17	Soluble	Soluble	0
17	513.84	2	1	0	93.69	9.23	4.08	Poorly soluble	Poorly soluble	2
18	551.69	2	0	1	92.25	12.03	3.62	Poorly soluble	Poorly soluble	2
19	551.69	2	0	1	92.25	12.03	3.62	Poorly soluble	Poorly soluble	2
20	551.69	2	0	1	92.25	12.03	3.56	Poorly soluble	Poorly soluble	2
21	684.79	6	5	0	117.43	46.15	4.46	Poorly soluble	Poorly soluble	2
22	577.84	4	5	4	100.68	90.15	2.87	Poorly soluble	Poorly soluble	1
23	297.93	1	3	3	52.02	60.69	1.86	Soluble	Soluble	0
24	340.01	4	3	0	65.68	27.69	3.04	Soluble	Soluble	0
25	563.82	1	5	4	94.64	90.15	0	Poorly soluble	Moderately soluble	1
26	607.87	3	6	3	106.41	88.38	0	Poorly soluble	Poorly soluble	1
27	649.95	6	6	0	119.81	55.38	0	Poorly soluble	Poorly soluble	1
28	261.11	4	3	0	57.98	27.69	2.87	Soluble	Soluble	0
29	219.03	1	3	3	44.32	60.69	1.65	Soluble	Very soluble	0
30	233.06	2	3	2	49.05	49.69	1.99	Soluble	Soluble	0
31	547.82	2	4	4	94.79	80.92	2.78	Poorly soluble	Poorly soluble	1
32	748.83	4	6	6	135.99	121.38	3.28	Poorly soluble	Poorly soluble	3
33	541.03	5	5	4	107.56	90.15	3.21	Poorly soluble	Poorly soluble	1
34	547.82	2	4	4	94.79	80.92	2.78	Poorly soluble	Poorly soluble	1
35	527	5	5	4	102.75	90.15	2.83	Moderately soluble	Poorly soluble	1
36	311.96	2	3	2	56.75	49.69	2.34	Soluble	Soluble	0
37	295.91	1	3	2	50.82	57.53	1.56	Soluble	Soluble	0

<b>38</b>	360.83	0	2	2	58.55	40.46	2.31	Moderately soluble	Moderately soluble	0
<b>39</b>	374.81	1	3	2	58.98	57.53	1.75	Moderately soluble	Soluble	0
<b>40</b>	376.82	1	3	2	60.08	49.69	2.31	Moderately soluble	Soluble	0
<b>41</b>	705.61	2	4	4	110.19	80.92	3.1	Poorly soluble	Poorly soluble	2
<b>42</b>	719.63	3	4	4	114.99	80.92	3.29	Poorly soluble	Poorly soluble	2
<b>43</b>	242.4	2	0	0	82.49	0	3.58	Moderately soluble	Moderately soluble	1
<b>44</b>	376.82	1	3	3	59.72	60.69	1.92	Soluble	Soluble	0
<b>45</b>	399.44	5	6	1	115.93	69.62	3.84	Moderately soluble	Moderately soluble	0
<b>46</b>	458.55	6	5	3	140.2	90.9	4.44	Poorly soluble	Poorly soluble	0
<b>47</b>	432.51	7	7	4	120.57	128.2	3.65	Moderately soluble	Moderately soluble	0
<b>48</b>	262.3	3	4	0	73.74	44.76	2.96	Soluble	Soluble	0
<b>49</b>	322.4	6	5	1	89.99	57.15	3.39	Soluble	Moderately soluble	0
<b>50</b>	296.36	1	3	1	87.93	54.37	3.11	Moderately soluble	Moderately soluble	0
<b>51</b>	316.43	5	3	1	94.27	54.37	3.04	Moderately soluble	Moderately soluble	0
<b>52</b>	298.42	3	2	0	93.91	18.46	3.89	Moderately soluble	Moderately soluble	0
<b>53</b>	316.31	4	6	0	85.96	67.13	2.94	Soluble	Soluble	0
<b>54</b>	354.44	6	4	1	103.78	47.92	3.98	Moderately soluble	Moderately soluble	0
<b>55</b>	328.32	4	6	1	89.42	78.13	2.87	Soluble	Moderately soluble	0
<b>56</b>	314.29	3	6	2	84.95	89.13	2.72	Moderately soluble	Moderately soluble	0
<b>57</b>	366.45	8	4	2	110.12	66.76	3.71	Moderately soluble	Poorly soluble	0
<b>58</b>	304.34	6	5	0	84.34	46.15	3.27	Soluble	Moderately soluble	0
<b>59</b>	476.6	6	5	4	144.5	97.99	4.69	Poorly soluble	Insoluble	0
<b>60</b>	298.38	1	3	1	87.16	54.37	3.06	Moderately soluble	Moderately soluble	0
<b>61</b>	312.36	1	4	1	87.59	71.44	2.44	Moderately soluble	Moderately soluble	0
<b>62</b>	360.31	4	8	3	93.46	118.59	2.66	Soluble	Moderately soluble	0
<b>63</b>	394.55	2	3	2	117.04	57.53	3.51	Moderately soluble	Moderately soluble	0
<b>64</b>	414.47	10	4	1	118.29	78.46	2.99	Moderately soluble	Poorly soluble	0
<b>65</b>	414.5	10	3	2	121.37	81.42	2.89	Moderately soluble	Poorly soluble	0
<b>66</b>	429.51	11	4	1	124.56	64.63	3.69	Poorly soluble	Poorly soluble	0
<b>67</b>	561.67	12	5	1	161.41	119.59	4.3	Poorly soluble	Insoluble	1
<b>68</b>	544.62	12	5	2	157.89	107.14	3.79	Poorly soluble	Poorly soluble	1
<b>69</b>	271.31	6	3	1	75.8	77	2.22	Very soluble	Very soluble	0

<b>70</b>	588.67	14	6	1	168.68	105.51	3.97	Poorly soluble	Poorly soluble	1
<b>71</b>	544.62	12	5	1	157.82	96.28	4.09	Poorly soluble	Poorly soluble	1
<b>72</b>	544.62	12	5	1	157.82	96.28	4.09	Poorly soluble	Poorly soluble	1
<b>73</b>	544.62	12	5	1	157.82	96.28	3.2	Poorly soluble	Poorly soluble	1
<b>74</b>	450.51	10	4	1	130.53	78.46	3.59	Poorly soluble	Poorly soluble	0
<b>75</b>	544.62	12	5	1	157.43	96.28	3.77	Poorly soluble	Poorly soluble	1
<b>76</b>	545.61	12	6	1	155.62	109.17	3.34	Poorly soluble	Poorly soluble	1
<b>77</b>	563.59	12	7	1	155.76	104.49	4.59	Poorly soluble	Poorly soluble	1
<b>78</b>	575.63	13	7	1	162.29	113.72	4.05	Poorly soluble	Insoluble	1
<b>79</b>	565.66	12	6	1	158.64	132.73	4.66	Poorly soluble	Insoluble	1
<b>80</b>	565.66	12	6	1	158.64	132.73	4.15	Poorly soluble	Insoluble	1
<b>81</b>	470.58	10	4	1	137.32	78.46	3.88	Poorly soluble	Poorly soluble	0
<b>82</b>	566.65	12	7	1	156.44	145.62	3.62	Poorly soluble	Insoluble	1
<b>83</b>	479.55	11	5	1	138.1	91.35	3.42	Poorly soluble	Poorly soluble	0
<b>84</b>	546.59	12	7	1	153.6	117.38	3.91	Poorly soluble	Poorly soluble	1
<b>85</b>	562.66	12	6	1	159.21	132.48	3.94	Poorly soluble	Poorly soluble	1
<b>86</b>	495.57	12	4	2	143.27	95.84	3.46	Poorly soluble	Poorly soluble	0
<b>87</b>	528.64	12	5	1	150.18	122.83	3.61	Poorly soluble	Poorly soluble	1
<b>88</b>	586.74	16	5	3	168.66	136.06	4.58	Poorly soluble	Poorly soluble	1
<b>89</b>	545.6	12	6	1	155.8	104.49	4.43	Poorly soluble	Poorly soluble	1
<b>90</b>	558.67	15	6	2	155.98	140.85	3.59	Poorly soluble	Poorly soluble	1
<b>91</b>	552.64	12	6	1	159.81	107.73	4.35	Poorly soluble	Poorly soluble	1
<b>92</b>	570.68	12	6	1	161.7	132.06	3.89	Poorly soluble	Poorly soluble	1
<b>93</b>	570.7	12	5	2	168.3	139.44	3.43	Moderately soluble	Poorly soluble	1
<b>94</b>	584.73	12	5	2	173.2	127.27	4.66	Poorly soluble	Poorly soluble	1
<b>95</b>	408.47	7	5	1	115.06	106.73	3.18	Moderately soluble	Poorly soluble	0
<b>96</b>	398.46	8	5	2	111.03	123.34	2.9	Moderately soluble	Moderately soluble	0
<b>97</b>	325.38	8	4	0	90.63	58.59	3.13	Moderately soluble	Moderately soluble	0
<b>98</b>	353.43	9	4	0	100.4	58.59	3.4	Moderately soluble	Moderately soluble	0
<b>99</b>	393.38	9	7	0	95.63	58.59	3.49	Moderately soluble	Moderately soluble	0
<b>100</b>	409.38	10	8	0	97.31	67.82	3.54	Moderately soluble	Poorly soluble	0
<b>101</b>	355.4	9	5	0	97.12	67.82	3.07	Moderately soluble	Moderately soluble	0

<b>102</b>	369.43	10	5	0	101.93	67.82	3.47	Moderately soluble	Moderately soluble	0
<b>103</b>	339.4	8	4	0	95.6	58.59	3.4	Moderately soluble	Moderately soluble	0
<b>104</b>	353.43	8	4	0	100.56	58.59	3.44	Moderately soluble	Moderately soluble	0
<b>105</b>	435.46	11	7	0	110.21	58.59	4.18	Poorly soluble	Poorly soluble	1
<b>106</b>	392.47	10	5	0	109.92	82.38	3.69	Moderately soluble	Poorly soluble	0
<b>107</b>	369.43	9	5	0	102.09	67.82	3.62	Moderately soluble	Moderately soluble	0
<b>108</b>	383.46	10	5	0	106.89	67.82	3.97	Moderately soluble	Moderately soluble	0
<b>109</b>	380.41	9	6	0	101.84	91.61	3.17	Moderately soluble	Moderately soluble	0
<b>110</b>	383.46	10	5	0	106.89	67.82	3.94	Moderately soluble	Moderately soluble	0
<b>111</b>	392.43	9	6	0	105.55	89.3	3.27	Moderately soluble	Moderately soluble	0
<b>112</b>	449.48	11	7	1	119.86	118.4	3.77	Soluble	Moderately soluble	0
<b>113</b>	426.87	9	6	0	110.56	89.3	2.96	Moderately soluble	Moderately soluble	0
<b>114</b>	491.5	10	9	0	119.5	112.61	4.03	Poorly soluble	Poorly soluble	0
<b>115</b>	407.5	9	4	0	113.94	86.83	3.89	Moderately soluble	Poorly soluble	0
<b>116</b>	391.44	9	5	0	108.33	71.73	3.66	Moderately soluble	Moderately soluble	0
<b>117</b>	440.51	9	4	1	127.92	74.38	3.68	Poorly soluble	Poorly soluble	0
<b>118</b>	402.46	9	5	0	113.86	71.48	3.54	Moderately soluble	Moderately soluble	0
<b>119</b>	432.49	10	6	0	120.35	80.71	3.84	Moderately soluble	Moderately soluble	0
<b>120</b>	403.45	9	6	0	111.66	84.37	3.37	Moderately soluble	Moderately soluble	0
<b>121</b>	463.5	11	8	0	124.64	102.83	4.24	Moderately soluble	Poorly soluble	0
<b>122</b>	416.49	9	5	0	118.83	71.48	3.98	Moderately soluble	Poorly soluble	0
<b>123</b>	470.46	10	8	0	118.86	71.48	3.82	Moderately soluble	Poorly soluble	0
<b>124</b>	450.57	11	5	0	126.32	99.72	3.6	Poorly soluble	Poorly soluble	0
<b>125</b>	507.6	12	7	0	136.3	139.85	3.51	Poorly soluble	Poorly soluble	1
<b>126</b>	495.54	11	8	0	128.21	149.08	3.55	Moderately soluble	Poorly soluble	0
<b>127</b>	438.52	10	6	0	118.23	108.95	4.14	Moderately soluble	Poorly soluble	0
<b>128</b>	468.54	11	7	0	124.72	118.18	4.35	Moderately soluble	Poorly soluble	0
<b>129</b>	452.54	11	6	0	123.04	108.95	4.18	Moderately soluble	Poorly soluble	0
<b>130</b>	494.62	13	6	0	137.62	108.95	4.78	Poorly soluble	Poorly soluble	0
<b>131</b>	508.65	13	6	0	142.42	108.95	4.96	Poorly soluble	Poorly soluble	1
<b>132</b>	496.6	12	7	0	134.34	118.18	4.57	Poorly soluble	Poorly soluble	0
<b>133</b>	438.52	10	6	0	118.23	108.95	3.92	Moderately soluble	Poorly soluble	0

134	452.54	10	6	0	123.2	108.95	4.22	Moderately soluble	Poorly soluble	0
135	436.54	10	5	0	121.51	99.72	4.25	Moderately soluble	Poorly soluble	0
13	507.68	11	5	0	145.06	105.71	5.48	Poorly soluble	Poorly soluble	1
137	518.57	12	8	0	131.32	99.72	4.55	Poorly soluble	Poorly soluble	2
138	436.54	9	5	0	121.67	99.72	4.27	Moderately soluble	Poorly soluble	0
139	466.57	10	6	0	128.16	108.95	4.35	Moderately soluble	Poorly soluble	0
140	448.55	9	5	0	124.36	99.72	4.03	Moderately soluble	Poorly soluble	0
141	504.66	11	5	0	143.75	99.72	4.41	Poorly soluble	Poorly soluble	2

**Table 6: Pharmacokinetics of best 10 candidates based on binding energy (E score).**

Comp	HBA	HBD	NoRB	logP(o/w)	TPSA	MW	LNV
6	18	12	6	2.56	36.95	756.58	3
7	12	10	5	1.39	220.76	496.38	2
12	5	2	5	4.78	37.3	438.69	1
32	6	6	4	3.28	121.38	748.83	3
66	4	1	11	3.69	64.63	429.51	0
80	6	1	12	4.15	132.73	565.66	1
89	6	1	12	4.43	104.49	545.60	1
121	8	0	11	4.24	102.83	463.50	0
138	5	0	9	4.27	99.72	436.54	0
139	6	0	10	4.35	108.95	466.57	0
Rosig	3	2	3	1.04	77.46	332.35	0

**KEY:** MW:molecularweight;HBA:hydrogenbondacceptor;HBD:hydrogenbonddonor;TPSA:totalpolarsurfacearea; NoRB:numberof rotatablebond;LNV:Lipinski'snumberofviolations;Rosig; rosiglitazone

## 5.0 DISCUSSION

### Molecular docking

The calculated free binding energy after molecular docking is given in Table 3 as compounds which showed strong binding affinity with the receptors. Compound **6** gave the lowest binding energy (the highest binding affinity) with **3VI8** (8.12 kcal/mol) while other compounds also had higher significant binding affinity for **3VI8** compared to rosiglitazone. Molecular operating environment (MOE) was used for docking studies. Table 2 shows the free binding energy, DG (kcal/mol) of the compounds against each selected drug receptor. These DG were compared to both the co-crystallized PPAR $\gamma$  and the standard drug. Virtual screening of the entire 141 compounds were done simultaneously using the MOE. 50 compounds showed good binding affinity based on their energy (E) scoring functions. 20 compounds were further selected and subjected to rigid docking and thereafter toxicity and physicochemical properties considerations including compounds; (6,7,12, 30,31,32,39,40,66,67,72,80,89,98,105,109,118,121,139 and 139). The result showed that 10 compounds (139, 138, 121, 80, 89, 66,6, 7, 12 and 32) of the 20 compounds showed high binding affinity (-7.4kcal/mol) for the target protein (3vi8, a PPAR $\gamma$ ) same as the standard reference drug (Rosiglitazone).

Prediction of the bioavailability of molecules in human body could be optimized using the results of the physicochemical properties. Hence, we explored and calculated the molecular descriptors including: molecular weight (MW), partition coefficient (log P), hydrogen bond acceptor (HBA), hydrogen bond donor (HBD), topological polar surface area (TPSA) and number of rotatable bond (NoRB), molar refractivity (MR) and number of atoms (nA). Lipophilicity is a property that has a major effect on solubility, absorption, distribution, metabolism, and excretion properties as well as pharmacological activity. Highly lipophilic molecules will partition into the lipid interior of membranes and retain there. When log P is higher than the upper limit, the drug molecule will have low solubility whereas in lower log P, the drug has difficulty to penetrate the lipid membranes. The pharmacokinetics results (Table 4)

showed that all the compounds reported have good balance between compound solubility and its penetration of the lipid bilayers. The empirical conditions to satisfy Lipinski's rule and manifest a good oral bioavailability involve a balance between the aqueous solubility of a compound and its ability to diffuse passively through the different biological barriers. Reckitt reported the modified Lipinski's rule of 5 (ro5), stating that a likely drug molecule

should have an octanol-water partition coefficient ( $\log P$ ) between  $-0.4$  and  $5.6$ , molar refractivity (AMR) between  $40$  and  $130$ , number of atoms ( $nA$ ) between  $20$  and  $70$ , hydrogen bond donor (HBD)<sub>5</sub> taken as equivalent to the number of  $-OH$  and  $-NH$  groups, hydrogen bond acceptor (HBA)<sub>10</sub> taken as equivalent to the number of oxygen and nitrogen atoms and molecular weight (MW) not more than  $500$ . A violation of more than one of these physicochemical parameters disqualifies a compound from being a likely drug. However, compounds that will serve as substrate for biological transporters do not obey this rule. They can have violations up to 435. This then imply that violations of more than one rule does not totally rule out a compound as a likely drug candidate.

Total polar surface area (TPSA) has often been used as a surrogate property for cell permeability. A molecule with TPSA  $\leq 140 \text{ \AA}^2$  would be able to permeate the cell. Only compounds 17g-1 had TPSA less than  $140$  and as such can permeate the cell membranes. The percentage solubility calculated from  $\% \text{ ABS} = 109 - 0.345 \cdot \text{TPSA}$ <sup>37</sup>, showed that only compounds 17g-1 had good solubility at  $74\%$  which is a designation of good bioavailability upon oral administration.

Toxic doses of chemical substances are often given as LD50 values in mg/kg body weight. The LD50 is the median lethal dose, a dose at which  $50\%$  of test subjects die upon exposure to a compound. Toxicity classes are defined according to the globally harmonized system of classification of labeling of chemicals (GHS).

- Class I: fatal if swallowed ( $\text{LD}_{50} \leq 5$ )
- Class II: fatal if swallowed ( $5 < \text{LD}_{50} \leq 50$ )
- Class III: toxic if swallowed ( $50 < \text{LD}_{50} \leq 300$ )
- Class IV: harmful if swallowed ( $300 < \text{LD}_{50} \leq 2000$ )
- Class V: may be harmful if swallowed ( $2000 < \text{LD}_{50} \leq 5000$ )

Class VI: non-toxic (LD50 > 5000)

The results from the toxicity assessment include the LD50 of the compounds and their activity or inactivity to hepatotoxicity, carcinogenicity, immunotoxicity, mutagenicity, cytotoxicity, Aryl hydrocarbon Receptor, Androgen Receptor, Androgen Receptor Ligand Binding Domain, Estrogen Receptor Alpha, Estrogen Receptor Ligand Binding Domain, Peroxisome Proliferator Activated Receptor Gamma, Nuclear factor (erythroid-derived 2)-like 2/antioxidant responsive element (nrf2/ARE), Heat shock factor response element, Mitochondrial Membrane Potential, Phosphoprotein (Tumor Suppressor) p53, and the ATPase family AAA domain-containing protein 5 (**Table3**). Special consideration was given to the LD50, hepatotoxicity, carcinogenicity and immunotoxicity of compounds that showed safety LD50 values.

#### 4.0 Conclusion

The findings of the study showed that ten (10) out of the 141 compounds analyzed are potential peroxisome proliferators-activated receptor gamma (PPAR  $\gamma$ ) agonists with possible antidiabetic activity. The promising compounds (6,7,12, 32, 66, 80, 89, 121, 138 and, 139) are of the marine origin and extracted from the species of: *Pelvitasiliquoso*, *Ecklonia cava*, *Eckloniastolonia*, *Eckloniacava*, *Laurenciasimili*, *Sargasumthumbegii*, *Ishigeokamurae*, *Sargassumringgoldianum*, *Laurenciasimili*, and *Laurenciasimili* respectively. These compounds showed higher binding affinity with the PPAR $\gamma$  protein target (PDB code: 3vi8) when compared with the standard PPAR $\gamma$  agonist, Rosiglitazone used as the test reference. Consequently, they have been selected for further assay as "hits" and lead-hopping candidates for possible development of a new PPAR  $\gamma$  agonist. The result is therefore, a precedent for deeper investigation into the potency of these compounds. We recommend that *in vitro* and *in vivo* studies could be proceeded on the selected "Hits" for further validation and possible development as new drug entities and perhaps novel therapeutic antidiabetic agents for the management of Diabetes Mellitus.

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