



# Establishing A Remedy Of Dyslexia From Indian Traditional Herbs By *in-silico* Methods

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**Abstract:** Dyslexia is a major childhood disorder. It causes learning disability because of which several behavior flaw is noticed like stammering, faulty writing, lack of attention and seen these children more energetic than normal children of same age. Recent studies suggest association of dyslexia with multiple gene mutations. In this work, we are identifying the genes responsible for dyslexia, establishing their 3d structure by using the structural coordinates of homologous templates. Several medicinal herbs from Ayurvedic, Siddha and Unani background are found to have a cure for neurological disorders. In this work we are targeting the compounds from these herbs against the gene receptors as potential ligands for the disorder.

**Keywords:** dyslexia, medicinal herbs, homology modelling, virtual screening

## I. INTRODUCTION

Dyslexia is a language-based learning disability. Dyslexia refers to an association of symptoms, which result in people having difficulties with specific language skills, particularly reading. Dyslexia, or developmental reading disorder, is characterized by difficulty with learning to read fluently and with accurate comprehension despite normal or above-average intelligence. Students with dyslexia usually experience difficulties with other language skills such as spelling, writing, and pronouncing words. Dyslexia affects individuals throughout their lives, however, its impact can change at different stages in a person's life. It is referred to as a learning disability because dyslexia can make it very difficult for a student to succeed academically in the typical instructional environment, and in its more severe forms, will qualify a student for special education, special accommodations, or extra support services [1, 2]. Genes associated to dyslexia referred in this work are DCDC2, KIAA0319, DYX1C1, CMIP [3].

**DCDC2: DOUBLECORTIN DOMAIN-CONTAINING 2**

This gene encodes a doublecortin domain-containing family member. The doublecortin domain has been demonstrated to bind tubulin and enhance microtubule polymerization. This family member is thought to function in neuronal migration where it may affect the signaling of primary cilia. Mutations in this gene have been associated with reading disability (RD) type 2, also referred to as developmental dyslexia. Alternatively spliced transcript variants encoding the same protein have been found for this gene. Mutations in this gene have been associated with reading disability (Rd) also referred to as developmental dyslexia.

Changes in the DCDC2 gene are frequently found among dyslexics. Altered alleles often occur among children with reading and writing difficulties. The gene appears to have a strong linkage with the processing of speech information when writing [4, 5].

### **KIAA0319:**

KIAA0319 is a protein which in humans is encoded by the KIAA0319 gene and its variants of the KIAA0319 gene have been associated with developmental dyslexia. Reading disability, also called dyslexia, is a major social, educational, and mental health problem associated with it. In spite of average intelligence and adequate educational opportunities, 5 to 10% of school children have been detected having substantial reading deficits. Twin have shown a substantial genetic component to this disorder, with heritable variation estimated at 50 to 70%. An NIDCD-supported investigator has identified a mutation in a gene on chromosome 6, called the KIAA0319 gene, that appears to play a key role in Specific Language Impairment. The KIAA0319 protein is expressed on the cell membrane



and may be involved in neuronal migration and furthermore, KIAA0319 follows a clathrin-mediated endocytic pathway [6, 7, 8].

**DYX1C1: DYSLEXIA SUSCEPTIBILITY 1 CANDIDATE 1**

This gene encodes a tetratricopeptide repeat domain-containing protein. The encoded protein interacts with estrogen receptors and the heat shock proteins, Hsp70 and Hsp90. An homologous protein in rat has been shown to function in neuronal migration in the developing neocortex. A chromosomal translocation involving this gene is associated with a susceptibility to developmental dyslexia. Mutations in this gene are associated with deficits in reading and spelling. Alternative splicing results in multiple transcript variants [9, 10].

**CMIP: C-MAF INDUCING PROTEIN**

This gene encodes a c-Maf inducing protein that plays a role in T-cell signaling pathway. Alternatively spliced transcript variants encoding different forms have been described for this gene. CMIP is a protein coding gene diseases associated with CMIP include speech and communication disorder specific language impairment [11].

**II METHODOLOGY**

Dyslexia receptor amino acid sequences were downloaded from NCBI (Table 1). Their 3d structure was modelled using swiss model.

Table 1: Dyslexia receptors with their NCBI Accession number

Gene name	Accession number
DCDC2	AAH50704.1
KIAA0319	BAA20777.3
DYX1C1	NP_570722.2
CMIP	AAH38113.1

Herbs useful in treating CNS disorders are used in this work [12, 13, 14, 15, 16]. The active compounds of the herbs used in this work are downloaded from PUBCHEM (Table 2)

Table 2: Herbs with their compounds used in this work.

Herb name	Scientific name	Compounds
Schizandra	Schisandra chinensis	Schizandrin Deonyschizandrin Gomisine
Kava kava	Piper	Kavain

	methysticum	Caffeic Acid Dihydrokavain Methysticin Dihydromethysticin Yangonin Desmethoxyangonin Allantoin Rosmaric Acid Methyl Oleanolate Stigmasterol Isoleucine Phnyylarine
Piperaceae	Piper caninum	Bornyll Caffeate
	Rheum officinale	Cinnanic Acid
	Forsythia suspensa	Pinoresinol
	Podophyllum peltatum	Pudophyllotonin
	Nardostachys jatamansi	Valeranone

The active compounds of the herbs are docked with the dyslexia receptors DCDC2, KIAA0319, DYX1C1 and CMIP in this work.

**III RESULTS & DISCUSSION**

Amino acid sequence of the gene responsible for Dyslexia was retrieved from Genbank/Uniprot database and their accession number was noted.

Structure modeling was done using Swiss Model. Unknown and reference ligand were identified and virtual screening and docking studies was performed on the ligands with the receptor proteins.

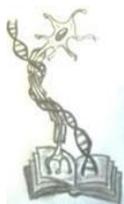
Table 3: Docking analysis of the dyslexia receptors with the active compounds of the CNS herbs

COMPOUN DS docking with DCDC2 receptor	DOC KING SCORE KCA L/MO L	INTER ACTIN G AMIN O ACIDS	NO. OF INTER ACTIO NS	DOC KIN G (YES /NO)
BRONYL CAFFEATE	2612 KCA L/MO L	LEV 190	1	YES
CINNAMIC ACID	2208 KCA	THR 189	1	YES



	L/MO L			
PINORESIN OL	3784 KCA L/MO L	ARG 180	1	YES
PODOPHYL LOTOXIN	3814 KCA L/MO L	GLN 136	1	YES
VALERANO NE	2208 KCA L/MO L	THR 189	1	YES
ALLANTOI N	1892 KCA L/MO L	PHE 206	1	YES
METHYSTI CIN	2552 KCA L/MO L	GLU 202	1	YES
DESMETHO XYANGONI N	3100 KCA L/MO L	GLY 204 PHE 206	1 1	YES
DIHYDROK AVAIN	2538 KCA L/MO L	ARG 180 THR 178	1 1	YES
ISOLUCINE	2032 KCA L/MO L	PHE 206	1	YES
KAVAIN	2470 KCA L/MO L	THR 178	1	YES
GOMISIN	2894 KCA L/MO L	TYR 188 PHE 215	1 1	YES
ROSMARIN IC ACID	3790 KCA L/MO L	THR 178	1	YES
DIHYDROM ETHYSTICI N	3354 KCA L/MO L	ARG 190	1	YES

PINORESIN OL	3784 KCA L/MO L	ARG 180	1	YES
STIGMAST EROL	4590 KCA L/MO L	ARG 180	1	YES
METHYLOL EANOLATE	NO DOC KING			
YANGONIN	NO DOC KING			
PHENYLAL ARINE	NO DOC KING			
<b>COMPOUN DS docking with KIAA receptor</b>	<b>DOC KING SCO RE KCA L/MO L</b>	<b>INTER ACTIN G AMIN O ACIDS</b>	<b>NO. OF INTER ACTIO NS</b>	<b>DOC KIN G (YES /NO)</b>
GOMISIN	3540 KCA L/MO L	LEV 684	1	YES
METHYSTI CIN	3222 KCA L/MO L	ASN 715	1	YES
ROSMARIN IC ACID	4938 KCA L/MO L	ASN 715	1	YES
CINNAMIC ACID	2910 KCA L/MO L	SER 648	1	YES
PODOPHYL LOTOXIN	NO DOC KING			
ALLANTOI N	2772 KCA L/MO L	ASN 780	1	YES
SCHIZAND RIN	3930 KCA	ASN 715	1 1	YES



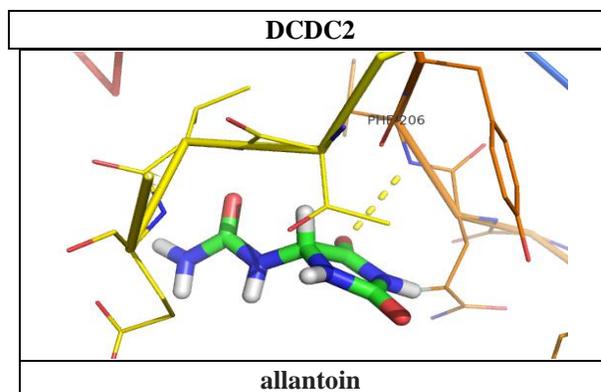
	L/MO L	SLR 795		
DESMETHOXYANGONIN	3752 KCA L/MO L	ASN 715	1	YES
BORNYLCAFFEATE	NO DOC KING			
DIHYDROMETHYSTICIN	4356 KCA L/MO L	GLN 685 ASN 715	1 2	YES
ISOLEUCINE	2794 KCA L/MO L	ASN 780 THR 779	1 1	YES
PINORESINOL	NO DOC KING			
STIGMASTEROL	NO DOC KING			
METHYLOLEANOLATE	NO DOC KING			
YANGONIN	NO DOC KING			
PHENYLALANINE	NO DOC KING			
<b>COMPOUNDS docking with CIMP receptor</b>	<b>DOC KING SCORE KCA L/MO L</b>	<b>INTERACTING AMINO ACIDS</b>	<b>NO. OF INTERACTIONS</b>	<b>DOC KING (YES /NO)</b>
GOMISIN	3526 KCA L/MO L	HIS 556	1	YES
METHYSTICIN	NO DOC KING			
ROSMARINIC ACID	NO DOC KING			

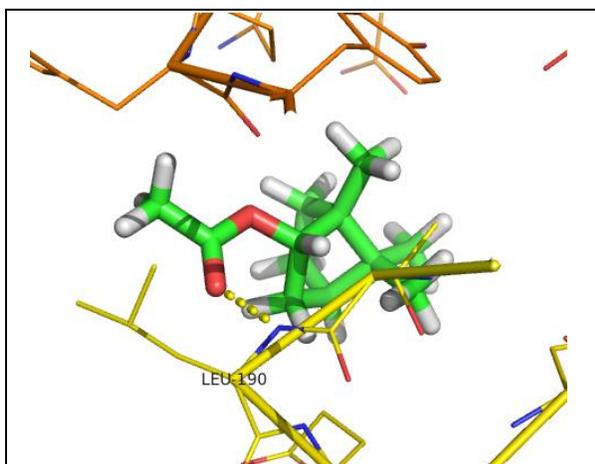
CINNAMIC ACID	NO DOC KING			
PODOPHYLLOTOXIN	4720 KCA L/MO L	ARG 501	1	YES
ALLANTONIN	NO DOC KING			
SCHIZANDRIN	3680 KCA L/MO L	GLU 527	1	YES
DESMETHOXYANGONIN	3754 KCA L/MO L	GLU 527	1	YES
BORNYLCAFFEATE	NO DOC KING			
DIHYDROMETHYSTICIN	4174 KCA L/MO L	ALA 526	1	YES
ISOLEUCINE	NO DOC KING			
PINORESINOL	4740 KCA L/MO L	LEU 514	1	YES
STIGMASTEROL	5100 KCA L/MO L	ARG 501	1	YES
METHYLOLEANOLATE	NO DOC KING			
YANGONIN	4040 KCA L/MO L	GLU 527	1	YES
PHENYLALANINE	3310 KCA L/MO L	LEU 540	1	YES
<b>COMPOUNDS docking with</b>	<b>DOC KING SCORE</b>	<b>INTERACTING</b>	<b>NO. OF INTERACTIONS</b>	<b>DOC KING</b>



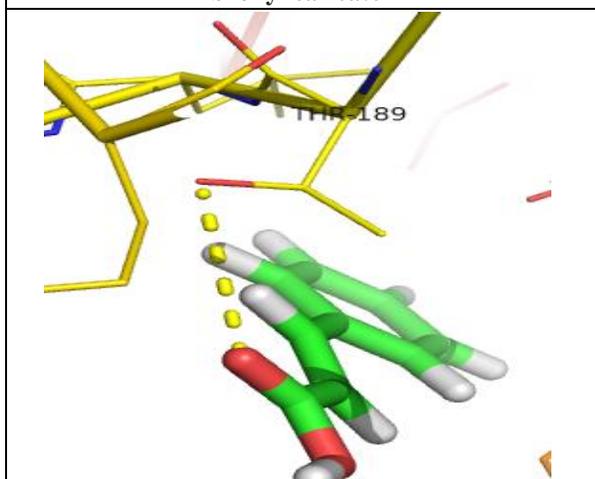
DYX1C1 receptor	RE KCA L/MO L	AMIN O ACIDS	NS	(YES /NO)
DIHYDROKAVAIN	NO DOC KING			
SCHIZANDRIN	3258 KCA L/MO L	ASN 339	1	YES
METHYSTICIN	2686 KCA L/MO L	ARG 365 ASN 363	1 1	YES
VALERANONE	NO DOC KING			
METHYLENEOLANOLATE	NO DOC KING			
DIHYDROMETHYSTICIN	3314 KCA L/MO L	LYS 408	1	YES
PODOPHYLLOTOXIN	3616 KCA L/MO L	ARG 371 THR 374	1 1	YES
KAVAIN	2528 KCA L/MO L	ASP 396	1	YES
DEOXYSCHIZANDRIN	NO DOC KING			
ISOLEUCINE	2046 KCA L/MO L	ARG 371	2	YES
GOMISIN	2866 KCA L/MO L	ASN 339	1	YES
YANGONIN	3202 KCA L/MO L	LYS 367	1	YES
ALLANTOIN	1980	ARG	1	YES

N	KCA L/MO L	365		
BORNYLCAFFEATE	2500 KCA L/MO L	ARG 365	1	YES
DESMETHOXYANGONIN	2934 KCA L/MO L	LYS 349	1	YES
CINNAMIC ACID	NO DOC KING			
STIGMASTEROL	4268 KCA L/MO L	ASN 361	1	YES
PHENYLALANINE	2212 KCA L/MO L	ASP 396	2	YES
ROSMARINIC ACID	3598 KCA L/MO L	TYR 306 LYS 408 ARG 371	1 1 1	YES
PINORESINOL	3854 KCA L/MO L	LYS 322	1	YES

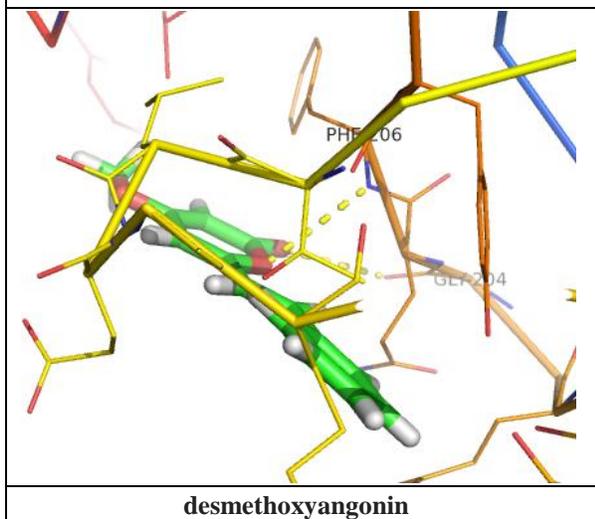




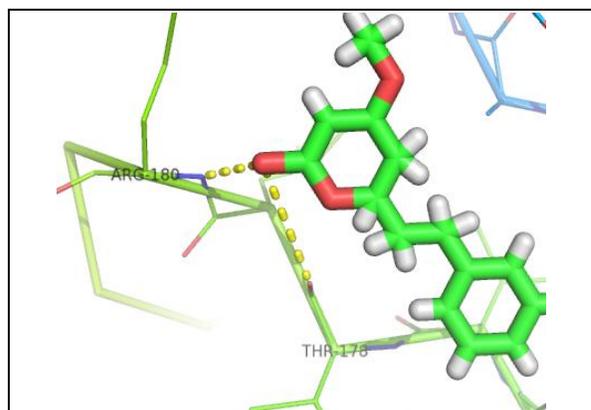
**bronyl caffeate**



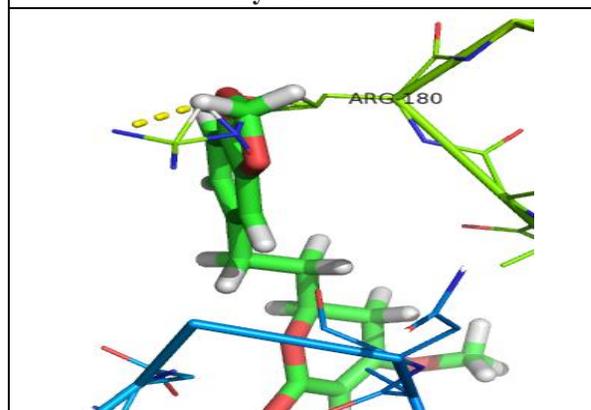
**cinnamic acid**



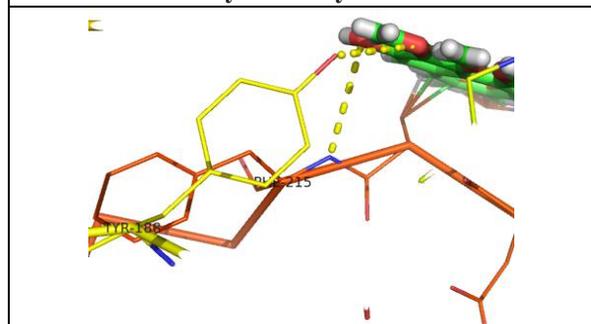
**desmethoxyangonin**



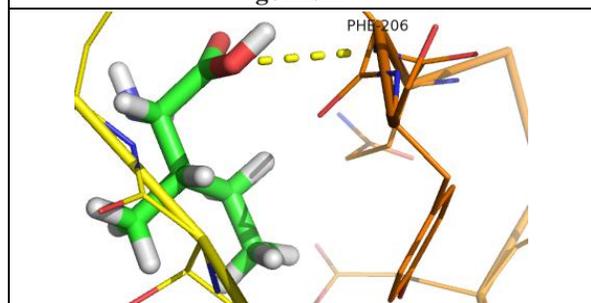
**dihydrokavain**

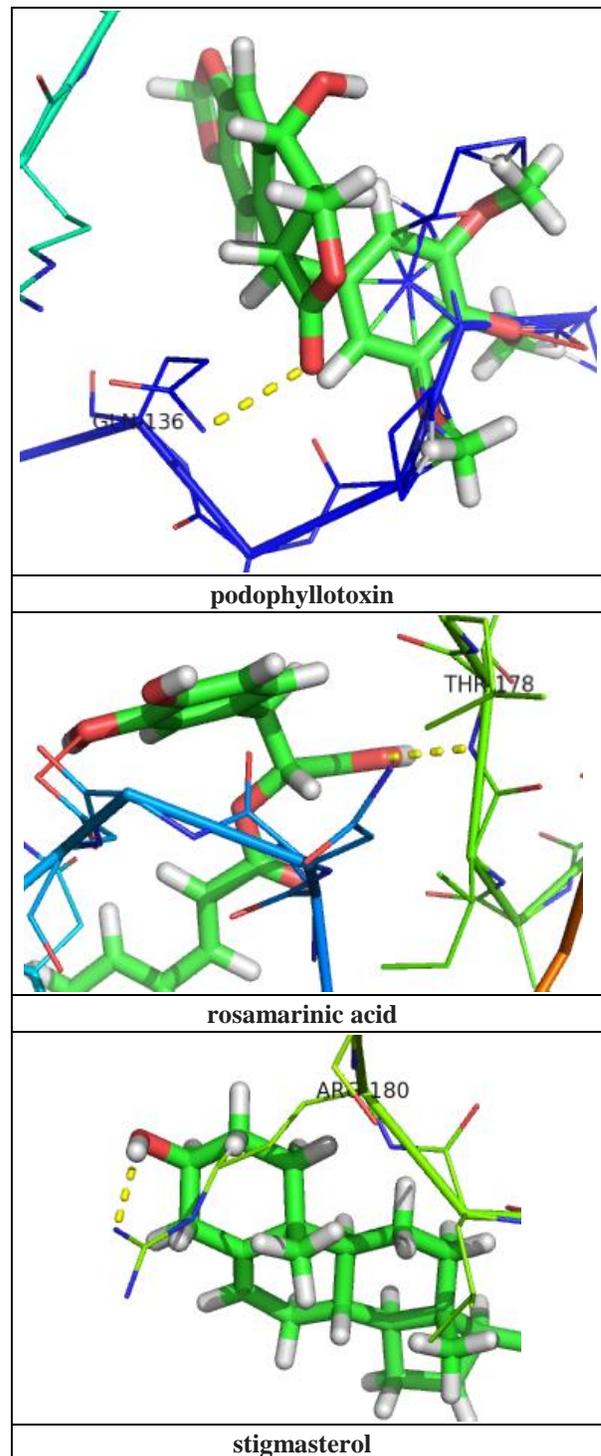
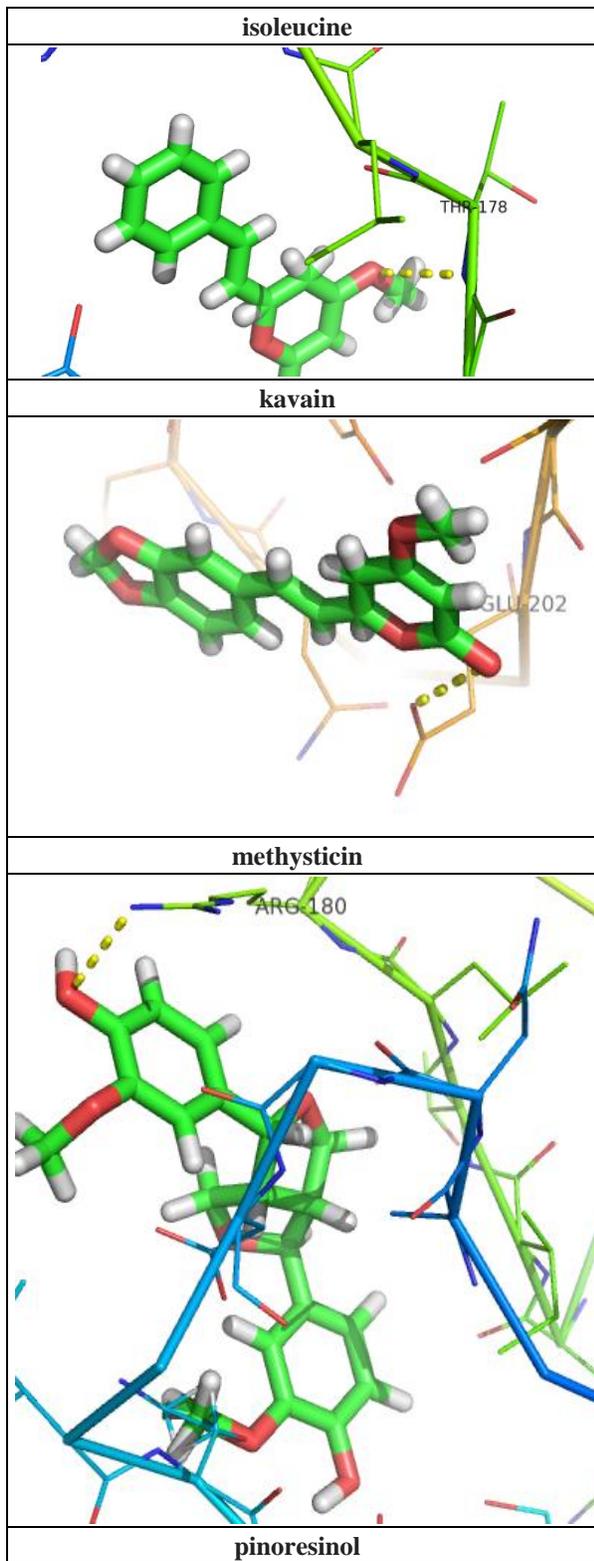


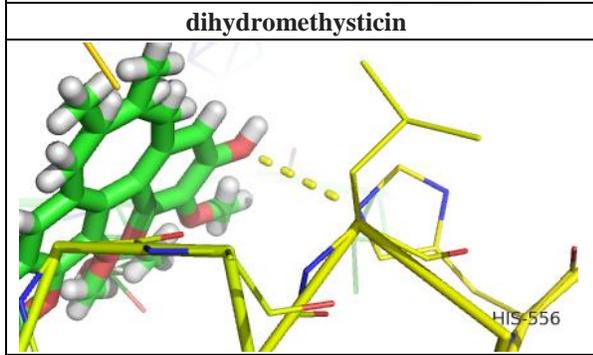
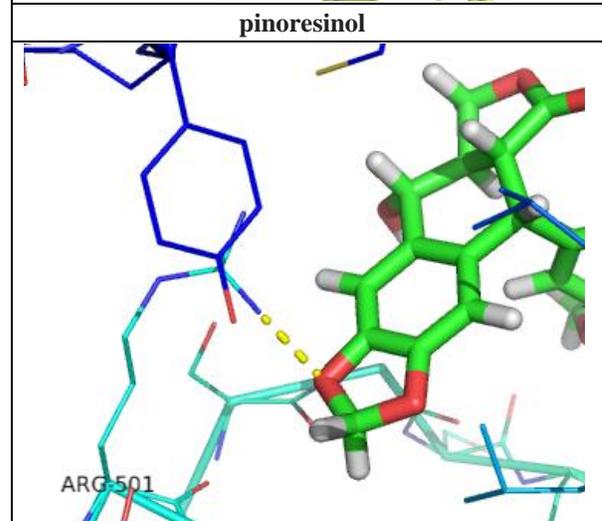
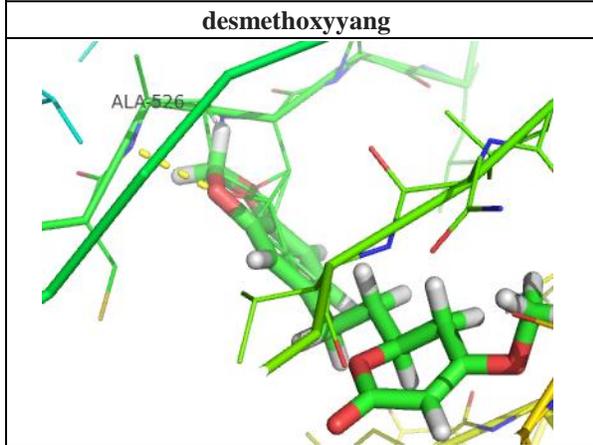
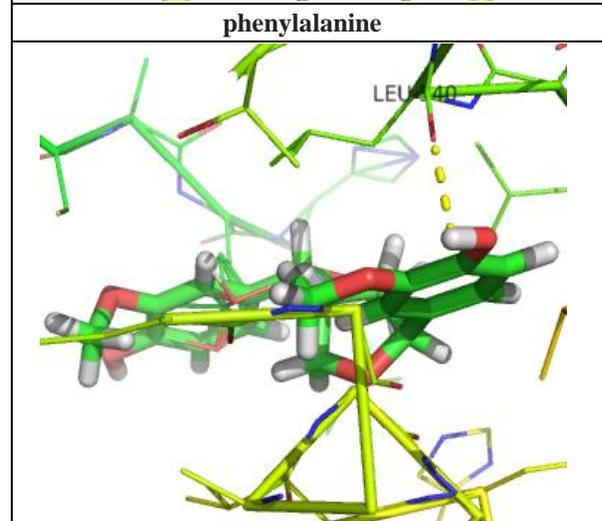
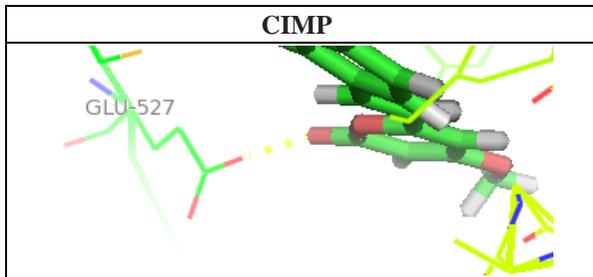
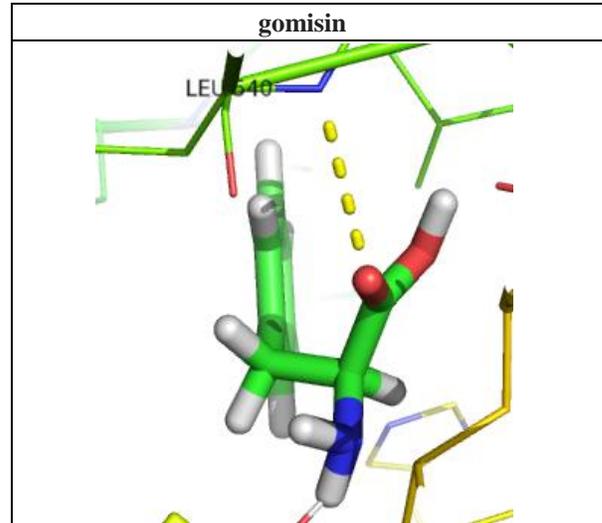
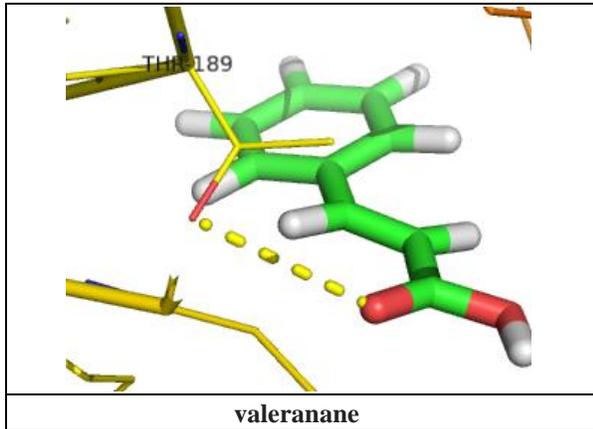
**dihydromethysticin**

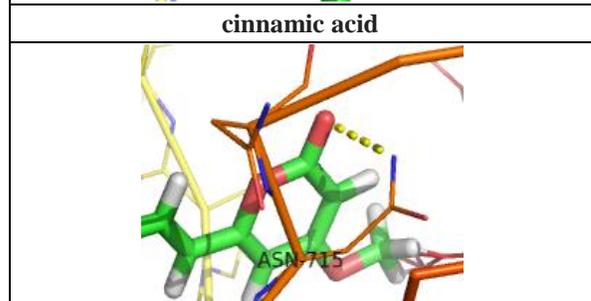
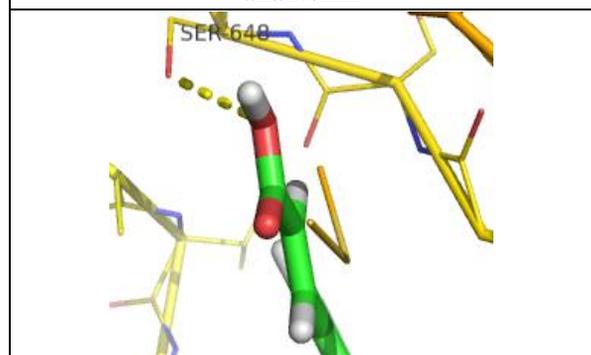
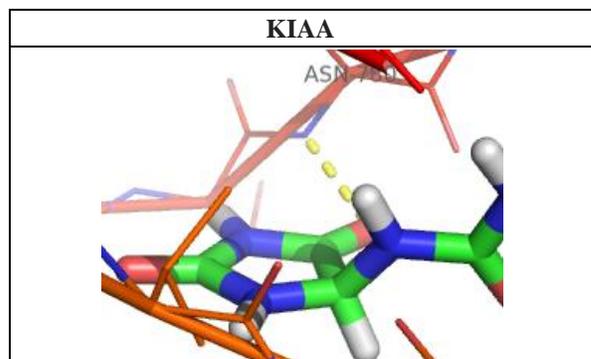
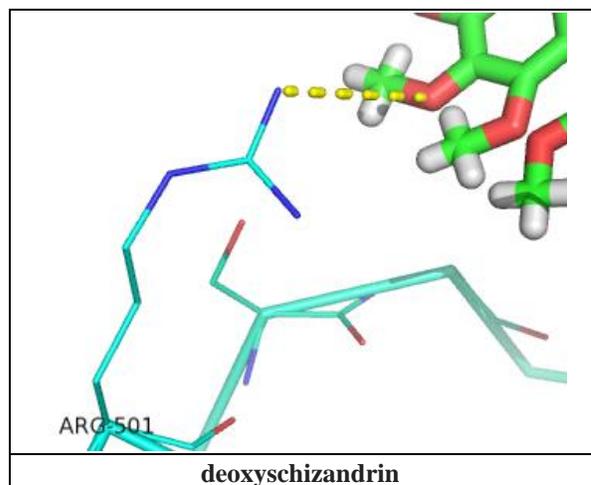
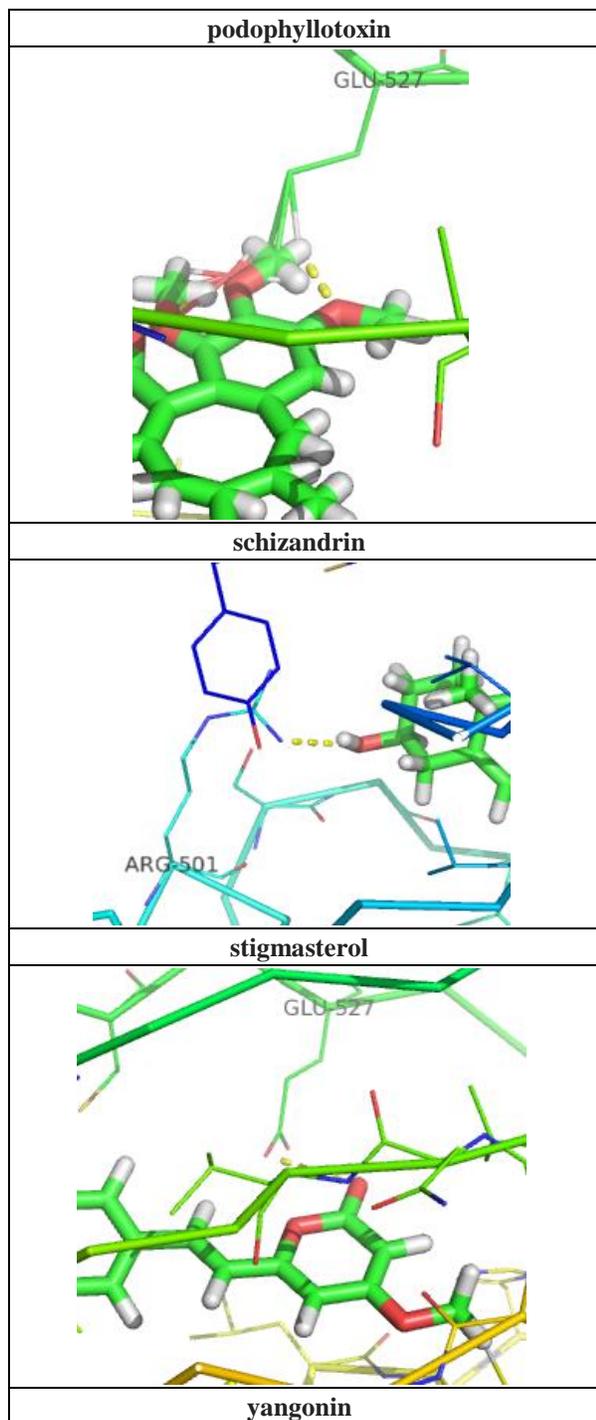


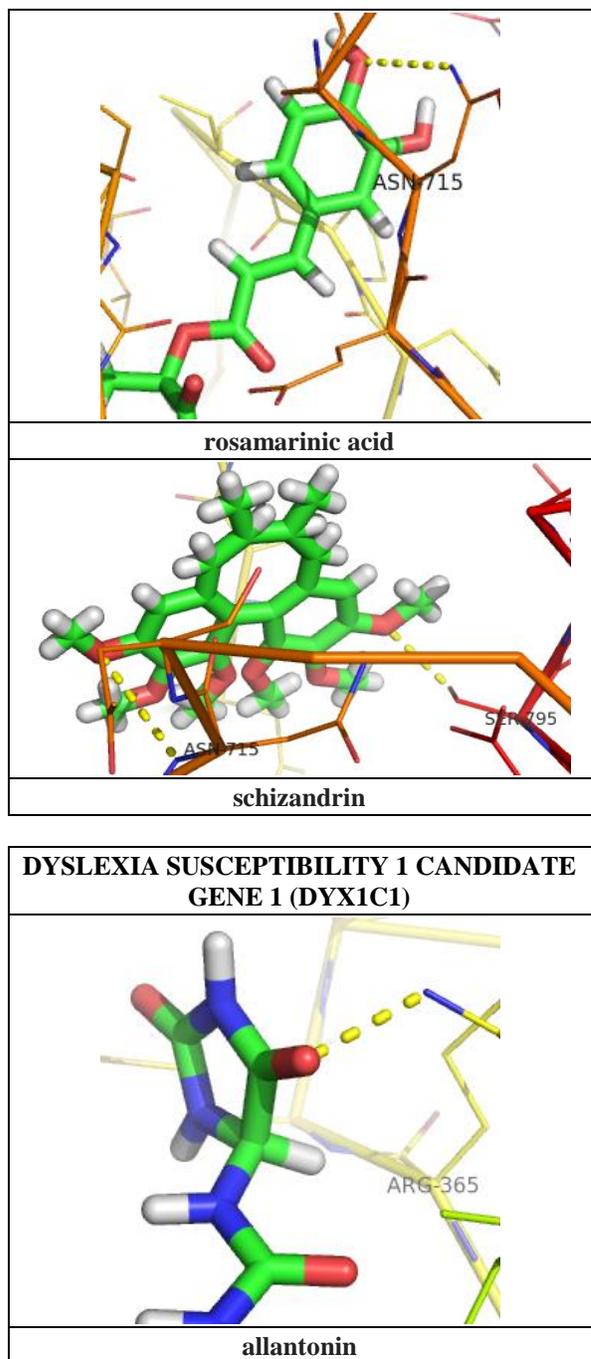
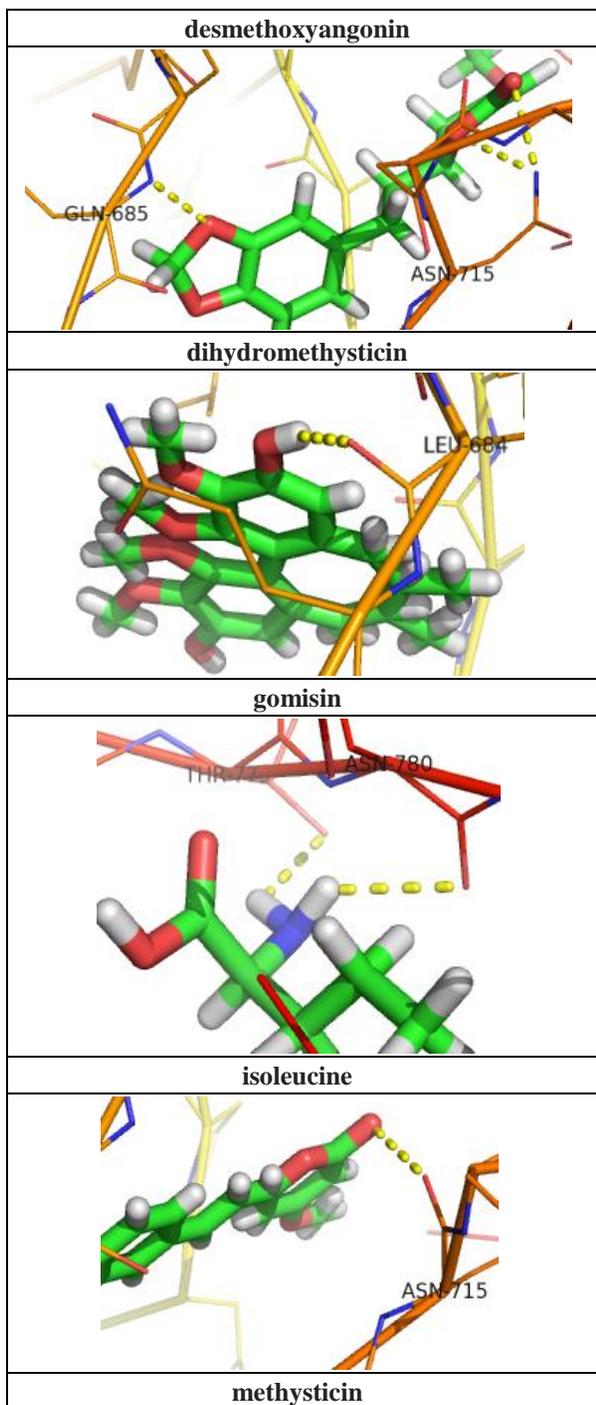
**gomisin**

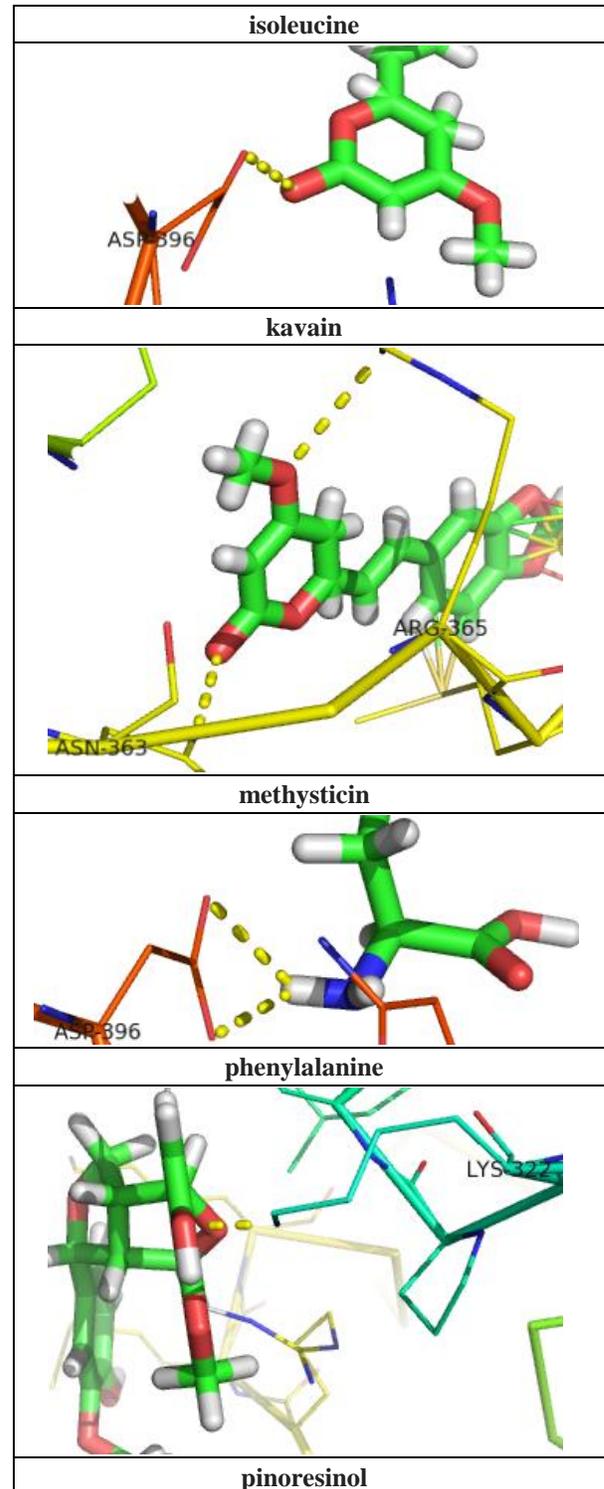
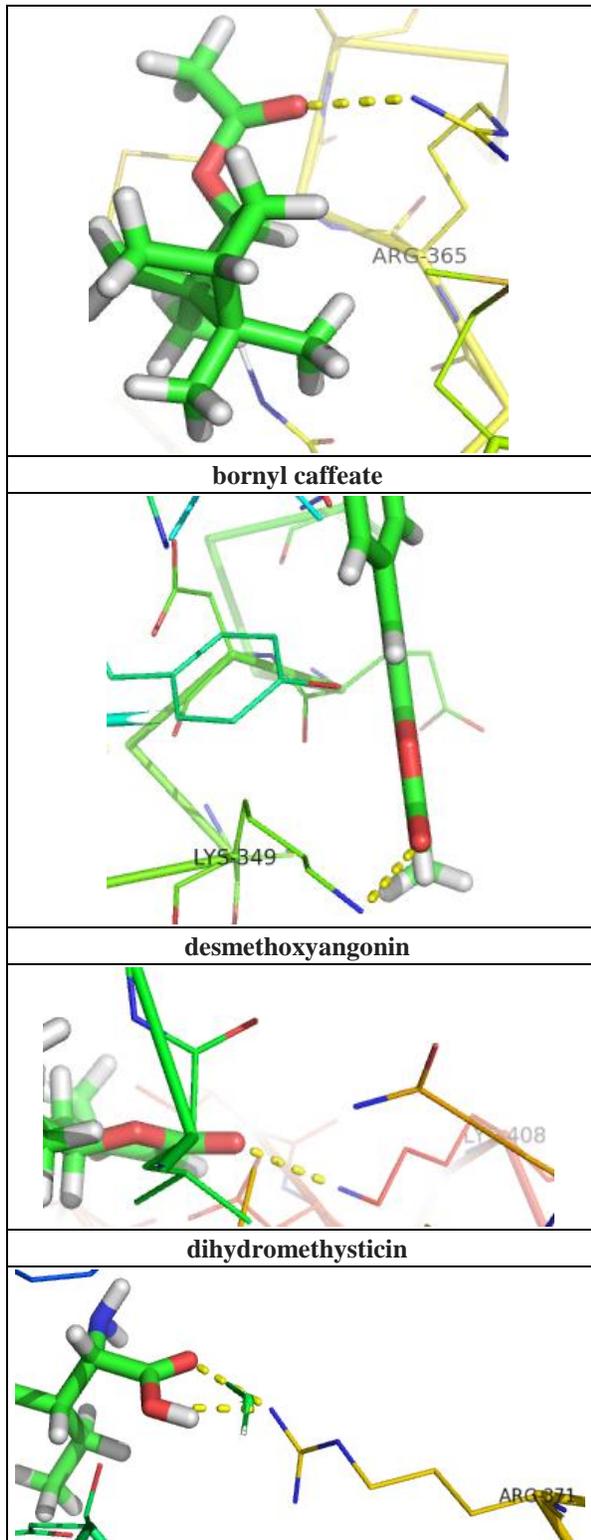


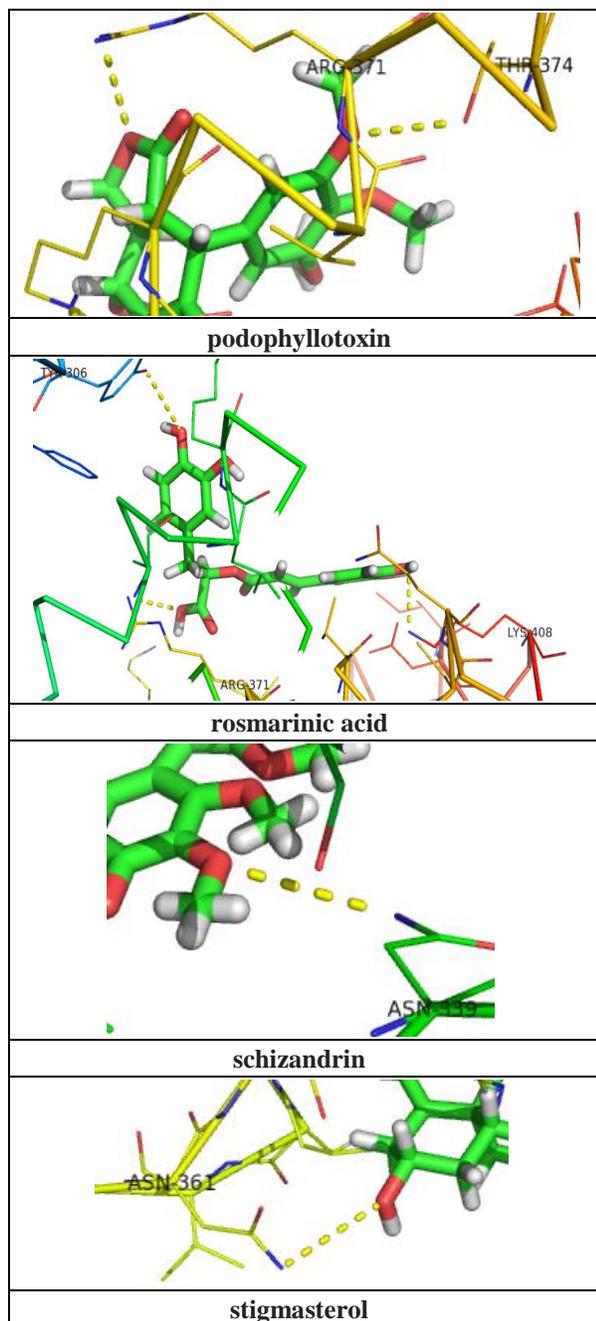












**Fig. 1:** Docking images of the dyslexia receptors with the active compounds of the CNS herbs

It is seen that DCDC2 best interacts with desmethoxyangonin, dihydrokavain and gomisin. It is seen that KIAA best docks with schizandrin, dihydromethysticin and isoleucine.

It is seen that CMIP docks with gomisin, podophyllotoxin, schizandrin, desmethoxyangonin, pinosresinol, stigmasterol, yangonin and phenylalanine.

It is seen that DYX1C1 docks best with methysticin, podophyllotoxin, isoleucine, phenylalanine and rosmarinic acid and has a good docking score and interaction with desmethoxyangonin.

#### IV CONCLUSION

It is seen that the compound desmethoxyangonin docks with all the receptors and has good interaction with all the receptors and hence selected as best ligand for dyslexia receptor proteins.

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