

From EST to structure models for functional inference of APP, BACE1, PSEN1, PSEN2 genes

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Abstract:

Successive oxidative stress and biochemical changes results in neuronal death and neuritic plaques growth in Alzheimer's disease (AD). Therefore, it is interest to analyze amyloid- β precursor protein (APP), beta-secretase 1 (BACE1), presenilin (PSEN1 and PSEN2) genes from brain tissues to gain insights. Development of potential inhibitors for these targets is of significance. EST sequences of 2898 (APP), 539 (BACE1), 786 (PSEN1) and 314 (PSEN2) genes were analyzed in this study. A contig sequences with APP (contigs 1-4), BACE1 (contigs 5-7), PSEN1 (contigs 8, 9, 10, 11), PSEN2 (contigs 13, 14) except PSEN1 (contigs 10) and PSEN2 (contigs 13) genes were identified. APP (contig 3 without translational error) was further analyzed using molecular modeling and docking to show its binding with curcumin (principal curcuminoid of turmeric) having -7.3 kcal/mol interaction energy for further consideration as a potential inhibitor.

Keywords: Alzheimer's disease, Curcumin, Hypothetical protein

Background:

Alzheimer's disease (AD) is caused due to the structural and functional loss of neurons which shows symptoms like cognitive and memory deterioration, progressive destruction of intellectual activities in day to life and behavioral abnormalities [1]. About 36 million people were found to be affected by AD worldwide in 2010 and it was anticipated to rise 66 million by 2030 and 115 million by 2050 [2]. In India, 3.7 million people were affected by AD [3] and the number of people having AD. Prevalence increases exponentially with age, affecting a little more than 1% in the population aged 65-69 years up to as much as 30-40% in the oldest old [4]. Alzheimer's disease is mainly caused due to the accumulation of β -amyloid peptides [5], which are formed by the action of sequential cleaving of the APP gene which plays an important role in the central nervous system. Proteolytic cleavage of APP by β - and γ -secretase enzymes resulting in the release of neurotoxic $A\beta$ peptides which can aggregate into oligomer is known. A mutation in the APP gene is likely to inhibit α -secretase cleavage which further enables preferential cleavage by β -secretase. Mutations in the PSEN1 and PSEN2 genes (which are components

of the γ -secretase complex) results in increased cleavage by γ -secretase at this site. Both these conditions result in the excess production of $A\beta$ peptide. Eventually, subsequent oxidative stress and biochemical changes result in the neuronal death and development of neuritic plaques in AD [6].

Expressed sequence tags are sequenced regions of complementary deoxyribonucleic acid (cDNA) imitates of messenger ribonucleic acid (mRNA) that are expressed in different states and represents element of the transcribed portion of the genome. The EST sequence information plays a vital role in human biology and disease, such as neurological disorders [7]. This helps to identify the functional genes expressed in diseased condition. Mutations in the alzheimer's susceptibility genes APP, BACE1, PSEN1 and PSEN2 greatly increase the risk of AD. The approved drugs for AD namely, tacrine, donepezil, rivastigmine and galantamine failed due to severe side effects and were abandoned. This work will help to identify the functional annotation of APP, BACE1, PSEN1,

PSEN2 genes and new discovery for the development of novel therapeutic approaches for the treatment of AD.

Methodology:

Retrieval of ESTs sequence and assembly:

In silico analysis of AD human genes APP, BACE1, PSEN1 and PSEN2 taken from UniGene database and those genes originating from brain tissues were taken. The 5' ESTs were considered, as the ESTs generated from the 3' end are most error prone as of the low base-call quality at the start of sequence reads. The 5' EST sequences were extracted using contig assembly program by CAP3 server [8]. The default parameters were used and each gene sequences were submitted to DNA sequence assembly program (CAP3) server in FASTA formatted text file and result was displayed in different output files e.g. contigs, single sequences, Assembly details and sequence file. We have selected contig sequence data set as it is useful functionality ascertained.

Database similarity search:

The contig sequences were obtained from clustering and similarity search using tools like nucleotide BLAST (BLASTN) and BLASTX (search protein). The contig sequence is aligned to the genome sequence of the organism using BLAT (BLAST like alignment tool) [9] to assist genome mapping and gene discovery. Each genes contig sequence was generated by BLAT analysis with parameters reading (genome: human, assembly: Dec. 2013 (GRCh38/hg38), query type: translated DNA, sort output: Score, output type: hyperlink).

Conceptual translation of ESTs and functional annotation:

ESTScan is a program that can identify the coding regions in DNA sequences and this was translated into amino acid sequences at either N- or C-terminus. Each contig sequence was generated by ESTScan2 tool [10]. Finally, the amino acid sequences were selected using multiple sequence alignment by CLC Genomics Workbench and further functional annotations were carried out. Our translated protein sequences for each sequence were generated by InterProScan 5.0 [11].

Molecular modelling of hypothetical protein:

Structural annotation of APP hypothetical amino acid sequence was used for build a 3D structure by Modeller v9.13 software [12]. The hypothetical protein sequence was aligned in BLASTP against the Protein Data Bank (PDB) database to select their appropriate templates. The template was selected for hypothetical protein query sequence aligning 18-199 amino acid residues, showing 97% sequence identity with 3KTM [13]; aligning 342-551 amino acid residues shows 99% sequence identity with 3NYL [14] and aligning

652-751 amino acid residues shows 100% sequence identify with 2LP1 [15].

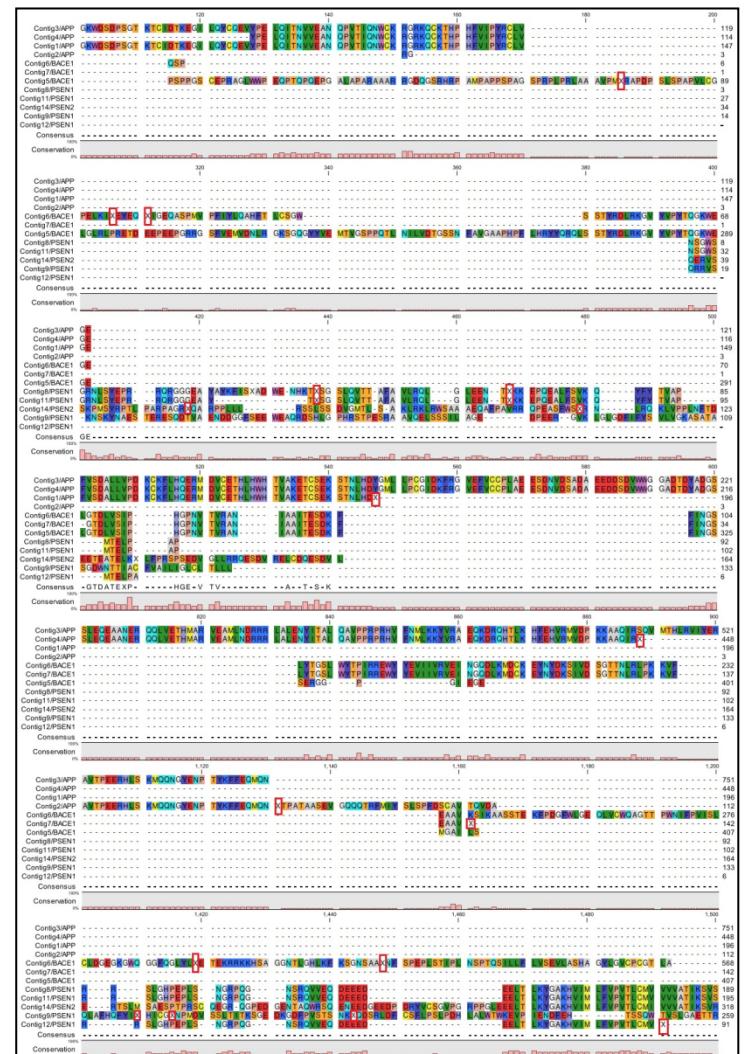


Figure 1: Graphical representation of contig protein sequences obtained from ESTScan2 translation sequences. Red color box represents X error translate level in APP (Contig 1, 2, 4), BACE1 (5, 6, 7), PSEN1 (8,9,11) and PSEN2 (14) except APP contig 3 sequence.

These templates were used to build a 3D structure for homology modelling. Modelled structure was energy minimized using Swiss-PDB viewer program (Gromos96 force field). Theoretically predicted structure was visualized using PyMol visualization

software. The amino acid constraint validation of the modeled APP protein was done by PROCHECK program (www.ebi.ac.uk/thornton-srv/software/PROCHECK/) [16]. Further, 3D profile of the modelled protein was computed by Verify3D program.

Selection of ligands:

The 2D structure of synthetic compounds tacrine, donepezil, rivastigmine, galantamine and natural remedy like compounds from plants such as *Rosmarinus officinalis* (α -Pinene, Camphene, β -Pinene, 1,8-Cineole, α -Thujone, β -Thujone, Chrysanthrone, Camphor, (+)-Borneol, Bornyl acetate, α -Copaene, Trans-Caryophyllene, α -Humulene, Germacrene-D and (+)- δ -Cadinene); *Ginkgo biloba* (Quercetin, Kaempferol, Isorhamnetin, Ginkgolide A, B, C, J, M); *Panax ginseng* (Ginsenoside Rb1 and Rg1); *Curcuma Longa* (Curcumin, Demethoxycurcumin and Bisdemethoxycurcumin); *Salvia officinalis* (Borneol, Caryophyllene, Linalool); *Huperzia serrata* (Huperzine A, B and Lycopodine) ; *Melissa officinalis* (1-Octen-3-ol, 6-Methyl-5-hepten-2-one, Myrcene, (Z)- β -Ocimene, (E)- β -Ocimene, n-Nonanal, Cis-Rose oxide, (+)-Trans-Rose oxide, (+)-Trans-Limonene oxide, Citronellal, Menthol, Isomenthol, Nerol, Neral, Piperitone, Geraniol, Geranial, α -Cubebene, Geranyl acetate, β -Cubebene, β -Caryophyllene, Valencene, Caryophyllene oxide, 1-Hexadecene, n-Eicosane, n-Heneicosane); *Withania somnifera* (Propane,1,1-diethoxy-2-methyl-2-Nonanone, PhenylethylAlcohol, Amyl nitrite, Dodecanoic acid, 3-tert-Butyl-4-hydroxyanisole, Tetradeanoic acid, n-Hexadecanoic acid, 9-Octadecenal, 1-tridecyne, Oleic acid); *Baccopa monnieri* (2-octanol, Dimethoxane, 2-Methyl-1-Phenyl-1-butanol, Phytol, Phytol acetate, Octadecanamide); *Centella asiatica* (Thujopsene, α -Thujene, Eucalyptol, 3-Nonen-2-one, β -Linalool, L-Camphor, trans-Borneol, α -Terpenol, Cis-Geraniol, Isobornyl acetate, 7-Tetradecene, β -Elemene, β -Gurjunene, γ -Elemene, Isocaryophyllene, Aromadendrene, β -Farnesene, β -Acoradiene, β -Selinene, α -Selinene, α -Chamigrene, α -Panasinsen, -(β)-Spathulenol, Viridiflorol, Valerenone, Isoaromadendrene epoxide, Aristolene epoxide, 1-Naphthalenol); *Celastrus paniculatus* (Palmitic acid, Erucic acid, γ -Murolene, Cubenol) were downloaded from PubChem databases as .sdf format. Further, the .sdf format converted into .pdb format using Openbabel 2.3.2.

Molecular docking:

Docking studies was carried out using Glide module from Schrodinger suite [17] to find the interaction between modeled APP protein with natural and synthetic compounds. All the compounds were prepared by LigPrep Module. The protein grids were prepared with the mutated residues and the size of the bounding box was set to 30 \AA . Modelled APP protein coordinates file of enclosing box was set as x=3.9023 \AA ; y=32.884 \AA ; z=30 \AA respectively. All the prepared inhibitory compounds were docked against the

grid generated APP modelled protein. The inhibitory compounds used for docking was screened using Virtual screening. Glide score was selected as the scoring function to rank the poses of each inhibitory compound. Validation of the docking is useful technique to identify best docked complex among number of docked complex.

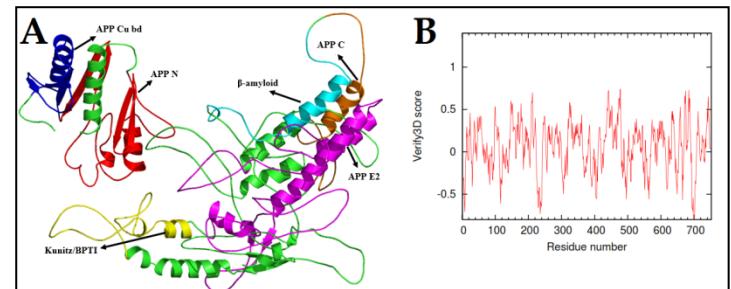


Figure 2: Graphical representation of modelled and validated hypothetical protein of APP. A) Domain regions of red color show APP N (Amyloid A4 N-terminal heparin binding); Blue color is APP Cu bd (Copper-binding of amyloid precursor, CuB); Yellow color is Kunitz BPTI (Kunitz/Bovine pancreatic trypsin inhibitor); Magenta color is APP E2 (E2 domain of amyloid precursor protein); Cyan color is β -amyloid and Orange color is APP C (APP-amyloid). B) Verify 3D plot showed score ranges in between -1.0 to 0.7.

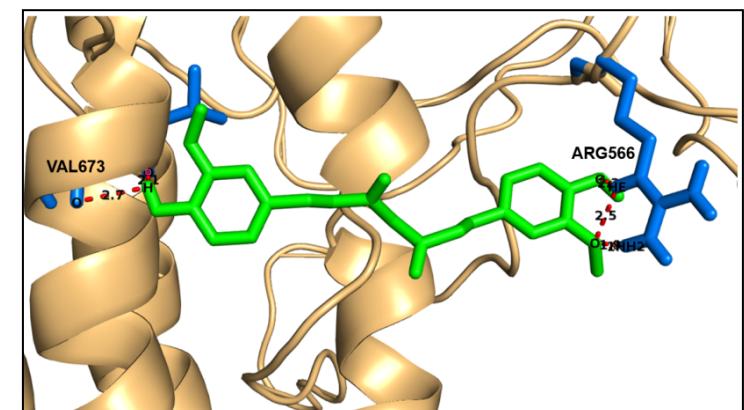


Figure 3: Interaction of modeled protein APP docked with curcumin. Light orange color represents protein; green color is curcumin compound and blue color is interacted residues.

Results and Discussion:

Retrieval of ESTs sequence:

The EST sequences for human AD genes APP, BACE1, PSEN1 and PSEN2 were searched from UniGene database. The gene entries

with their mRNA and ESTs information are listed in **Table 1**. ESTs of four gene entries originating from brain tissue were used for further analysis.

Table 1: UniGene information on human Alzheimer's disease

S. No	Name of the Genes	Source	mRNA	ESTs
1.	APP	Homo sapiens	38	2898
2.	BACE1	Homo sapiens	27	539
3.	PSEN1	Homo sapiens	16	786
4.	PSEN2	Homo sapiens	10	314

It shows the list of mRNA and ESTs entries.

Table 2: BLAT output showing the alignment of APP, BACE1, PSEN1 and PSEN2 contigs sorted by score

Query	Score	Start	End	Qsize	Identity (%)	Chromosome	Strand
APP							
Contig1	547	30	582	583	99.9	21	-
Contig1	515	31	550	583	99.9	21	+
Contig2	734	6	752	780	99.5	21	-
Contig2	734	7	757	780	99.2	21	+
Contig3	3838	1	3876	4579	99.8	21	-
Contig3	3838	1	3876	4579	99.8	21	+
Contig4	1331	2	1340	1340	100	21	+
Contig4	1330	1	1340	1340	100	21	-
BACE1							
Contig5	1605	3	1616	1616	99.9	11	+
Contig5	1604	2	1616	1616	99.8	11	-
Contig6	4916	1	5092	5184	99.4	11	-
Contig6	4916	1	5100	5184	99.3	11	+
Contig7	563	1	572	572	99.7	11	-
Contig7	563	1	572	572	99.7	11	+
PSEN1							
Contig8	546	1	589	589	99.7	14	-
Contig8	545	2	589	589	99.7	14	+
Contig9	4161	1	4265	4265	99.6	14	+
Contig9	4159	1	4265	4265	99.6	14	-
Contig10	465	1	477	604	98.8	14	-
Contig10	464	1	478	604	98.6	14	+
Contig11	1491	1	1501	1680	100	14	-
Contig11	1491	1	1505	1680	99.9	14	+
Contig12	501	44	638	638	99.5	14	+
Contig12	499	38	638	638	98.7	14	-
PSEN2							
Contig13	577	1	580	580	99.9	1	-
Contig13	577	1	580	580	99.9	1	+
Contig14	1499	1	1602	1918	98.4	1	-
Contig14	1467	1	1603	1918	97.9	1	+

Note: (+) given segment and (-) reverse complement. It shows the contig sequence of APP, BACE1, PSEN1 and PSEN2 similarity score.

EST clustering and assembly:

Each gene sequence of ESTs from brain tissue was retrieved. The 5' ESTs were analyzed, as the ESTs created from the 3' end are most error prone because of the low base-call quality at the start of sequence reads. The subjected ESTs along with their resulting contigs found a total of 988 ESTs from four reported gene entries as listed in **Table 5** (Supplementary Material at the bottom of the article). The tissue-based ESTs from four reported genes were subjected to cluster analysis by CAP3 Server. 14 contigs of four genes were found and further analysis was under taken.

Database similarity searches:

The database similarity search by querying these contigs in BLAT against human genome revealed that alzheimer's contig of APP shows good matches with chromosomes 21. The BACE1, PSEN1 and PSEN2 contigs were showing good matches with chromosomes

11, 14 and 1 respectively and are shown in **Table 2**. The conceptual translation of 14 contigs sequences in ESTScan2 provides 12 protein sequences from APP, BACE1, PSEN1 and PSEN2, as presented in this analysis and protein sequences were not available for the rest of two contig nucleotide sequences contig 10 and contig 13. Multiple sequence alignment was done for these 12 protein sequences obtained by ESTScan2 tool. The entire alignment shows contig 3 sequence of APP protein alone with no error at translate level and rest of the 11 protein sequences were left due to some erroneous readings (X, which does not code for somewhat amino acids or refers to a stop codon) in their sequence as shown in **Figure 1**, obtained by CLC Genomics Workbench 7.6. The APP protein sequence of contig 3 is 751 amino acids with a molecular weight of 84818.77 Daltons and this sequence was named as hypothetical protein for further annotation.

Table 3: The InterProScan annotations for Hypothetical protein

Protein	GENE3D	PANTHER	PFAM	PRINTS	PROFILE	SMART	SUPER FAMILY
Contig3 /APP	G3DSA: 3.90.570.10; 1.10.287.510; 4.10.410.10	PTHR23103	PF12925; PF02177; PF12924 and PF00014	PR00204 PS00319; 00280; 00320	PS00319; 000131	SM000006; 56491; 89811; 57362	SF10984;

Conceptual translation of ESTs

The APP protein sequence was reported from 5' ESTs of brain tissues and it belongs to the APP amyloid and beta-APP families of proteins with a distinct N-terminal and C-terminal. The major part of the amyloid plaques found in the brains of AD and peptide regions of 36-43 amino acids are fatefully involved in amyloid precursor protein. A β molecules can aggregate to form oligomers and the resulting amyloid plaques are toxic to nerve cells [18]. N-terminal region of the APP is a member of the heparin-binding class of GFLDs (Growth Factor-Like Domain) and may itself have growth factor function, neuronal development. It contains four structurally similar domains represented by PFAM families PF12925 [14], PF02177 [19], PF12924 [20] and PF00014 [21] as shown in **Table 3**. In structural classification by CATH, the classification lineage of hierarchy 3.90.570.10, 3.30.1490.140, 4.10.230.10 is amyloid beta A4 protein; 1.10.287.510 is amyloid protein and 4.10.410.10 is protease inhibitor IX.

Table 4: Molecular docking analysis of modeled APP protein with synthetic and medicinal compounds

S. No	Compound Name	Glide Score (Kcal/mol)	No. of Hydrogen Bonds	Interacting Residues	Ligand Atom	Distance Length (Å)
APP						
1.	Curcumin	-8.7	5	ARG566:HE	O	2.3
				ARG566:HE	O	2.5
				ARG566:1HH2	O	1.9
				VAL673:O	H	2.7
				VAL673:O	H	2.1
2.	Ginsenoside Rb1	-6.1	2	VAL673:O	H	1.9
3.	Aristolene epoxide	-5.2	2	LYS680:H	O	1.9
4.	Phytol acetate	-3.8	1	PHE745:O	H	2.41
5.	Dimethoxane	-3.6	1	GLU747:O	H	2.21
6.	Valerenone	-3.6	1	SER711:O	H	1.98
				GLY677:O	H	2.01
				LYS680:O	H	1.69

7.	Erucic acid	-3.5	2	LYS680:H	O	2.37
8.	Rivastigmine	-2.9	1	ASP720:O	H	1.80
9.	Tacrine	-1.4	1	LYS600:O	H	1.99
10.	Galantamine	-1.0	1	ASP720:O	H	1.66
11.	Donepezil	-	-	ASP720:O	H	1.51

Note: Hyphen sign (-) denotes no interaction between protein and ligand. Highlighted compound curcumin shows best glide score and more number of hydrogen bonds, best interaction with mutated residues among other compounds.

Molecular modelling of hypothetical protein:

The 3D structure of hypothetical protein of human APP was predicted using MODELLER v9.13. This program was generated ten different 3D modeled structures and validating these structures was considered based on the scoring percentage of the favored regions. Finally, we selected the best modeled structure for hypothetical protein (model 3) as depicted in **Figure 2A**. Validation of Ramachandran plot showed >96% of the residues in most favored and additional allowed regions and the structure of our modeled protein was found to be stable. Verify3D methods evaluate protein structure using 3D profiles and this program analyzed the compatibility of an atomic model (3D) with their possess amino acid sequence (1D). Each residue is allocated a structural class based on the scores ranges from -1 to +1. In our results verify3D score value of modeled APP protein is -1.0 to 0.7 (**Figure 2B**). Validation results showed stero chemical properties and geometrical arrangements of the atoms of the protein was stable. The root-mean-square deviation value of modeled APP protein 3D structure was higher (0.439Å) than the existing crystal structure PDB IDs: 3KTM (2.70Å) and 3NYL (2.80Å) with an energy value of -30227.773KJ/mol.

Molecular docking:

Molecular docking studies were performed for modeled complete sequences of APP protein with current drugs and medicinal compounds. Various synthetic drugs are available against AD such as tacrine, donepezil, rivastigmine and galantamine, but causing side effects like diarrhea, nausea, vomiting etc [22]. Hence, a new drug development is important to cure AD without these side effects. In our study, we have selected 11 medicinal plants such as *R. officinalis* (α -Pinene, Camphene, β -Pinene, 1,8-Cineole, α -Thujone, β -Thujone, Chrysanthenone, Camphor, (+)-Borneol, Bornyl acetate, α -Copaene, Trans-Caryophyllene, α -Humulene, Germacrene-D and (+)- δ -Cadinene) plant essential oils have a potent effect in patients with symptoms of AD [23] and mentioned 15 natural compounds were identified in this plant using GC-MS analysis. *G. biloba* extract from leaves has been found to improve the symptoms of AD [24] and this plant compounds like Quercetin, Kaempferol, Isorhamnetin, Ginkgolide A, B, C, J, M. *P. ginseng* plant extract from root has a potential role in the treatment AD²⁹ and compounds like Ginsenoside Rb1 and Rg1. *C. Longa Linn* plant extract from root have been used to treat of AD and compounds like Curcumin,

Demethoxycurcumin and Bisdemethoxycurcumin [25]. *S. officinalis* extract from leaf has been found a significant benefit in cognition to the patients with mild to moderate AD and compounds like Borneol, Caryophyllene, Linalool. *H. serrata* has been studied extensively for it is role in treating AD and this plant leaves had been extracted to identify compounds like Huperzine A, B and Lycopodine. The essential oil is obtained from leaves of *M. officinalis* compounds 1-Octen-3-ol, 6-Methyl-5-hepten-2-one, Myrcene, (Z)- β -Ocimene, (E)- β -Ocimene, n-Nonanal, Cis-Rose oxide, (+)-Trans-Rose oxide, (+)-Trans-Limonene oxide, Citronellal, Menthol, Isomenthol, Nerol, Neral, Piperitone, Geraniol, Geranial, α -Cubebene, Geranyl acetate, β -Cubebene, β -Caryophyllene, Valencene, Caryophyllene oxide, 1-Hexadecene, n-Eicosane, n-Heneicosane [26] and this plant has been modulate mood and cognitive performance in AD. Compounds like Propane, 1,1-diethoxy-2-methyl-, 2-Nonanone, PhenylethylAlcohol, Amyl nitrite, Dodecanoic acid, 3-tert-Butyl-4-hydroxyanisole, Tetradecanoic acid, n-Hexadecanoic acid, 9-Octadecenal, 1-tridecyne, Oleic acid extracted from *W. somnifera* root [27] are mainly used to treat AD. *B. monnierii* leaf extract has been used to promote memory increasing activity and treat psycho neurological disorders. GC-MS analysis of this plant identified compounds such as 2-octanol, Dimethoxane, 2-Methyl-1-Phenyl-1-butanol, Phytol, Phytol acetate, Octadecanamide [28]. *C. asiatica* plant essential oil extract from leaves and GC-MS analysis compounds like Thujopsene, α -Thujene, Eucalyptol, 3-Nonen-2-one, β -Linalool, L-Camphor, trans-Borneol, α -Terpeneol, Cis-Geraniol, Isobornyl acetate, 7-Tetradecene, β -Elemene, β -Gurjunene, γ -Elemene, Isocaryophyllene, Aromadendrene, β -Farnesene, β -Acoradiene, β -Selinene, α -Selinene, α -Chamigrene, α -Panasinsen, -(β)Spathulenol, Viridiflorol, Valeranone, Isoaromadendrene epoxide, Aristolene epoxide, 1-Naphthalenol. This plant has ability to prevent cognitive deficits treatment for AD. *C. paniculatus* plant contains essential oil extract from seeds and GC-MS analysis compounds like Palmitic acid, Erucic acid, γ -Murolene, Cubenol. The seed oil is studied as best nervine tonic and used in treatment of various neurological disorders [29]. We validated the efficacy of synthetic and medicinal plants based compounds with modeled APP protein using molecular docking approach to identify the best inhibitor for AD.

APP is a transmembrane protein without known function that is constitutively cleaved into peptides during cell metabolism. The amyloidogenic 40 or 42 amino acid A β peptide is released after cleavage by β -secretase and γ -secretase. Familial alzheimer's disease (FAD) mutations have been identified in APP, PSEN1 and PSEN2 genes, which are essential for the generation of A β peptides [30]. Reported APP mutation sequences include A673V [31], V717I [32]. **Figure 3** shows the interaction of modeled APP protein with curcumin having least glide score value of -7.3Kcal/mol and more

number of hydrogen bonds (ARG566, VAL673) were formed than other compounds. From the results of docking study, out of 11 medicinal plant compounds only six medicinal plants such as *P. ginseng* (Ginsenoside Rb1), *C. longa Linn* (Curcumin), *C. asiatica* (Aristolene epoxide, Valerenone), *B. monnierii* (Phytol acetate), *B. monnierii* (Dimethoxane), *C. paniculatus* (Erucic acid) and synthetic (Rivastigmine, Tacrine, Galantamine) compounds showed proper interaction but mutated residues docked with ginsenoside rb1 and curcumin compounds (**Table 4**). Tang and Taghibiglou 2017 [33] has reported curcumin compound to be more effective than current treatment of AD. Alcigir *et al.* [34] found that positive results in new-born rodent pups, curcumin compound as a natural therapy for permanent treatment based on neuronal impairment. Abdolahi *et al.* [35] has considered curcumin compound as a novel promising therapy in migraine prevention. From the molecular interaction study, we conclude that, natural compound curcumin shows better interaction than synthetic, other natural screened compounds and AD approved drugs. Hence we suggested as an alternative lead compound of curcumin in alzheimer's disease research.

Conclusion:

EST analysis of the four genes associated with AD produced 14 contig sequences. APP contig 3, the only contig with no error of translation was annotated using functional and structural data. APP was further analyzed using molecular modeling and docking with natural compound of curcumin, it shows the best glide score of -7.3kcal/mol into mutated residues unlike the synthetic and other natural compounds. Hence to avoid the side effects of synthetic drugs and natural compound, curcumin is suggested for the treatment of AD.

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Conflict of Interest:

The authors confirm that this article content has no conflict of interest.

References:

- [1] Cummings JL *et al.* *New England Journal of Medicine* 2004 **351**:56. [PMID:15229308]
- [2] Singh A *et al.* *CNS Neurol Disord Drug Targets* 2018 **17**:571. [PMID: 29952273]
- [3] Peter KP & Emily RA. *Neuroscience & Medicine* 2011 **2**:120.
- [4] Gatz M *et al.* *Neurobiology of aging* 2005 **26**:439. [PMID: 15653172]
- [5] Tharp WG & Sarkar IN. *BMC genomics* 2013 **14**:1. [PMID: 23627794]
- [6] Patterson C *et al.* *Canadian Medical Association Journal* 2008 **178**:548. [PMID: 18299540]
- [7] Lemere CA *et al.* *The American journal of pathology* 2004 **165**:283. [PMID: 15215183]
- [8] Huang X & Madan A. *Genome research* 1999 **9**:868. [PMID: 10508846]
- [9] Kent WJ. *Genome research* 2002 **12**:656. [PMID: 11932250]
- [10] Iseli C *et al.* *In ISMB* 1999 **99**:138. [PMID: 10786296]
- [11] Quevillon E *et al.* *Nucleic acids research* 2005 **33**:116. [PMID: 15980438]
- [12] Šali A & Blundell TL. *Journal of molecular biology* 1993 **234**:779. [PMID: 8254673]
- [13] Dahms SO *et al.* *Proceedings of the National Academy of Sciences* 2010 **107**:5381. [PMID: 20212142]
- [14] Wang Y & Ha Y. *Molecular cell* 2004 **15**:343. [PMID: 15304215]
- [15] Barrett PJ *et al.* *Science* 2012 **336**:1168. [PMID: 22654059]
- [16] Laskowski RA *et al.* *Journal of applied crystallography* 1993 **26**:283.
- [17] GLIDE, Version 5.8, Schrodinger, LLC, New York, 2012
- [18] Pulawski W *et al.* *Applied biochemistry and biotechnology* 2012 **166**:1626. [PMID: 22350870]
- [19] Rossjohn J *et al.* *Nature Structural & Molecular Biology* 1999 **6**:327. [PMID: 10201399]
- [20] Kong GKW *et al.* *Journal of molecular biology* 2007 **367**:148. [PMID: 17239395]
- [21] Arnoux B *et al.* *Acta Crystallographica Section D: Biological Crystallography* 2002 **58**:1252. [PMID: 12077460]
- [22] Hake AM. *Cleveland Clinic journal of medicine* 2001 **68**:608. [PMID: 11453078]
- [23] Ozarowski M *et al.* *Herba Polonica* 2009 **55**:148.
- [24] Li H *et al.* *Nutrients* 2018 **10**:589. [PMID: 29747475]
- [25] Iriti M *et al.* *Molecules* 2010 **15**:3517. [PMID: 20657497]
- [26] Abdellatif F *et al.* *EXCLI Journal* 2014 **13**:772. [PMID: 26417300]
- [27] Kumar S *et al.* *Int. J. Pharm. Phytochem. Res* 2011 **3**:31.
- [28] Subashri B & Pillai YJ. *Int J Pharm Pharm Sci* 2014 **6**:494.
- [29] Arora N & Pandey-Rai S. *Industrial Crops and Products* 2014 **61**:345.
- [30] De Strooper B *et al.* *Nature Reviews Neurology* 2010 **6**:99. [PMID: 20139999]
- [31] Di Fede G *et al.* *Science* 2009 **323**:1473. [PMID: 19286555]
- [32] Goate A *et al.* *Nature* 1991 **349**:704. [PMID: 1671712]
- [33] Tang M & Changiz T. *Journal of Alzheimer's Disease* 2017 **58**:1003. [PMID: 28527218]
- [34] Alcigir ME *et al.* *CNS Neurol Disord Drug Targets* 2018 **17**:132. [PMID: 29546838]

[35] Abdolahi M *et al.* *CNS Neurol Disord Drug Targets* 2018

17:430. [PMID: 29938621]



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Supplementary material: Table 5 ESTs data for APP, BACE1, PSEN1 and PSEN2 genes (Page 1)

No.	GB accession No.	Description (clone)	Tissue Type	EST Type	Code	S. No.	GB accession No.	Description (clone)	Tissue Type	EST Type	Code	S. No.	GB accession No.	Description (clone)	Tissue Type	EST Type	Code
1.	RG1959	IMAGE:34389	brain	5' read	P	94.	DA1169013	BRACE0306402	brain	5' read	P	190.	DA3288171	BRHP1018158	brain	5' read	P
2.	RG2570	IMAGE:36360	brain	5' read	P	95.	DA2404941	BRAH1030525	brain	5' read	P	191.	DA2788071	BRCA2006262	brain	5' read	P
3.	RG2913	IMAGE:36445	brain	5' read	-	96.	DA3916614	BRAWH1030540	brain	5' read	P	192.	DA3751611	BRAMY1018150	brain	5' read	P
4.	CDS15501A1	CD288079005	brain	5' read	P	97.	DA288079005	BRAWH1030579	brain	5' read	P	193.	DA3720497	BRAWH10304073	brain	5' read	P
5.	RG2914	IMAGE:42589	brain	5' read	-	98.	DA3808911	BRAWH1030598	brain	5' read	P	194.	DA3720497	BRAWH10304159	brain	5' read	P
6.	AL056477	DKRZ:554/B1262	brain	5' read	P	99.	DA0554113	BRAMY1000405	brain	5' read	P	195.	DA3720497	BRAMY10203599	brain	5' read	P
7.	RG348162	CS09W0327001	brain	5' read	P	100.	DA1929931	BRAMY1000465	brain	5' read	P	196.	DA2588013	BRAMY10304469	brain	5' read	P
8.	RG348863	CS09W0317015	brain	5' read	P	101.	DA1329041	BRHP10302211	brain	5' read	P	197.	DA0553933	BRAC15003794	brain	5' read	P
9.	RG348163	CS09W0317015	brain	5' read	P	102.	DA1054501	BRAMy0200083	brain	5' read	P	198.	DA3233153	BRAL23000912	brain	5' read	P
10.	RG411752	CS09W0317015	brain	5' read	P	103.	DA1054761	BRAMY0200028	brain	5' read	P	199.	DA0553933	BRAC13004688	brain	5' read	P
11.	RG3426012	CS09W0327001	brain	5' read	P	104.	DA3677941	BRAC20212159	brain	5' read	-	200.	DA1321741	BRAC13051605	brain	5' read	P
12.	RG410063	CS09W0317015	brain	5' read	P	105.	DA3455761	BRAWH10303541	brain	5' read	P	201.	DA1328251	BRAC13051778	brain	5' read	P
13.	RG4184112	CS09W0401116	brain	5' read	P	106.	DA1899611	BRAMY20103170	brain	5' read	P	202.	DA1328301	BRAC13051853	brain	5' read	P
14.	RG4105812	CS09N005YC22	brain	5' read	P	107.	DA3443611	BRAWH10306191	brain	5' read	P	203.	DA1328721	BRAC13051927	brain	5' read	P
15.	CDS154463	IMAGE:30176636	brain	5' read	-	108.	DA1891841	BRAMY3010051	brain	5' read	P	204.	DA1328101	BRAC13052122	brain	5' read	P
16.	RG4120123	IMAGE:30374500	brain	5' read	P	109.	DA1890311	BRAMY30101070	brain	5' read	P	205.	DA1328651	BRH102003440	brain	5' read	P
17.	RG462052	CS09R0387013	brain	5' read	P	110.	DA1628441	BRAY2023095	brain	5' read	P	206.	DA1289271	BRAC13052651	brain	5' read	P
18.	RG413112	CS09D0525A20	brain	5' read	P	111.	DA2784031	BRAWH103022731	brain	5' read	P	207.	DA1802021	BRAMY20404844	brain	5' read	P
19.	CDS145301	IMAGE:30195742	brain	5' read	P	112.	DA1890641	BRAMY3012122	brain	5' read	P	208.	DA0739751	ASTRO2006466	brain	5' read	P
20.	CDS162981	IMAGE:30408890	brain	5' read	P	113.	DA2827061	BRAWH10302307	brain	5' read	P	209.	DA3656531	BRWH103045531	brain	5' read	P
21.	CDS167691	IMAGE:30198693	brain	5' read	P	114.	DA2787851	BRAMY10302728	brain	5' read	P	210.	DA3668571	BRWH103045894	brain	5' read	P
22.	CD518169	IMAGE:30987272	brain	5' read	P	115.	DA2019371	BRHW10302781	brain	5' read	P	211.	DA1746821	BRAMY2037864	brain	5' read	P
23.	CDS187601	IMAGE:30198647	brain	5' read	-	116.	DA1730201	BRAC3045131	brain	5' read	P	212.	DA1324651	BRWH10303271	brain	5' read	P
24.	CG000751	IMAGE:30721155	brain	5' read	P	117.	DA1323561	BRAC3045205	brain	5' read	P	213.	DA0681051	BRACE3006174	brain	5' read	P
25.	CK003561	IMAGE:30720873	brain	5' read	P	118.	DA1292971	BRAC306669	brain	5' read	P	214.	DA1689631	BRAMY2030435	brain	5' read	P
26.	CK003991	IMAGE:30719468	brain	5' read	-	119.	DA1257151	BRAC3045706	brain	5' read	P	215.	DA1689481	BRAMY20302724	brain	5' read	P
27.	CK00407021	IMAGE:30723098	brain	5' read	P	120.	DA1127211	BRAC3070318	brain	5' read	P	216.	DA0277121	BRWH12014849	brain	5' read	P
28.	CDS053241	IMAGE:30722169	brain	5' read	A	121.	DA1128271	BRAC30703042	brain	5' read	P	217.	DA2096411	BRWH102015212	brain	5' read	P
29.	CK0058771	IMAGE:30721285	brain	5' read	P	122.	DA1128931	BRAC30703736	brain	5' read	P	218.	DA0302021	ASTRO201322	brain	5' read	P
30.	CDS204561	IMAGE:307201921	brain	5' read	-	123.	DA1326511	BRHP10302712	brain	5' read	P	219.	DA1052651	BRACE3012521	brain	5' read	P
31.	H282321	IMAGE:51381	brain	5' read	P	124.	DA3127271	BRHP10302726	brain	5' read	P	220.	DA3251921	BRH102002063	brain	5' read	P
32.	RP1901815	AD070016	brain	5' read	P	125.	DA2748031	BRCC20008999	brain	5' read	P	221.	DA2890311	BRCCOC2017878	brain	5' read	P
33.	CK0018613	AD070016	brain	5' read	P	126.	DA1461621	BRAC306669	brain	5' read	P	222.	DA0204971	ASTRO2007682	brain	5' read	P
34.	RP1900401	AMR042423	brain	5' read	P	127.	DA3138881	BRHP10300301	brain	5' read	P	223.	DA2077411	BRWH13012625	brain	5' read	P
35.	BP2045631	AMR02345	brain	5' read	P	128.	DA1961581	BRAMY001764	brain	5' read	P	224.	DA1308841	BRHP1021231	brain	5' read	P
36.	BP2040431	AMR04293	brain	5' read	P	129.	DA3072921	BRHP102042092	brain	5' read	P	225.	DA2132301	BRWH103044315	brain	5' read	P
37.	BP2025701	AMR080808	brain	5' read	P	130.	DA1963631	BRAMY0020014	brain	5' read	P	226.	DA1244841	BRWH103004525	brain	5' read	P
38.	BP2028031	AMR06620	brain	5' read	P	131.	DA3531891	BRAWH10301578	brain	5' read	P	227.	DA2749091	BRWH10302761	brain	5' read	P
39.	BP228671	FBR00557	brain	5' read	P	132.	DA1962428	BRAC30020090	brain	5' read	P	228.	DA1030921	BRACE3015891	brain	5' read	P
40.	BP2496531	H1R01011	brain	5' read	P	133.	DA1963821	BRAC30020290	brain	5' read	P	229.	DA0988931	BRCE3006663	brain	5' read	P
41.	BP2510151	H1R01849	brain	5' read	P	134.	DA3553511	BRAWH10303203	brain	5' read	P	230.	DA1036121	BRACE3016663	brain	5' read	P
42.	BP258671	H1R04503	brain	5' read	P	135.	DA1633991	BRAMY20200406	brain	5' read	P	231.	DA3333911	BRHP10304095	brain	5' read	P
43.	BP2590011	H1R05341	brain	5' read	P	136.	DA3577011	BRAY10302217	brain	5' read	P	232.	DA2870811	BRCCOC2018089	brain	5' read	P
44.	BP2510441	H1R06069	brain	5' read	P	137.	DA0853931	BRAC20139695	brain	5' read	P	233.	DA2609011	BRCAN2012243	brain	5' read	P
45.	BP2515484	H1R08028	brain	5' read	P	138.	DA1890511	BRAC30307953	brain	5' read	P	234.	DA3261611	BRHP1021491	brain	5' read	P
46.	BP2516061	H1R08878	brain	5' read	P	139.	DA2464141	BRAC30402748	brain	5' read	P	235.	DA2210421	BRWH103012957	brain	5' read	P
47.	BP2101871	CBL04159	brain	5' read	P	140.	DA3181821	BRCC2009860	brain	5' read	P	236.	DA2111121	BRWH103013042	brain	5' read	P
48.	BP2105241	CB004258	brain	5' read	P	141.	DA1866001	BRAC3080726	brain	5' read	P	237.	DA2482531	BRWH10307637	brain	5' read	P
49.	BP2145711	CC001118	brain	5' read	P	142.	DA2409811	BRAWH10309946	brain	5' read	P	238.	DA2162621	BRAWH10301371	brain	5' read	P
50.	BP2166751	CCR0755	brain	5' read	P	143.	DA3426711	BRAY103040222	brain	5' read	P	239.	DA2105761	BRWH103005331	brain	5' read	P
51.	BP2176741	CRN01789	brain	5' read	P	144.	DA1701161	BRAY103023090	brain	5' read	P	240.	DA2486011	BRAWH10300809	brain	5' read	P
52.	BP2180301	CRN02054	brain	5' read	P	145.	DA2757471	BRCC2002473	brain	5' read	P	241.	DA1749441	BRAMY20204056	brain	5' read	P
53.	BP2192841	CRN02575	brain	5' read	P	146.	DA3465011	BRHP10304586	brain	5' read	P	242.	DA2488011	BRAWH103048341	brain	5' read	P
54.	BP2192894	CRN0974	brain	5' read	P	147.	DA3693421	BRAC2011472	brain	5' read	P	243.	DA3104241	BRWH102005772	brain	5' read	P
55.	BP2142441	CRN04373	brain	5' read	P	148.	DA3427451	BRAWH10304801	brain	5' read	P	244.	DA1434651	BRALZ101944	brain	5' read	P
56.	BP1881331	CRN05756	brain	5' read	P	149.	DA1253011	BRAC3004035	brain	5' read	P	245.	DA2608231	BRACE30107246	brain	5' read	P
57.	BP2188271	CRN05940	brain	5' read	P	150.	DA1581751	BRAC30040355	brain	5' read	P	246.	DA2941101	BRHP102006484	brain	5' read	P
58.	BP1891991	CRN06941	brain	5' read	P	151.	DA1889761	BRAMY2018659	brain	5' read	P	247.	DA0319031	ASTRO2016830	brain	5' read	P
59.	BP2191761	CRN07124	brain	5' read	P	152.	DA1910101	BRAC3039240	brain	5' read	P	248.	DA2483531	BRHP10302135	brain	5' read	P
60.	BP2194161	CRN07868	brain	5' read	P	153.	DA2007614	ASTRO201340	brain	5' read	P	249.	DA3204211	BRHP10201435	brain	5' read	P
61.	BP2661614	TH1R01187	brain	5' read	P	154.	DA2823194	BRAC2011472	brain	5' read	P	250.	DA2509241	BRHP10202942	brain	5' read	P
62.	BP2661221	TH1R01275	brain	5' read	P	155.	DA2295921	BRAC3040748	brain	5' read	P	251.	DA2584841	BRCCOC2003035	brain	5' read	P
63.	BP2661154	TH1R01345	brain	5' read	P	156.	DA1918951	BRAC3040402	brain	5' read	P	252.	DA3007421	BRHP10215238	brain	5' read	P
64.	BP2662811	TH1R01522	brain	5' read	P	157.	DA2212231	BRAMY2034535	brain	5' read	P	253.	DA3008521	BRHP10215395	brain	5' read	P
65.	BP2661911	KG015956	brain	5' read	P	158.	DA2664241	BRAC3049937	brain	5' read	P	254.	DA1834041</				

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286.	DA317310_1	BRAW1H3008184	brain	5' read	P	382	DA311857_1	BRAH1P3000628	brain	5' read	P	478	DA411093_1	BRTTHA3018133	brain	5' read	P
287.	DA317305_1	BRAW1H3008217	brain	5' read	P	383	DA309502_1	BRAH1P3000615	brain	5' read	P	479	DA410923_1	BRTTHA3018705	brain	5' read	P
288.	DA317305_1	BRAW1H3008283	brain	5' read	P	384	DA309502_1	BRAH1P3000619	brain	5' read	P	480	DA410924_1	BRTTHA3018869	brain	5' read	P
289.	DA317305_1	BRAW1H3008488	brain	5' read	P	385	DA309502_1	BRAH1P3000624	brain	5' read	P	481	DA410924_1	BRTTHA3019144	brain	5' read	P
290.	DA317305_1	BRAW1H3008508	brain	5' read	P	386	DA309502_1	BRAH1P3000624	brain	5' read	P	482	DA410924_1	BRTTHA3019152	brain	5' read	P
291.	DA3113129_1	BRACT3028973	brain	5' read	P	387	DA309502_1	BRAH1P3000624	brain	5' read	P	483	DA410924_1	BRTTHA3019164	brain	5' read	P
292.	DA319520_1	BRAH1P3002929	brain	5' read	P	388	DA309502_1	BRAH1P3000624	brain	5' read	P	484	DA410924_1	BRTTHA3019709	brain	5' read	P
293.	DA319520_1	BRAH1P3005936	brain	5' read	P	389	DA309502_1	BRAH1P3000624	brain	5' read	P	485	DA410924_1	BRTTHA3019752	brain	5' read	P
294.	DA319520_1	BRAH1P301073	brain	5' read	P	390	DA309502_1	BRAH1P3000624	brain	5' read	P	486	DA410924_1	BRTTHA3019806	brain	5' read	P
295.	DA319520_1	BRAH1P301426	brain	5' read	P	391	DA309502_1	BRAH1P3000624	brain	5' read	P	487	DA410924_1	BCBHF3001099	brain	5' read	P
296.	DA319520_1	BRAH1P3015083	brain	5' read	P	392	DA309502_1	BRAH1P3000624	brain	5' read	P	488	DA410924_1	BCBHF3001331	brain	5' read	P
297.	DA319520_1	BRAH1P3016512	brain	5' read	P	393	DA309502_1	BRAH1P3000624	brain	5' read	P	489	DA410924_1	FEHRFA201645	brain	5' read	P
298.	DA319520_1	BRAH1P3016512	brain	5' read	P	394	DA309502_1	BRAH1P3000624	brain	5' read	P	490	DA410924_1	BRTTHA3019944	brain	5' read	P
299.	DA319520_1	BRAH1P3016513	brain	5' read	P	395	DA309502_1	BRAH1P3000624	brain	5' read	P	491	DA410924_1	BRTTHA3019951	brain	5' read	P
300.	DA319520_1	BRAH1P3016513	brain	5' read	P	396	DA309502_1	BRAH1P3000624	brain	5' read	P	492	DA410924_1	BRTTHA3020370	brain	5' read	P
301.	DA319520_1	BRAH1P3016513	brain	5' read	P	397	DA309502_1	BRAH1P3000624	brain	5' read	P	493	DA410924_1	BRTTHA3020436	brain	5' read	P
302.	DA319520_1	BRAH1P3016513	brain	5' read	P	398	DA309502_1	BRAH1P3000624	brain	5' read	P	494	DA410924_1	BRTTHA3021792	brain	5' read	P
303.	DA319520_1	BRAH1P3016515	brain	5' read	P	399	DA309502_1	BRAH1P3000624	brain	5' read	P	495	DA410924_1	BRTTHA3021880	brain	5' read	P
304.	DA319520_1	BRAH1P3016560	brain	5' read	P	400	DA309502_1	BRAH1P3000624	brain	5' read	P	496	DA410924_1	FCBHF3020221	brain	5' read	P
305.	DA319520_1	BRACT2013042	brain	5' read	P	401	DA309502_1	BRACT205589	brain	5' read	P	497	DA410924_1	FEHRFA202785	brain	5' read	P
306.	DA319520_1	BRACT2013052	brain	5' read	P	402	DA309502_1	BRAH1P3006013	brain	5' read	P	498	DA410924_1	BRTTHA3021010	brain	5' read	P
307.	DA319520_1	BRAH1H3009194	brain	5' read	P	403	DA309502_1	BRAH1P3000618	brain	5' read	P	499	DA410924_1	BRTTHA3021206	brain	5' read	P
308.	DA319520_1	BRAH1H3009224	brain	5' read	P	404	DA309502_1	BRAH1P3000910	brain	5' read	P	500	DA410924_1	BRTTHA3021620	brain	5' read	P
309.	DA319520_1	BRAH1P3005440	brain	5' read	P	405	DA309502_1	FCBHF300978	brain	5' read	P	501	DA410924_1	BRTTHA3021740	brain	5' read	P
310.	DA319520_1	BRAH1P3005440	brain	5' read	P	406	DA309502_1	FCBHF3016894	brain	5' read	P	502	DA410924_1	BRTTHA3022124	brain	5' read	P
311.	DA319520_1	BRAH1P3013052	brain	5' read	P	407	DA309502_1	FCBHF3017033	brain	5' read	P	503	DA410924_1	BRTTHA3023067	brain	5' read	P
312.	DA319520_1	BRAH1H3010147	brain	5' read	P	408	DA309502_1	BRAH1H3009584	brain	5' read	P	504	DA410924_1	H1D9X312	brain	5' read	P
313.	DA319520_1	BRAH1H3010149	brain	5' read	P	409	DA309502_1	BRAHSN2015364	brain	5' read	P	505	DA410924_1	H1D9X328	brain	5' read	P
314.	DA319520_1	BRAH1H3010149	brain	5' read	P	410	DA309502_1	BRAH1P3027842	brain	5' read	P	506	DA410924_1	H1D9U710	brain	5' read	P
315.	DA319520_1	BRAH1H3010149	brain	5' read	P	411	DA309502_1	BRAH1P3027972	brain	5' read	P	507	DA410924_1	H1D9AQ65	brain	5' read	P
316.	DA319520_1	BRAH1H3010175	brain	5' read	P	412	DA309502_1	BRAH1P3028616	brain	5' read	P	508	DA410924_1	H1TCAD305	brain	5' read	P
317.	DA319520_1	BRAH1P3013179	brain	5' read	P	413	DA309502_1	BRAH1P3028661	brain	5' read	P	509	DA410924_1	BRTTHA3015682	brain	5' read	P
318.	DA319520_1	BRAH1P3027818	brain	5' read	P	414	DA309502_1	BRAH1P3010714	brain	5' read	P	510	DA410924_1	FEHRFA2023446	brain	5' read	P
319.	DA319520_1	BRAH1P301816	brain	5' read	P	415	DA309502_1	BRAH1P3010761	brain	5' read	P	511	DA410924_1	FCBHF3020849	brain	5' read	P
320.	DA319520_1	BRAH1P3002885	brain	5' read	P	416	DA309502_1	BRAH1A2030863	brain	5' read	P	512	DA410924_1	BRAH1A2030863	brain	5' read	P
321.	DA319520_1	BRACT3022737	brain	5' read	P	417	DA309502_1	BCHB1H3018897	brain	5' read	P	513	DA410924_1	FEHRFA2036519	brain	5' read	P
322.	DA319520_1	BRACT3022737	brain	5' read	P	418	DA309502_1	BRAH1P3023624	brain	5' read	P	514	DA410924_1	BRTTHA3022693	brain	5' read	P
323.	DA319520_1	BRACT3022738	brain	5' read	P	419	DA309502_1	BRAH1P3022089	brain	5' read	P	515	DA410924_1	BRTTHA3023267	brain	5' read	P
324.	DA319520_1	BRAH1H3012630	brain	5' read	P	420	DA309502_1	BRAH1P3023032	brain	5' read	P	516	DA410924_1	IMR520207845	brain	5' read	P
325.	DA319520_1	BRAH1P3005626	brain	5' read	P	421	DA309502_1	BRAH1P3026664	brain	5' read	P	517	DA410924_1	BRTTHA3023445	brain	5' read	P
326.	DA319520_1	BRAH1P3010955	brain	5' read	P	422	DA309502_1	BRAH1P3027134	brain	5' read	P	518	DA410924_1	BRTTHA3021424	brain	5' read	P
327.	DA319520_1	BRAH1P3008977	brain	5' read	P	423	DA309502_1	BRAH1P3027135	brain	5' read	P	519	DA410924_1	FCBHF3020870	brain	5' read	P
328.	DA319520_1	BRAH1P3008154	brain	5' read	P	424	DA309502_1	BRAH1P3021165	brain	5' read	P	520	DA410924_1	BRTTHA3021513	brain	5' read	P
329.	DA319520_1	BRAH1P3008159	brain	5' read	P	425	DA309502_1	BRAH1P3021063	brain	5' read	P	521	DA410924_1	BRTTHA3022693	brain	5' read	P
330.	DA319520_1	BRAH1P3008159	brain	5' read	P	426	DA309502_1	BRAH1P3021761	brain	5' read	P	522	DA410924_1	BRTTHA3022739	brain	5' read	P
331.	DA319520_1	BRAH1H30102230	brain	5' read	P	427	DA309502_1	BRAH1P3021910	brain	5' read	P	523	DA410924_1	BRAH1P3020640	brain	5' read	P
332.	DA319520_1	BRAH1H30102230	brain	5' read	P	428	DA309502_1	BRAH1P3023922	brain	5' read	P	524	DA410924_1	FCBHF30205198	brain	5' read	P
333.	DA319520_1	BRAH1H30101587	brain	5' read	P	429	DA309502_1	BRAH1P3011688	brain	5' read	P	525	DA410924_1	IMR320210575	brain	5' read	P
334.	DA319520_1	BRAH1H30101587	brain	5' read	P	430	DA309502_1	BRAH1P3020297	brain	5' read	P	526	DA410924_1	BRAH1P3020297	brain	5' read	P
335.	DA319520_1	BRAH1P3012161	brain	5' read	P	431	DA309502_1	BRAH1P3003161	brain	5' read	P	527	DA410924_1	BRTTHA3026868	brain	5' read	P
336.	DA319520_1	BRAH1P3014275	brain	5' read	P	432	DA309502_1	BRAH1P3003115	brain	5' read	P	528	DA410924_1	FCBHF3021115	brain	5' read	P
337.	DA319520_1	BRAH1P3006179	brain	5' read	P	433	DA309502_1	BRAH1P3004243	brain	5' read	P	529	DA410924_1	FEHRFA2035226	brain	5' read	P
338.	DA319520_1	BRAH1P3006194	brain	5' read	P	434	DA309502_1	BRAH1P3003232	brain	5' read	P	530	DA410924_1	FCBHF3003089	brain	5' read	P
339.	DA319520_1	BRAH1H3019103	brain	5' read	P	435	DA309502_1	BRAH1P3024425	brain	5' read	P	531	DA410924_1	BRTTHA3015501	brain	5' read	P
340.	DA319520_1	BRAH1P3006001	brain	5' read	P	436	DA309502_1	BRAH1P3003814	brain	5' read	P	532	DA410924_1	BRAH1P3015858	brain	5' read	P
341.	DA319520_1	BRACT3016260	brain	5' read	P	437	DA309502_1	BRAH1P3003120	brain	5' read	P	533	DA410924_1	BRTTHA3016026	brain	5' read	P
342.	DA319520_1	BRAH1H3011127	brain	5' read	P	438	DA309502_1	BRAH1P3004081	brain	5' read	P	534	DA410924_1	BRTTHA3016079	brain	5' read	P
343.	DA319520_1	BRAH1P3003340	brain	5' read	P	439	DA309502_1	BRAH1P3024867	brain	5' read	P	535	DA410924_1	FCBHF3020327	brain	5' read	P
344.	DA319520_1	BRAH1P3003340	brain	5' read	P	440	DA309502_1	BRAH1P3003277	brain	5' read	P	536	DA410924_1	FCBHF3003027	brain	5' read	P
345.	DA319520_1	BRAH1P3003341	brain	5' read	P	441	DA309502_1	BRAH1P3003940	brain	5' read	P	537	DA410924_1	FCBHF3003712	brain	5' read	P
346.	DA319520_1	BRAH1P3003341	brain	5' read	P	442	DA309502_1	BRAH1P3025411	brain	5' read	P	538	DA410924_1	FCBHF300307304	brain	5' read	P
347.	DA319520_1	BRAH1P3012620	brain	5' read	P	443	DA309502_1	BRAH1P3005646	brain	5' read	P	539	DA410924_1	FCBHF300309681	brain	5' read	P
348.	DA319520_1	BRAH1P3003971	brain	5' read													

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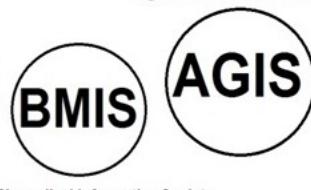
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586.	GB075573_1	OCCBBE3021600	brain	5 read	P	682.	BG328542_1	IMAGE:53000177	brain	5 read	P	82.	BG328774_1	14032026M02	brain	5 read	P
587.	GB075574_1	OCCBBE3023933	brain	5 read	P	683.	BG328543_1	IMAGE:5301766	brain	5 read	P	83.	BG328775_1	14032026M02	brain	5 read	P
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592.	GB075579_1	IMAG:4327534	brain	5 read	P	688.	BG328548_1	IMAGE:5307974	brain	5 read	P	88.	BG328780_1	14032026M02	brain	5 read	P
593.	GB075580_1	OCCBBE3025952	brain	5 read	P	689.	BG328549_1	IMAGE:5326214	brain	5 read	P	89.	BG328781_1	14032026M02	brain	5 read	P
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595.	GB075582_1	OCCBBE3025895	brain	5 read	P	691.	BG328551_1	IMAGE:5327438	brain	5 read	P	91.	BG328783_1	14032026M02	brain	5 read	P
596.	GB075583_1	OCCBBE3025895	brain	5 read	P	692.	BG328552_1	IMAGE:53198424	brain	5 read	P	92.	BG328784_1	14032026M02	brain	5 read	P
597.	GB075584_1	OCCBBE3025959	brain	5 read	P	693.	BG328553_1	IMAGE:5327236	brain	5 read	P	93.	BG328785_1	14032026M02	brain	5 read	P
598.	GB075585_1	OCCBBE3025899	brain	5 read	P	694.	BG328554_1	IMAGE:5330712	brain	5 read	P	94.	BG328786_1	14032026M02	brain	5 read	P
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600.	GB075587_1	OCCBBE3021514	brain	5 read	P	696.	BG328556_1	IMAGE:5327211	brain	5 read	P	96.	BG328788_1	14032026M02	brain	5 read	P
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602.	GB075589_1	OCCBBE3014446	brain	5 read	P	698.	BG328558_1	IMAGE:5327153	brain	5 read	P	98.	BG328790_1	14032026M02	brain	5 read	P
603.	GB075590_1	OCCBBE30214773	brain	5 read	P	699.	BG328559_1	IMAGE:5309317	brain	5 read	P	99.	BG328791_1	14032026M02	brain	5 read	P
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606.	GB075593_1	OCCBBE30219444	brain	5 read	P	702.	BG328562_1	HIBHU95	brain	5 read	P	102.	BG328794_1	14032026M02	brain	5 read	P
607.	GB075594_1	OCCBBE30203495	brain	5 read	P	703.	BG328563_1	PSEN1	brain	5 read	P	103.	BG328795_1	14032026M02	brain	5 read	P
608.	GB075595_1	OCCBBE3023867	brain	5 read	P	704.	BG328564_1	IMAGE:5309391	brain	5 read	P	104.	BG328796_1	14032026M02	brain	5 read	P
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611.	GB075598_1	OCCBBE3025895	brain	5 read	P	707.	BG328567_1	IMAGE:5326014	brain	5 read	P	107.	BG328799_1	14032026M02	brain	5 read	P
612.	GB075599_1	OCCBBE3027204	brain	5 read	P	708.	BG328568_1	IMAG:4302635	brain	5 read	P	108.	BG328800_1	14032026M02	brain	5 read	P
613.	GB075600_1	OCCBBE30301500	brain	5 read	P	709.	BG328569_1	IMAGE:5317509	brain	5 read	P	109.	BG328801_1	14032026M02	brain	5 read	P
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616.	GB075603_1	OCCBBE30264554	brain	5 read	P	712.	BG328572_1	IMAGE:43184140	brain	5 read	P	112.	BG328804_1	14032026M02	brain	5 read	P
617.	GB075604_1	OCCBBE30264559	brain	5 read	P	713.	BG328573_1	IMAGE:4315661	brain	5 read	P	113.	BG328805_1	14032026M02	brain	5 read	P
618.	GB075605_1	OCCBBE30264596	brain	5 read	P	714.	BG328574_1	IMAGE:4320321	brain	5 read	P	114.	BG328806_1	14032026M02	brain	5 read	P
619.	GB075606_1	OCCBBE30264596	brain	5 read	P	715.	BG328575_1	IMAGE:4320321	brain	5 read	P	115.	BG328807_1	14032026M02	brain	5 read	P
620.	GB075607_1	OCCBBE30264596	brain	5 read	P	716.	BG328576_1	IMAGE:4320321	brain	5 read	P	116.	BG328808_1	14032026M02	brain	5 read	P
621.	GB075608_1	OCCBBE30264596	brain	5 read	P	717.	BG328577_1	IMAGE:4320321	brain	5 read	P	117.	BG328809_1	14032026M02	brain	5 read	P
622.	GB075609_1	OCCBBE30264596	brain	5 read	P	718.	BG328578_1	IMAGE:4320321	brain	5 read	P	118.	BG328810_1	14032026M02	brain	5 read	P
623.	GB075610_1	OCCBBE30264596	brain	5 read	P	719.	BG328579_1	IMAGE:4320321	brain	5 read	P	119.	BG328811_1	14032026M02	brain	5 read	P
624.	GB075611_1	OCCBBE30264596	brain	5 read	P	720.	BG328580_1	IMAGE:4320321	brain	5 read	P	120.	BG328812_1	14032026M02	brain	5 read	P
625.	GB075612_1	OCCBBE3000009	brain	5 read	P	721.	BG328581_1	IMAGE:432040207	brain	5 read	P	121.	BG328813_1	14032026M02	brain	5 read	P
626.	GB075613_1	OCCBBE3027218	brain	5 read	P	722.	BG328582_1	IMAG:432014575	brain	5 read	P	122.	BG328814_1	14032026M02	brain	5 read	P
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628.	GB075615_1	OCCBBE302903112	brain	5 read	P	724.	BG328584_1	IMAGE:432032152	uterus	5 read	P	124.	BG328816_1	14032026M02	uterus	5 read	P
629.	GB075616_1	OCCBBE3029040103	brain	5 read	P	725.	BG328585_1	IMAGE:432021166	uterus	5 read	P	125.	BG328817_1	14032026M02	uterus	5 read	P
630.	GB075617_1	OCCBBE302904103	brain	5 read	P	726.	BG328586_1	IMAGE:432021166	uterus	5 read	P	126.	BG328818_1	14032026M02	uterus	5 read	P
631.	GB075618_1	OCCBBE302904103	brain	5 read	P	727.	BG328587_1	IMAGE:432021166	uterus	5 read	P	127.	BG328819_1	14032026M02	uterus	5 read	P
632.	GB075619_1	OCCBBE302904103	brain	5 read	P	728.	BG328588_1	IMAGE:432021166	uterus	5 read	P	128.	BG328820_1	14032026M02	uterus	5 read	P
633.	GB075620_1	OCCBBE302904103	brain	5 read	P	729.	BG328589_1	IMAGE:432021166	uterus	5 read	P	129.	BG328821_1</a				

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64.	DA2099901.1	BRAW120170959	brain	5' read	P	2	C9003561.1	IMAGE-30720023	brain	5' read	
65.	BR055101.1	BRCOC2019518	brain	5' read	P	3	BR055102.1	BRAW20235304	brain	5' read	P
66.	BR0424700.1	BRIH120172359	brain	5' read	P	4	BR04247002.1	BRAW20235379	brain	5' read	
67.	BR04247002.1	BRCOC2020217	brain	5' read	P	5	BR04247003.1	BRAZ1201203	brain	5' read	P
68.	BR0501452.1	IMAGE-2481578	brain	5' read	-	6	BR05014522.1	BNSN2009483	brain	5' read	P
69.	BR0370205.1	IMAGE-2486283	brain	5' read	-	7	BR03702053.1	BRIH12025890	brain	5' read	
70.	BR037205.1	IMAGE-2486717	brain	5' read	-	8	BR0372053.1	BRAW2033218	brain	5' read	
71.	QA144712.1	BRAMY2000995	brain	5' read	P	9	QA144712.1	BRAW12002949	brain	5' read	
72.	QA144712.1	BRAW12001292	brain	5' read	-	10	QA144712.1	BRAW12002949	brain	5' read	
73.	QA250504.1	BRAH1201208393	brain	5' read	P	11	QA2505042.1	BNGH12001566	brain	5' read	
74.	QA2505042.1	BRAH1201208399	brain	5' read	P	12	QA2505043.1	BNGH12001601	brain	5' read	
75.	QA337327.1	BRSN2000475	brain	5' read	P	13	QA337327.1	BRCOC2013488	brain	5' read	P
76.	QA2505043.1	BRAH1201209200	brain	5' read	P	14	QA2505043.1	BNGH12001672	brain	5' read	
77.	QA2505043.1	BRCAN2015700	brain	5' read	P	15	QA2505043.1	BRAZ20009922	brain	5' read	P
78.	QA2505043.1	IMAGE-2486528	brain	5' read	-	16	QA2505043.1	BRIH12003271	brain	5' read	
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83.	QA2505043.1	BRCAN2009051	brain	5' read	P	21	QA2505043.1	BRSN2002870	brain	5' read	
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85.	QA1446372.1	BRAW12002449	brain	5' read	P	23	QA3531453.1	BRSN2017180	brain	5' read	
86.	QA207854.1	BRAH12011089	brain	5' read	P	24	QA3531453.1	BRIH1202032125	brain	5' read	
87.	QA337327.1	BRAH1201205073	brain	5' read	P	25	QA337327.1	BRIH120204472	brain	5' read	
88.	QA2081004.1	BRAH1201201193	brain	5' read	P	26	QA2081004.1	FCBH03015262	brain	5' read	
89.	QA1446983.1	BRAZC2002635	brain	5' read	P	27	QA1446983.1	BRSN2013890	brain	5' read	P
90.	QA208351.1	BRAH1201201204	brain	5' read	P	28	QA208351.1	OCB012002935	brain	5' read	
91.	QA208351.1	BRAH1201202807	brain	5' read	-	29	QA208351.1	OCB012028454	brain	5' read	
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93.	QA2505043.1	BRAH120120208	brain	5' read	P	31	QA2505043.1	OCB01204545	brain	5' read	
94.	QA1446023.1	IMAGE-24865177	brain	5' read	P	32	QA1446023.1	OCB012028040	brain	5' read	
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96.	QA311925.1	BRAH12012003738	brain	5' read	P	34	QA311925.1	BRAC2001617	brain	5' read	P
97.	QA309824.1	BRAY1200305289	brain	5' read	P	35	QA309824.1	IMAG65732941	brain	5' read	P
98.	QA3011263.1	BRAY12003052890	brain	5' read	P	36	QA3011263.1	IMAGE-5477670	brain	5' read	P
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101.	QA408657.1	IMR32206160	brain	5' read	P						
102.	QA3011256.1	BRIH120230022	brain	5' read	P						
103.	QA3026614.1	BRAH1201203097	brain	5' read	P						
104.	QA353823.1	BRSN20117529	brain	5' read	P						
105.	QA353823.1	IC3903022740	brain	5' read	P						
106.	QA509415.1	FCB0120192031	brain	5' read	P						
107.	QA509415.1	FCB012020099	brain	5' read	P						
108.	QA301846.1	BRIH1202032476	brain	5' read	P						
109.	QA387460.1	BRIH12012023207	brain	5' read	P						
110.	QA387528.1	BRIH12012023308	brain	5' read	P						
111.	QA354215.1	BRSN2018169	brain	5' read	P						
112.	QA509600.1	FCB013022442	brain	5' read	P						
113.	QA385231.1	BRIH12012035318	brain	5' read	P						
114.	QA363387.1	BRAH120009016	brain	5' read	P						
115.	QA352652.1	BRSN2001446	brain	5' read	P						
116.	QA400202.1	BRIH120305441	brain	5' read	P						
117.	QA526272.1	F33R012010987	brain	5' read	P						
118.	QA550881.1	FCB01302518	brain	5' read	P						
119.	QA359810.1	BRSN2005743	brain	5' read	P						
120.	QA481320.1	FCB011000375	brain	5' read	P						
121.	QA365554.1	FCB0120014506	brain	5' read	P						
122.	QA511849.1	FEBR20207680	brain	5' read	P						
123.	QA369214.1	BRSN2016635	brain	5' read	P						
124.	QA525966.1	FEBR2017611	brain	5' read	P						
125.	QA487608.1	FCB012005641	brain	5' read	P						
126.	QA5090037.1	FCB012013875	brain	5' read	P						
127.	QA532732.1	FEBR2027807	brain	5' read	P						
128.	QA509341.1	FCB012014362	brain	5' read	P						
129.	QA351221.1	BRSN2014138	brain	5' read	P						
130.	QA511862.1	OCB012027591	brain	5' read	P						
131.	QA511862.1	OCB012016429	brain	5' read	P						
132.	QA382433.1	OCB012018252	brain	5' read	P						
133.	QA528137.1	OCB012019166	brain	5' read	P						
134.	QA277912.1	OCB0120172441	brain	5' read	P						
135.	QA781496.1	OCB012019008	brain	5' read	P						
136.	QA785260.1	OCB012018162	brain	5' read	P						
137.	QA278212.1	OCB012014163	brain	5' read	P						
138.	QA292203.1	OCB012015159	brain	5' read	P						
139.	QA786627.1	OCB012025726	brain	5' read	P						
140.	QA787493.1	OCB012025047	brain	5' read	P						
141.	QA786200.1	OCB012026041	brain	5' read	P						
142.	QA786801.1	OCB012026934	brain	5' read	P						
143.	QA509538.1	SKNNC2007468	brain	5' read	P						
144.	QA785445.1	OCB0120196316	brain	5' read	P						
145.	QA809653.1	SKNNC-3008312	brain	5' read	P						
146.	QA789610.1	OCB012027054	brain	5' read	P						
147.	QA786664.1	OCB012017986	brain	5' read	P						
148.	QA786870.1	OCB012019855	brain	5' read	P						
149.	QA848186.1	H03209116	brain	5' read	P						
150.	QA848186.1	H03300424	brain	5' read	P						
151.	QA846793.1	H02309008	brain	5' read	P						
152.	QA629849.1	IMAGE-4297305	brain	5' read	P						
153.	QA351385.1	EERBA2008325	brain	5' read	P						
154.	QA725252.1	IMAGE-2196165	brain	5' read	P						
155.	QA725205.1	IMAGE-2193181	brain	5' read	P						
156.	QA831006.1	IMAGE-5272563	brain	5' read	P						
157.	QA913201.1	IMAGE-5243215	brain	5' read	P						
PSEN2	1.	AM16292931	IMAGE-2785548	brain	5' read	-					



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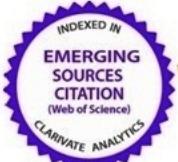
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