

# NEUROINFLAMMATION IN NEURODEGENERATIVE DISORDERS: MECHANISMS, BIOMARKERS, AND THERAPEUTIC TARGETS

**Dr Nandita Prabhat**

DM Neurology

ASST PROFESSOR

Neurology Department Hind Institute of Medical Science Barabanki

**Dr Chandra Shekhar Tiwari**

MD Medicine

Associate Professor

Medicine Department

Hind Institute of Medical Science Barabanki

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## Abstract

This top to bottom examination inspects the complex associations among neuroinflammation and neurodegeneration, researching central components, recognizing biomarkers, and distinguishing potential new treatment targets. In view of broad bioinformatics, this work looked to find center point qualities and pathogenic pathways associated with neuroinflammation in irregular Creutzfeldt-Jakob sickness (SCJD). Tests of SCJD and sound people were taken from GSE160208. Key qualities were found utilizing the Limma R bundle and Weighted Quality Co-Articulation Organization Examination (WGCNA), which were then utilized for improvement and invulnerable cell penetration examinations. The primary SCJD qualities were screened utilizing the protein communication (PPI) organization, cytoHubba, and AI. By utilizing sub-atomic docking, the particles related with center qualities were anticipated and examined. 88 potential qualities were analyzed. They were essentially associated with bacterial and viral disease and safe cell enactment, as indicated by improvement examinations.

**Keywords:** Neuroinflammation, Neurodegenerative, Disorders, Therapeutic

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## 1. INTRODUCTION

A major global health problem is the spread of neurodegenerative illnesses, a group of incapacitating conditions characterized by the progressive destruction of neurons. The importance of neuroinflammation in the pathophysiology and development of various disorders is becoming more and more clear as our knowledge of the complex workings of the nervous

system expands. Insights into the mechanisms behind disease progression, the discovery of potential biomarkers, and the creation of novel therapeutic approaches are all made possible by the interaction between chronic inflammation and neurodegeneration. The constant loss of neuronal structure and function is a defining feature of neurodegenerative diseases like Alzheimer's disease, Parkinson's disease, amyotrophic lateral sclerosis (ALS), and Huntington's disease. Although the underlying causes of these ailments are diverse, neuroinflammation—a sophisticated immune response that takes place within the central nervous system—is a common factor that connects them. The release of pro-inflammatory cytokines, chemokines, and reactive oxygen species is a result of complex interactions between invading immune cells and resident immune cells, including microglia and astrocytes.

The backdrop of neuroinflammation in relation to neurodegenerative diseases is established by this introduction. Researchers hope to find new approaches for therapeutic intervention by elucidating the complex pathways through which neuroinflammation affects the course of disease. Furthermore, the discovery of trustworthy biomarkers that can spot neuroinflammatory processes early in the course of the disease holds promise for enhancing diagnostic precision and disease monitoring. Parallel to this, new therapeutic approaches targeted at modifying the inflammatory response to delay disease progression and improve symptoms have been developed as a result of our growing understanding of the role of neuroinflammation in neurodegenerative illnesses. Targeting neuroinflammation offers patients and carers alike a glimmer of hope as research into immune-modulating drugs, biologics, and gene treatments advances.

- **Role of Neuroinflammation Emergence:** Our understanding of neurodegenerative illnesses has changed as a result of the discovery that neuroinflammation plays a significant role in their development. Studying disease mechanisms has taken on a new dimension as a result of the connection between immune responses and brain degeneration.
- **Diverse Spectrum of Neurodegenerative Disorders:** There are many different types of neurodegenerative disorders, each with its own clinical features and underlying processes. Despite these variations, recent evidence suggests that neuroinflammation contributes to the advancement of a variety of diseases, including Alzheimer's, Parkinson's, and ALS.
- **Core Inflammatory Mechanisms:** The activation of in-house immune cells, like microglia and astrocytes, is a core feature of neuroinflammation. These cells react to different stimuli

by releasing cytokines, chemokines, and inflammatory mediators. This inflammatory cascade prolongs neuronal injury by upsetting the delicate microenvironmental balance of the brain.

- **Relationship:** The effects of neuroinflammation on neurodegeneration go beyond local immune responses. The existence of misfolded protein aggregates, which are common in many neurodegenerative illnesses, may cause immune activation, aggravating neuroinflammatory processes, according to the evidence. This back-and-forth interaction highlights how intricately neurodegeneration and inflammation interact.

## 2. REVIEW OF LITREATURE

T. L. Spires-Jones (2016) The core viewpoint on the relationship between synaptic pathology and other neurological illnesses is provided by this review article. The authors highlight the continuous disturbance of synapse structure and function as a common factor by looking at studies spanning a variety of illnesses. The review highlights how synaptic abnormalities, such as loss and dysfunction, affect a variety of neurological diseases, including those that affect motor function and cognition.

B. A. Barres (2017) This ground-breaking research reveals an important role for microglia and astrocytes in neuropathology linked to neuroinflammation. The authors show that activated microglia cause the development of neurotoxic reactive astrocytes, which aid in the progression of neurodegenerative diseases. The work highlights the deep interaction between many cellular elements of the central nervous system and their functions in causing synaptic dysfunction brought on by neuroinflammation.

Xu, H. (2020) This investigation examines the function of microglia in Alzheimer's illness and how soluble TREM2 affects those cells. The researchers discover a potential therapeutic route for treating the pathogenic features of Alzheimer's by examining the effects of this soluble protein on microglial activities. The results of the study highlight the possibility of reducing synaptic dysfunction in neurodegenerative illnesses by addressing microglial activation.

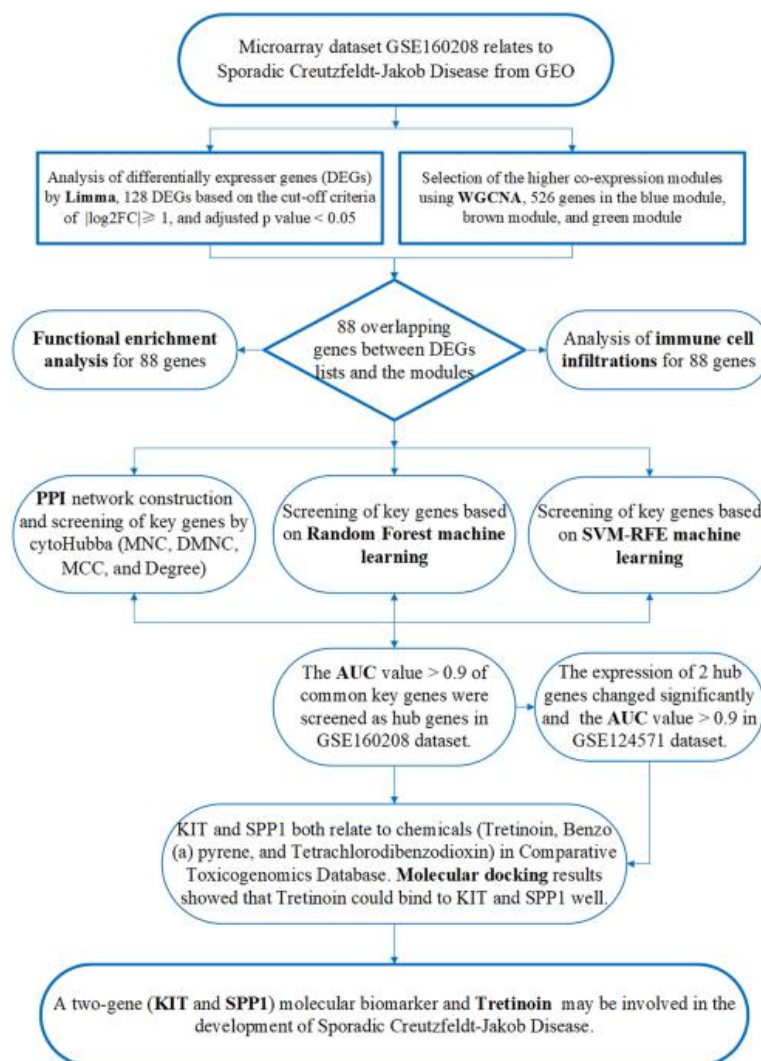
P. Edison (2021) Leng and Edison objectively evaluate our present knowledge of neuroinflammation and microglial activation in Alzheimer's disease in this extensive review. They go over the difficulties in converting research discoveries into efficient therapy approaches, the complexity of these processes, and its implications for the development of diseases. The

review provides a road map for future research directions in understanding and treating synaptic dysfunction caused by neuroinflammation.

T. O'Connor (2010) By examining protein aggregation diseases, a category of neurological conditions marked by the buildup of misfolded proteins, this review gives a wider context. The authors explore prospective therapy approaches as well as the pathological factors underlying these illnesses. The review adds to the overall knowledge of how neuroinflammation affects neurological illnesses by elucidating the role of protein aggregation in synaptic dysfunction.

### 3. MATERIALS AND METHODS

A multistage coordinated examination of this examination is introduced in the flowchart in Fig. 1.



**Figure 1:**The means of the bioinformatics examination, AI, and atomic docking in this study are portrayed in this flowchart.

### 3.1 Data collection and data preprocessing

In the GEO (Quality Articulation Omnibus) data set, the expressions "Irregular Creutzfeldt-Jakob Illness" and "Neuroinflammation" were looked, and eventually the GSE160208 dataset relating to neuroinflammation in SCJD was picked for extra analysis<sup>7</sup>. It contains data on the declaration of 800 qualities related with neuroinflammation in the mind tissues of 27 SCJD patients and 20 solid individuals. Moreover, the dataset contains gathering for each example (Control and SCJD) (Table 1), mind tissue parcel (cerebrum and cerebellum), orientation (Female and Male), codon-129 change (MM, MV, and VV), and SCJD subtype (MM1, MM2, MV1, MV2, VV1, and VV2).

**Table 1:**GSE160208 gives a portrayal of the examples used in this review.

Group	Brain. region	Gender	Codon-129 Mutation	SCJD Sub type
27 SCJD	14 FC 13 CB	16 Female 15 Male	16 MM 7 MV 8 VV	13 MM1 3 MM 1+2 5 MV1 3 MV2 8 VV2
30 Normal	15 FC 15 CB	20 Female 10 Male	9 MM 9 MM 5 VV	-

### 3.2 Gene co-expression network construction and identification of key modules

Telemedicine administrations are turning out to be increasingly more typical in the healthcare framework as data innovation propels. In this paper, we look at three healthcare frameworks and dissect how telemedicine has impacted the delivery of healthcare administrations. We center

around the patients' inclination for the first analytic among disconnected and online channels. As per our information, telemedicine sometimes can help with bringing down both the general expense of the healthcare framework and the hanging tight times for patients.

In certain cases, we find that the hospital shouldn't offer telemedicine administrations, which is in accordance with Tarakci et al's. finding. Their discoveries, notwithstanding, that treating each patient utilizing telemedicine is rarely all that strategy, can be extended in specific conditions. Our discoveries show that the watchman framework might be the best strategy in specific conditions. The double channel administration framework, which joins the guard framework and the regular outpatient framework, is for the most part the best strategy. By adjusting the disconnected and online assistance limits, the hospital can choose how to portion the market while permitting customers to pick clinical benefits in light of their utility. Our review's discoveries show that there is a decent market parted among on the web and disconnected arrangements. To get the double channel administration framework's most minimal complete expense, the hospital can shift the on the web and disconnected help limit.

### **3.3 Identification of DEGs**

DEGs were separated the mind tissues of patients with inconsistent Creutzfeldt-Jakob infection (SCJD) and sound people utilizing the "limma" programming. The screening necessities were  $\text{adj.P } 0.05$  and  $|\log_2\text{FC}| > 1$ . The R language's "ggplot2" program was utilized to show the DEGs. The quality is communicated all the more exceptionally in the cerebrum tissue of SCJD patients assuming that the  $\log_2\text{FC}$  reciprocal to the DEG is  $> 0$ ; alternately, assuming the  $\log_2\text{FC}$  corresponding to the DEG is 0, the DEGs are communicated less profoundly in the mind tissue of SCJD patients.

The DEGs among SCJD and significant modules were portrayed in a Venn chart by the "Venn Outline" programming. Picturing the articulation levels of normal qualities across the significant modules and DEGs in SCJD was finished utilizing the "Complex Heatmap" bundle in R.

### **3.4 Functional enrichment analysis**

The intricacy of the examination can be diminished utilizing utilitarian enhancement investigation, which can dole out hundreds or thousands of qualities to different pathways. On possible qualities for SCJD, KEGG pathway investigation and GO improvement examination were completed utilizing the R language's "bunch Profiler" and "Portion" programs. Three

organic points are covered by GO enhancement investigation: natural cycles, cell parts, and atomic capabilities. Adj.P0.05 was used as the determination rule in this examination, and the main ten positioned pathways were decided to research the sub-atomic systems basic neuroinflammation in SCJD.

### **3.5 Analysis of immune cell infiltrations**

To survey the general substance and dynamic guideline cycle of 22 safe cells, the CIBERSORT technique is habitually used. It is more viable than different strategies at finding human-safe cell morphologies in foundation clamor and unidentified blends. The examination utilizes the R programming language related to the CIBERSORT deconvolution strategy to process the dispersion of 22 safe cells in the example, including Lymphocytes, memory and credulous B cells, resting and enacting NK cells, and others. The resistant cell extent score of each example in the not set in stone after the calculation has been performed multiple times. The stacked bar outline shows the circulation of resistant cells in the example. The relationship between's particular insusceptible cells is likewise determined utilizing the Pearson connection coefficient and showed as an air pocket graph. The position aggregate test is then performed to analyze resistant cell articulation in the SCJD gathering and control bunch.

## **4. HUB GENE GENE EXPRESSION VALUES FROM THE GSE124571 DATASET**

The GSE124571 dataset related with SCJD11, which incorporates ten examples of SCJD and ten examples of reference mind tissue, was utilized to approve the declaration of center point qualities at the RNA level in human cerebrum tissue. t-tests and the "ggplot2" bundle from the R language were utilized to dissect and show the declaration of potential center point quality competitors. The GSE124571 dataset was additionally used to evaluate the symptomatic adequacy of center qualities in SCJD utilizing ROC (AUC > 0.9).

### **4.1 Exploring the functions of hub genes and molecular docking analysis**

The Near Toxicogenomics Data set (CTD), which was distributed in 2004, has formed into a fundamental wellspring of toxicological information. The CTD data set makes sense of the associations among synthetic compounds and qualities/proteins, infections and synthetic substances, illnesses and qualities, aggregates and species, and openness data. It is feasible to find new or complete openness qualities and atomic components of synthetic compounds through month to month updates to the CTD information base, which can help create testable thoughts on

what openness means for human health<sup>12,13</sup>. Synthetics related with center point qualities were anticipated in CTD to explore their organic exercises, and sub-atomic docking examination was used to produce novel treatment choices for SCJD by looking at the relationship between center qualities and chemicals<sup>14</sup>.

The center qualities' protein structures were taken from the Protein Information Bank (PDB) and filtered utilizing PyMOL (V2.5.4) to avoid water particles and heteroatoms. Furthermore, Chem3D and AutoDockTools programming was utilized to change the 3D synthetic designs of the substances from SDF to "pdb" design after they were recovered in SDF design from the PubChem database. The proteins and synthetic substances were then changed via AutoDockTools (V1.5.7) to "pdbqt" design records. Explicit docking pockets in the picked proteins were characterized utilizing the matrix box component of AutoDockTools, where medications could tie. In the wake of social affair every one of the essential information, sub-atomic docking analysis<sup>15,16</sup> was done at the order brief, and PyMOL was used to show the docking results.

## 4.2 Statistical analyses

Running in R (variant 4.0.2) with the default measurements boundary and cut-off values expressed in each part were the accompanying projects: WGCNA (form 1.69), Limma (adaptation 1.9.6), ggplot2 (variant 3.3.3), ClusterProfiler (variant 3.16), Proc (form 1.18.0), e1071, randomForest (adaptation 4.7.1.1), Portion (form 3.14.0), and GO plot (adaptation 1.0.2). This work furthermore utilized Cytoscape (form 3.8.0), PyMOL (rendition 2.5.4), Chem3D, and AutoDockTools (variant 1.5.7). Factual not entirely set in stone by the accompanying edges: \*p 0.05, \*\*p 0.01, and \*\*\*p 0.001.

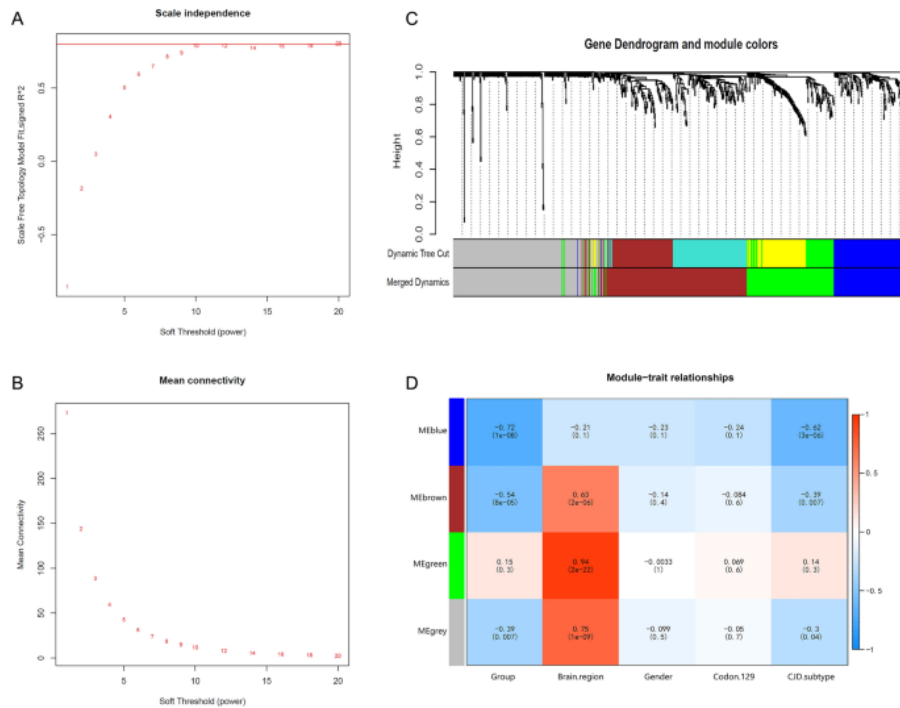
## 5. RESULTS

### 5.1 Construction of a weighted expression network and identification of important modules

Utilizing the WGCNA approach, 800 qualities related with neuroinflammation were thought about between the Control and SCJD gatherings. The worth of 20 under the  $R^2 = 0.88$ , figured by pick Delicate Limit, was chosen as the delicate edge for building an unweighted network. Four modules (blue, brown, green, and dim) were found after the module was gathered and divided in light of the qualities of quality articulation (Fig. 1). Utilizing the trademark vector upsides of every module, the connection among modules and clinical aggregates was determined (Fig. 1). The blue and earthy colored modules showed a higher significance with groupings (cor>



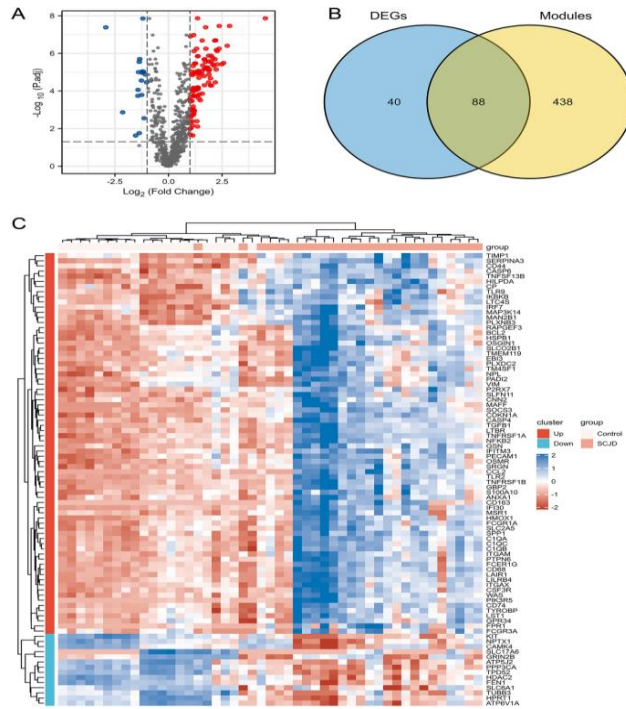
0.5, P 0.01), and the brown, green, and dim modules were exceptionally connected with mind locales ( $cor > 0.5$ , P 0.01). Also, we picked the 526 qualities remembered for the blue, brown, and green modules for extra examination in light of the fact that the qualities in the dark module couldn't be gathered.



**Figure 1:** Important gene modules that are connected to neuroinflammation in SCJD with WGCNA have been identified

## 5.2 Identification of DEGs

The "limma" bundle's differential examination created 128 SCJD DEGs, containing 110 upregulated DEGs and 18 downregulated DEGs, as displayed in Fig. 2. Dim means non-DEGs, though red indicates DEGs that are upregulated, blue signifies DEGs that are downregulated. The DEGs and basic module qualities in SCJD shared 88 qualities, as per the Venn chart bundle (Fig. 2). The "Complex Heatmap" bundle was utilized to show the articulation levels of these 88 qualities (Fig. 2).

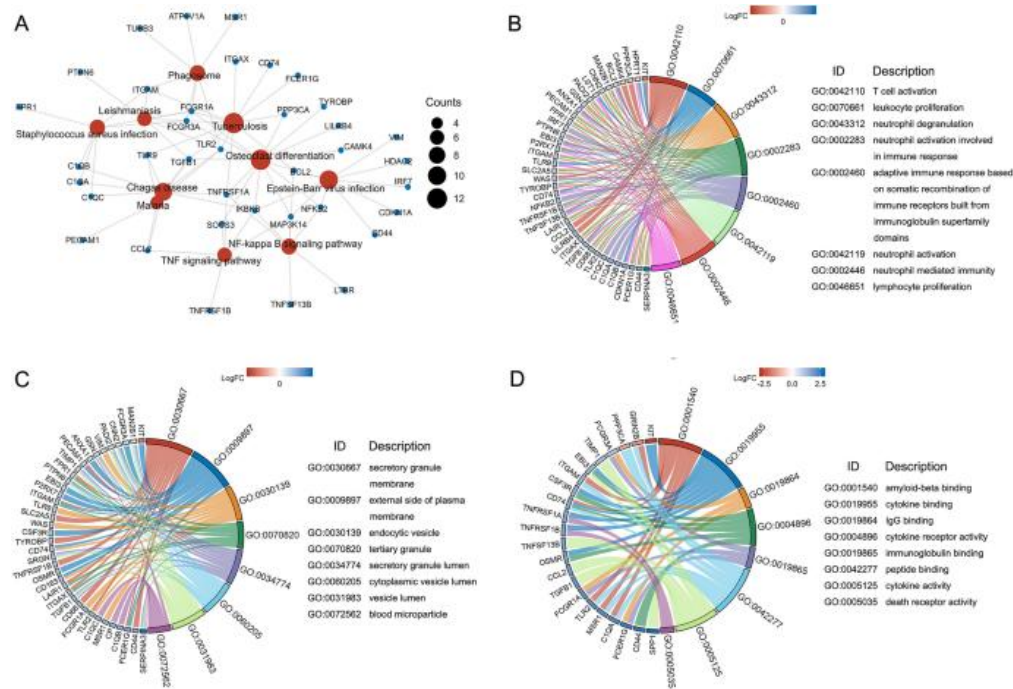


**Figure 2:** Identification of Key Genes Associated with Neuroinflammation in SCJD and Differentially Expressed Genes (DEGs)

### 5.3 Functional enrichment analysis

As per the KEGG pathway examination results the 88 applicant qualities are essentially advanced in natural flagging pathways like Osteoclast separation, Tuberculosis, Chagas sickness, Epstein-Barr infection contamination, Staphylococcus aureus disease, NF-kappa B flagging pathway, Leishmaniasis, Jungle fever, Phagosome, and TNF flagging pathway (Fig. Furthermore, the 88 up-and-comer qualities' GO examinations uncovered that these qualities are principally improved in organic cycles connected with the enactment of resistant cells and the limiting of safe proteins, for example, Lymphocyte actuation, leukocyte expansion, neutrophil degranulation, neutrophil initiation associated with resistant reaction, neutrophil enactment, neutrophil interceded resistance, lymphocyte multiplication, mononuclear cell expansion, and reaction to interferon-gam. As found in Fig. 4C, the CC examination shows that the quality items are basically tracked down in the secretory granule layer, outer side of the plasma film, endocytic vesicle, tertiary granule, secretory granule lumen, cytoplasmic vesicle lumen, and vesicle lumen. As found in Fig. 4D, these qualities are principally enhanced in exercises, for example, 1-phosphatidylinositol-3-kinase controller movement, 1-phosphatidylinositol-3-kinase restricting,

amyloid-beta restricting, cytokine restricting, IgG restricting, cytokine receptor action, immunoglobulin restricting, peptide restricting, and cytokine action.



**Figure 3:** Analysis of 88 genes' functional enrichment in SCJD neuroinflammation

## 6. DISCUSSION

The beginning of Creutzfeldt-Jakob sickness (CJD), as per connected concentrate on discoveries, includes organic cycles such invulnerable framework reaction, digestion, formative science, and vesicle-intervened transport11. The exact interaction is obscure, and there are not many distributions of the neuroinflammatory pathways connected to the pathophysiology of SCJD. To examine the system and significant qualities engaged with neuroinflammation during the pathogenesis of SCJD, we utilized a dataset connected to SCJD neuroinflammation. Initial, 800 neuroinflammatory-related qualities in SCJD were examined utilizing the WGCNA approach, and 526 qualities were tracked down in the significant modules. Another technique delivered a sum of 128 DEGs by running DEGS examination on two gatherings of tests utilizing the lima customized. The 88 most significant DEGs in the fundamental module qualities and DEGs were the subject of improvement and safe penetration examination. As per the aftereffects of the advancement study, these qualities are for the most part engaged with bacterial and viral diseases, safe cell enactment and state changes, and other related processes. The movement of SCJD might be affected by invulnerable cell initiation and modified resistant framework action,

as indicated by research on safe cell invasion. Unit (CD117) and SPP1 were demonstrated to be urgent qualities embroiled in SCJD neuroinflammation subsequent to examining and affirming the 88 DEGs with various procedures. Utilizing CTD, it was found that Pack (CD117) and SPP1 have associations with tretinoin, tetrachlorodibenzodioxin, and benzo(a)pyrene. At long last, sub-atomic docking methods checked that SPP1 and Unit (CD117) connect well with tretinoin. Thus, Tretinoin, SPP1, and Unit (CD117) are fundamental for the neuroinflammation part of SCJD.

The contribution of neuroinflammation in the advancement of SCJD has been talked about, in spite of the way that there haven't been numerous examinations on the neuroinflammation related with SCJD previously. An illness explicit marker of neuroinflammation called YKL-40 (otherwise called Chitinase 3-like 1) was demonstrated to be extensively higher in SCJD17 patients with neurodegenerative dementia in a few examinations on cerebrospinal liquid pointers. The outflow of sTREM2 (the dissolvable rendition of TREM2), an intrinsic safe cell surface receptor that controls microglial movement, was connected to PRNP codon 129 and subtypes, including CSF14-3-3 positive, all out tau, and YKL-40, and it expanded as the infection progressed<sup>18</sup>. Also, the cerebral liquid of people with SCJD showed raised articulation of irritation related qualities as chitotriosidase 1 (CHIT1), glial fibrillary acidic protein (GFAP)<sup>19</sup>, and SERPINA1 (- 1 antitrypsin)<sup>20</sup>. There have additionally been a few examinations on the SCJD's mechanism. A concentrate on mice contaminated with the astrocyte-and neuron-related PrPSc strains 22L and ME7 found that the elements of PrPSc quality articulation are inconsequential to cerebrum district or cell affinity<sup>21</sup>. As indicated by the latest investigations on SCJD neuroinflammation, the impact of SCJD subtypes may not be the greatest or just variable influencing how firmly neuroinflammatory qualities are expressed<sup>7</sup>. Nonetheless, it was found that specific quality articulations were controlled diversely in different pieces of the cerebrum. Preceding this, a few papers guaranteed that varieties in local quality guideline are impacted by the genotype of the patient at the PRNP codon.

## 7. CONCLUSION

The muddled cooperation among neuroinflammation and neurodegenerative illnesses has uncovered a relationship that is diverse and complex, and it considerably affects the

pathophysiology and improvement of sickness. The evaluated writing underscores the pivotal part that neuroinflammation plays in heightening neuronal harm and regularly fills in as a connection between different neurological illnesses. An outpouring of favorable to provocative middle people and receptive oxygen species are delivered because of the enactment of microglia, astrocytes, and immunological reactions, and this interaction eventually prompts synaptic brokenness and neuronal demise. The meaning of neuroinflammation is additionally featured by the common system of synaptic harm, as upset neurotransmitters are a figure mental degradation, engine shortfalls, and other clinical side effects found in different disorders. Investigations of cytokines, chemokines, and other sub-atomic signs have shown intriguing possibilities with regards to the quest for biomarkers to distinguish and follow neuroinflammation. These biomarkers have the ability to upset analyze, empower early mediations, and improve patient results through individualized treatment plans. In preclinical and clinical examination, a few treatment moves toward that mean to balance neuroinflammatory reactions have showed guarantee. These methodologies, which range from inventive biologics and quality treatments to insusceptible designated drugs, are intended to diminish ongoing aggravation, shield the strength of the sensory system, and maybe even stop the course of neurodegenerative sicknesses. Notwithstanding, the hardships in formulating effective therapeutic procedures are featured by the intricacy of the focal sensory system, complex safe reactions, and likely unforeseen outcomes.

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