Review

# Contribution of Inhibitor of Differentiation (ID) and Estrogenic Endocrine Disruptors to Neurocognitive Disorders

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Abstract: The devastating growth in the worldwide frequency of neurocognitive disorders and its allied difficulties such as decline in memory, spatial competency, and ability to focus poses a significant psychological public health problem. Inhibitor of Differentiation (ID) proteins are members of a family of helix-loop-helix (HLH) transcription factors. ID proteins have been demonstrated to be involved in neurodevelopmental & depressive diseases and thus may influence neurocognitive deficiencies due to environmental exposure. Previously, it has been demonstrated that environmental factors such as estrogenic endocrine disruptors (EEDs) have played an essential role in the influence of various neurocognitive disorders such as Alzheimer's, Dementia, and Parkinson's disease. Based on this increasing number of reports, we consider the impact of these environmental pollutants on ID proteins. Better understanding of how these ID proteins by which EED exposure can affect neurocognitive disorders in populations will prospectively deliver valuable information in the impediment and regulation of these diseases linked with environmental factor exposure.

**Keywords:** endocrine disruptor; environmental health sciences; gene-environment; inhibitor of differentiation; neurocognitive disorders

## 1. Introduction

Neurocognitive disorders, which was earlier identified as dementia, denotes a range of disorders that affect the brain and cause worsening in one or more cognitive areas. Deterioration in cognition need not only be damage to cognitive abilities, but also visible by others and tested by cognitive assessment. This type of disorder frequently affects older people, however is not part of the standard aging development and has the capability to affect young people as well. Neurocognitive disorders can affect language, memory, attention, social cognition, learning, and perception [1-2]. Today, there are main and trivial neurocognitive disorders that are reliant on how severely the symptoms influence an individual's capability to function self-sufficiently in daily activities such as Alzheimer's, Parkinson's, Creutzfeldt-Jakob disease, Huntington's disease, and Vascular Dementia. Factors inducing various forms of neurocognitive ailments can include endocrine & metabolic ailments, substance abuse, infections, trauma, and nutritional deficiencies [2]. Presently, there is a necessity to recognize how factors such as estrogenic endocrine disruptors contribute to neurocognitive disorder susceptibility.

Estrogen, which belongs to a class of hormones, has been shown to have various functions including the regulation of metabolism and endocrine growth & development [2]. Furthermore, estrogen has been previously demonstrated to affect neurocognitive outcomes [3-8]. Due to this, neurocognitive disorders may be susceptible to estrogenic endocrine disruptors or EEDs. Endocrine



2 of 16

disruptors modify hormone function or production including anthropogenic chemicals, heavy metals, and phytoestrogens. More particularly, EEDs consist of arsenic, DES, phthalates, bisphenol A (BPA), and polychlorinated biphenyls (PCBs) [9-13]. Studies have reported links between EED exposure and neurocognitive disorders [14-16]. Based on findings that demonstrate that a family of proteins, Inhibitor of Differentiation (ID) has been associated with neurocognitive disorders [16], we will also highlight how EED exposure from the environment may potentiate neurocognitive outcomes via ID proteins. This review is concentrated on connecting EEDs to ID interactions leading to altered outcomes in neurocognitive disorders. More study in these capacities may uncover innovative or additional applicable beneficial modalities and deliver approaches for neurocognitive outcomes with EED exposure.

# 2. Inhibitor of Differentiation (ID)

#### 2.1 Structure & Function

Inhibitor of Differentiation is a family of proteins that consist of four genes (ID1, ID2, ID3, & ID4). The members of the ID family share an extensive amino acid sequence homology within their HLH (helix-loop-helix) domain (69-78%) [17-18]. It has been reported that ID protein acts as a transcription regulator, which regulate transcription in a dominant-negative manner by dimerizing with basic HLH transcription factors like HEB, E47, and E12 [19-21]. ID proteins are pleiotropic proteins involved in the modulation of biological processes such as cell proliferation & differentiation, cell cycle control, senescence, apoptosis or angiogenesis, and metastasis [22-23]. In case of the central nervous system, ID1 and ID3 are greatly expressed in the premature stages of nervous tissue development, but their levels decline in later stages [24-25]. Although ID1 and ID3 expression cease as tissue matures, ID2 and ID4 expression remain constant throughout the nervous tissue development [26-28]. Therefore, ID proteins play a very important role in nervous tissue biology. Intriguingly, reactivation of ID proteins in adult tissues is held responsible for the involvement of various cancers [29-30]. Researchers have shown the reactive oxygen species (ROS) induced ID protein-mediated cell proliferation and dysregulation of tissue biology in vitro and in vivo conditions [31-33]. Similarly, in another study Das et al have shown the exposure of 17-β estradiol (E2) and estrogenic endocrine disruptors (EED) like polychlorinated biphenyl 153 (PCB153) to vascular endothelial cells (ECs) increase ROS. While ID3 is a redox-sensitive gene, it acts as an important determinant of the ROS-induced proliferation of E2 and ECs exposed to PCB153 [34-36]. Neurodegenerative disorders like Alzheimer's and Parkinson's are the main causes of dementia, and their symptoms worsen slowly over the time. The causes of neurodegenerative diseases apart from genetic mutations are mainly environmental factors, head injuries, depression, and hypertension. Researchers have shown that these causes are strongly interconnected with elevated levels of ROS [31, 37-38]. The increased levels of ROS affect human tissues at a molecular level over time. As age increases, longer exposure to ROS may result in increasing tissue injury and severe disease symptoms. Since ID proteins are shown to be redox sensitive, we predict EEDs exposure boosts ROS-induced ID proteins levels, which may be responsible for the onset of neurocognitive deficiencies.

# 2.2 ID Proteins and Neurocognitive Disorders

There has been accumulative evidence demonstrating the role of ID proteins in various neurological deficiencies & disorders. One essential pathological hallmark of Alzheimer's disease (AD) is the buildup of senile plaques largely comprised of amyloid beta-peptide (A $\beta$ ) in the patients' brains. Hung et al investigated if A $\beta$  may stimulate Sonic hedgehog (SHH) expression and its essential mechanisms. A $\beta$ 25-35 induced ID1, which has been exhibited to stabilize HIF-1 $\alpha$ . Further, A $\beta$ 25-35-mediated induction of HIF-1 $\alpha$  and SHH was both suppressed by ID1 siRNA. Taken together, the pathway inhibitor cyclopamine SHH and its antibody reduced A $\beta$  cytotoxicity. Based

3 of 16

on these lines of evidence, results showed a signaling pathway of  $A\beta \rightarrow ID1 \rightarrow HIF-1 \rightarrow SHH$  [39]. Kitajima et al investigated ID2 mRNA-expressing cells in the adult mouse brain. Results showed that ID2 mRNA is identified in more diverse brain areas, including the amygdaloidal complex, globus pallidus, substantia nigra pars reticulata, suprachiasmatic nucleus caudate putamen, and the frontal part of the sub-ventricular zone. These suggest that ID2 may have a function in cognitive and neural activity. Additionally, expression of ID3 (moderate or low) was demonstrated alongside high expression in some specific areas such as the molecular layer of the dentate gyrus. ID4 mRNA was detected in the regions such as the lateral amygdaloidal nucleus. Based on these lines of evidence, the ID2 pattern expression is distinctive from those of partnering ID proteins [40]. Donepezil, is a common medication for Alzheimer's disease. Acetylcholinesterase (AChE) has been demonstrated to play a role in osteoblast function, however the mechanism of AChE on osteoclastogenesis nevertheless remains uncertain. Donepezil reduced receptor activator of nuclear factor-kappa B ligand (RANKL) expression in bone marrow macrophages (BMMs), resulting in the up-regulation of ID2 and inhibition of osteoclast differentiation with down-regulation of c-Fos. These particular results show that inhibition of osteoclast function due to Donepezil prevents bone loss, which may suggest the chance that donepezil reduces fracture risk in patients with Alzheimer's disease [41].

Uncertainties in rhythm circadian-related developments are recurrently established in anxiety and depressed-driven patients. Several genes have been recognized as factors for the progression of mood disorders. It was demonstrated that mild stress-stimulated chronic anhedonic behavior is connected amongst distressed diurnal alternation of the expression of CLOCK, CRY2, PER3 PER1, REV-ERB $\alpha$ , ID2, ROR- $\beta$  and ROR- $\gamma$  in the mouse basolateral amygdala [42]. Furthermore, premature life abandonment increases risk for the psychopathological development through both childhood and adulthood, including anxiety disorders and depression. It was recently reported that epigenetic changes in DNA resulted in three genes predicting depression in maltreated children: GRIN1, ID3, & TPPP. Behavioral tests demonstrated that GRIN1, ID3, and TPPP gene expression were established to significantly predict behavioral alterations. These lines of evidence support the role of these genes in the etiology of anxiety and depressive phenotypes succeeding premature life stress [43]. Transcriptome wide association studies have furthermore been used to predict various epigenetic markers of depression. Methylation in 3 genes developed as predictors of depression: ID3, NMDA, GRIN1, and TPPP [44].

Similarly, Becker et al targeted a strategy to uncover foundations of variability in subcortical brain areas. Results demonstrated important improvement of genomic loci that affect the area of the hippocampus, a result that strongly passed the adjusted threshold for testing of multiple brain phenotypes. Investigation of individual single nucleotide polymorphisms (SNPs) also revealed a connection upstream of the ID2 gene with rs7588305. Results show that targeting recognized regulatory regions indicates ways to comprehend the biology that links genotypes to phenotypes. [45]. Previously, Kepa et al identified a network of HLH transcriptional regulators controlled by myelin transcription factor 1-like (MYT1L), as specified in the human brain and neural stem cells. It was demonstrated that MYT1L is essential for neuronal differentiation and identified ID1 as a target. Furthermore, MYT1L prohibited ID1 expression and induced expression of a large quantity of terminal differentiation genes. Consistent expression of MYT1L corresponded with neuronal maturation and linked ID1 and ID3 during the lifecycle. Additionally, genetic polymorphisms that abridged expression of MYT1L in the hippocampus caused enlarged ID1 and ID3 expression, reduced TCF4 & NEUROD6 levels and reduced gene expression involved in cancer, neurodegeneration, long-term potentiation, and synaptic transmission. As a result, these outcomes indicate that MYT1L controls memory-related developments by regulating a neuronal proliferation & differentiation mechanism of ID family proteins [46]. Additionally, six genes displaying at least differential expression among hemispheres (BAIAP2, DAPPER1, LMO4, NEUROD6, ATP2B3, and ID2) in a case-control association study in an initial Spanish sample of ADHD patients and control subjects. Outcomes support the contribution of genomic factors in the stability of ADHD in some of the populations examined and may supply influencing deviant cerebral lateralization in this condition [47].

4 of 16

Fetal Alcohol Spectrum Disorders, or FASD, signify a variety of antagonistic developmental ailments triggered by prenatal ethanol exposure (PrEE) from parental intake of alcohol. A mouse model of FASD demonstrated stable phenotypes transmitting by the male germline to the unexposed third generation. Global DNA methylation levels, modifications in ectopic intraneocortical connectivity, and up-regulation of neocortical Rzr $\beta$  & ID2 was seen. These phenotypes may contribute to sensorimotor, cerebral, and communicative insufficiencies seen in individuals with FASD. Therefore, understanding the conceivable epigenetic mechanisms may uncover innovative targets for beneficial mediation of FASD [48].

#### 2.3 ID Proteins & Estrogenic Endocrine Disruptors

Previously, we demonstrated how ID3 influence obesity and metabolic health in reaction to environmental influences. We also highlighted the understanding of how ID3 may contribute to multifaceted diseases through metabolic perturbations [16, 20]. PCBs are a class of organochlorine compounds that are persistent and have the potential to disrupt the homeostasis of thyroid hormones (THs). In this study, thyroid histology, plasma TH levels, and iodothyronine deiodinase (ID proteins including ID1, ID2, and ID3) gene expression patterns were examined in juvenile Japanese flounder (Paralichthys olivaceus) following exposure to environmentally relevant concentrations of a commercial PCB mixture, Aroclor 1254. Exposure to Aroclor 1254 for 50 days increased follicular cell height, colloid depletion, and hyperplasia. Profiles of the changes in mRNA expression levels of IDs were detected in the liver and kidney, which may be associated with a reduction in plasma THs levels. ID2 mRNA expression in the liver exhibited an increase based on dose-dependence, signifying that this isotype may function as a constant marker for thyroid-disrupting chemical (TDC) exposure. Overall, results suggest that applicable levels of Aroclor 1254 produce noteworthy thyroid disruption [49].

Micro-vascular lesions from endothelial cell dysfunction are formed in various areas of patients with complex chronic diseases such as the lung, brain, retina, and kidney. The mechanisms dependable for starting micro-vascular injury remain weakly distinct, while factors have been suggested, including oxidative stress induced by environmental chemicals. Heightened neovascularization has been associated in the progression or development of proliferative vascular lesions. Previously, support for how ROS via PCBs may contribute to neo-vascular phenotype development with the aim of revealing the function of environmental chemicals in endothelial dysfunction with a concentration on ID3 has been shown. Results demonstrated that PCB-induced ROS intermediated neo-vascular phenotype are additionally depended on ID3 and Pyk2. Additionally, treatment of PCB153 enlarged the dimension of endothelial spheroids with circumstances that function on behalf of stem cell spheroid clonal selection. Elevated ID3 protein expression compared with a larger amount of oxidative DNA damage marker 8-OHdG in blood vessels. The results provide the potential role of ID3 in regulating development of micro-vascular lesions and vascular endothelial cell survival prompted by environmental chemicals such as PCB153 [34-35]. Furthermore, another study determined if in utero exposure to Bisphenol A (BPA) stimulated reproductive tract irregularities in the adult male testis. Adult males including hospathology, anogenital distance, and sex-organ weights were exposure in utero through oral gavage to sesame oil, 50 µg/kg BPA, 1000 µg/kg BPA, or 2 µg/kg diethylstilbestrol (DES) from gestational days 10 to 16 was examined in C57/BI6 mice. Adult mRNA levels of genes connected with differentiation and sexual maturation, GATA4 and ID2, were lower only in testes exposed to DES. At the molecular level, DES exposure via in utero, not BPA, leads to reduced mRNA expression of genes linked with Sertoli cell differentiation [50].

5 of 16

# 3. Influence of EED Exposure on Neurocognitive Disorders

Exposure to estrogenic endocrine disruptors has been previously shown in various population and animal studies. Previously Bell et al tested the effects of PCBs on prenatal or juvenile individuals. The effects had differential results on age-dependent and sex behaviors. Females demonstrated different social and anxiety behavior in adolescence, while males exhibited small but significant changes in socio-sexual preferences in adulthood [51]. Exposure to low levels of PCBs is known to lead to anxious behavior in offspring mice, both young and adult. At further advanced life stages, an effect on the mouse brain of neuronal stress induced by the AB peptide was evaluated. Significant impairment in long-duration memory was identified in the mice treated lacatational with non dioxin-like PCBs (NDL-PCBs). Early exposure to low levels of NDL-PCBs stimulates late neuronal susceptibility to amyloid tension [52]. Exposure to PCBs may result in changed procreative behaviors in adulthood. Rat dams (pregnant) were injected on gestational days with PCB mixture Aroclor 1221 at one of two doses. Females were unaffected but males treated with A1221 presented decreased indicators of anxiety [53]. Researchers have also studied sub-chronic embryonic exposure to PCBs concerning anxiety-associated components. Exposure induced behavioral deficiencies at 7 days post-fertilization was detected. Outcomes demonstrated that exposed larvae had enhanced edge preference relative to the control. Furthermore, larvae that were exposed reacted contrarily to a graphical risk comparative to control larvae [54]. Comprehensive gene expression profiles in cerebellar exhibiting the highest suggestive stimulation of anxiety-like behavior has been warranted via male mice. Outcomes demonstrated alterations in the expression of genes in the neurons of the PCB-exposed mice [55]. Furthermore, studies have analyzed the relationship between work-related PCB exposures. Analysis including individuals with their plasma PCBs collected via bio monitoring and the psychological syndromes assessed with a consistent screening method. Results showed greater PCB burdened individuals had greater depression but not anxiety syndrome [56].

Chronic exposure to PCBs has also been demonstrated to cause neurocognitive abnormalities. Repressor element 1-silencing transcription factor (REST) plays an important role in neuronal phenotype expansion in both neural progenitor cells and non-neuronal cells. Chronic exposure to the PCB mixture Aroclor-1254 caused cell death thru the initiation of calpain but not caspase-3. The mixture of PCBs reduced acetylation of the histone proteins H3 and H4. Together, these outcomes indicate that A-1254 employs its toxic influence via REST by down-regulating synapsin-1 and diminishing H3 and H4 acetylation [57]. Alterations in calcium signaling, thyroid hormones, and neurotransmitters have been hypothesized as candidate mechanisms for developmental neurotoxicity in animal models via PCBs. Expression of gene levels in the cerebellum and hippocampus from postnatal days (PNDs) 7 and 14 animals were determined. In the cerebellum, transcripts demonstrated modification at PND7 compared to transcripts at PND14 by Aroclor-1254 exposure, with only one transcript disturbed at both ages. Examination indicates that pathways associated to calcium homeostasis (GNG3, RYR2, TRDN, CACNA1A), intracellular signaling (CAMK2D, STK17B, PACSIN2, RYR2, TRIO, FERT2, PTK2B), axonal guidance (LUM, MXD3, AKAP11, GUCY1B3), aryl hydrocarbon receptor signaling (NFIA, 2COL1A2), involvement in cell proliferation (GSPT2, CDKN1C, PTK2B), and differentiation (LFITM31, HPCA, ZFP260, LGSF4A, HES5) leading to nervous system development were suggestively changed by Aroclor-1254 exposure. Aroclor-1254-induced genomic changes were greater in the hippocampus than the cerebellum. The outcomes indicate that stimulated neurotoxic effects via PCB were due to disturbance of standard ontogenetic design of nervous system growth and development by adjusting signaling pathways [58].

Individuals were exposed to PCBs and polychlorinated dibenzofurans (PCDFs) due to absorption of polluted cooking oil in Taiwan. Neurocognitive performance in individuals exposed to PCDFs and PCBs with that of unexposed was compared. Evaluation of neurocognitive examinations was directed. In exposed men, outcomes were comparable to the reference group; conversely, exposed women had diminished functioning in attention and digit span visual memory span, and verbal memory recalls. The study demonstrated neurocognitive deficits in certain dose-dependent

6 of 16

aspects of attention, learning ability, and visual memory in women previously exposed to PCBs and PCDFs however, not in exposed men [59]. Samples taken from women and blood samples during birth, pregnancy, from the umbilical cord, and breast milk were tested for PCB congeners and organochlorine pesticides. PCB153 levels in these media were comparatively low in relation to other studies. Measurements of PCBs in samples taken during the second trimester of pregnancy, at birth, and in the umbilical cord were strongly associated. Particular measurements of PCB153 and PCB180 among those subjects with concluded sampling of blood samples from mothers and cord samples were significantly correlated. Maternal blood measurement can reliably estimate the fetal exposure to PCBs during the second trimester. This assessment is consistent for PCB 153 and PCB 180 [60].

Additionally, BPA was examined in urine samples from women in the 1st and 3rd trimester of pregnancy. Psychomotor alongside cognitive growth was measured using psychologist-based scales. BPA exposure concentrations in the highest tectile were associated with a reduction of psychomotor scores at 1 year of age, however no associations with psychomotor outcomes at 4 years were shown. Exposure to BPA was linked with a higher risk of ADHD symptoms at 4 years old. Overall, the results indicate that BPA exposure prenatally does not disturb cognitive growth up to 4 years old. Links are demonstrated with ADHD-correlated signs and psychomotor development at initial ages but do not appear to remain until later ages [61]. Furthermore, hormone-induced alterations in brain composition and role indicate that EED exposure may be connected with sex-specific modifications in behavior. BPA has shown to alter androgen, estrogen, and thyroid hormone signaling pathways. Mothers with measurable prenatal urinary BPA were correlated with relatively higher affecting and expressing depressed behavior, somatic problems, and ODD behaviors in boys. The results indicate higher behavior problems in school aged boys but not girls [62]. Additionally, it has been shown that communicative effects of developmental exposure to a low dose of BPA with detail to the maternal environment, time of exposure, and sex & age. During both testing ages, females whom were exposed presented suggestion of higher anxiety and were less disposed to search a new environment. The results specify that sexually dimorphic behaviors are delicate to endocrine disruption during important developmental periods, specifically during important early neonatal stages [63]. BPA exposure during gestation has been additionally suggested as a risk element of neurobehavioral disorders. Exposure to a regular low-dose of BPA during pregnancy tested offspring of mice. Mice who were exposed had offspring that had increased anxiety-like behavior [64].

# 4. Interaction of ID proteins & EEDs in neurocognitive disorders

In order to investigate how ID proteins and environmental exposures affect neurocognitive deficiencies at a genetic level, we used various publicly accessible databases to help support our understanding. First we used Comparative Toxicogenomic Database (CTD), a research tool that determines chemical-gene and chemical-disease association, to help decipher gene-environment or gene-EED interactions involved in the production of various neurocognitive deficiencies [65]. To demonstrate an overlapping interaction between neurocognitive disorders, we established a list of neurocognitive-interacting genes (36,367), mood disorder interacting-genes (31,182), and neurodegenerative-interacting genes (30,127). As shown in Figure 1, 26,081 genes are commonly interacting. We also selected interacting genes with estrogenic endocrine disruptors PCB & BPA. We established a gene list for both: PCBs having 7,825 interacting genes and BPA having 20,504 interacting genes. Because ID proteins are our candidate gene group, we also generated a list of overlapping genes between each of the ID proteins (ID1, ID2, ID3, & ID4) and established interacting genes between the two chosen EEDs and neurocognitive disorder categories. We established that 63 genes interact with EEDs, our ID proteins, and neurocognitive disorders as shown in Figure 2 and Table 1 [65]. To demonstrate an interaction between these 63 genes; we furthermore inputted them

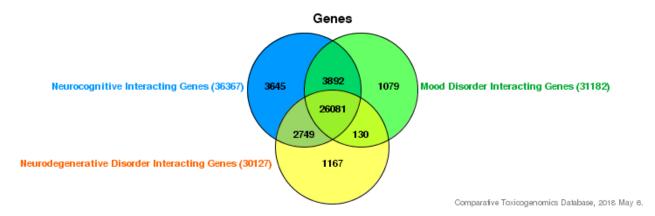
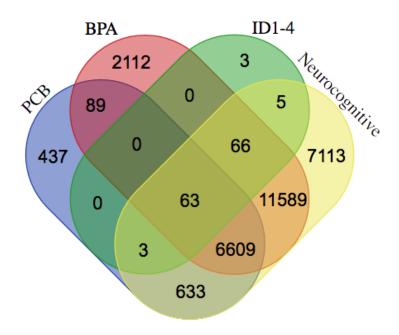


Figure 1. Interacting genes of neurocognitive categories. Shown are 26,081 genes that interact within the three categories.



**Figure 2.** Venn diagram displaying interacting genes between EEDs (PCBs & BPA), ID1-4, and neurocognitive disorders. Results show 63 overlapping genes.

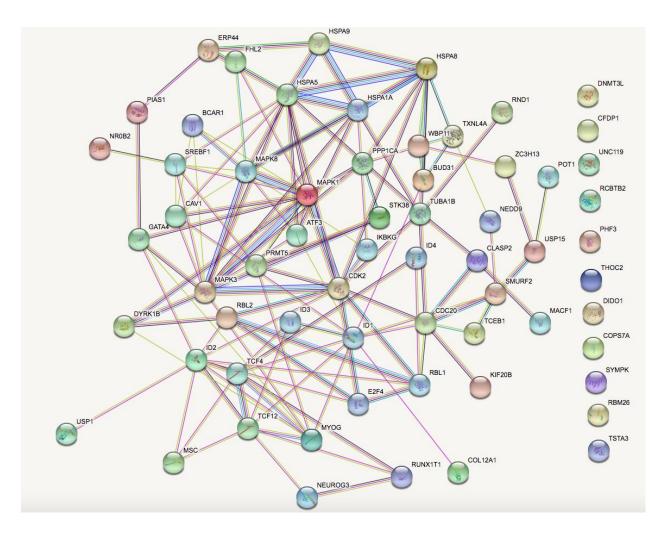
 $\textbf{Table 1.} \ \ \textbf{Gene symbols \& gene names of 63 overlapping EED-ID protein-neurocognitive genes}.$ 

Gene Symbol	Gene Name		
ATF3	Activating transcription factor 3 (ATF3)		
BCAR1	BCAR1, Cas family scaffolding protein (BCAR1)		
BUD31	BUD31 homolog (BUD31)		
CAV1	caveolin 1 (CAV1)		
CDC20	cell division cycle 20 (CDC20)		
CDK2	cyclin-dependent kinase 2 (CDK2)		
CFDP1	craniofacial development protein 1 (CFDP1)		
CLASP2	cytoplasmic linker associated protein 2 (CLASP2)		
COL12A1	collagen type XII alpha 1 chain (COL12A1)		
COPS7A	COP9 signalosome subunit 7A (COPS7A)		

DIDO1	death inducer-obliterator 1 (DIDO1)			
DNMT3L	DNA methyltransferase 3 like (DNMT3L)			
DYRK1B	dual specificity tyrosine phosphorylation regulated kinase 1B (DYRK1B)			
E2F4	E2F transcription factor 4 (E2F4)			
ELOC	elongin c (ELOC)			
ERP44	endoplasmic reticulum protein 44 (ERP44)			
FHL2	four and a half LIM domains 2 (FHL2)			
GATA4	GATA binding protein 4 (GATA4)			
HSPA1A	heat shock protein family A member 1A (HSPA1A)			
HSPA5	heat shock protein family A member 5 (HSPA5)			
HSPA8	heat shock protein family A member 8 (HSPA8)			
HSPA9	heat shock protein family A member 9 (HSPA9)			
ID1	inhibitor of DNA binding 1, HLH protein (ID1)			
ID2	inhibitor of DNA binding 2, HLH protein (ID2)			
ID3	inhibitor of DNA binding 3, HLH protein (ID3)			
ID4	inhibitor of DNA binding 4, HLH protein (ID4)			
IKBKG	inhibitor of kappa light polypeptide gene enhancer in B-cells, kinase gamma (IKBKG)			
KIF20B	kinesin family member 20B (KIF20B)			
MACF1	microtubule-actin crosslinking factor 1 (MACF1)			
MAPK1	mitogen-activated protein kinase 1 (MAPK1)			
MAPK3	mitogen-activated protein kinase 3 (MAPK3)			
MAPK8	mitogen-activated protein kinase 8 (MAPK8)			
MSC	Musculin (MSC)			
MYOG	Myogenin (MYOG)			
NEDD9	neural precursor cell expressed, developmentally down-regulated 9 (NEDD9)			
NEUROG3	neurogenin 3 (NEUROG3)			
NR0B2	nuclear receptor subfamily 0 group B member 2 (NR0B2)			
PHF3	PHD finger protein 3 (PHF3)			
PIAS1	protein inhibitor of activated STAT 1 (PIAS1)			
POT1	protection of telomeres 1 (POT1)			
PPP1CA	protein phosphatase 1 catalytic subunit alpha (PPP1CA)			
PRMT5	protein arginine methyltransferase 5 (PRMT5)			
RBL1	RB transcriptional corepressor like 1 (RBL1)			
RBL2	RB transcriptional corepressor like 2 (RBL2)			
RBM26	RNA binding motif protein 26 (RBM26)			
RCBTB2	RCC1 and BTB domain containing protein 2 (RCBTB2)			
RND1	RNA binding motif protein 26(RBM26) Rho family GTPase 1 (RND1)			
RUNX1T1	RUNX1 translocation partner 1 (RUNX1T1)			
SMURF2	SMAD specific E3 ubiquitin protein ligase 2 (SMURF2)			
SREBF1	sterol regulatory element binding transcription factor 1 (SREBF1)			
STK38	serine/threonine kinase 38 (STK38)			
SYMPK	Symplekin (SYMPK)			
TCF12	transcription factor 12 (TCF12)			
TCF4	transcription factor 4 (TCF4)			
THOC2	THO complex 2(THOC2)			
TSTA3	tissue specific transplantation antigen P35B (TSTA3)			

TUBA1B	tubulin alpha 1b (TUBA1B)			
TXNL4A	thioredoxin like 4A (TXNL4A)			
UNC119	unc-119 Lipid Bind Chaperone (UNC119)			
USP1	ubiquitin specific peptidase 1 (USP1)			
USP15	ubiquitin specific peptidase 15 (USP15)			
WBP11	WW domain binding protein 11 (WBP11)			
ZC3H13	zinc finger CCCH-type containing 13 (ZC3H13)			

STRING, a recognized database that can provide protein-to-protein interactions. Overall, the interactions contain either indirect (functional) or direct (physical) connections, which stem from interactions gathered from various databases, computational prediction, and from transfer of knowledge between organisms. As seen in Figure 3, STRING provides a network of these common 63 genes and additionally provides connections between all the genes [66-67]. Furthermore, to demonstrate importance of these genes we also used KEGG pathway, a collection of pathways representing knowledge for molecular reaction, interaction, and relation networks. We established that these genes were involved in 71 biological/molecular pathways [68]. The top 5 are represented in Table 2 and it is revealed that these pathways play functionality in neurocognitive disorders [69-74].



**Figure 3.** Gene network of 63 common interacting genes between EEDs, ID proteins, & neurocognitive disorders.

**Table 2.** Top 5 pathways with common 63 interacting genes with EEDs, ID, & neurocognitive disorders.

Pathway Name	Gene Count	Matching Genes in Network (Nodes)
TGF-beta signaling pathway	9	E2F4, ID1, ID2, ID3, ID4, MAPK1, MAPK3, RBL1, SMURF2
Toxoplasmosis	6	HSPA1A, HSPA8, IKBKG, MAPK1, MAPK3, MAPK8
Focal adhesion	6	HSPA1A, HSPA5, MAPK1, MAPK3
Viral carcinogenesis	6	FHL2, IKBKG, MAPK1, MAPK3, MAPK8
MAPK signaling pathway	6	BCAR1, CAV1, MAPK1, MAPK3, MAPK8, PPP1CA

### 5. Conclusion

ID proteins have been demonstrated to be involved with neurocognitive disorders. Studies have reported links between neurocognitive disorders and exposure to EEDs such as PCBs & BPA. Based on the evidence discussed in this review, we have demonstrated that exposure to EEDs may activate ID proteins to alter molecular mechanisms, additionally changing neurocognitive disorder outcomes. It is important to understand how the influence of EEDs and ID proteins affects neurocognitive perturbations. This essential information will allow research scientists, toxicologists, and public health professionals to discover novel avenues for the prevention and treatment of these types of disorders.

**Conflict of Interest:** The authors declare no conflict of interest.

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