

The role of ALDH2 and its substrates in central nervous system disorders

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Yashuang Chen^a, Fengyuan Tiana^b, Shu Meng^c, Jiayu Wang^d

ABSTRACT

Background: Aldehyde Dehydrogenase 2 (ALDH2) is a mitochondrial dehydrogenase enzyme primarily tasked with the detoxification of acetaldehyde produced from alcohol metabolism and endogenous aldehydes. These aldehyde compounds, such as 4-Hydroxynonenal (4-HNE) and 3,4-dihydroxyphenylacetaldehyde (DOPAL), are predominantly generated through lipid peroxidation and are known to form adducts with proteins, DNA, and lipids, thereby inducing neurotoxicity.

Methods: This review examines the role of ALDH2 in central nervous system (CNS) diseases by analyzing epidemiological studies and disease models. The focus is on understanding the impact of the ALDH2 rs671 G>A polymorphism, which reduces or eliminates enzyme activity.

Results: This genetic polymorphism of ALDH2 is closely associated with the onset of various central nervous system (CNS) diseases such as Alzheimer's disease (AD), Parkinson's disease (PD), schizophrenia, and stroke potentially linked to the accumulation of aldehyde compounds within the CNS.

Conclusions: A deeper understanding of the mechanisms by which ALDH2 influences these diseases will enhance therapeutic strategies for patients carrying the ALDH2 rs671 polymorphism and also offer new insights for the prevention and diagnosis of these conditions.

Keywords: ALDH2; Alzheimer's disease; Parkinson's disease; Schizophrenia; Stroke; 4-HNE.

^a Department of Immunology, Institute of Basic Medical Sciences, Chinese Academy of Medical Sciences, Beijing, China. Contributed equally.

^b Department of Immunology, Institute of Basic Medical Sciences, Chinese Academy of Medical Sciences, Beijing, China. Contributed equally.

^c Department of Immunology, Institute of Basic Medical Sciences, Chinese Academy of Medical Sciences, Beijing, China. Corresponding Author: mengshu129@163.com

^d Department of Immunology, Institute of Basic Medical Sciences, Chinese Academy of Medical Sciences, Beijing, China. Corresponding Author: wang1208jiau@163.com

However, at times, the production of ROS may surpass the compensatory capacity of mitochondria, leading to mitochondrial damage, subsequent initiation of mitochondrial dysfunction, apoptosis, and cell death(4, 5).

In the realm of neurobiology, the dysregulation of ROS and the resultant augmented oxidative stress, specifically manifesting as lipid peroxidation in mitochondria, lead to the formation of various aldehyde species that exhibit reactivity and toxicity(6). These aldehydes are prevalent not only in external environments, such as automobile emissions, chemical manufacturing processes, cosmetics, and food and beverages(7), but also originate within the body through the metabolic processes of neurotransmitters, amino acids, and lipids. Among them, 4-hydroxynonenal (4-HNE), acetaldehyde, acrolein, formaldehyde, 3,4-dihydroxyphenylacetaldehyde (DOPAL), and 3,4-dihydroxyphenylglycolaldehyde (PDPEGAL) are particularly crucial high-reactivity aldehyde species. The overabundance of 4-HNE, which has been the subject of extensive research, can disrupt cellular operations and precipitate apoptosis. These aldehydes are known to form conjugates with vital biomolecules, including DNA, proteins, and enzymes(8, 9), including mitochondrial electron transport chain proteins, proteasome proteins, and cytoskeletal proteins, leading to their functional impairment(6). The aggregation of these aldehydes, coupled with mitochondrial dysfunction, is implicated in the etiology of numerous neurological conditions such as Huntington's disease (HD), Alzheimer's disease (AD), Parkinson's disease (PD), various forms of dementia, ataxia, seizures, cerebral pathologies associated with hypertension, and ischemic stroke(10-12).

Mitochondrial aldehyde dehydrogenase 2 (ALDH2), as a critical detoxifying enzyme, is primarily responsible for metabolizing endogenous and exogenous aldehyde species. It not only functions in alcohol metabolism but also plays a crucial role in processes associated with oxidative stress(10, 13, 14). ALDH2 deficiency arises from a polymorphism in the 487th amino acid structure, where the substitution of glutamate with lysine results in an allelic gene change from G (wild-type allele *1) to A (mutant allele *2), significantly reducing the enzyme's catalytic activity, leading to the accumulation of toxic aldehyde species. However, the association between ALDH2 enzyme activity deficiency and the risk assessment of CNS disorders has not been extensively studied, and

research models are lacking. Therefore, gaining a deeper understanding of the function and molecular regulatory mechanisms of ALDH2 in the CNS is crucial for unraveling the complexity of neurological disorders. This review aims to amalgamate existing studies, specifically concentrating on the involvement of ALDH2 in conditions such as AD, PD, schizophrenia, and stroke.

2. ALDH2 AND ENDOGENOUS ALDEHYDE METABOLISM

The ALDH2 enzyme, localized in mitochondria, is encoded by a gene situated on the 12q24 chromosome region. This gene is structured into 13 exons and translates into a 517 amino acid long polypeptide(15). The enzyme is targeted to the mitochondrial matrix by a mitochondrial targeting sequence at its N-terminus, comprising 17 amino acids. Upon arrival, this sequence is cleaved by mitochondrial proteases, a step critical for the enzyme's maturation(16). Within the genetic structure of ALDH2, researchers have identified a total of 535 single nucleotide polymorphisms (SNPs), with the rs671 SNP in exon 12 receiving significant scientific focus. The rs671 SNP encompasses a G to A nucleotide transition, resulting in an amino acid change from glutamate to lysine at the 487th position (E487K or E504K, accounting for the cleavage of the N-terminal sequence). This specific mutation leads to an alteration in hydrogen bonding, which adversely affects the binding site for NAD⁺, a cofactor essential for aldehyde oxidation. Consequently, the rs671 variant is associated with reduced enzymatic activity of the dehydrogenase(17).

ALDH2 plays a pivotal role in metabolizing toxic aldehydes generated directly or indirectly from ethanol metabolism. Ethanol can be oxidized to acetaldehyde via the following pathways: in most cases, ethanol is oxidized to acetaldehyde by alcohol dehydrogenases (ADHs) in the cytoplasm; ethanol is oxidized through the microsomal ethanol-oxidizing system (MEOS); in the presence of catalase (CAT), ethanol is converted to acetaldehyde(18). ALDH2 further metabolizes acetaldehyde into acetate, which can either exit the cell through a carrier or undergo enzymatic transformation into acetyl-coenzyme A (acetyl-CoA) by cytosolic acetyl-CoA synthetase 2 (ACSS2)(19). Acetyl-CoA subsequently participates in several metabolic pathways, including the tricarboxylic acid (TCA) cycle and the oxidative phosphorylation (OXPHOS)

pathway(20), eventually being metabolized into water and carbon dioxide. Moreover, malondialdehyde (MDA), generated through lipid peroxidation, undergoes conversion to malonic semialdehyde facilitated by ALDH2. Subsequently, malonic semialdehyde further metabolizes into malonic acid (MOA). Alternatively, MDA can undergo a process where a carboxyl group is removed, resulting in its conversion into acetaldehyde. Under the influence of ALDH2, a portion of the 4-NHE generated during lipid peroxidation converts into 4-hydroxy-2-nonenic acid (HNA). Concurrently, the remaining portion is catalyzed by other reductase enzymes, like glutathione-S-transferase (GST) and aldo-keto reductases (AKRs)/ADHs(21). The reduction or functional impairment of ALDH2 activity disrupts its protective role against aldehydes, resulting in escalated production of ROS and enhanced oxidative cellular damage, which may culminate in various human pathologies.

ALDH2, a vital enzyme in aldehyde detoxification, experiences a reduction in its dehydrogenase function due to the rs671 G>A polymorphism. This genetic variation results in an increased buildup of acetaldehyde and other endogenous aldehydes, often evidenced by the facial flushing observed post-alcohol intake. Intriguingly, this polymorphism is predominantly found in East Asian populations, with a prevalence of 30-50%, in stark contrast to less than 5% in individuals of European ancestry (22). Due to the activation of ALDH2 through multiple pathways and its close association with various pathophysiological processes, including its involvement in and impact on the occurrence and development of diverse diseases and metabolic processes, ALDH2, as a key regulatory enzyme in oxidative stress, has garnered increasing attention. Given its exceptionally high mutation rate in Asian populations, gaining a deeper understanding of the role of ALDH2 in the diseases of the Chinese population holds significant significance.

3. ALDH2 AND NEURODEGENERATIVE DISEASES

The cerebral environment, characterized by an abundance of polyunsaturated fatty acids, engenders a heightened oxygen presence within its lipid membranes. This aspect renders the brain more vulnerable to lipid peroxidation, a consequence of oxidative stress, as well as mitochondrial impairments and subsequent accumulation of aldehyde

derivatives. These elements are crucial contributors to the deterioration of memory functions, cognitive deficits, and the pathogenesis of neurodegenerative disorders(23). This section reviews epidemiological evidences and findings from model-based studies related to ALDH2 in AD and PD, both of which are key neurodegenerative conditions, and assesses the implications of aldehyde buildup in these neurodegenerative states.

3.1. ALDH2 and Alzheimer's disease

Currently, Alzheimer's disease (AD) stands as the most widespread neurodegenerative disorder, yet efficacious treatment modalities remain elusive. Data reveals that in the United States alone, over 4.5 million individuals are afflicted with AD, a figure anticipated to escalate to 13.8 million by 2060 owing to increased longevity(24). The development of AD is closely associated with oxidative stress, a phenomenon commonly observed in neurodegenerative events(25). Studies have identified signs of oxidative stress in the blood, cerebrospinal fluid, and brain tissues of AD patients, highlighting its critical role in the disease's pathophysiology(26). ROS facilitate the production of various aldehyde by-products, among which 4-HNE is considered a fundamental signaling molecule in the pathology of AD. Similar to other aldehydes, the detoxification of 4-HNE depends on the enzymatic activity of ALDH2. When ALDH2 activity is inhibited, cells become more susceptible to damage induced by 4-HNE. Therefore, an enhanced comprehension of the role of ALDH2 in AD, especially its impact on oxidative stress and 4-HNE metabolism, could provide crucial insights for developing strategies to prevent or reverse the progression of AD.

The relationship between ALDH2 polymorphism and AD remains inconclusive, as evidenced by varied research outcomes summarized in Table 1. Two separate investigations conducted in South Korea, with participant pools of 690 and 510, respectively, did not establish a significant link between the ALDH2 polymorphism and AD(27, 28). Similarly, research within the Mongolian demographic of China also failed to recognize the ALDH2 gene as a contributing factor for AD(29). Conversely, outcomes from Japanese research present a contrasting picture. One study involving 447 individuals indicated the genotype frequency for the ALDH2*2 allele was significantly higher in the patients with AD than individuals without AD(30).

A cohort study in Japan, encompassing 271 AD patients, echoed these observations(31). Yet, another case-control study from Japan did not find any notable connection between the ALDH2*2 variant and AD(32).

Further investigations in China, particularly a study focusing on individuals over 90 years old, found a link between ALDH2*2 polymorphism and cognitive impairment(33). Additionally, two independent Chinese studies suggested that ALDH2*2 polymorphism is associated with an increased risk of AD(34, 35). A comprehensive meta-analysis, aggregating data from 1824 AD cases and 4300 controls across six Asian case-control studies, inferred that possession of the ALDH2*2 allele might escalate AD risk(29, 32, 34-37). Recently, an analysis of 469 human brain tissue samples from a Chinese brain tissue repository revealed that the ALDH2 rs671 variant increases amyloid-beta pathology, unveiling its molecular mechanism at the chemical bond and organelle levels for the first time. The connection between ALDH2 and AD, therefore, continues to be a topic of debate, underscoring the need for expanded research endeavors involving larger and more heterogeneous cohorts to arrive at more conclusive insights.

In AD research, the use of cellular and animal models has significantly contributed to elucidating the disease's underlying mechanisms. Initial investigations utilizing the PC12 cell line demonstrated that the presence of the ALDH2*2 allele results in diminished mitochondrial ALDH2 enzyme activity. These cells, when subjected to external 4-HNE, displayed an increased susceptibility, highlighting their vulnerability under oxidative stress conditions(38). Furthermore, research conducted on rat primary neuronal cells has shown that an elevation in ALDH2 enzymatic activity serves as a defensive mechanism, countering synaptic damage and oxidative stress triggered by 4-HNE(39).

In terms of animal models, transgenic AD mouse models have been pivotal for investigating the impact of oxidative stress on AD's pathological and behavioral aspects(40). These animal models are engineered to express mutated versions of the human amyloid precursor protein (APP) and harbor a detrimental mutation in mitochondrial ALDH2. Observations from this model include a reduction in lifespan, expedited amyloid plaque formation, enhanced tau protein phosphorylation, and increased gliosis. Furthermore, the DAL101 mouse model, distinguished by its deficient ALDH2

activity, represents an age-related dementia model. It displays progressively worsening pathological characteristics, mirroring chronic oxidative stress exposure(41).

The ALDH2^{-/-} mouse model has proven to be exceptionally valuable in mimicking disease mechanisms and in the exploration of potential therapeutic approaches. In these models, the lack of ALDH2 activity results in an accumulation of 4-HNE, which in turn initiates a series of pathological changes characteristic of AD. These changes include an increase in A β and phosphorylated tau, a decline in synaptic proteins, cerebral atrophy, and cognitive deficits(42). Similar findings were also reported by Ohta *et al.* in 18-month-old ALDH2 knockout mice(43). Consequently, ALDH2^{-/-} mice are increasingly recognized as an accurate animal model for sporadic AD, effectively replicating the oxidative stress-induced cognitive impairments observed in AD pathology(44-46). Intriguingly, dietary intervention with D-PUFAs in ALDH2^{-/-} mice has shown a reversal in cognitive impairments, highlighting D-PUFAs as a promising approach for countering AD-related cognitive decline(47). Furthermore, the overexpression of ALDH2 in APP/PS1 mice resulted in significant cognitive improvements, emphasizing the critical importance of ALDH2 activity in AD pathology(48). These models have not only facilitated a more profound understanding of AD pathology but also opened avenues for the development of novel therapeutic strategies.

3.2. ALDH2 and Parkinson's disease

Parkinson's disease (PD) is a degenerative neurological condition marked by the progressive loss of dopaminergic neurons in the substantia nigra, resulting in involuntary movements and varying degrees of dementia. Epidemiologically, PD affects approximately 1% of the population aged over 65, with rates increasing to 4-5% in individuals aged over 85(49). Given the increasing life expectancy in Asia, projections suggest that by 2030, the number of PD patients might reach approximately 5 million, accounting for nearly 60% of PD cases worldwide(50). The etiology of PD involves familial genetic defects, including mutations in α -synuclein, PARKIN, PINK1, DJ-1, and LRRK2 genes, as well as hereditary and environmental toxins(51). Additionally, oxidative stress in mitochondria is considered a significant influencing factor in

PD(52). Research has observed a decline in mitochondrial numbers in the dopaminergic neurons of the substantia nigra in PD patients, emphasizing the crucial role of mitochondrial integrity in these neurons and their vulnerability to the detrimental effects of aldehydes.

As previously mentioned, the mutation rate of ALDH2 rs671 can reach up to 50% in the East Asian population, making it crucial to understand the impact of rs671 polymorphism on the Asian PD population. Epidemiological studies, including one by Zhao *et al.* using polymerase chain reaction-restriction fragment length polymorphism (PCR-RFLP) analysis, have shown that the risk of PD in the Chinese population carrying the ALDH2 gene GA and AA genotypes is significantly higher than those with the GG genotype(34). However, a cohort study based on the Chinese population indicated no correlation between ALDH2 rs671 polymorphism and PD(53). A comprehensive meta-analysis encompassing 5315 samples and 9 case-control studies concluded that there is no notable correlation observed between the ALDH2 rs671 polymorphism and the risk of PD(54). A case-control study comparing 93 Parkinson's disease patients with different genotypes to 297 healthy controls found no association between the ALDH2 genotype and PD. Nonetheless, the patients with PD carrying ALDH2*2 allele are more likely to develop heightened daytime sleepiness and have difficulty in maintaining asleep. These results support the hypothesis that the accumulation of neurotoxic monoamine neurotransmitter aldehyde metabolites, resulting from decreased ALDH2 enzyme activity, could contribute to more severe loss of monoaminergic neurons, thus exhibiting more severe symptoms in wakefulness and sleep regulation(55). Moreover, PD patients with the ALDH2*1 allele genotype are more prone to depressive moods(56).

In the context of animal model research, α -synuclein has been identified as highly susceptible to modification by reactive aldehydes, potentially leading to the formation of neurotoxic oligomers. Specifically, in rat PD models, the administration of 6-hydroxydopamine (6-OHDA) has shown to significantly reduce ALDH activity in the striatum(57). Furthering this, Stott's research also found that 6-OHDA-induced striatal damage in mice led to a decrease in the number of dopaminergic neurons in the substantia nigra, accompanied by an increase in 4-HNE and DOPAL. Similarly, the ratio

of DOPAL to dopamine significantly increased in the striatum of double gene knockout mice, indicating increased neurotoxicity, which mirrors the situation in the striatum of PD patients' post-mortem. The neurotoxin 1-methyl-4-phenylpyridinium ion (MPP⁺) is commonly used to establish PD models, and in PC12 cells, treatment with MPP⁺ increased the aldehyde load of MDA and 4-HNE. The specific inhibitor of ALDH2, daidzin, prevented the reduction in cell viability and the increase in apoptosis, oxidative stress, and aldehyde stress induced by MPP⁺, indicating a protective role of ALDH2 against neurotoxicity and PD(58). Through the study of these models, we not only deepen our understanding of the pathophysiological mechanisms of PD but also provide possible directions for developing new therapeutic methods.

4. ALDH2 AND SCHIZOPHRENIA

Schizophrenia, a complex psychiatric condition manifesting in behavioral, emotional, and cognitive impairments, often follows a chronic trajectory(59). Since the 1930s, when Roy Hoskins first posited the influence of oxidative stress in the genesis and progression of schizophrenia, substantial attention has been focused on patients, whether untreated, in the early stages of the disease, or in its chronic phases(60-62). In the research conducted by Wang *et al.* immunohistochemical techniques were employed to measure HNE-protein adducts, revealing a 47% increase in HNE levels in the anterior cingulate cortex of post-mortem schizophrenic patients compared to healthy controls, thereby suggesting a correlation between schizophrenia and oxidative stress(63). Additionally, a tight association between ALDH2 and oxidative stress was observed; research involving 1316 schizophrenic patients and 1349 healthy controls indicated a positive correlation between the ALDH2 rs671 G A mutation and schizophrenia(64). Mounting evidence underscores the significant role of inflammation in schizophrenia. For instance, a study demonstrated that treatments with risperidone and amisulpride alleviated negative symptoms such as affective flattening and social withdrawal in schizophrenic patients with ALDH2 polymorphism(65). Proteome-wide association studies (PWAS), combined with MR and genetic colocalization analysis, further confirmed that an increase in ALDH2 protein abundance helps reduce feelings of loneliness in schizophrenic patients(66).

Current studies demonstrate significant differences between male and female schizophrenia patients in terms of disease onset, progression, treatment response, and brain structural abnormalities(67). In the study by Ramos-Loy *et al.*, a gender-based analysis of serum HNE levels in patients suffering from paranoid schizophrenia was performed. The findings revealed elevated 4-HNE concentrations in male patients compared to their female counterparts. This disparity could potentially be ascribed to the neuroprotective role of estrogens against oxidative stress, which are more prevalent in females(68). Additionally, the incidence of schizophrenia in women has been observed to rise during menopause, a phase characterized by a reduction in estrogen's protective influence(69). However, contrary results were reported by Wang *et al.* (63), whose immunohistochemical study did not identify any significant gender-based differences in HNE levels among schizophrenia patients. Thus, the existence of gender-specific variations in 4-HNE levels within schizophrenia remains an open question, highlighting the need for further focused investigations to clarify this important dimension.

5. ALDH2 AND STROKE

Stroke stands as a predominant contributor to chronic disability and mortality worldwide, a statement corroborated by robust evidence(70). Remarkably, China experiences the highest incidence and fatality rates associated with stroke globally, with staggering figures exceeding 2.4 million new cases and 1.1 million deaths each year. Significantly, ischemic stroke constitutes approximately 87% of these cases(71). This alarming prevalence underscores the imperative need for comprehensive genetic research in stroke susceptibility. Such studies are pivotal for the identification of high-risk demographics and the elucidation of the intricate etiological pathways of stroke. Additionally, these researches are instrumental in advancing predictive methodologies for stroke progression, thereby facilitating the development and application of efficacious preventative strategies within clinical paradigms(72, 73).

Ischemic stroke, characterized as a life-threatening cerebrovascular condition, has been the focus of recent scientific inquiry. Two groundbreaking genome-wide association studies (GWAS) have shed light on the genetic underpinnings of

this ailment. These studies have pinpointed a correlation between genetic markers near the ALDH2 gene located on chromosome 12q24 and the onset of stroke(74). Intriguingly, the ALDH2 rs671 polymorphism has been identified as a genetic determinant in East Asian populations, primarily linked to the phenomenon of facial flushing post-alcohol consumption(75). An expansive study involving a cohort of over half a million participants further explored this link, indicating a heightened risk of stroke associated with alcohol consumption, especially in individuals with the ALDH2 rs671 G>A variant(10, 76, 77). Concurrently, dedicated research within the Han Chinese demographic, coupled with comprehensive meta-analyses focusing on the broader Chinese population, unanimously echoes these findings. They affirm the ALDH2 rs671 variant as a significant risk factor for ischemic stroke(78-80).

Expanding the scope of this research, a distinct study targeting the Hakka population in China, involving 329 hemorrhagic stroke patients alongside 515 control subjects, meticulously analyzed the ALDH2 rs671 genotypes. The study unveiled notable statistical disparities in the genotype and allele distributions between the patient cohort and control group. Crucially, it revealed a markedly amplified risk of hemorrhagic stroke in individuals carrying the ALDH2*2 allele. This revelation propels the hypothesis that the ALDH2 rs671 polymorphism might serve as a salient risk factor for hemorrhagic stroke, further enriching our understanding of stroke pathophysiology(81).

The role of chronic inflammation, tissue hypoxia, and oxidative stress in cerebrovascular diseases is substantial(82). ALDH2 emerges as a pivotal biological agent in this context, exerting protective effects against oxidative damage. It achieves this function by metabolizing aldehydes, compounds known for their toxicity, thereby mitigating oxidative stress. This function of ALDH2 is particularly pronounced in the case of ischemic stroke, where its impact is most palpable(83, 84). Additionally, research focusing on hemorrhagic stroke has unveiled a correlation between ALDH2 activity and key cardiovascular markers, namely systolic and diastolic blood pressure, as well as its association with serum levels of HDL and LDL cholesterol(85, 86). These findings collectively offer a more profound insight into the complex interplay between genetic factors and stroke, paving the way for targeted therapeutic interventions and preventive measures.

Cellular and animal model studies have indicated a potential link between ALDH2 and stroke. Through an unbiased proteomic search, the study revealed ALDH2 deficiency in stroke-prone spontaneously hypertensive rats (SHR-SP) compared to spontaneously hypertensive rats. It was concluded that the deficiency of ALDH2 plays a causative role in neuronal injury, as evidenced by in vitro studies showing that overexpression or activation of ALDH2 provides neuroprotection by clearing 4-HNE(87). Some studies suggest that ROS may act as potential activators of the JNK signaling pathway, triggering neuronal apoptosis during cerebral ischemia. Overexpression of ALDH2, by inhibiting JNK-induced caspase 3 activation and transcription, has been shown to reduce mitochondrial apoptosis

in both in vivo rat focal cerebral ischemic model and in vitro neuronal cell oxygen-glucose deprivation/re-oxygenation (OGD/R) models of cerebral ischemia(88). These experimental findings suggest that ALDH2 might play a significant role in the development of ischemic stroke.

6. CONCLUSION

ALDH2 plays a key role not only in preventing and managing alcohol addiction but also in the intricate regulatory pathways of neurodegenerative ailments and cerebrovascular disorders (Figure 1) Diminished ALDH2 enzymatic activity leads to the accumulation of aldehydes, which are inherently volatile and strong electrophilic reagents. These toxic

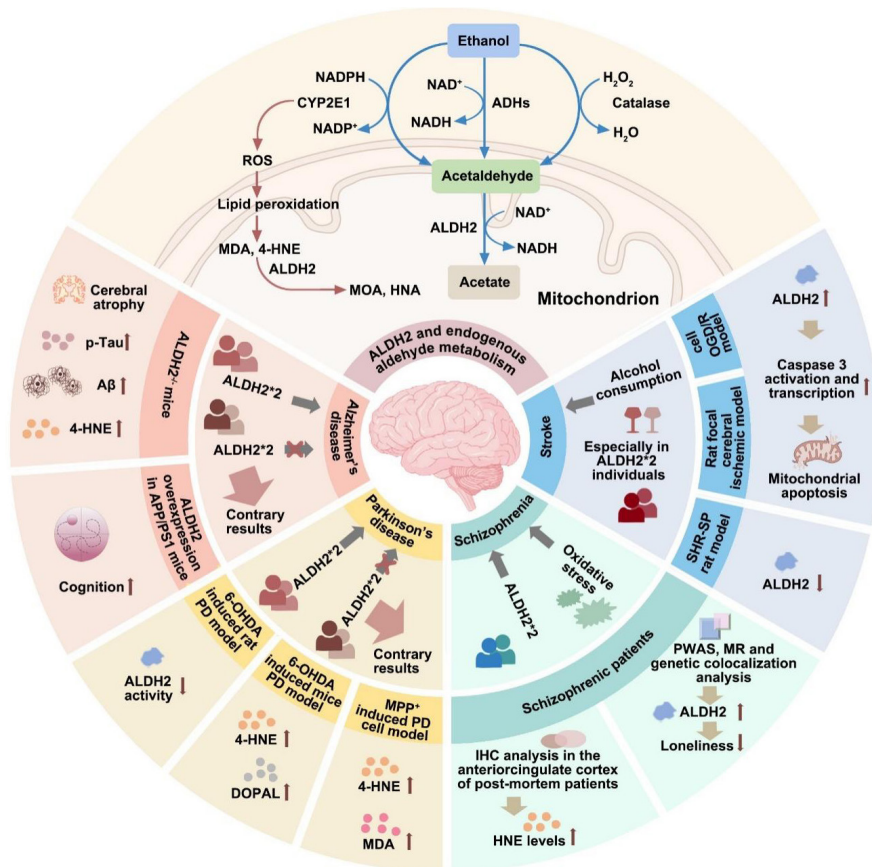


Figure 1. The effect of reduced ALDH2 enzyme activity and the role of ALDH2 in central nervous system diseases. 4-HNE, 4-Hydroxynonenal; 6-OHDA, 6-Hydroxydopamine; ADHs, Alcohol dehydrogenases; ALDH2, Aldehyde dehydrogenase 2; APP, Amyloid precursor protein; Aβ, Amyloid-beta peptides; CYP2E1, Cytochrome P450 2E1; DOPAL, 3,4-Dihydroxyphenylacetaldehyde; HNA, 4-Hydroxy-2-nonenic acid; MDA, Malondialdehyde; MOA, Malonic acid; MPP+, 1-Methyl-4-phenylpyridinium ion; OGD/R, Oxygen-glucose deprivation/re-oxygenation; MR, Magnetic resonance; PS1, Presenilin-1; p-Tau, Phosphorylated-tau protein; PWAS, Proteome-wide association studies; ROS, Reactive oxygen species; SHR-SP, Stroke-prone spontaneously hypertensive rats.

aldehydic substances readily interact with macromolecules, perturbing normal biological functions and leading to cellular dysregulation. Epidemiological studies, genetic analyses, and animal model research have pinpointed the excessive buildup of aldehydes as a key molecular mechanism implicated in the pathogenesis of neurodegenerative conditions and cerebrovascular incidents. Considering the crucial role of mitochondrial ALDH2 in the detoxification of aldehydes within the brain, a deeper exploration of its function and impact in neurodegenerative diseases, schizophrenia and stroke becomes imperative, particularly for East Asian populations where ALDH2 deficiency is widespread. The insights

provided in this review illuminate the complex relationship among ALDH2, AD, PD, schizophrenia, and stroke, offering a nuanced understanding of these intricate interconnections.

Conflict of Interest Disclosures

The authors declare no conflicts of interest to declare.

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Nonstandard abbreviations and acronyms

Abbreviations

Full name

4-HNE	4-Hydroxynonenal
6-OHDA	6-Hydroxydopamine
Acetyl-CoA	Acetyl-coenzyme A
ACSS2	Acetyl-CoA synthetase 2
AD	Alzheimer's disease
ADHs	Alcohol dehydrogenases
AKRs	Aldo-keto reductases
ALDH2	Aldehyde dehydrogenase 2
APP	Amyloid precursor protein
CAT	Catalase
CNS	Central nervous system
DOPAL	3,4-Dihydroxyphenylacetaldehyde
GST	Glutathione-S-transferase
GWAS	Genome-wide association studies
HD	Huntington's disease
HNA	4-Hydroxy-2-nonenic acid
MDA	Malondialdehyde
MEOS	Microsomal ethanol-oxidizing system
MOA	Malonic acid
MPP ⁺	1-Methyl-4-phenylpyridinium ion
OGD/R	Oxygen-glucose deprivation/re-oxygenation
OXPHOS	Oxidative phosphorylation
PCR-RFLP	Polymerase chain reaction-restriction fragment length polymorphism
PD	Parkinson's disease
PDPEGAL	3,4-Dihydroxyphenylglycolaldehyde
PWAS	Proteome-wide association studies
ROS	Reactive oxygen species
SHR-SP	Stroke-prone spontaneously hypertensive rats
SNPs	Single nucleotide polymorphisms
TCA	Tricarboxylic acid

References	Published time	Region	Case	Control	Association between ALDH2 rs671 polymorphism and disease	Disease type
Kim, <i>et al.</i> (89)	2004	South Korea	80	229	No association	AD
Shin, <i>et al.</i> (27)	2005	South Korea	37	473	No association	AD
Zhou, <i>et al.</i> (29)	2010	Mongolia	106	100	No association	AD
Komatsu, <i>et al.</i> (32)	2014	Japan	130	201	No association	AD
Kamino, <i>et al.</i> (36)	2000	Japan	477	477	Increases AD risk	AD
Ueno, <i>et al.</i> (31)	2022	Japan	271	492	Increases AD risk	AD
Wang, <i>et al.</i> (35)	2008	China	188	223	Increases AD risk	AD
Ma, <i>et al.</i> (34)	2016	China	115	236	Increases AD risk	AD
Jin, <i>et al.</i> (33)	2021	China	1114	805	Increases AD risk	AD
Chen, <i>et al.</i> (54)	2019	Asia	Meta		Increases AD risk	AD
Yu, <i>et al.</i> (56)	2021	China	114	—	Associated with PD non-motor symptoms	PD
Michel, <i>et al.</i> (90)	2014	Germany	9	12	ALDH2 activity was significantly increased in the putamen of PD patients	PD
Zhang, <i>et al.</i> (53)	2015	China	584	582	No association	PD
Zhao, <i>et al.</i> (91)	2016	China	115	214	Increases PD risk	PD
Lin, <i>et al.</i> (55)	2019	China	83	—	PD patients carrying allele rs671 (A) are more likely to experience impairment in the regulation of arousal and sleep	PD
Chen, <i>et al.</i> (54)	2019	—	Meta		No association	PD
Zheng, <i>et al.</i> (64)	2017	China	1316	1349	Related to risk for schizophrenia	Schizophrenia
Lee, <i>et al.</i> (65)	2015	China	95	—	The ALDH2*2 genotype predicts that DM drug therapy reduces negative symptoms in schizophrenia patients	Schizophrenia
Zhang, <i>et al.</i> (81)	2023	China	329	515	Increases hemorrhagic stroke	Stroke
Shi, <i>et al.</i> (80)	2022	China	4727	5029	Increases ischemic stroke	Stroke
Jiang, <i>et al.</i> (79)	2020	China	2762	2503	Increases ischemic stroke	Stroke
Xu, <i>et al.</i> (78)	2019	China	3236	2893	Increases ischemic stroke	Stroke

AD: Alzheimer's disease, PD: Parkinson's disease, DM: Dextromethorphan

Table 1. Association between ALDH2 rs671 polymorphism and diseases.

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