

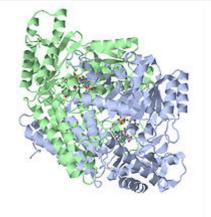
Glutamate decarboxylase

Glutamate decarboxylase or glutamic acid decarboxylase (GAD) is an enzyme that catalyzes the decarboxylation of glutamate to gamma-aminobutyric acid (GABA) and carbon dioxide (CO $_2$). GAD uses pyridoxal-phosphate (PLP) as a cofactor. The reaction proceeds as follows:

glutamate decarboxylase	
	Identifiers
EC no.	4.1.1.15 (https://www.en zyme-database.org/quer y.php?ec=4.1.1.15)
CAS no.	9024-58-2m
	Databases
IntEnz	IntEnz view (https://www.ebi.ac.uk/intenz/query?cmd=SearchEC&ec=4.1.1.15)
BRENDA	BRENDA entry (http://www.brenda-enzymes.org/enzyme.php?ecno=4.1.1.
ExPASy	NiceZyme view (https://e nzyme.expasy.org/EC/4. 1.1.15)
KEGG	KEGG entry (https://www.genome.jp/dbget-bin/www_bget?enzyme+4.1.1.
MetaCyc	metabolic pathway (http s://biocyc.org/META/sub string-search?type=NIL& object=4.1.1.15)
PRIAM	profile (http://priam.prab i.fr/cgi-bin/PRIAM_profile s_CurrentRelease.pl?EC =4.1.1.15)
PDB structures	RCSB PDB (https://www.rcsb.org/search?q=rcsb_polymer_entity.rcsb_ec_lineage.id:4.1.1.15) PDBe (https://www.ebi.ac.uk/pdbe/entry/search/index?ec_number:4.1.1.15) PDBsum (https://www.ebi.ac.uk/thornton-srv/

Gene Ontolog	у	databases/cgi-bin/enzym es/GetPage.pl?ec_numb er=4.1.1.15) AmiGO (http://amigo.gen eontology.org/amigo/ter m/GO:0004351) /
		QuickGO (https://www.e bi.ac.uk/QuickGO/term/ GO:0004351)
	Search	
PMC	h.g me N% 20	icles (https://www.ncbi.nlm.ni jov/entrez/query.fcgi?db=pub rd&term=4.1.1.15%5BEC/R %20Number%5D%20AND% pubmed%20pmc%20local% sb%5D)
PubMed	h.g	icles (https://www.ncbi.nlm.ni jov/entrez/query.fcgi?db=pub id&term=4.1.1.15%5BEC/R %20Number%5D)
NCBI	ih.	oteins (https://www.ncbi.nlm.n gov/protein?term=4.1.1.15% EC/RN%20Number%5D)

Glutamic acid decarboxylase 1



GAD67 derived from PDB: 2okj (http s://www.rcsb.org/structure/2okj)

Identifiers		
Symbol	GAD1	
Alt. symbols	glutamate decarboxylase 1	
	(brain, 67kD); GAD67	
NCBI	2571 (https://www.ncbi.nl	
gene	m.nih.gov/gene?cmd=retri	

	eve&dopt=default&list_uid
	s=2571&rn=1)
HGNC	4092 (https://www.genena
	mes.org/data/gene-symbo
	<pre>I-report/#!/hgnc_id/HGNC:</pre>
	4092)
OMIM	605363 (https://omim.org/
	605363)
RefSeq	NM 000817 (https://www.
<u></u>	ncbi.nlm.nih.gov/protein/N
	M_000817)
UniProt	Q99259 (https://www.unip
	rot.org/uniprot/Q99259)
	Other data
EC	4.1.1.15 (https://www.gen
number	ome.jp/dbget-bin/www_bg
	et?enzyme+4.1.1.15)
Locus	Chr. 2 q31 (https://omim.o
	rg/search/?index=geneMa
	p&search=2q31)
Search for	
Structures	Swiss-model (https://swissm
	odel.expasy.org/repository/u
	niprot/Q99259)
Domains	InterPro (https://www.ebi.ac.u

glutamic acid decarboxylase 2			
	Identifiers		
Symbol	GAD2		
Alt.	GAD65		
symbols			
NCBI	2572 (https://www.ncbi.nl		
gene	m.nih.gov/gene?cmd=retri		
	eve&dopt=default&list_uid		
	s=2572&rn=1)		
HGNC	11284 (https://www.genen		
	ames.org/data/gene-symb		
	ol-report/#!/hgnc_id/HGN		
	<u>C:11284)</u>		
ОМІМ	4093 (https://omim.org/40		
	93)		

RefSeq	NM_001047 (https://www.ncbi.nlm.nih.gov/protein/NM_001047) Q05329 (https://www.unip	
	rot.org/uniprot/Q05329)	
Other data		
EC number	4.1.1.15 (https://www.gen ome.jp/dbget-bin/www_bg et?enzyme+4.1.1.15)	
Locus	Chr. 10 p11.23 (https://om im.org/search/?index=gen eMap&search=10p11.23)	
Search for		
Structures	Swiss-model (https://swissm odel.expasy.org/repository/u niprot/Q05329)	
Domains	InterPro (https://www.ebi.ac.u k/interpro/protein/Q05329)	

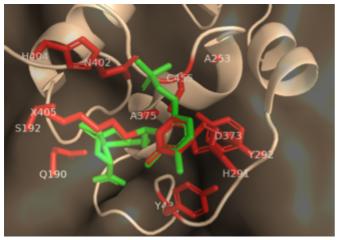
 $HOOC-CH_2-CH_2-CH(NH_2)-COOH \rightarrow CO_2 + HOOC-CH_2-CH_2-CH_2NH_2$

In mammals, GAD exists in two <u>isoforms</u> with molecular weights of 67 and 65 <u>kDa</u> (GAD₆₇ and GAD₆₅), which are encoded by two different <u>genes</u> on different <u>chromosomes</u> (GAD1 and <u>GAD2</u> genes, <u>chromosomes 2</u> and <u>10</u> in humans, respectively). [1][2] GAD₆₇ and GAD₆₅ are expressed in the brain where GABA is used as a <u>neurotransmitter</u>, and they are also expressed in the <u>insulin-producing β -cells</u> of the <u>pancreas</u>, in varying ratios depending upon the species. [3] Together, these two enzymes maintain the major physiological supply of GABA in mammals, [2] though it may also be synthesized from <u>putrescine</u> in the <u>enteric nervous system</u>, [4] brain, [5][6] and elsewhere by the actions of <u>diamine oxidase</u> and <u>aldehyde</u> dehydrogenase 1a1. [4][6]

Several truncated transcripts and <u>polypeptides</u> of GAD_{67} are detectable in the developing brain, $\overline{^{[7]}}$ however their function, if any, is unknown.

Structure and Mechanism

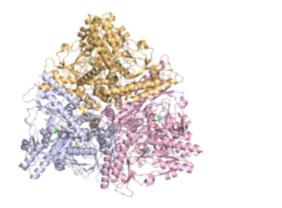
Both isoforms of GAD are homodimeric structures, consisting of 3 primary domains: the PLP, C-terminal and N-terminal domains. The PLP-binding domain of this enzyme adopts a type I PLP-dependent transferase-like fold. The reaction proceeds via the canonical mechanism, involving Schiff base linkage between PLP and Lys405. PLP is held in place through base-stacking with an adjacent histidine residue, and GABA is positioned such that its carboxyl group forms a salt bridge with arginine and a hydrogen bond with glutamine.

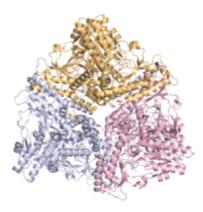


GAD67 active site containing PLP-glutamate complex (shown in green), with Schiff base linkage at Lys405. Side chain residues shown in red.

Dimerization is essential to maintaining function as the active site is found at this interface, and mutations interfering with optimal association between the 2 chains has been linked to pathology, such as schizophrenia. [9][10] Interference of dimerization by GAD inhibitors such as 2-keto-4-pentenoic acid (KPA) and ethyl ketopentenoate (EKP) were also shown to lead to dramatic reductions in GABA production and incidence of seizures. [11][8]

Catalytic activity is mediated by a short flexible loop at the dimer interface (residues 432–442 in GAD67, and 423–433 in GAD65). In GAD67 this loop remains tethered, covering the active site and providing a catalytic environment to sustain GABA production; its mobility in GAD65 promotes a side reaction that results in release of PLP, leading to autoinactivation. The conformation of this loop is intimately linked to the C-terminal domain, which also affects the rate of autoinactivation. Moreover, GABA-bound GAD65 is intrinsically more flexible and exists as an ensemble of states, thus providing more opportunities for autoantigenicity as seen in Type 1 diabetes. GAD derived from *Escherichia coli* shows additional structural intricacies, including a pH-dependent conformational change. This behavior is defined by the presence of a triple helical bundle formed by the N-termini of the hexameric protein in acidic environments. 16]





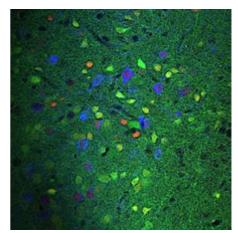
Hexameric E. coli GAD conformational transition: low-pH (left), neutral pH (right).

Regulation of GAD65 and GAD67

Despite an extensive sequence similarity between the two genes, GAD65 and GAD67 fulfill very different roles within the human body. Additionally, research suggests that GAD65 and GAD67 are regulated by distinctly different cellular mechanisms.

 ${\rm GAD}_{65}$ and ${\rm GAD}_{67}$ synthesize ${\rm GABA}$ at different locations in the cell, at different developmental times, and for functionally different purposes. ${}^{[17][18]}_{}$ ${\rm GAD}_{67}$ is spread evenly throughout the cell while ${\rm GAD}_{65}$ is localized to nerve terminals. ${}^{[17][19][20]}_{}$ ${\rm GAD}_{67}$ synthesizes ${\rm GABA}$ for neuron activity unrelated to neurotransmission, such as synaptogenesis and protection from neural injury. ${}^{[17][18]}_{}$ This function requires widespread, ubiquitous presence of ${\rm GABA}$. ${\rm GAD}_{65}$, however, synthesizes ${\rm GABA}$ for neurotransmission, ${}^{[17]}_{}$ and therefore is only necessary at nerve terminals and synapses. In order to aid in neurotransmission, ${\rm GAD}_{65}$ forms a complex with heat shock cognate 70 (HSC $_{70}$), cysteine string protein (CSP) and vesicular ${\rm GABA}$ transporter VGAT, which, as a complex, helps package ${\rm GABA}$ into vesicles for release during neurotransmission. ${}^{[21]}_{}$ ${\rm GAD}_{67}_{67}$ is transcribed during early development, while ${\rm GAD}_{65}_{65}$ is not transcribed until later in life. ${}^{[17]}_{}$ This developmental difference in ${\rm GAD}_{67}$ and ${\rm GAD}_{65}_{65}$ reflects the functional properties of each isoform; ${\rm GAD}_{67}_{67}$ is needed throughout development for normal cellular functioning, while ${\rm GAD}_{65}_{5}$ is not needed until slightly later in development when synaptic inhibition is more prevalent. ${}^{[17]}_{}$

 GAD_{67} and GAD_{65} are also regulated differently posttranslationally. Both GAD₆₅ and GAD₆₇ are regulated via phosphorylation of a dynamic catalytic loop, [22][12] but the regulation of these isoforms differs; GAD₆₅ is activated by phosphorylation while GAD₆₇ is inhibited by phosphorylation. GAD67 is predominantly found activated (~92%), whereas GAD65 is predominantly found inactivated (\sim 72%). [23] GAD₆₇ is phosphorylated at threonine 91 by protein kinase A (PKA), while GAD₆₅ is phosphorylated, and therefore regulated by, protein kinase C (PKC). Both GAD₆₇ and GAD₆₅ are also regulated posttranslationally by pyridoxal 5'-phosphate (PLP); GAD is activated when bound to PLP and inactive when not bound to PLP.[23] Majority of GAD_{67} is bound to PLP at any given time, whereas binds PLP when GABA neurotransmission. [23] This reflects the functional properties of the two isoforms; GAD₆₇ must be active at all times for normal cellular



Gad65 in red, Gad67 in green, and tyrosine hydroxylase (blue) in the ventral tegmental area of the mouse brain

functioning, and is therefore constantly activated by PLP, while GAD_{65} must only be activated when GABA neurotransmission occurs, and is therefore regulated according to the synaptic environment.

Studies with mice also show functional differences between Gad67 and Gad65. GAD67-/- mice are born with cleft palate and die within a day after birth while GAD65-/- mice survive with a slightly increased tendency in seizures. Additionally, GAD65+/- have symptoms defined similarly to attention deficith/peractivity disorder (ADHD) in humans. attention deficith/peractivity disorder (ADHD) in humans. attention deficith/peractivity disorder (ADHD) in humans.

Role in the nervous system

Both GAD67 and GAD65 are present in all types of synapses within the human nervous system. This includes <u>dendrodendritic</u>, axosomatic, and axodendritic synapses. Preliminary evidence suggests that GAD65 is dominant in the visual and neuroendocrine systems, which undergo more phasic changes. It is also believed that GAD67 is present at higher amounts in tonically active neurons. [25]

Role in pathology

Autism

Both GAD65 and GAD67 experience significant downregulation in cases of <u>autism</u>. In a comparison of autistic versus control brains, GAD65 and GAD67 experienced a downregulation average of 50% in parietal and cerebellar cortices of autistic brains. Cerebellar <u>Purkinje cells</u> also reported a 40% downregulation, suggesting that affected cerebellar nuclei may disrupt output to higher order motor and cognitive areas of the brain. [18]

Diabetes

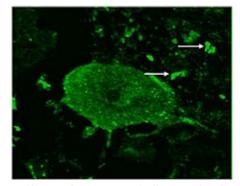
Both GAD_{67} and GAD_{65} are targets of <u>autoantibodies</u> in people who later develop type 1 <u>diabetes mellitus</u> or <u>latent autoimmune diabetes</u>. [27][28] Injections with GAD_{65} in ways that induce immune tolerance have been shown to prevent type 1 diabetes in rodent models. [29][30][31] In clinical trials, injections with GAD_{65} have been shown to preserve some insulin production for 30 months in humans with type 1 diabetes. [32][33] A <u>Cochrane systematic review</u> also examined 1 study showing improvement of C-peptide levels in cases of Latent Autoimmune Diabetes in adults, 5 years following treatment with GAD_{65} . Still, it is important to highlight that the studies available to be included in this review presented considerable flaws in quality and design. [34]

Stiff person syndrome

High <u>titers</u> of autoantibodies to glutamic acid decarboxylase (GAD) are well documented in association with <u>stiff person syndrome</u> (SPS). Glutamic acid decarboxylase is the rate-limiting enzyme in the synthesis of γ -aminobutyric acid (GABA), and impaired function of GABAergic neurons has been implicated in the pathogenesis of SPS. Autoantibodies to GAD might be the causative agent or a disease marker. GAD

Schizophrenia and bipolar disorder

Substantial dysregulation of GAD \underline{mRNA} expression, coupled with downregulation of \underline{reelin} , is observed in $\underline{schizophrenia}$ and $\underline{bipolar}$ $\underline{disorder}.^{\underline{[37][38]}}$ The most pronounced downregulation of GAD_{67} was found in hippocampal $\underline{stratum}$ oriens layer in both disorders and in other layers and structures of hippocampus with varying degrees. $\underline{[39]}$



Stiff man human cerebellum stained with a reference anti-GAD65 monoclonal antibody. Thin arrows show presynaptic terminals staining only with the anti-GAD65 monoclonal antibody

 ${
m GAD_{67}}$ is a key enzyme involved in the synthesis of inhibitory neurotransmitter ${
m \underline{GABA}}$ and people with schizophrenia have been shown to express lower amounts of ${
m GAD_{67}}$ in the <u>dorsolateral prefrontal cortex</u> compared to healthy controls. The mechanism underlying the decreased levels of ${
m GAD_{67}}$ in people with schizophrenia remains unclear. Some have proposed that an immediate early gene, Zif268, which normally binds to the <u>promoter</u> region of ${
m GAD_{67}}$ and increases transcription of ${
m GAD_{67}}$, is lower in schizophrenic patients, thus contributing to decreased levels of ${
m GAD_{67}}$. Since the dorsolateral prefrontal cortex (DLPFC) is involved in working memory, and ${
m GAD_{67}}$ and Zif268 mRNA levels are lower in the DLPFC of schizophrenic patients, this molecular alteration may account, at least in part, for the working memory impairments associated with the disease.

Parkinson disease

The bilateral delivery of glutamic acid decarboxylase (GAD) by an <u>adeno-associated viral vector</u> into the subthalamic nucleus of patients between 30 and 75 years of age with advanced, progressive, levodoparesponsive <u>Parkinson disease</u> resulted in significant improvement over baseline during the course of a sixmonth study. [42]

Cerebellar disorders

Intracerebellar administration of GAD autoantibodies to animals increases the excitability of motoneurons and impairs the production of <u>nitric oxide</u> (NO), a molecule involved in learning. Epitope recognition contributes to cerebellar involvement. Reduced GABA levels increase glutamate levels as a consequence of lower inhibition of subtypes of GABA receptors. Higher glutamate levels activate microglia and activation of xc(-) increases the extracellular glutamate release.

Neuropathic pain

<u>Peripheral nerve injury</u> of the sciatic nerve (a <u>neuropathic pain</u> model) induces a transient loss of GAD₆₅ immunoreactive terminals in the <u>spinal cord</u> <u>dorsal horn</u> and suggests a potential involvement for these alterations in the development and amelioration of pain behaviour. [45]

Other Anti-GAD-associated neurologic disorders

Antibodies directed against glutamic acid decarboxylase (GAD) are increasingly found in patients with other symptoms indicative of central nervous system (CNS) dysfunction, such as <u>ataxia</u>, <u>progressive encephalomyelitis</u> with <u>rigidity and myoclonus</u> (PERM), <u>limbic encephalitis</u>, and <u>epilepsy</u>. The pattern of anti-GAD antibodies in epilepsy differs from type 1 diabetes and stiff-person syndrome.

Role of glutamate decarboxylase in other organisms

Besides the synthesis of GABA, GAD has additional functions and structural variations that are organism-dependent. In <u>Saccharomyces cerevisiae</u>, GAD binds the Ca²⁺ regulatory protein <u>calmodulin</u> (CaM) and is also involved in responding to oxidative stress. [48] Similarly, GAD in plants binds calmodulin as well. [49] This interaction occurs at the 30-50bp CAM-binding domain (CaMBD) in its C terminus and is necessary for proper regulation of GABA production. [50] Unlike vertebrates and invertebrates, the GABA produced by GAD is used in plants to signal abiotic stress by controlling levels of intracellular Ca²⁺ via CaM.

Binding to CaM opens Ca^{2+} channels and leads to an increase in Ca^{2+} concentrations in the cytosol, allowing Ca^{2+} to act as a secondary messenger and activate downstream pathways. When GAD is not bound to CaM, the CaMBD acts as an autoinhibitory domain, thus deactivating GAD in the absence of stress. [50] Interesting, in two plant species, rice and apples, Ca^{2+} /CAM-independent GAD isoforms have been discovered. [51][52] The C-terminus of these isoforms contain substitutions at key residues necessary to interact with CaM in the CaMBD, preventing the protein from binding to GAD. Whereas CaMBD of the isoform in rice still functions as an autoinhibitory domain, [51] the C-terminus in the isoform in apples does not. [52] Finally, the structure of plant GAD is a hexamer and has pH-dependent activity, with the optimal pH of 5.8 in multiple species. [50][53] but also significant activity at pH 7.3 in the presence of CaM [16]

It is also believed that the control of glutamate decarboxylase has the prospect of improving citrus produce quality post-harvest. In Citrus plants, research has shown that glutamate decarboxylase plays a key role in citrate metabolism. With the increase of glutamate decarboxylase via direct exposure, citrate levels have been seen to significantly increase within plants, and in conjunction post-harvest quality maintenance was significantly improved, and rot rates decreased. [54]

Just like GAD in plants, GAD in $E.\ coli$ has a hexamer structure and is more active under acidic pH; the pH optimum for $E.\ coli$ GAD is 3.8-4.6. However, unlike plants and yeast, GAD in $E.\ coli$ does not require calmodulin binding to function. There are also two isoforms of GAD, namely GadA and GadB, encoded by separate genes in $E.\ coli$, [55] although both isoforms are biochemically identical. The enzyme plays a major role in conferring acid resistance and allows bacteria to temporarily survive in highly acidic environments (pH < 2.5) like the stomach. This is done by GAD decarboxylating glutamate to GABA, which requires H+ to be uptaken as a reactant and raises the pH inside the bacteria. GABA can then be exported out of $E.\ coli$ cells and contribute to increasing the pH of the nearby extracellular environments.

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- Genetics, Expression Profiling Support GABA Deficits in Schizophrenia (https://web.archive.org/web/20070815162255/http://www.schizophreniaforum.org/new/detail.asp?id=1358) Schizophrenia Research Forum, 25 June 2007.
- Overview of all the structural information available in the PDB for UniProt: Q99259 (https://www.ebi.ac.uk/pdbe/pdbe-kb/proteins/Q99259) (Glutamate decarboxylase 1) at the PDBe-KB.
- Overview of all the structural information available in the PDB for UniProt: Q05329 (https://www.ebi.ac.uk/pdbe/pdbe-kb/proteins/Q05329) (Glutamate decarboxylase 2) at the PDBe-KB.

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