γ-Aminobutyrate (GABA) Regulated Defense Tolerance: Mechanisms and Opportunities

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Supplementary Table S1. Important *Arabidopsis thaliana* genes associated with GABA metabolism and signaling.

Gene name	Common gene name	Gene I.D.
ALDH10A8 (*AMADH2*)	Aminoaldehyde dehydrogenase/4-aminobutanal dehydrogenase	At1g74920
ALDH10A9 (*AMADH1)*	Aminoaldehyde dehydrogenase/4-aminobutanal dehydrogenase	At3g48170
ALMT1	Aluminum-activated malate transporter	At1G08430
ALMT4	Aluminum-activated malate transporter	At1g25480
ALMT6	Aluminum-activated malate transporter	At2G17470
ALMT9	Aluminum-activated malate transporter	At3G18440
ALMT12	Aluminum-activated malate transporter	At4g17970
CAT9	Cationic amino acid transporter	At1g05940
CuAOα1	Copper amine oxidase	At1g31670
CuAOα3	Copper amine oxidase	At1g31710
CuAOδ	Copper amine oxidase	At4g12290
CuAOζ (previously *CuAO3* or *CuAO1*)	Copper amine oxidase	At2g42490
CuAOα3 (previously *CuAO2)*	Copper amine oxidase	At1g31710
DIT1	Dicarboxylate transporter	At5g12860
DIT2.1	Dicarboxylate transporter	At5g64290
DIT2.2,	Dicarboxylate transporter	At5g64280
DTC	Dicarboxylate-tricarboxylate transporter	At5G19760
GABA-TOG	2-Oxogluarate-dependent GABA transaminase	uncertain
GABA-TP,	Pyruvate-dependent GABA transaminase	At3g22200
GABP/BAT1,	GABA permease	At2g01170
GAD1	Glutamate decarboxylase	At5g17330
GAD2	Glutamate decarboxylase	At1g65960
GAD4	Glutamate decarboxylase	At2g02010
GAT1	GABA transporter	At1g15040

GLYR1/SSR1	Glyoxylate/succinic semialdehyde reductase	At3g25530
GLYR2/SSR2	Glyoxylate/succinic semialdehyde reductase	At1g17650
GORK	Guard cell outward rectifying channel	At5g37600
OMT (mitochondrial)	2-oxoglutarate/malate transporter	uncertain
PAO2,	Polyamine oxidase	At2g43020
PAO3	Polyamine oxidase	At3g59050
PAO4	Polyamine oxidase	At1g65840
PROT1	Proline transporter	At2g39890
PROT2	Proline transporter	At3g55740
PROT3	Proline transporter	At2g36590
SSADH (ALDH5F1)	Succinic semialdehyde dehydrogenase	At1g79440
UCP1	Uncoupling protein	At3g5411
UCP2	Uncoupling protein	At5g58970