

**Table 2. *Arabidopsis* genotypes and references (see list below)**

Genotypes	Code	Comment on phenotype	Ecotype background	Primary reference or NASC No.
Col-0	Co0	Wild ecotype: Columbia	WT	N1093
Col-2	Co2	Variant of Columbia	WT	N907
C24	C24	Wild ecotype: C24	WT	N906
Ak1	Ak1	Wild ecotype: Achkarren	WT	N939
Ler-0	Le0	Wild ecotype: Landsberg erecta	WT	NW20
Nd-0	Nd0	Wild ecotype: Niederzenz	WT	N1391
Bch-1	Bch	Wild ecotype: Büchen/Lauenburg	WT	N957
Ws-0	Ws0	Wild ecotype: Vasljevići	WT	N1603
<b>Transgenics</b>				
35S-NahG	nah	Constitutive salicylate hydroxylase, defence suppressed	Ler	1
CPD2	C2	Brassinosteroid synthesis gene antisense	C24	2, 3
CPD9	C9	Brassinosteroid synthesis gene antisense	C24	2, 3
CPD31	C31	Brassinosteroid synthesis gene antisense	C24	2, 3
<b>Mutants</b>				
etr1-1	etr	Insensitive to ethylene	Col-1	4, 5
lsd-1	ls1	Lesion mimic, spontaneous cell death	Ws	6
lsd-5	ls5	Lesion mimic, spontaneous cell death	Ws	6
uvr2-1	uvr	Sensitive to UV light, mutant type II CPD photolyase	Ler	7, 8
vtc1-2	vtc	Ascorbic acid deficient, UV sensitive	Col-0	9, 10
fah1-2	fah	Defective in ferulate-5-hydroxylase, light sensitive	Col-2	11, 12
axr3-1	axr	Auxin resistant, transcription factor mutation with pleiotrophic phenotype	Col-1	13-15
eds1	eds1	Enhanced disease susceptibility	Ler	16
pgm1-1	pgm	Chloroplast phosphoglucomutase deficient, starch deficient	Col	17, 18
amt1-1	a11	NH <sub>4</sub> transporter defective	Col-0	19, 20
amt1-2	a12	NH <sub>4</sub> transporter defective	Col-0	20
amt1-4	a14	NH <sub>4</sub> transporter defective	Col-0	21
amt2	a2	NH <sub>4</sub> transporter defective	Col-0	22, 23

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