

Crop	Genome Editing Technique	Targeted Gene	Molecular Function	Type of Editing	Effect	Reference
Wheat	ZFN	AHAS	Role in branched amino acid formation	Insertion and replacement	Resistance against herbicide	[141]
	ZFN	IPK1	Phytate formation	Deletion	Removal of antinutritional phytate, mineral accumulation against abiotic stress (Fe, Zn)	[142]
	CRISPR/Cas9	IPK1	Phytic acid biosynthesis	Deletion and insertion	Reduced phytic acid and enhanced Fe and Zn in wheat grains	[143]
	CRISPR/Cas9	HRC	Encodes a putative histidine-rich calcium-binding protein	Deletion and insertion	Reduced fusarium head blight severity	[144]
	CRISPR/Cas9	GW2	Genetic determinant of grain weight	Deletion	Increase in thousand-grain weight and grain protein content	[145]
Rice	ZFN	SSIVa	Involved in the starch biosynthesis pathway	Deletion	Improve eating quality	[146]
	TALENs	11N3	Rice bacterial blight susceptibility gene	Deletion and insertion	Increase resistance to rice bacterial blight	[147]
	CRISPR/Cas9	SAIPK2	Regulate drought response	Deletion	Improved drought tolerance	[148]
	CRISPR/Cas9	OsPIN5b GS3 OsMYB30	Regulate panicle length, grain size, and cold tolerance	Mutation deletion	Increased panicle length, enlarged grain size, and increased cold tolerance	[149]
	CRISPR/Cas9	NRT1.1B	Control yield and early maturation	Base editing	Increased yield	[150]
Maize	ZFN	IPK1	Catalyses the final step in phytate biosynthesis in maize seeds	Deletion and insertion	Removal of antinutritional phytate herbicide tolerance	[151]
	TALENs	g2	Cuticular lipid functions	Deletion	Reduce epicuticular wax	[152]
	CRISPR/Cas9	RR22	Salinity tolerance	Deletion, insertion, substitution	Regulation of salt tolerance	[153]
	CRISPR/Cas9	NC1 QTL and, HKT1	Encodes an HKT-type transporter	Deletion	Reduce salt tolerance	[154]
	CRISPR/Cas9	AOC	Jasmonic acid biosynthesis pathway	Deletion, insertion, substitution	Efficient coordination with the environment	[155]
Barley	CRISPR/Cas9	HPT HGT	Biosynthesis of tocotrienols and tocopherol	Deletion and insertion	Decreased grain size and weight Shrunken phenotype Lower total starch content in grains	[156]
		ARE1	Involved in nitrogen use efficiency	Missense and/or frameshift mutations	Increase in plant height, tiller number, grain protein content, yield, and chlorophyll content	[157]
	CRISPR/Cas9	ENGase	Production of GN1 type FNGs (Free N Glycans	Indels and deletions	Increased abiotic tolerance	[158]

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Charles N. Bebee



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