WT	AGGGCCGTGCCGGGAGGCGCCACCTCGGCC-GAAGCGGCCCTACTC	Plant ID
D1(×9) D1,+1	AGGGCCGTGCCGGGAGGCGCCACCTCGGCC-AA-GCGGCCCTACTC AGGGCCGTGCCGGGAGGCGCCCACCTCGGCC-GGA-GCGGCCCTACTC	1
D1(×5)	AGGGCCGTGCCGGGAGGCGCCCACCTCGGCCAAGCGGCCCTACTC	2
D1(×8)	AGGGCCGTGCCGGGAGGCGCCCACCTCGGCCAAGCGGCCCTACTC	3
D1,+1 D1(×3) +1	AGGGCCGTGCCGGGAGGCGCCACCTCGGCC-GGAGCGGCCCTACTC AGGGCCGTGCCGGGAGGCGCCACCTCGGCC-AAGCGGCCCTACTC AGGGCCGTGCCGGGAGGCGCCACCTCGGCC-GAAA-GCGGCCCTACTC	4
D1(×8) +1(×7)	AGGGCCGTGCCGGGAGGCGCCACCTCGGCC-AA-GCGGCCCTACTC AGGGCCGTGCCGGGAGGCGCCCACCTCGGCC-GAAA-GCGGCCCTACTC	5
D1(×3)	AGGGCCGTGCCGGGAGGCGCCACCTCGGCC-GAGCGGCCCTACTC	6
D1(×20)	AGGGCCGTGCCGGGAGGCGCCACCTCGGCCAAGCGGCCCTACTC	7
D1(×5)	AGGGCCGTGCCGGGAGGCGCCACCTCGGCCAAGCGGCCCTACTC	8
D1(×7) +1(×3)	AGGGCCGTGCCGGGAGGCGCCACCTCGGCC-AA-GCGGCCCTACTC AGGGCCGTGCCGGAGGCGCCCACCTCGGCC-GAAA-GCGGCCCTACTC	9
D3,+7(×3)	AGGGCCGTGCCGGGAGGCGCCCACCTCGGCATGGACTGCGGCCCTACTC	10
D1(×12)	AGGGCCGTGCCGGGAGGCGCCCACCTCGGCCAAGCGGCCCTACTC	11
D1(×13) D1	AGGGCCGTGCCGGGAGGCGCCACCTCGGCC-AA-GCGGCCCTACTC AGGGCCGTGCCGGGAGGCGCCCCCCCCCGCC-GAGCGGCCCTACTC	12

Supplementary information, Figure S10 Targeted indel mutations induced by engineered sgRNA:Cas9 at the *YSA* gene sgRNA1 site in rice.

Alleles shown were amplified from genomic DNA isolated from 12 independent T0 transgenic plants separately and sequenced after cloned into vectors. The wild type sequence is shown at the top with the PAM sequence highlighted in magenta and the target sequence in cyan. Red dashes, deleted bases; red bases, insertions or mutations. The net change in length is to the right of each sequence (+, insertion; D, deletion). The number of clones representing each mutant allele is shown in brackets. The plants #5 and #7 correspond to the two plants showing albino leaf phenotype.