

## Published data sample

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### Gene

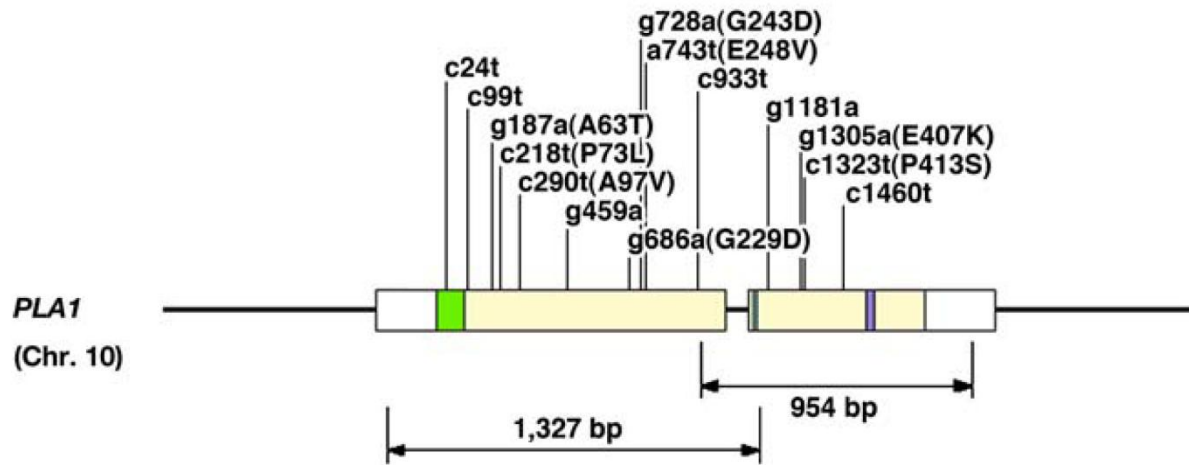
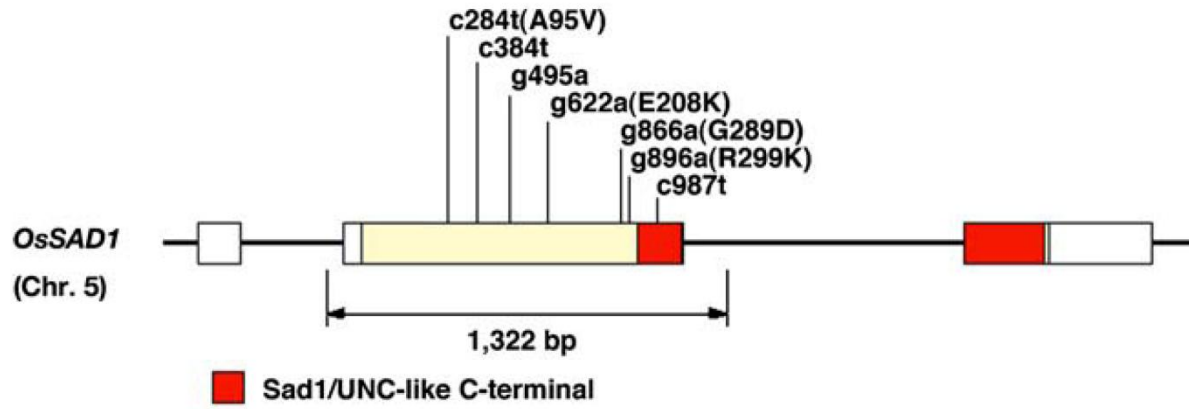
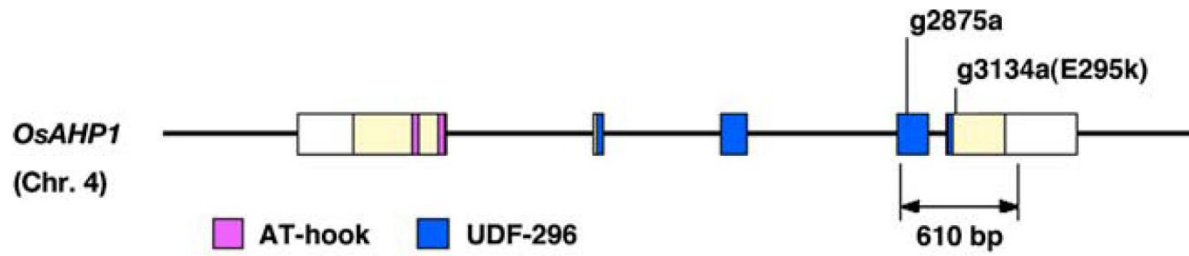
Os04g.....

Os05g.....

Os10g.....

### Reference:

Suzuki T., M. Eiguchi, T. Kumamaru, H. Satoh, H. Matsusaka, K. Moriguchi, Y. Nagato, N. Kurata. MNU-induced mutant pools and high performance TILLING enable finding of any gene mutation in rice. *Mol. Gen. Gen.*, 279: 213-223 (2008)



Hydrophobic region Oxygen binding motif  
Heme binding motif

UTR CDS

Gene	Line	Genotype of individuals			Base change <sup>a</sup>	Position from atg	Amino acid change	Region
		Wt	Hetero	Homo				
<i>OsAHP1</i>	1S31	0	3	3	G/A	3134	E295K	cds(exon5)
	7S102	5	1	0	T/A	3002		intron
	8S65	2	3	1	G/A	2875		cds(exon4)
<i>OsSAD1</i>	2S50	0	5	0	G/A	622	E208K	cds
	3S03	3	3	0	C/T	284	A95V	cds
	4S82	2	0	4	G/A	495		cds
	5S88	2	2	0	G/A	866	G289D	cds
	6S40	0	5	0	G/A	896	R299K	cds
	7S17	3	2	0	C/T	987		cds
	8S05	3	2	0	C/T	384		cds
	<i>PLA1</i>	1S32	1	3	1	C/T	1460	
2S03		1	2	1	G/A	459		cds(exon1)
2S91		4	2	0	G/A	187	A63T	cds(exon1)
3S81		2	3	0	C/T	933		cds(exon1)
4S17		1	3	2	C/T	24		cds(exon1)
4S35		2	2	1	C/T	290	A97V	cds(exon1)
4S111		2	2	2	A/T	743	E248V	cds(exon1)
5S43		2	3	1	G/A	1181		cds(exon2)
5S75		1	3	2	C/T	1323	P413S	cds(exon2)
5S84		2	2	1	G/A	728	G243D	cds(exon1)
5S91		1	3	0	G/A	1305	E407K	cds(exon2)
6S05		0	2	4	C/T	99		cds(exon1)
6S74		4	2	0	C/T	218	P73L	cds(exon1)
8S05		1	0	2	G/A	686	G229D	cds(exon1)