

Table S3: Candidate genes for *GN2* in the fine mapping region according to the annotation system of Nipponbare

Gene ID	Start	Stop	Annotation
LOC_Os02g56630	2214	4707	OsWAK receptor-like protein kinase
LOC_Os02g56640	7952	11394	retrotransposon protein, putative, unclassified
LOC_Os02g56650	12092	13870	retrotransposon protein, putative, unclassified
LOC_Os02g56660	14491	17909	retrotransposon protein, putative, unclassified
LOC_Os02g56670	22161	22482	hypothetical protein
LOC_Os02g56680	23842	25630	dehydrogenase, putative