

Table S3. Number of QTLs identified from different phenotypes, populations, and *M. oryzae* isolates

Phenotype	Population	<i>M. oryzae</i> isolates		Total ^a
		D41-2	12YL-DL-3-2	
LT	Full	1	3	6
	<i>indica</i>	3	1	
	<i>japonica</i>	-	1	
DLA	Full	13	-	30
	<i>indica</i>	1	12	
	<i>japonica</i>	-	4	
Total ^a		16	18	32

^a Total number of non-overlapped QTLs