

Additional information

Table S1. List of primers used in this study

| Primer Name | Sequence |
|----------------|-------------------------------|
| MoXYL1A-AF | 5' CGACCGAACCAAGGATTGC 3' |
| MoXYL1A-AR | 5' GAAGATTTGAACCGCACC 3' |
| MoXYL1A-BF | 5' CGCGAGTTGATGTTGA 3' |
| MoXYL1A-BR | 5' CAAGGGCATAGAACGGAAT 3' |
| MoXYL1A-OF | 5' GCAGCTCATGCGCGACAGT 3' |
| MoXYL1A-OR | 5' CCCTCGGTGGCGACAATC 3' |
| MoXYL1A-UF | 5' CACTTGCAGTTAGCCTGAG 3' |
| MoXYL1A-UR | 5' CACAGTTGCCAGTGATACA 3' |
| MoXYL1A-COMP-F | 5' AACATCAGGCAGCTCGTA 3' |
| MoXYL1A-COMP-R | 5' CGCCGGAGTCTGAACGTTGA 3' |
| MoXYL1A-qPCR-F | 5' ATGGCAACATGACGGGATTTC 3' |
| MoXYL1A-qPCR-R | 5' ACTGCCACTGAACGCTGTAG 3' |
| MoXYL1A-pGDG-F | 5' ATGGTCTCCTCACCTCCATC 3' |
| MoXYL1A-pGDG-R | 5' TCACGCCGGAGTCTGAA 3' |
| MoXYL1A-Y2H-F | 5' ATGGTCTCCTCACCTCCATC 3' |
| MoXYL1A-Y2H-R | 5' TCACGCCGGAGTCTGAA 3' |
| MoXYL1B-AF | 5' ACGAGGTGATTACTTTGGG 3' |
| MoXYL1B-AR | 5' TCTGGCTCTATGATGTATGTCTT 3' |
| MoXYL1B-BF | 5' GCGAGATCGACTGTAACAAACC 3' |
| MoXYL1B-BR | 5' ACGGAGGAAAGTGACCTACCATT 3' |
| MoXYL1B-OF | 5' GGGGAGTGCTCGGTATGTT 3' |
| MoXYL1B-OR | 5' GGCGACGATCTGGTAGTCGT 3' |
| MoXYL1B-UF | 5' TTCAGACAAGACGCAACCC 3' |
| MoXYL1B-UR | 5' GCTCCATACAAGCCAACCAC 3' |
| MoXYL1B-COMP-F | 5' AACATCAGGCAGCTCGTA 3' |
| MoXYL1B-COMP-R | 5' CGCCGGAGTCTGAACGTTGA 3' |

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| MoXYL1B-qPCR-F | 5' ATGGCAACATGACGGGATT C 3' |
| MoXYL1B-qPCR-R | 5' ACTGCCACTGAACGCTGTAG 3' |
| MoXYL1B-pGDG-F | 5' ATGGTCTCCTTCACCAC 3' |
| MoXYL1B-pGDG-R | 5' TCACCGCTGGATGGTGA 3' |
| MoXYL1B-Y2H-F | 5' ATGGTCTCCTTCACCAC 3' |
| MoXYL1B-Y2H-R | 5' TCACCGCTGGATGGTGA 3' |
| Os-CH-pGDR-F | 5' ATGTCGTCGGTGCAGCT 3' |
| Os-CH-pGDR-R | 5' TCAAGATAGGACAGCCATCA 3' |
| G418-F2 | 5' CAACAACACGCATCATCCC A 3' |
| G418-R2 | 5' TCAGAAGAACTCGTCAAGAA 3' |
| rice-actin-F | 5' GCGTGGACAAAGTTTCAACCG 3' |
| rice-actin-R | 5' TCTGGTACCCCTCATCAGGCATC 3' |
| OsPR1a-RT-F | 5' GGAAGTACGGCGAGAACATC 3' |
| OsPR1a-RT-R | 5' GGCGAGTAGTTGCAGGTGAT 3' |
| OsCht1-RT-F | 5' CGTGGTGACCAACATCATCA 3' |
| OsCht1-RT-R | 5' GAGTTGAAAGGCCTCTGGTTGT 3' |
| OsPBZ1-F | 5' CTGTGGAAGGTCTGCTTGG A 3' |
| OsPBZ1-R | 5' TCTTGTATACGCTCCCTGCG 3' |
| OsPAL-F | 5' GAGATCAACTCCGTCAAC 3' |
| OsPAL-R | 5' TGTAGAAGTCGTTCACCA 3' |
| OsPR1a-F | 5' GTCGGAGAACGCAGTGGTA 3' |
| OsPR1a-R | 5' CGAGTAGTTGCAGGTGATG 3' |
| OsPR5-F | 5' CAGCCAGGACTTCTACGA 3' |
| OsPR5-R | 5' TGTGTCTTGGTGTGTTCTTC 3' |
| OsKSL7-F | 5' CCATATCCACAGCCAACA 3' |
| OsKSL7-R | 5' ACTTAGCCTCTCCTGATGT 3' |
| OsKSL10-F | 5' TAACCCTTGCCTCTGGGATG 3' |
| OsKSL10-R | 5' ATTGTTCCACCGGAATCCCT 3' |
| OsKOL4-F | 5' GTGCACAGCTGACAGATGAC 3' |

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|-------------|-----------------------------|
| OsKOL4-R | 5' TCGGATCTCTGGTAGAGTAGC 3' |
| OsCYP76M8-F | 5' ATGGAGAATAGCCAGATGTG 3' |
| OsCYP76M8-R | 5' AAGTGTAGGTTGCCGATG 3' |
| OsNOMT-F | 5' GAAGGTGTGGAGCATATCGG 3' |
| OsNOMT-R | 5' GTGCAGAACCCACTTGAGC 3' |

Table S2. Predicted location of pathogenicity-related protein in rice.

| Pathogenicity-related genes in rice | Abbreviated annotation | Gene accession number (Gene ID) | Predicted localization |
|---|------------------------|----------------------------------|---|
| Probenazole-inducible protein | OsPBZ1 | Os12g0555200 | Extracellular, Mitochondrial, ER, Peroxisomal, Chloroplast |
| Pathogenesis-related protein 1a | OsPR1a | Os07g0129200-01 | Extracellular, and Mitochondrial |
| Thaumatin-like pathogenesis-related protein 3 precursor | OsPR5 | Os12g0628600 | Vacuolar, and Extracellular |
| Ent kaurene synthase 7 | OsKSL7 | Os02g0570400 | Extracellular, and Vacuolar |
| kaurene synthase like 10 | OsKSL10 | Os12g0491800 | Cytoplasmic, Mitochondrial, and Chloroplast |
| Ent-kaurene oxidase 4 | OsKOL4 | Os06g0569500 | Mitochondrial, Peroxisomal, and Chloroplast |
| Cytochrome P450 family protein | OsCYP76M8 | Os02g0569400 | Plasma membrane, Mitochondrial, and ER |
| naringenin 7-O-methyltransferase | OsNOMT | Os12g0240900 | Extracellular, Mitochondrial, ER, Peroxisomal, and Chloroplast |
| Chitinase 1 | OsCHT1 | Os06g0726200 | Extracellular, Vacuolar |

| | | | |
|--------------------------------|-------|--------------|--------------------------|
| phenylalanine ammonia-lyase | OsPAL | Os05g0427400 | Cytoplasmic, Chloroplast |
|--------------------------------|-------|--------------|--------------------------|

Additional figures

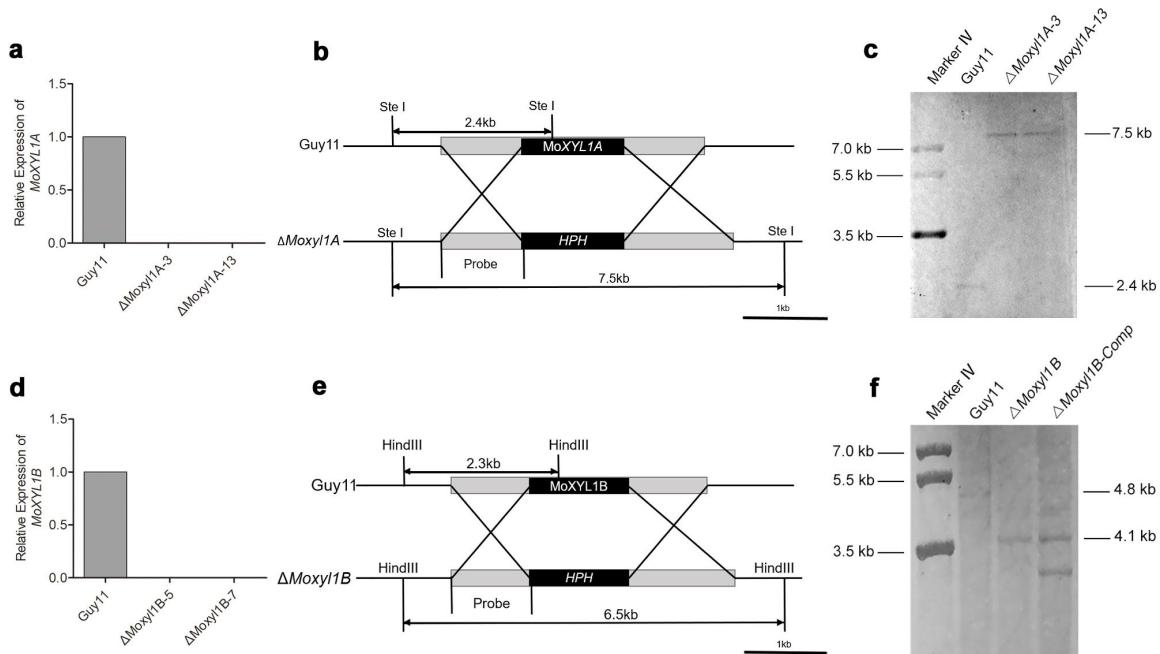


Figure S1. Single copy insertions confirmed by Southern Blot and relative expression of *MoXYL1A* gene in Guy11 and the two putative mutant strains was assessed by qRT-PCR. (a) Schematic of probe design for Southern blot for *MoXYL1A* gene. **(b)** Successful gene replacement of *MoXYL1A* by single insertion of *hph* at the *MoXYL1A* locus by Southern blot analysis. **(c)** The relative expression of *XYL1B* gene in Guy11 and the putative mutant strains was assessed by qRT-PCR. **(d)** Schematic of probe design for Southern blot for *MoXYL1B* gene **(e)** Successful gene replacement of *MoXYL1B* by single insertion of *hph* at the *MoXYL1B* locus by Southern blot analysis.

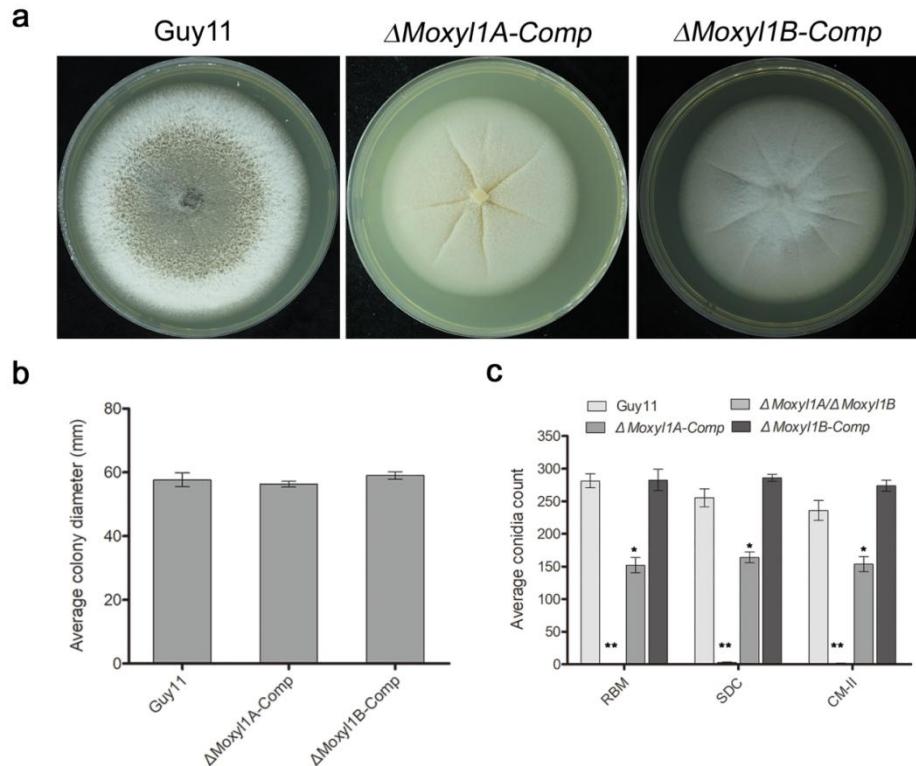


Figure S2. Complementation of MoXYL1A and MoXYL1B rescued the defects exhibited by the mutant strains

(a) Photographs showing colony growth diameter (mm) of complemented deletion strains for *MoXYL1A&B* compared to WT. (b-c) Quantification of colony growth in (a) Quantification of conidiation in the complemented strains and the double mutant.

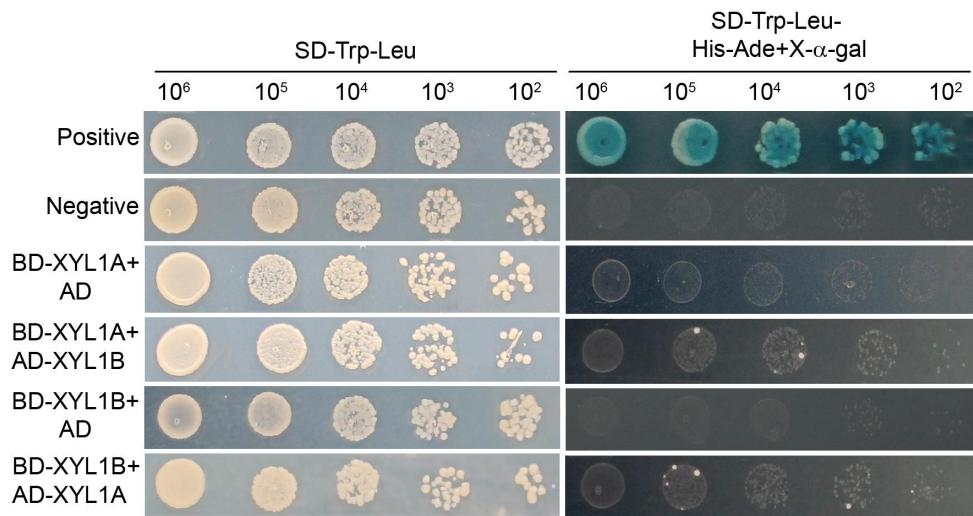


Figure S3. Interaction in Y2H and in-planta expression of putative *M. oryzae* xylanases 1A and B

(a) Yeast-2-hybrid experiments to assess interaction between the xylanases 1A and B in *M. oryzae* did not show physical interaction between the two proteins.