## MoErv29 promotes apoplastic effector secretion contributing to virulence of the rice blast fungus *Magnaporthe oryzae*

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Table S1. Primers used in this study

| Primer name | Sequence (5'-3') | Remark |
|---|---|---|
| ERV29 F1 | TAAATCGATCGAGTGGGAATCGTCTATCCAAG | amplify *MoERV29* 5' flank sequence |
| ERV29 F2 | TAAGAATTCGTTGCTAGCTCTACCCGGAG | amplify *MoERV29* 5' flank sequence |
| ERV29 F3 | TAATCTAGAGAAAAAACGACGCATCACACAATTG | amplify *MoERV29* 3' flank sequence |
| ERV29 F4 | TAACCGCGGGAGAGACAAGACTTCCTTCTGC | amplify *MoERV29* 3' flank sequence |
| ERV29 KO-F | GGAGACTAGTTTACGTGGCCGCTAAC | amplify *MoERV29* probe sequence |
| ERV29 KO-R | GAAGTACATCTTGCGGTCCTTCTCCTC | amplify *MoERV29* probe sequence |
| ERV29 BY | CTACATATGCAGTGTCTGATGTG | validation of *MoERV29* deletion (HPH) |
| HPH R | GCTGATCTGACCAGTTGCCTA | |
| ERV29 HB-F1 | ACTCACTATAGGGCGAATTGGGTACTCAAATTGGTTCAACCTTCACACCCAACCTTAG | *MoERV29* complementation |
| ERV29 HB-R1 | CACCACCCCGGTGAACAGCTCCTCGCCCTTGCTCACGTAAACCTTCTTCTCGTCG | *MoERV29* complementation |
| AD-ERV29 F1 | TACCAGATTACGCTCATATGATGTCGCAACGAGGACCTTCGGGC | Construction of AD-ERV29 |
| AD-ERV29 R1 | TGCCCACCCGGGTGGAATTCTTAGTAAACCTTCTTCTTCTCGTCG | Construction of AD-ERV29 |
| BD-ERV29 F1 | TGATCTCAGAGGAGGACCTGCATATGATGTCGCAACGAGGACCTTCGGGC | Construction of BD-ERV29 |
| BD-ERV29 R1 | GCAGGTCGACGGATCCCCGGGAATTCTTAGTAAACCTTCTTCTTCTCGTCG | Construction of BD-ERV29 |
| MoHac1 32a F1: | CAAGGCCATGGCTGATATCATGAACACTTGGAGCTCCACCCCAG | Construction of MoHac1 32a |
| MoHac1 32a F2: | GTGCGGCCGCAAGCTTGTCGACCTATAGACTACCTTGCTCGTTGC | Construction of MoHac1 32a |
| Slp1-Stag F1 | TTTCGTAGGAACCCAATCTTCAAAATGCAGTTCGCTACCATCACC | Construction of Slp1-S |
| Slp1-Stag R1 | GCTCCTCGCCCTTGCTCACTTAGCTATCCACATGCTGACGTTCAAATTTCGCCGCCGCGGTTTCTTTGTTCTTGCAGATGGGGATG | Construction of Slp1-S |
| Slp1-GFP F1 | ACTCACTATAGGGCGAATTGGGTACTCAAATTGGTTCGCACAATTGAACAGTGCGATG | Construction of Slp1-GFP |
| Slp1-GFP R1 | CACCACCCCGGTGAACAGCTCCTCGCCCTTGCTCACGTTCTTGCAGATGGGGATG | Construction of Slp1-GFP |
| Bas4-GFP F1 | TTTCGTAGGAACCCAATCTTCAAAATGCAGCTCTCATTCTCAGCAATC | Construction of Bas4-GFP |
| Bas4-GFP R1 | CACCACCCCGGTGAACAGCTCCTCGCCCTTGCTCACAGCAGGGGGGATAGACGAGC | Construction of Bas4-GFP |
| 28S rDNA LL | TACGAGAGGAACCGCTCATTCAGATAATTA | qRT-PCR |

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|---|---|---|
| 28S rDNA RR | TCAGCAGATCGTAACGATAAAGCTACTC | qRT-PCR |
| Rubq1 LL | GTGGTGGCCAGTAAGTCCTC | qRT-PCR |
| Rubq1 RR | GGACACAATGATTAGGGATCA | qRT-PCR |
| MGG_00227 F1 | GTATCGATAAGCTTGATATCGCCAAACCCATACCGCGATTG | amplify MGG_00227 5' flank sequence |
| MGG_00227 R1 | GGTGGAGGATGGGCAAGGTAG | amplify MGG_00227 5' flank sequence |
| Hph F: | CTACCTTGCCCATCCTCCACCGGAGGTCAACACATCAATG | amplify Hph sequence |
| Hph R: | CTCTATTCCTTTGCCCTCG | amplify Hph sequence |

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|---|---|---|
| MGG_00227 F2: | CGAGGGCAAAGGAATAGAGCGTGTCTTGATAGGTAGGG | amplify MGG_00227 3' flank sequence |
| MGG_00227 R2: | GGTGGCGGCCGCTCTAGAACTAGTGAGATCAACTGTGAACT | amplify MGG_00227 3' flank sequence |
| MGG_00227Con F: | GCGCAATCCGGCTACGGCACCTGC | amplify MGG_00227 probe sequence |
| MGG_00227Con R: | CGTTGTCACGGTAACGGTAG | amplify MGG_00227 probe sequence |
| MGG_00227BY: | CTGGTCCAACTACATCATTC | validation of MGG_00227 deletion (HPH) |
| MGG_00227 GFPF1 | TTTCGTAGGAACCCAATCTTCAAAATGCGCAACACGCTCGTTAAC | Construction of MGG_00227 GFPF1 |
| MGG_00227 GFPR1: | CACCACCCCGGTGAACAGCTCCTCGCCCTTGCTCACATCAAAATCAACGATGGTTG | Construction of MGG_00227 GFPR1 |
| MGG_00259 F1: | GTATCGATAAGCTTGATATCCACGGATATGTTGAAACCTTG | amplify MGG_00259 5' flank sequence |
| MGG_00259 R1: | GTTTACTTGGGGAGTTGGATG | amplify MGG_00259 5' flank sequence |
| Hph F 00259: | CATCCAACTCCCCAAGTAAACGGAGGTCAACACATCAATG | amplify Hph sequence |
| Hph R00259: | CTCTATTCCTTTGCCCTCG | amplify Hph sequence |
| MGG_00259 F2: | CGAGGGCAAAGGAATAGAGGGTTGGTTGGGAAGAGATG | amplify MGG_00259 3' flank sequence |
| MGG_00259 R2: | GGTGGCGGCCGCTCTAGAACTAGTCTGGCCAAGGAGGGCAAGCT | amplify MGG_00259 3' flank sequence |
| MGG_00259 Con F: | GGCGCCGTGGCCGACCTGTC | amplify MGG_00259 probe sequence |
| MGG_00259 Con R: | GCACGGTCGAGTTGTCTGCC | amplify MGG_00259 probe sequence |
| MGG_00259 BY: | GGCACTACTTTGCCATACCAC | validation of MGG_00229 deletion (HPH) |
| MGG_00259 GFPF1: | TTTCGTAGGAACCCAATCTTCAAAATGAAGGTCTCACTAAGCATC | Construction of MGG_00259 GFPF1 |
| MGG_00259 GFPR1: | CACCACCCCGGTGAACAGCTCCTCGCCCTTGCTCACCGGCAGAGCCAACGGGCTCAG | Construction of MGG_00259 GFPR1 |
| MGG_01655 F1: | GTATCGATAAGCTTGATATCCAGCCCCATCAGGCCACTAG | amplify MGG_01655 5' flank sequence |
| MGG_01655 R1: | CTTGGCGAGTATATACAGCC | amplify MGG_01655 5' flank sequence |
| Hph 01655 F: | GGCTGTATATACTCGCCAAGGGAGGTCAACACATCAATG | amplify Hph sequence |
| Hph 01655 R: | CTCTATTCCTTTGCCCTCG | amplify Hph sequence |
| MGG_01655 F2: | CGAGGGCAAAGGAATAGAGATTGACGGCACGCATTTTCGG | amplify MGG_01655 3' flank sequence |
| MGG_01655 R2: | GGTGGCGGCCGCTCTAGAACTAGTGTAGAGCAGCGATGATTGCC | amplify MGG_01655 3' flank sequence |
| MGG_01655 Con F: | CGGCAATCATTCCAAGATTGC | amplify MGG_01655 probe sequence |
| MGG_01655 Con R: | GTTTGGCTGCGGACCCACTG | amplify MGG_01655 probe sequence |
| MGG_01655 BY: | CCGCGGATTACGGTCCCGCC | validation of MGG_01655 deletion (HPH) |
| MGG_01655 GFPF1: | TTTCGTAGGAACCCAATCTTCAAAATGACCTTGTTATCCCT | Construction of MGG_01655 GFPF1 |

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| MGG_01655 GFPR1: | CACCACCCCGGTGAACAGCTCCTCGCCCTTGCTCACGACA GGGCCGACACCGGGC | Construction of MGG_01655 GFPR1 |
| MGG_05970 F1: | GTATCGATAAGCTTGATATCCTGCCGATCGTGTACCCTTAC | amplify MGG_05970 5' flank sequence |
| MGG_05970 R1: | GTGCAAGATAACGAGTGATG | amplify MGG_05970 5' flank sequence |
| Hph 05970 F: | CATCACTCGTTATCTTGCACGGAGGTCAACACATCAATG | amplify Hph sequence |
| Hph 05790 R: | CTCTATTCCTTTGCCCTCG | amplify Hph sequence |
| MGG_05970 F2: | CGAGGGCAAAGGAATAGAGGCTGTCTGGTCTGGGTGAGC | amplify MGG_05970 3' flank sequence |
| MGG_05970 R2: | GGTGGCGGCCGCTCTAGAACTAGTCATGGCGCTTAACCACT GCG | amplify MGG_05970 3' flank sequence |
| MGG_05970 Con F: | CAATTCTAATCATGACTCACAG | amplify MGG_05970 probe sequence |
| MGG_05970 Con R: | CAGACCTTGGTCCAGAGTGG | amplify MGG_05970 probe sequence |
| MGG_05970 BY: | CAAACGACCGTAACAATCGC | validation of MGG_05970 deletion (HPH) |
| MGG_05970 GFPF1： | TTTCGTAGGAACCCAATCTTCAAAATGGCTCGTCACGCACT CGC | Construction of MGG_05970 GFPF1 |
| MGG_05970 GFPR1: | CACCACCCCGGTGAACAGCTCCTCGCCCTTGCTCACAATGC CCGAGTCCTCCTGCG | Construction of MGG_05970 GFPR1 |
| MGG_08127 F1: | GTATCGATAAGCTTGATATCCACAGAACCCCGGGTAAGTC | amplify MGG_08127 5' flank sequence |
| MGG_08127 R1: | AATGGCGGGGTCTTTGAGAG | amplify MGG_08127 5' flank sequence |
| Hph 08127 F: | CTCTCAAAGACCCCGCCATTGGAGGTCAACACATCAATG | amplify Hph sequence |
| Hph 08127 R: | CTCTATTCCTTTGCCCTCG | amplify Hph sequence |
| MGG_08127 F2: | CGAGGGCAAAGGAATAGAGACTGGTCGTGCTTGAAAGGTG | amplify MGG_08127 3' flank sequence |
| MGG_08127 R2: | GGTGGCGGCCGCTCTAGAACTAGTGATGACCATTTTCCTAG TATC | amplify MGG_08127 3' flank sequence |
| MGG_08127 Con F: | GAGGTCGTTATAACCGCGTC | amplify MGG_08127 probe sequence |
| MGG_08127 Con R: | CGTCAATGGCGGATCCGTTGAC | amplify MGG_08127 probe sequence |
| MGG_08127 BY: | GGTATCCCTGCCTTTTATGC | validation of MGG_08127 deletion (HPH) |
| MGG_08127GFPF1： | TTTCGTAGGAACCCAATCTTCAAAATGAATCTTCGGGACAC CATC | Construction of MGG_08127 GFP |
| MGG_08127GFPR1: | CACCACCCCGGTGAACAGCTCCTCGCCCTTGCTCACTCTCC TCAAACCAGAGTCCAG | Construction of MGG_08127 GFP |
| MGG_09139 GFPF1: | TTTCGTAGGAACCCAATCTTCAAAATGGTTCAATCTGTGATA TTTG | Construction of MGG_09139 GFP |
| MGG_09139 GFPF1: | CCCGGTGAACAGCTCCTCGCCCTTGCTCACGACACCAGAAT CCGCCAGCAC | Construction of MGG_09139 GFP |
| MGG_13464 GFPF1: | TTTCGTAGGAACCCAATCTTCAAAATGAAGGTCACCAGCTT CTTC | Construction of MGG_13464 GFP |
| MGG_13464 GFPR1: | CCCGGTGAACAGCTCCTCGCCCTTGCTCACGACTGCGCGA GCAGTGCTGTTAC | Construction of MGG_13464 GFP |