

# Supplementary material

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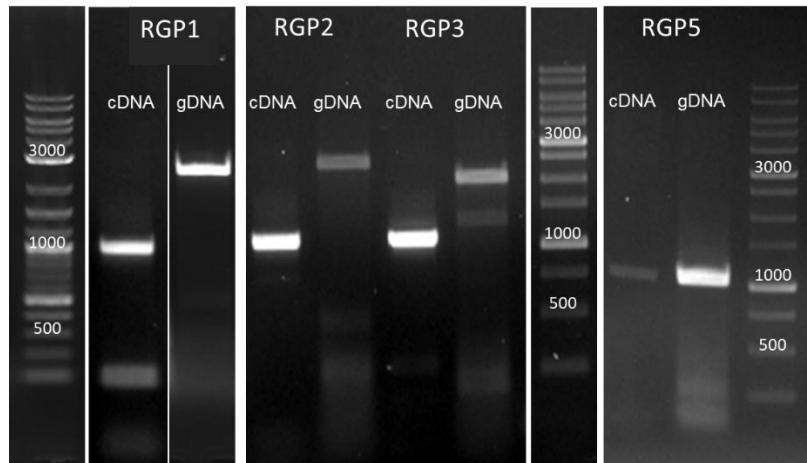
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NbRGP3 274 NPFVNLKKEVKGIVYVQEEIIPFFGOSVALPKDCTSVVKCYLELSKVVAALKGVDDYFNVLAADAMVTVIEAEDELNPATATA-VKIPNGSSK--- 364
NbRGP4 268 NPFVNLKKEINGIFWEEIIPFFGUNLNLSKEETTVVKCIVEMARQVNEKLGKIDDPYFVVLAEAMVTVIEAEDELNPAKDTA-KIPNGAA--- 357
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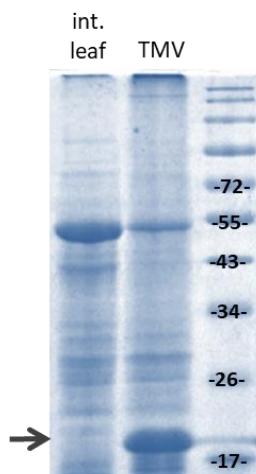
## Amino Acid Sequence Identities of NbRGP Proteins

	NbRGP1	NbRGP2	NbRGP3	NbRGP5
NbRGP1	-	94.72%	87.43%	48.80%
NbRGP2	94.72%	-	92.46%	48.87%
NbRGP3	87.43%	92.46%	-	50.30%
NbRGP5	48.80%	48.87%	50.30%	-

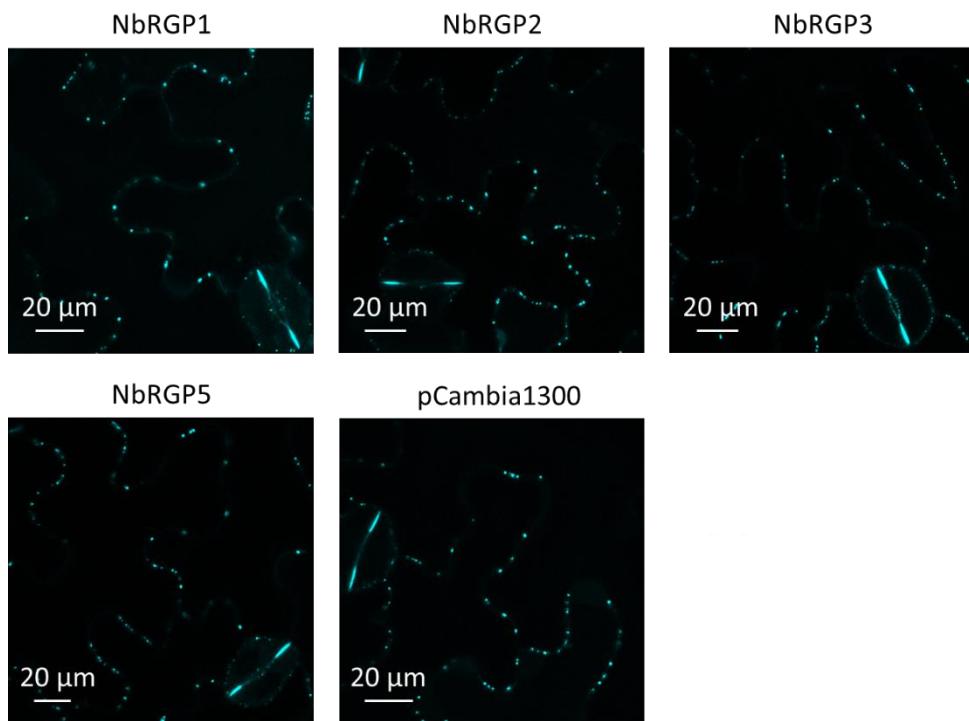
**Figure S1.** Multiple alignment of NbRGP amino acid sequences (top) and the percent of similarity (bottom).



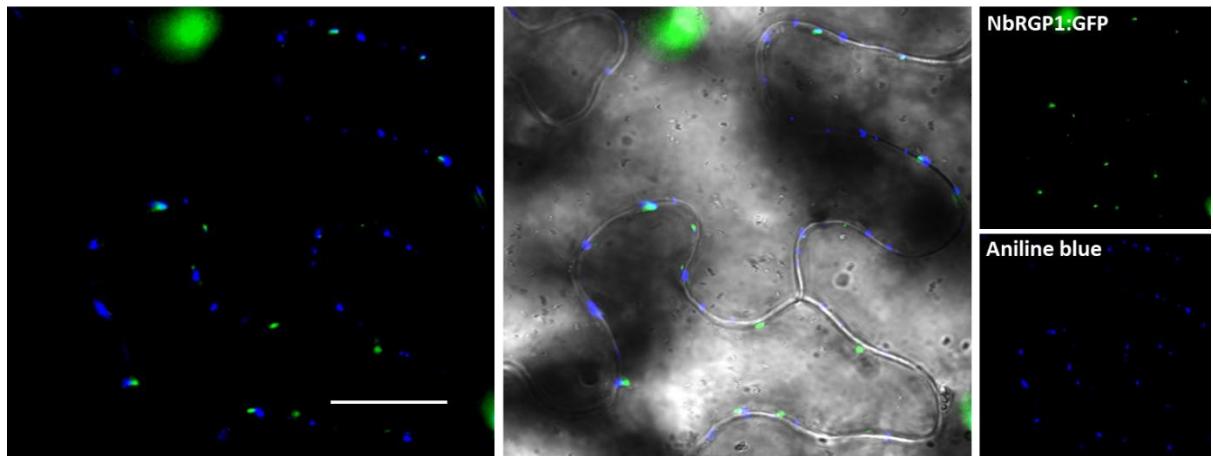
**Figure S2.** Comparison of PCR products obtained from cDNA and genomic DNA (gDNA) for each NbRGP.



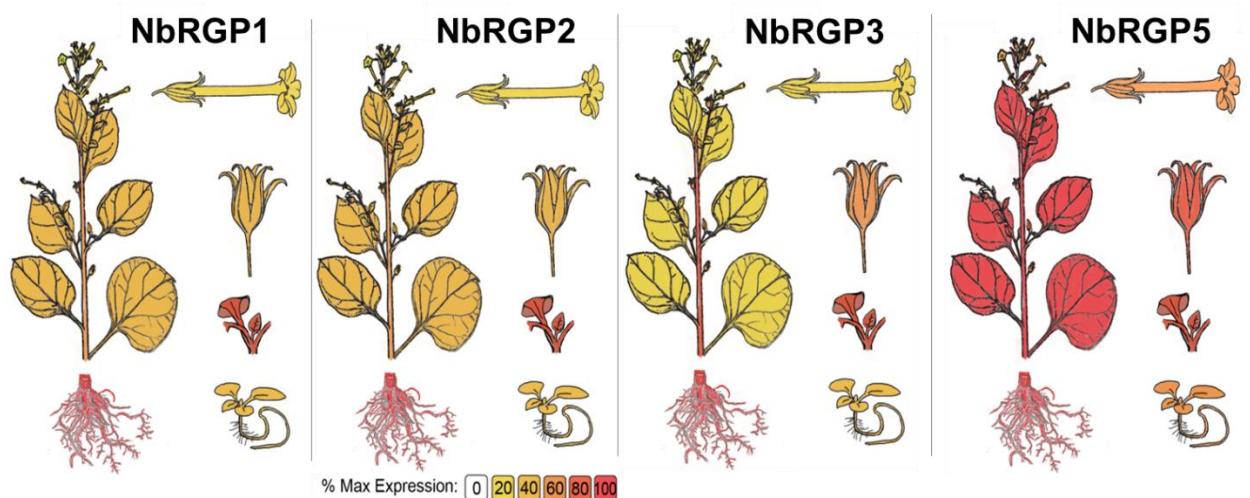
**Figure S3. TMV systemic infection.** Total soluble protein extracted from the intact leaf (int. leaf) and leaf with systemic TMV infection separated in PAAG and stained with Coomassie blue. The arrowhead indicates the band corresponding to TMV coat protein.



**Figure S4.** Confocal images of epidermal cells with *NbRGPs* overexpression after callose staining with aniline blue.



**Figure S5.** Confocal images of epidermal cells expressing 35S-NbRGP1:GFP (green) and stained with aniline blue (blue).



**Figure S6.** *NbRGPs* patterns of expression according to Version 6 Gene expression Atlas <https://sefapps02.qut.edu.au/atlas/tREX6.php>

**Table S1. Oligonucleotides used for cloning**

F1	GGTACCATGGCAGCAGCAACGCCATTG
F2	GGTACCATGGTAGGACGATCAATG
F3	TCCGATGGCAACGGCTTC
F4	GGGCCGTGGTGAGCAAGGGCGAGGAG
F5	GGTACCATGTCGCTAGTCAGTG
F6	GGGCCGCCGACAAGCAGAAGAACGG
F7	GGTACCATGGCTCTAGTTGTTAAAGG
F8	GGTACCGCAGCAGCAACGCCATTG
F9	GGATCCATAAGAACGGGGCCCAGAGTGAG
F10	GGATCCATAAGAACGGGGCCCATCGCCTCCTC
F11	GGTACCATGTCGCTCTCGATTGATTG
F12	GGATCCATAAGAACGGGGCCCATCGCCTCCTC
F13	GGTACCATGAAGATCATATCAAGGA
R2	GTCGACCTACTTGCCGGGCCATTG
R3	GTCGACTATTTGAGCTGCCATTGG
R4	GGATCCCAGCTTGAGCCTTGCAGG
R5	GTCGACTTAAGTTGGGTTTGACAG
R9	GTCGACTTACCTGTACAGCTCGTCCA
R10	GTCGACTTAGGCAGCCGGTGGAGTG
R11	GTCGACTTACATGATATAGACGTTGTGGCTGTTGAG
R12	GTCGACTTAGGACTTGTACAGCTCGTCCATGCC
R13	GGATCCAAACGAATCCGATTGGC
R14	GGATCCTACAAACAGAACGCTAGTTCCCCG
R15	GTCGACTTAGGCAGCCGGTGGAGTG
R16	GGATCCAACCTTCTTGAACACAAT

**Table S2. Oligonucleotides used for qRT-PCR**

Gene	Forward primer	Reverse primer
18S rRNA	ACGGCTACCACATCCAAG	ACTCATTCCAATTACCAGACTC
PP2A	ATTGCTGCCTGTGGTTATTAC	ATAGACTGAAGTGCTTGATTGG
NbRGP1	GGAATGAACCTGGCCTTGACC	GCCCACATATCATCGTAACGAC
NbRGP2	GATTCTGCTTGTGGTGCTT	CACGAACGAAATCTGCACCT
NbRGP3	TCGAGGATTCTGCTTGCCGT	TCCTGCCAGTTGGGTCTTT
NbRGP5	CTGGCGCTGAGGAACACT	ATGGTAGCAAAGCAGGTCCC
MP	GGTGTGAGCGTGTCTGG	GCGTCCTGGGTGGTTATAGC
GFP	GCAGAAGAACGGCATCAAG	GCTCAGGTAGTGGTTGTCG