

# Supplemental data

## Impact of *Paraburkholderia phytofirmans* PsJN on grapevine phenolics metabolism

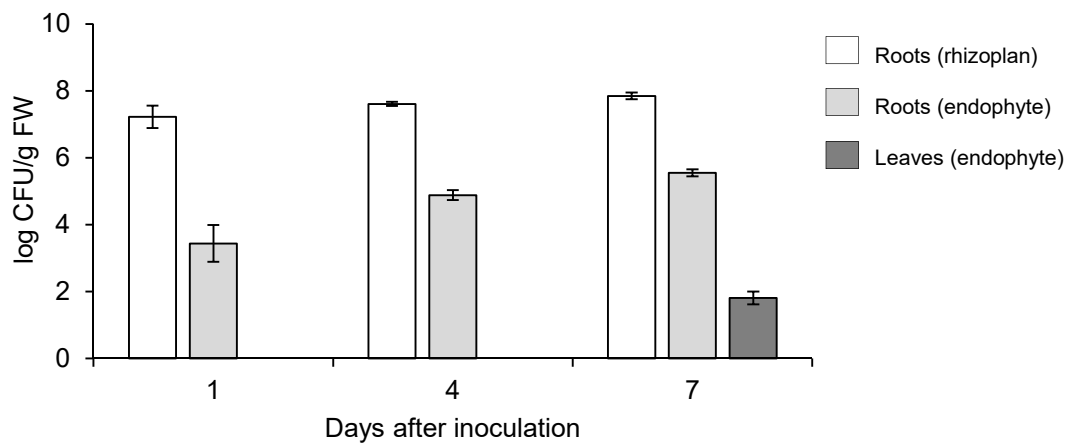
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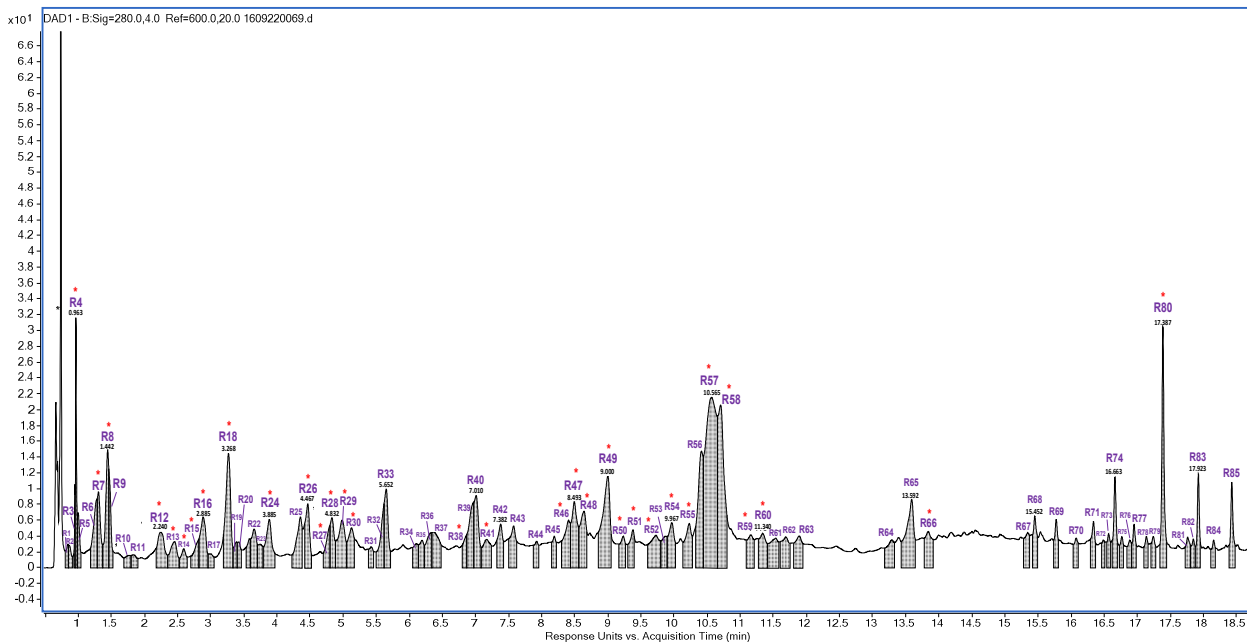
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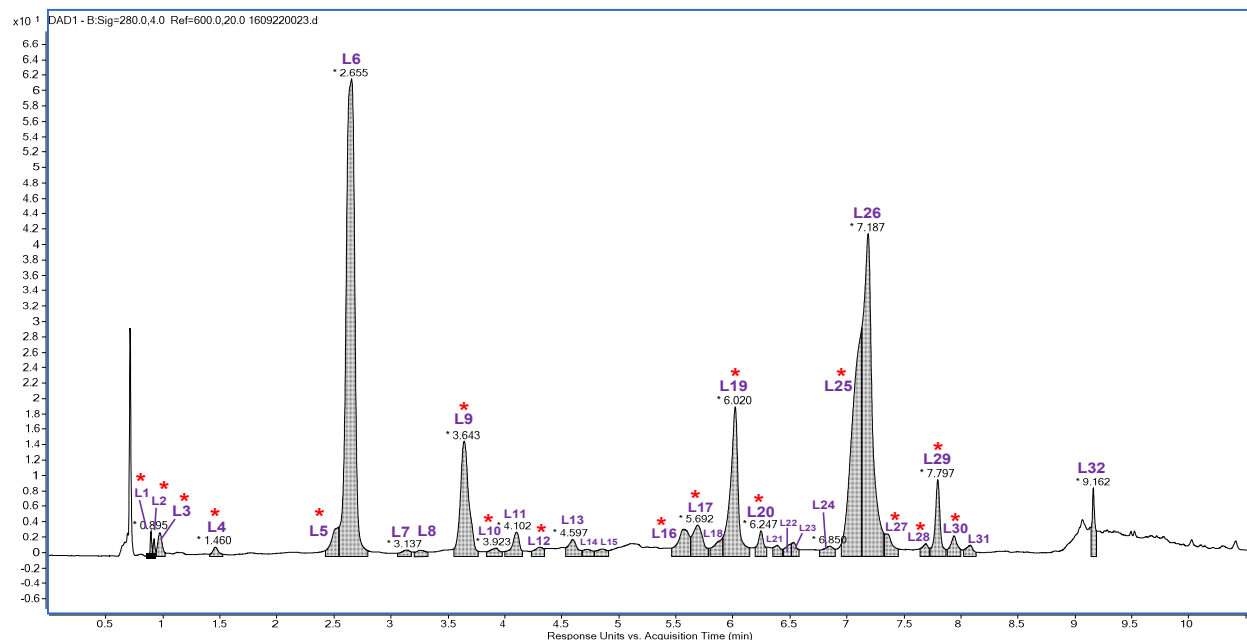


Supplemental figure S1: Grapevine colonization by *P. phytofirmans* PsJN. Values shown are means +/- SD of three independent repetitions (each repetition was realized in triplicates)

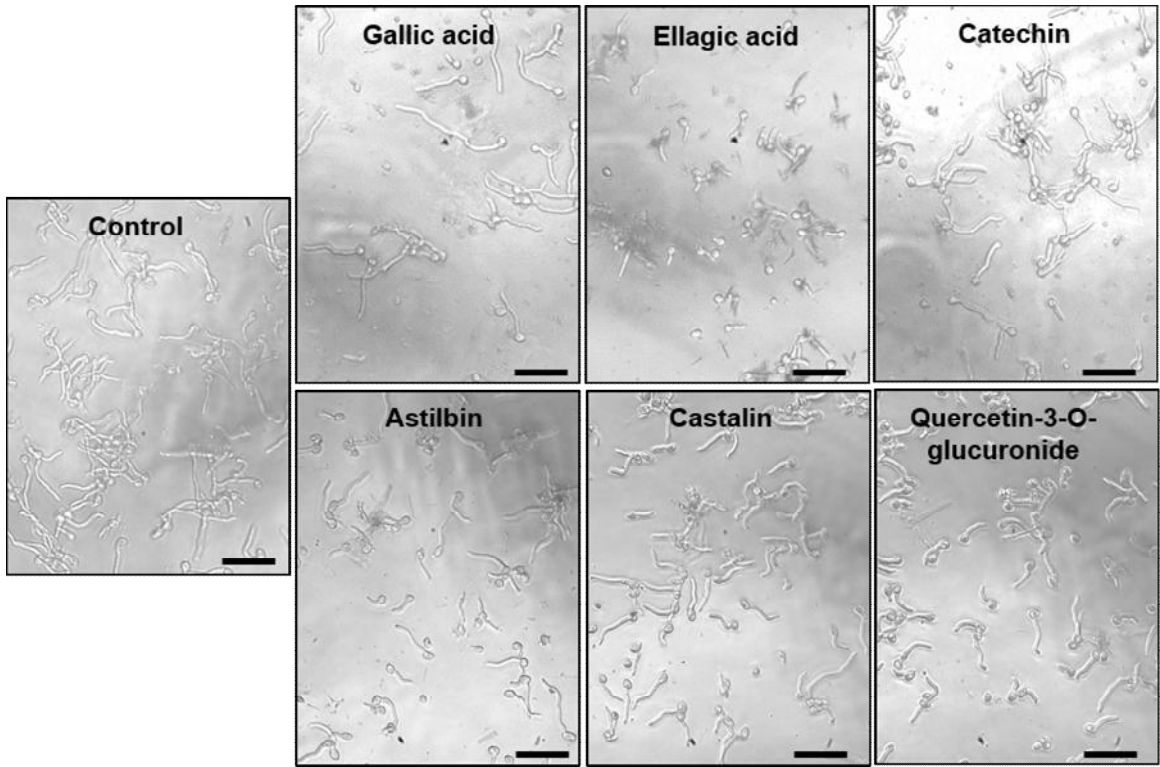
(a)



(b)



Supplemental figure S2: Analyses by UHPLC-UV/DAD-MS QTOF of phenolic compounds extracted from grapevine roots (a) and leaves (b) inoculated or not with *P. phytofirmans* strain PsJN. The chromatogram at 280 nm presented corresponds to the analysis of the QC (Quality control) sample for (a) roots and (b) leaves of the first biological repeat. The integrated and numbered chromatographic peaks were taken into account for the comparison of the profiles between the two conditions (control and bacterized). \* indicates the peak with a significant difference ( $P < 0.05$ , Student t-test) between the two conditions.



Supplemental figure S3: Effect of molecules on *Botrytis cinerea* spore germination. Conidia were placed in growth medium supplemented with solutions of molecules at 0.1mg/mL. Germ tubes were observed by inverted light microscopy 8h later. Scale: 50  $\mu$ m.

Supplemental Table S1: Primers used in this study

Gene symbol	Gene name	Forward primer sequence	Reverse primer sequence	Reference
<i>PAL</i>	phenylalanine-ammonia lyase	TCCTCCCGGAAAACAGCTG	TCCTCCAATGCCTCAAATCA	Varnier et al. 2009
<i>STS</i>	stilbene synthase	AGGAAGCAGCATTGAAGGCTC	TGCACCAGGCATTTCTACACC	Varnier et al. 2010
<i>CHS1</i>	chalcone synthase	CGAAGGAGCAATCGACGGA	GTCGCTGATGCCTATCGGAG	this study
<i>CHS2</i>	chalcone synthase	GGAAGATGGGAATGGCTGCT	GAGAGAAGGCACAGGGACAC	this study
<i>CHS3</i>	chalcone synthase	GCCCTAAAGCCCAGAAAGTT	AGCCGACTTCTCCTCATCT	this study
<i>CHI1</i>	chalcone isomerase 1	GCAGAAGCCAAAGCCATTGA	GCCGATGATGGACTCCAGTAC	this study
<i>CHI2</i>	chalcone isomerase 2	TCCAGATCAAGTTCACAGCA	GAAACAAGAGCCTCAAAGAA	Gutha et al. 2010
<i>FLS1</i>	flavonol synthase	CAGGGCTTGCAGGTTTTTAG	GGGTCTTCTCCTTGTTACAG	Gutha et al. 2010
<i>LAR1</i>	leucoanthocyanidin reductase 1	AAATGAACTCGCATCTGTGT	CTGTGGGATGATGTTTTCTC	Gutha et al. 2010
<i>LAR2</i>	leucoanthocyanidin reductase 2	TGATATCAGCTGTGGGTGGA	CCCAAATTCTGATGGAAGGA	Gutha et al. 2010
<i>LDOX</i>	leucoanthocyanidin dioxygenase	ATGAGGGCAAGTGGGTGACA	TTGACCAGTCCCCTGTGAAGA	this study
<i>ANR</i>	anthocyanidin reductase	GCTGCTGTTACCATCAATCA	GCAGGATAGCCCCAAGTAGG	Gutha et al. 2010
<i>UFGT</i>	UDP-glucose:flavonoid 3-O-glucosyltransferase	GGGATGGTAATGGCTGTGG	ACATGGGTGGAGAGTGAGTT	Gutha et al. 2010

- Gutha, L.R.; Casassa, L.F.; Harbertson, J.F.; Naidu, R.A. Modulation of flavonoid biosynthetic pathway genes and anthocyanins due to virus infection in grapevine (*Vitis vinifera* L.) leaves. *BMC Plant Biol* **2010**, *10*, 187. doi: 10.1186/1471-2229-10-187.

- Varnier, A.L.; Sanchez, L.; Vatsa, P.; Boudesocque, L.; Garcia-Brugger, A.; Rabenoelina, F.; Sorokin, A.; Renault, J.H.; Kauffmann S.; Pugin, A.; Clément, C.; Baillieul, F.; Dorey, S. Bacterial rhamnolipids are novel MAMPs conferring resistance to *Botrytis cinerea* in grapevine. *Plant Cell Environ* **2009**, *32*, 178-193. doi: 10.1111/j.1365-3040.2008.01911.x.