

TABLE S4 A: OD 30-75 Values: Means \pm Standard Deviations

<i>Agrobacterium</i> strains	N	OD ₃₀₋₇₅ Mean \pm SE		
HP1836	12	0.66883333 \pm	0.04599769	
HP1837	12	0.15591667 \pm	0.02306693	
HP1838	12	0.81525000 \pm	0.06015000	
HP1839	12	0.03091667 \pm	0.01625623	
HP1840	12	0.55350000 \pm	0.14573232	
HP1841	12	0.43513333 \pm	0.06826824	
HP1842	12	0.79608333 \pm	0.07247502	
HP1843	12	0.79933333 \pm	0.05940054	
SZ1	12	0.04375000 \pm	0.01183312	
SZL2	12	0.33241667 \pm	0.04599769	
SZL3	12	0.66883333 \pm	0.06230199	
SZL4	12	0.06183333 \pm	0.0172459	

TABLE S4 B: Duncan’s Multiple Range test of the OD 30-75 VALUES											
Alpha = 0.05; Error Degrees of Freedom: 22; Error Mean Square = 40.23677;											
No of Means				2			3			4	
Critical Range				6.201			6.512			6.710	
No of Means	2	3	4	5	6	7	8	9	10	11	12
Critical Range	10.74	11.28	11.62	11.86	12.04	12.18	12.29	12.38	12.45	12.50	12.55
		DC									
Duncan’s Groups											
Super group		Duncan’s Groups			STRAINS		N		MEAN		
EMA RESISTANT		DC			STRAINS		N		MEAN		
		A			HP1838		12		0.81525		
		A			HP1843		12		0.79933		
		A			SZL004		12		0.79658		
		B			HP1836		12		0.66883		
		C			HP1840		12		0.55350		
		D			HP1841		15		0.43513		
		E			SZL002		12		0.33242		
EMA SENSITIVE		F			HP1837		12		0.15592		
		G			SZL003		12		0.06183		
		G			SZ001		12		0.04375		
		G			HP1839		12		0.03092		

Footnotes to Table S4: The data analysis was performed using [SAS/STAT] software, Version [9.4] of the SAS System for [Windows X 64 Based Systems]; (Copyright © [2013 of copyright]; SAS Institute Inc. SAS, Cary, NC, USA, see Footnotes to Table S3. The significance of differences of the means ($\alpha = 0.05$) were determined here by using Duncan's Multiple Range Tests, depending upon the experiment as a part of the Anova Procedure. Duncan's Multiple Range Test of OD₃₀₋₇₅ values measured in Liquid Culture Bioassay of EMA PF on *Agrobacterium* strains. **Abbreviations:** EMA PF: Antimicrobially Peptid Rich Fraction from the cell-free culture media (CFCM) of *Xenorhabdus budapestensis* (EMA) (AF13), see Vozik et al., 2015. MIC: minimal inactivation concentration. For HP1837: MIC₇₅; for HP1839, SZL1 & SZL3: MIC₉₀. We have been considering the Duncan's Multiple Range test as the most accurate to distinguish between experimental groups reacting differently to the same treatments. The means within a given Duncan's Group labelled with a letter, say, with letter A, may differ from each other, but the SD values overlap; but differ significantly from those belonging to another Duncan's Group, labelled, say, letter B, are significantly different at P=0.05 level. We overchecked each case with t(LSD) tests as well (data are not given), and found that the Duncan's Multiple Range Tests were completely fair.

The ANOVA-based Duncan's Multiple Range test of the OD (30-75) of the EMA_PF treated *Agrobacterium* strains scored them to **7 Duncan's Groups (DG)**; and we scored the 7 Duncan's Groups to 2 clearly unambiguously separable "Super-Groups", (**Table S3B**).

Four *Agrobacterium* strains of S phenotype, including HP1837 of Duncan's Group F); HP1839, SZL1, and SZL3 of Duncan's Group G) were scored to **Duncan's Super-Group I**.

The rest of the *Agrobacterium* strains are of the R phenotype, including the wild-type (TDNA) (+) AGR strain HP1838 (of Duncan's Group A); the pMP90 helper-plasmid harboring SZL4 and plasmid-cured (HP1836, HP1840; HP1841; HP1842; HP 1843)), all nopaline catabolizing strains belonging to Duncan's Groups A-D); as well as one (SZL2) of the two octopine catabolizing strains examined, were scored to **Duncan's Super-Group II**.