

Table S4. Cellular fatty acids content of *Photobacterium sanctipauli* sp. nov. and related taxa of the genus *Photobacterium*. Taxa: **1**, *P. sanctipauli* (range profile of five strains); **2**, *P. rosenbergii* LMG 22223^T (Thompson et al. 2005); **3**, *P. gaetbulicola* Gung47^T; and **4**, *P. lutimaris* LMG 25278^T (Kim et al. 2010). Summed feature 2 comprises C_{12:0} ALDE, iso-C_{16:1} I and/or C_{14:0} 3-OH and/or an unidentified fatty acid with equivalent chain length of 10.928. Summed feature 3 comprises C_{16:1}ω7c and/or iso-C₁₅ 2-OH. Data are expressed as percentages of total fatty acids. Fatty acids representing <1 % are not shown, except for C_{18:0}, because it was represented in all strains with approximately the same intensity

Fatty acids	1	2	3	4
C _{12:0}	3.7-4.5	2	1.7	2.4
C _{12:0} 3-OH	2.5-3.6	2-3	2.8	2.6
C _{14:0}	4.7-5.8	3-4	4.2	2.4
C _{15:0} iso	0.6-1.1	2-4	6.9	5
C _{16:0} iso	-	1.9	0.4	-
C _{16:0}	18.5-21.4	10-15	14	13.1
C _{17:0}	1.2-1.6	1	1	-
C _{17:0} iso	0.7-1.5	3-6	3.5	1.8
C _{17:1} ω8c	0.3-1.1	1	0.6	-
C _{18:0}	0.5-0.7	-	-	-
C _{18:1} ω6c	0-3.3	-	3.1	-
C _{18:1} ω7c	7.9-12.7	17-19	10.3	13
Summed feature 2	3.7-4.7	2-3	3.7	3
Summed feature 3	43-45.5	41-44	33.6	43.3

References

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