**Table S5.** Cellular fatty acid compositions (%) of strains D3-12<sup>T</sup> and G2-2<sup>T</sup> and closely related taxa of the family *Roseobacteraceae* 

Taxa: 1, strain D3-12<sup>T</sup>; 2, strain G2-2<sup>T</sup>; 3, *Rhodalgimonas zhirmunskyi* KCTC 72611<sup>T</sup>; 4, *Ponticoccus litoralis* KCCM 90028<sup>T</sup>; 5, *Aquicoccus porphyridii* KACC 18806<sup>T</sup>; 6, *Marimonas lutisalis* KCTC 62376<sup>T</sup>; 7, *Marimonas arenosa* KCTC 52189<sup>T</sup>; 8, *Cognatishimia maritima* KCTC 23347<sup>T</sup>. All data were obtained from this study. Data are presented as percentages of total fatty acids, with fatty acids accounting for less than 0.5% in all strains excluded from the results. Major components (> 5.0%) are highlighted in bold. tr, trace amount (< 0.5%); –, not detected.

Fatty acid	1	2	3	4	5	6	7	8
Saturated:								
$C_{10:0}$	0.6	0.6	0.7	0.8	tr	0.5	tr	tr
$C_{12:0}$	0.8	0.8	0.7	1.4	tr	1.2	tr	2.0
$C_{16:0}$	5.9	17.8	4.4	8.1	6.9	2.8	12.3	3.0
$C_{17:0}$	tr	1.4	1.2	tr	1.0	tr	1.8	1.0
$C_{18:0}$	2.3	1.2	0.8	6.1	1.7	3.4	4.2	3.3
Unsaturated:								
iso- $C_{17:1} \omega 5c$	_	tr	_	0.6	_	_	_	0.5
11-methyl- $C_{18:1} \omega 7c$	10.1	_	_	_	2.1	4.9	_	_
cyclo- $C_{19:0} \omega 8c$	_	1.6	_	_	10.8	1.1	0.7	_
$C_{20:1} \omega 7c$	tr	_	_	_	tr	0.7	_	_
Branched:								
iso- $C_{10:0}$	_	tr	_	0.7	_	_	_	0.7
iso- $C_{18:0}$	1.3	_	_	_	_	_	_	_
$iso-C_{19:0}$	_	tr	_	tr	_	_	_	0.5
Hydroxy:								
$C_{10:0}$ 3-OH	tr	tr	tr	_	tr	0.6	4.4	tr
$C_{11:0}$ 3-OH	_	_	0.7	_	_	_	_	_
$C_{12:0}$ 3-OH	2.1	2.2	2.2	0.9	0.7	0.8	tr	3.5
iso-C <sub>12:0</sub> 3-OH	_	0.6	tr	1.9	_	_	_	3.3
C <sub>12:1</sub> 3-OH	tr	tr	_	3.3	1.8	1.6	1.2	_
iso-C <sub>17:0</sub> 3-OH	_	_	0.6	_	_	_	_	_
Summed features*:								
3	tr	3.3	0.5	0.7	0.6	tr	0.7	0.7
7	_	_	tr	_	_	_	_	0.5
8	75.1	67.6	85.6	73.9	69.9	78.2	72.7	78.5

<sup>\*</sup>Summed features are groups of fatty acids that cannot be resolved reliably using the chromatographic conditions chosen. The MIDI system groups these fatty acids together as one feature with a single percentage of the total. Summed features 3, 7, and 8 comprise  $C_{16:1} \omega 7c$  and/or  $C_{16:1} \omega 6c$ ,  $C_{19:1} \omega 7c$  and/or  $C_{19:1} \omega 6c$ , and  $C_{18:1} \omega 7c$  and/or  $C_{18:1} \omega 6c$ , respectively.