

**Supplementary Table S1:** DDH and ANI values between *M. rosea* and selected myxobacteria. Color intensity changes from green to orange corresponding with higher to lower values, respectively.

Query	Sub order	Family	Members	Genome size (Mbp)	DDH	ANI value (%)
<i>Minicystis roseae</i> DSM 2400	Sorangineae	<i>Labilitrichaceae</i>	<i>Labilithrix luteola</i> DSM 27648	12.19	19.1	70.26
		<i>Polyangiaceae</i>	<i>Chondromyces apiculatus</i> DSM 436	11.49	19.8	72.93
			<i>Polyangium fumosum</i> DSM 14668	12.88	20.1	73.42
			<i>Polyangium</i> sp. SDU3-1	12.28	19.7	73.6
			<i>Sorangium cellulosum</i> So ce26	14.56	20.7	74.34
			<i>Sorangium cellulosum</i> So ce56	13.03	20.9	74.64
			<i>Sorangium cellulosum</i> So ce836	14.59	21.1	74.73
			<i>Sorangium cellulosum</i> So ceGT47	11.26	20.9	75
			<i>Sorangium cellulosum</i> So0003-19-2	10.09	21.4	75
			<i>Sorangium cellulosum</i> So0007-03	12.39	21.4	74.97
			<i>Sorangium cellulosum</i> So0008-312	11.21	20.8	75
			<i>Sorangium cellulosum</i> So0157-2	14.78	21	75
			<i>Sorangium cellulosum</i> So0163	12.46	21.2	74.86
	<i>Sandaracinaceae</i>	<i>Sandaracinus amyloyticus</i> DSM 53668	10.33	18.3	70.05	
	Cystobacterineae	<i>Anaeromyxobacteraceae</i>	<i>Anaeromyxobacter dehalogenans</i> 2CP-1	5.03	18.8	69.52
		<i>Archangiaceae</i>	<i>Archangium gephyra</i> DSM 2261	12.49	18.4	68.51
			<i>Cystobacter fuscus</i> DSM 52655	12.35	19.1	68.72
			<i>Hyalangium minutum</i> DSM 14724	10.12	18.8	66.39
		<i>Myxococcaceae</i>	<i>Myxococcus hansupus mixupus</i>	9.49	18.5	68.19
	<i>Vulgatibacteraceae</i>	<i>Vulgatibacter incomptus</i>		4.35	18.1	68.56
	<i>Nannocystineae</i>	<i>Nannocystaceae</i>	<i>Enhygromyxa salina</i> DSM 15201	10.26	17.3	67.32

**Supplementary Table S2:** Excel file has been provided.