Ε	DMSP degradation:	SP degradation: Step Summaries for Genomes above 75% Complete															
Verrucomicrobiota -																	
SAR324 -																	
Planctomycetota -																	
Patescibacteria -																	
Other -																	
Marinisomatota -																	
Gemmatimonadota -																	
Gamma-SAR92 -																	
Gamma-SAR86 -																	
Gamma-Other -																	
Gamma-Alteromonas -																	
Firmicutes -																	
Dadabacteria -																	
Cyano–Syn -																	Bro
Cyano–Pro -																value 1.00	ыо
dnoo Cyano-Other -																0.75 0.50	
Chloroflexota -																0.25	
Bacteroidota -																	
Bacteria-Other -																	
Archaea-Poseidoniia MGIIb -																	
Archaea-Poseidoniia MGIIa -																	
Archaea-Other -																	
Archaea-Nitrosopumilaceae -																	
Archaea–Halobacteria -																	
Alpha-Rhodobacter-LowGC -																	
Alpha-Rhodobacter -																	
Alpha-Puniceispirilla -																	
Alpha-Pelagibacter -																	
Alpha-Other -																	
Actinobacteriota -																	
Acidobacteriota -																	
	Step1	Step2	Step3	Step4	Step5	Step6	Step7	Subpathway1 Step or Calculation	Subpathway2	Subpathway3	Subpathway4	Subpathway5	Subpathway6	Subpathway7	MaxCompletion		

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