

Table S2.PCA analysis of SEED Subsystems level 1 (Meyer *et al.*, 2008).

Correlation of each variable with factor 1

	Correlation	P-value
Cluster system	0.99922	2.95E-42
Cofactors and pigments	0.997675	2.18E-35
Miscellaneous	0.997555	4.52E-35
Amino acids	0.997344	1.50E-34
Carbohydrates	0.996973	9.97E-34
Protein metabolism	0.996624	4.83E-33
Cell division	0.995114	1.02E-30
Cell wall	0.995105	1.05E-30
Lipids	0.99492	1.79E-30
Nucleotides	0.994764	2.77E-30
DNA metabolism	0.994473	6.06E-30
RNA metabolism	0.994238	1.11E-29
Respiration	0.993927	2.37E-29
Stress response	0.993802	3.18E-29
Transport	0.993616	4.88E-29
Sulfur	0.992958	2.02E-28
Aromatic	0.992208	8.69E-28
Nitrogen	0.991532	2.89E-27
Secondary metabolism	0.990146	2.58E-26
Virulence	0.987861	5.24E-25
Cell signaling	0.987054	1.33E-24
Spore	0.980752	4.00E-22
Motility	0.97793	2.86E-21
Potassium	0.977141	4.73E-21
Phosphorus	0.97681	5.82E-21
Phages	0.969652	2.75E-19
Iron	0.969093	3.57E-19
Photosynthesis	0.927324	6.60E-14