

Significantly enriched (FDR<0.05) GO Cellular Component terms and groups in the 80 differential proteins (p<0.05 or 0.5>FC>2 between Control and Inoculated)								
GO_ID	GO_Term	Source	Term FDR	Group FDR	Enrichment score	GO groups	% Associated proteins	Number of proteins
GO:0044445	Cytosolic part	CC	4.83E-27	1.96E-25	82.1	4	7.20	25.00
GO:0022626	Cytosolic ribosome	CC	1.02E-26	1.96E-25	82.1	4	7.57	24.00
GO:0005840	Ribosome	CC	1.42E-24	1.96E-25	82.1	4	5.49	25.00
GO:0044391	Ribosomal subunit	CC	2.01E-20	1.96E-25	82.1	4	6.83	19.00
GO:0022627	Cytosolic small ribosomal subunit	CC	2.85E-12	1.96E-25	82.1	4	9.17	10.00
GO:0015935	Small ribosomal subunit	CC	7.46E-12	1.96E-25	82.1	4	8.20	10.00
GO:0022625	Cytosolic large ribosomal subunit	CC	3.10E-10	1.96E-25	82.1	4	7.09	9.00
GO:0015934	Large ribosomal subunit	CC	1.61E-09	1.96E-25	82.1	4	5.81	9.00
GO:0010319	Stromule	CC	1.17E-07	1.41E-07	22.8	1	13.89	5.00
GO:0000786	Nucleosome	CC	5.74E-07	3.60E-05	14.8	2	10.00	5.00
GO:0000785	Chromatin	CC	3.38E-05	3.60E-05	14.8	2	4.24	5.00
GO:0000790	Nuclear chromatin	CC	3.65E-05	3.60E-05	14.8	2	6.90	4.00
GO:0009521	Photosystem	CC	6.51E-05	5.64E-05	14.1	3	5.71	4.00
GO:0009522	Photosystem I	CC	1.64E-04	5.64E-05	14.1	3	8.57	3.00
GO:0009523	Photosystem II	CC	4.71E-04	5.64E-05	14.1	3	5.88	3.00

The leading term of each enriched group is that with the lowest term FDR (highest enrichment score), highlighted in bold and used to name the respective group.