Supplementary Figure: 8 :Snapshot of PAGE SEACOMPARE. It is Cross comparison of PAGE of DETs at all 6 timepoints .

Bottom panel

GO Analysis Toolkit and Database for Agricultural Community

IOME ANALYSIS TOOL SEARCH DOWNLOADS MANUAL FAQ

Select terms to [Draw graphical results 击] [Create bar chart 🏭] [Suppress term number ந] [Use NO-adjusted Pvalue 🎲] Θ

GO Information		СМ			Ohr			12hr			24hr			48hr			120hr			240hr		
No GO Term Onto Number	Description	1 2 3	4 5	6 Z-score	Mean	FDRnbsp;	Z-score	Mean	FDRnbsp;	Z-score	Mean	FDRnbsp	; Z-score	Mean	FDRnbsp;	Z-score	Mean	FDRnbsp;	Z-score	Mean	FDRnbsp;	
74 GO:0043436 P 18	oxoacid metabolic process			-2	-1.8	0.17	-2.6	-2.4	0.08	-1.5	-1.7	0.41	-3.1	-2.2	0.044	-2.4	-1.8	0.13	-3.3	-2.7	0.016	
75 GO:0042180 P 18	cellular ketone metabolic process			-2	-1.8	0.17	-2.6	-2.4	0.08	-1.5	-1.7	0.41	-3.1	-2.2	0.044	-2.4	-1.8	0.13	-3.3	-2.7	0.016	
76 GO:0019752 P 18	carboxylic acid metabolic process			-2	-1.8	0.17	-2.6	-2.4	0.08	-1.5	-1.7	0.41	-3.1	-2.2	0.044	-2.4	-1.8	0.13	-3.3	-2.7	0.016	
77 GO:0006082 P 18	organic acid metabolic process			-2	-1.8	0.17	-2.6	-2.4	0.08	-1.5	-1.7	0.41	-3.1	-2.2	0.044	-2.4	-1.8	0.13	-3.3	-2.7	0.016	
78 GO:0032787 P 14	monocarboxylic acid metabolic process			-1.5	-1.5	0.37	-2.6	-2.7	0.08	-1.5	-1.9	0.41	-3.1	-2.6	0.044	-2.6	-2.1	0.13	-3.4	-3.2	0.016	
79 GO:0015979 P 13	photosynthesis			-1.2	-1.3	0.52	-1.5	-1.7	0.37	-1.6	-2	0.41	-3.2	-2.7	0.044	-2.7	-2.3	0.13	-2.8	-2.7	0.051	
80 GO:0044281 P 37	small molecule metabolic process			-2.3	-1.5	0.12	-2.6	-1.7	0.08	-1.8	-1.4	0.39	-3.5	-1.7	0.044	-2.7	-1.3	0.13	-4	-2.3	0.01	
81 GO:0003676 F 227	nucleic acid binding		8	1.8	0.4	0.15	1.2	0.068	0.45	1.4	0.0066	0.32	2.8	0.82	0.019	1.8	0.47	0.16	2	0.62	0.12	
82 GO:0003700 F 182	transcription factor activity			1.6	0.39	0.22	1.3	0.15	0.36	1.1	-0.03	0.46	2.5	0.81	0.037	1.1	0.34	0.42	1.4	0.5	0.28	
83 GO:0030246 F 14	carbohydrate binding			2.2	2.2	0.058	1	0.74	0.48	1.6	1.3	0.22	2.4	2.4	0.039	2.4	2.1	0.048	2.1	2.2	0.087	
84 GO:0030528 F 193	transcription regulator activity			1.5	0.35	0.26	1.2	0.11	0.42	1.1	-0.027	0.45	2.4	0.77	0.042	1	0.32	0.46	1.4	0.51	0.27	
85 GO:0009505 C 23	plant-type cell wall			-1.7	-1.4	0.22	-2.4	-2	0.06	-1.8	-1.7	0.18	-2.5	-1.5	0.022	-2	-1.3	0.074	-2.4	-1.6	0.036	
86 GO:0043234 C 48	protein complex			-0.56	-0.34	0.76	-0.22	-0.32	0.91	-0.75	-0.75	0.61	-2.6	-1	0.019	-2.3	-1	0.04	-1.8	-0.78	0.12	
87 GO:0044464 C 661	cell part			-1.3	-0.23	0.37	-1.2	-0.37	0.41	-1.6	-0.58	0.22	-2.9	-0.19	0.0082	-2.3	-0.2	0.043	-2.2	-0.18	0.046	
88 GO:0005623 C 661	cell			-1.3	-0.23	0.37	-1.2	-0.37	0.41	-1.6	-0.58	0.22	-2.9	-0.19	0.0082	-2.3	-0.2	0.043	-2.2	-0.18	0.046	
89 GO:0043229 C 391	intracellular organelle			-0.31	-0.1	0.86	-0.15	-0.23	0.94	-0.84	-0.5	0.6	-3.1	-0.33	0.0043	-3.1	-0.41	0.0048	-2.4	-0.31	0.032	
90 GO:0043226 C 391	organelle			-0.31	-0.1	0.86	-0.15	-0.23	0.94	-0.84	-0.5	0.6	-3.1	-0.33	0.0043	-3.1	-0.41	0.0048	-2.4	-0.31	0.032	
91 GO:0043231 C 379	intracellular membrane-bounded organelle			-0.36	-0.11	0.86	-0.26	-0.25	0.9	-1	-0.54	0.5	-3.2	-0.36	0.0033	-3.2	-0.44	0.0034	-2.5	-0.33	0.031	
92 GO:0005622 C 448	intracellular			-0.17	-0.073	0.94	-0.046	-0.22	0.96	-0.67	-0.46	0.61	-3.2	-0.32	0.0033	-3.2	-0.39	0.004	-2.6	-0.3	0.026	
93 GO:0043227 C 380	membrane-bounded organelle			-0.36	-0.11	0.86	-0.25	-0.25	0.9	-1.1	-0.54	0.49	-3.3	-0.37	0.0031	-3.3	-0.45	0.0031	-2.5	-0.33	0.029	
94 GO:0009570 C 31	chloroplast stroma			-1.3	-0.88	0.4	-1.2	-1	0.4	-1.4	-1.3	0.31	-3.4	-1.8	0.0022	-3.5	-1.9	0.0014	-2.8	-1.7	0.014	
95 GO:0044424 C 428	intracellular part			-0.43	-0.12	0.83	-0.29	-0.26	0.9	-0.9	-0.5	0.59	-3.6	-0.38	0.0011	-3.5	-0.45	0.0014	-2.9	-0.37	0.013	
96 GO:0009532 C 37	plastid stroma			-1.5	-0.96	0.28	-1.5	-1.1	0.33	-1.6	-1.3	0.23	-3.9	-1.9	0.00037	-3.9	-2	0.00035	-3.2	-1.8	0.0039	
97 GO:0044446 C 104	intracellular organelle part			-1.5	-0.6	0.27	-1.1	-0.59	0.44	-2	-1.1	0.13	-4.1	-1.2	0.00013	-4.1	-1.2	0.00014	-3.3	-1	0.0034	
98 GO:0044422 C 104	organelle part			-1.5	-0.6	0.27	-1.1	-0.59	0.44	-2	-1.1	0.13	-4.1	-1.2	0.00013	-4.1	-1.2	0.00014	-3.3	-1	0.0034	
99 GO:0005737 C 300	cytoplasm			-1.1	-0.28	0.44	-0.58	-0.33	0.75	-1.3	-0.62	0.35	-4.8	-0.73	6e-06	-4.2	-0.7	8.7e-05	-3.6	-0.61	0.0014	
100 GO:0044444 C 283	cytoplasmic part			-1.6	-0.4	0.26	-1.1	-0.43	0.45	-1.9	-0.76	0.15	-5.3	-0.86	9.2e-07	-4.8	-0.84	5.1e-06	-4	-0.73	0.00026	
101 GO:0009507 C 188	chloroplast			-1.9	-0.55	0.16	-1.6	-0.61	0.29	-2.2	-0.95	0.074	-5.8	-1.2	2.2e-07	-5.5	-1.2	1.9e-07	-4.5	-1	3.4e-05	
102 GO:0009536 C 193	plastid			-2	-0.58	0.13	-1.7	-0.65	0.23	-2.3	-0.97	0.057	-6	-1.2	1.2e-07	-5.7	-1.2	8.2e-08	-4.6	-1.1	1.6e-05	
103 GO:0046483 P 12	heterocycle metabolic process			-1.4	-1.5	0.4	-1	-1.2	0.59	-2.6	-3.2	0.11	-2.4	-2.1	0.1	-3.8	-3.5	0.026	-2.9	-2.9	0.051	
104 GO:0009266 P 25	response to temperature stimulus			-0.56	-0.46	0.92	-1.3	-1.1	0.43	-1.9	-1.8	0.32	-2.1	-1.2	0.19	-3	-1.9	0.13	-3.5	-2.4	0.016	
105 GO:0004674 F 17	protein serine/threonine kinase activity			1.9	1.6	0.13	1.8	1.3	0.17	2.2	1.7	0.068	1.9	1.7	0.12	1.9	1.5	0.15	2.9	2.7	0.013	