

Supplementary data

Metabolite profiling reveals the effect of drought on sorghum (*Sorghum bicolor* L. Moench) metabolism

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Table 1. Identified metabolites of sweet sorghum leaves and roots harvested from plants grown under normal or drought stress conditions at two stress levels. Drought stress was induced by polyethylene glycol (PEG) at two levels of stress 2.5 % and 5 % PEG.

Suppl. Table 1a. Amino acids

	m/z	RT	Loadings		Control	2,5% PEG	5% PEG	P
			PC-1 (68%)	PC-2 (14%)	Average	Average	Average	
LEAVES								
Aspartic acid	232	17.6199	0.003212	-0.008828	2.75E-05	0.000159	0.00014	0.01
b-alanine	248	15.2100	0.008034	-0.003557	4.89E-06	0.000142	0.00029	0.00
DL-isoleucine	158	11.8931	0.005819	-0.006800	5.07E-05	0.000187	0.00026	0.05
glycine	174	12.1789	0.010352	-0.005670	2.4E-05	0.000252	0.00034	0.00
L-alanine	116	7.2591	0.038672	-0.030306	0.000199	0.001129	0.00138	0.00
L-asparagine	231	21.0587	0.180705	-0.259678	0.00026	0.005044	0.0072	0.00
L-cysteine	220	18.3804	n/d	n/d	n/d	n/d	n/d	-
L-glutamic acid	201	18.6980	0.022087	-0.019053	9.19E-05	0.000682	0.00087	0.00
L-homoserine	218	15.9019	0.000254	-0.000432	5.11E-07	7.04E-06	1.1E-05	0.03
L-lysine	317	26.2457	0.005667	0.006311	0	0	0.00013	0.14
L-methionine	176	17.4389	0.000496	-0.001212	1.21E-05	2.2E-05	4.7E-05	0.07
L-ornithine	174	24.1784	0.002399	-0.000744	0	3.93E-05	7.7E-05	0.02
L-serine	204	13.6984	0.057229	-0.066447	0.00025	0.001609	0.00229	0.00
L-threonine	218	14.3861	0.023377	-0.016571	0.000103	0.000629	0.00087	0.00
L-tryptophan	202	31.6634	0.028823	-0.033023	8.08E-05	0.000707	0.00114	0.00
Tyrosine	218	26.5646	0.050153	-0.044486	9.94E-06	0.001044	0.0017	0.00
L-valine	144	9.9049	0.037676	-0.005024	5.96E-05	0.000489	0.00141	0.00
ROOTS								
Aspartic acid	232	17.6199	0.004640	0.003836	8.11E-05	0.000399	0.00024	0.02
b-alanine	248	15.2100	0.000158	0.002001	5.16E-06	3.57E-05	5.3E-06	0.01
DL-isoleucine	158	11.8931	0.000018	-0.002300	0.000191	0.000138	0.0001	0.35
glycine	174	12.1789	0.006208	0.003953	3.98E-05	0.00027	0.00033	0.04

L-alanine	116	7.2591	0.013118	0.014621	4.69E-05	0.000667	0.00053	0.05
L-asparagine	231	21.0587	0.000071	0.010315	1.14E-05	0.000136	1E-05	0.08
L-cysteine	220	18.3804	-0.000086	-0.000033	6.41E-06	0	0	0.00
L-glutamic acid	201	18.6980	0.020389	0.002293	0.000195	0.000884	0.00109	0.05
L-homoserine	218	15.9019	0.000213	0.000226	8.35E-08	5.86E-06	1.6E-05	0.46
L-lysine	317	26.2457	0.002038	0.008083	9.12E-05	0.000289	0.00011	0.01
L-methionine	176	17.4389	-0.000107	-0.000077	9.2E-06	2.78E-06	0	0.01
L-ornithine	174	24.1784	0.001466	0.002243	1.47E-05	0.000113	6.4E-05	0.05
L-serine	204	13.6984	0.012675	0.021249	0.000283	0.001001	0.00069	0.04
L-threonine	218	14.3861	0.001968	0.007839	9.76E-05	0.000283	0.00013	0.01
L-tryptophan	202	31.6634	-0.001360	0.005405	0.00011	6.91E-05	1E-05	0.08
Tyrosine	218	26.5646	0.000529	0.003624	7.95E-05	0.000108	6.5E-05	0.42
L-valine	144	9.9049	0.000516	0.007929	0.000172	0.000238	0.00014	0.22

Table 1b. Organic acids

	m/z	RT	Loadings		Control	2,5% PEG	5% PEG	<i>P</i>
			PC-1 (68%)	PC-2 (14%)	Average	Average	Average	
LEAVES								
a ketoglutaric acid	147	18.9074	-0.000110	0.000185	4.71E-06	0	0	0.00
Caffeic acid	396	30.1434	0.003445	0.001521	3.21E-05	5.46E-05	0.00014	0.00
Citric acid	347	24.3955	0.006900	-0.020258	6.63E-05	0.000414	0.00025	0.01
Dehydroascorbic acid	173	24.9093	-0.000033	0.000053	1.41E-06	0	0	0.01
4-hydroxycinnamic acid	293	26.3507	0.002564	-0.000184	6.4E-06	3.02E-05	8.1E-05	0.01
DL-4-hydroxymandelic acid	267	23.4074	0.026514	-0.014107	1.6E-05	0.00063	0.00076	0.02
D-malic acid	147	16.8912	0.022863	-0.045825	0.000168	0.001097	0.00072	0.04
Fumaric acid	245	13.1481	0.000215	-0.000318	9.18E-07	6.24E-06	7.4E-06	0.03
Gluconic acid	333	28.1861	0.013507	-0.021143	2.69E-05	0.000578	0.0004	0.02
Glycolic acid	147	6.7007	0.022164	0.031249	1.18E-05	6.67E-05	0.00062	0.06
3,4-dihydroxymandelic acid	355	28.5997	0.000452	-0.000264	0	1.07E-05	1.4E-05	0.02
Mucic acid	333	28.9875	0.001692	-0.004968	1.22E-05	0.000105	5.3E-05	0.03
Pipecolic acid	156	13.5854	0.000401	-0.000711	5.87E-06	2.2E-05	1.7E-05	0.02
Pyruvic acid	174	6.2359	-0.000770	0.000946	3.8E-05	1.09E-05	3.4E-06	0.00
Saccharic acid	333	28.7452	0.001044	-0.002392	2.54E-06	5E-05	3.4E-05	0.01
Succinic acid	148	12.3228	0.001887	-0.002617	1.21E-05	6.9E-05	7.6E-05	0.00
Trans-aconitic acid	229	22.7348	0.037849	-0.088046	2.15E-05	0.001431	0.00127	0.01
Uric acid	441	29.8723	0.000130	-0.000055	7.7E-07	3.52E-06	4.5E-06	0.01
ROOTS								

a ketoglutaric acid	147	18.9074	n/d	n/d	n/d	n/d	n/d	-
Caffeic acid	396	30.1434	0.000063	0.000090	1.2E-06	4.12E-06	3.7E-06	0.21
Citric acid	347	24.3955	0.000402	0.005524	0.000185	0.000364	3.9E-05	0.00
Dehydroascorbic acid	173	24.9093	n/d	n/d	n/d	n/d	n/d	-
4-hydroxycinnamic acid	293	26.3507	0.001561	0.002647	2.25E-05	0.000122	0.00012	0.03
DL-4-hydroxymandelic acid	267	23.4074	0.004116	0.003047	4.63E-05	0.00029	0.0001	0.33
D-malic acid	147	16.8912	-0.006469	0.008983	0.000779	0.000362	0.00019	0.01
Fumaric acid	245	13.1481	0.000557	0.000598	1.27E-05	3.22E-05	3.5E-05	0.14
Gluconic acid	333	28.1861	0.001480	0.002879	5.89E-05	0.000194	0.0001	0.00
Glycolic acid	147	6.7007	0.021758	-0.015682	5.86E-05	0.000237	0.0016	0.00
3,4-dihydroxymandelic acid	355	28.5997	-0.000002	-0.000001	1.63E-07	0	0	0.12
Mucic acid	333	28.9875	-0.000036	0.001079	7.15E-06	1.72E-05	0	0.11
Pipecolic acid	156	13.5854	-0.000166	0.000339	1.91E-05	8.01E-06	2.5E-06	0.11
Pyruvic acid	174	6.2359	0.000033	-0.000133	5.87E-06	6.77E-06	8.4E-06	0.72
Saccharic acid	333	28.7452	-0.000010	0.000251	2.04E-06	5.93E-06	0	0.00
Succinic acid	148	12.3228	0.002841	0.000598	4.02E-05	0.000139	0.00019	0.01
Trans-aconitic acid	229	22.7348	0.000074	0.001552	3.76E-06	1.1E-05	6.7E-06	0.80
Uric acid	441	29.8723	n/d	n/d	n/d	n/d	n/d	-

Table 1c. Fatty acids

	m/z	RT	Loadings		Control Average	2,5% PEG Average	5% PEG Average	P
			PC-1 (68%)	PC-2 (14%)				
LEAVES								
Capric acid	229	15.7761	n/d	n/d	n/d	n/d	n/d	-
4-guanidinobutyric acid	174	17.6937	0.013202	-0.004076	1.65E-05	0.000277	0.00044	0.00
2-hydroxybutyric acid	147	7.7499	0.000037	0.000018	0	6.38E-08	1.2E-06	0.03
ROOTS								
Capric acid	229	15.7761	0.000092	0.000019	8.25E-07	1E-05	5.3E-06	0.04
4-guanidinobutyric acid	174	17.6937	0.003566	0.005331	6.19E-05	0.000318	0.0002	0.01
2-hydroxybutyric acid	147	7.7499	n/d	n/d	n/d	n/d	n/d	-

Table 1d. Sugars

	m/z	RT	Loadings		Control Average	2,5% PEG Average	5% PEG Average	P
			PC-1 (68%)	PC-2 (14%)				
LEAVES								
D-glucose	319	26.2202	0.731198	-0.104959	0.001403	0.007798	0.02154	0.01

D-lyxose	307	21.2187	0.001893	0.000188	3.36E-05	6.68E-05	8.5E-05	0.05
D-(+) trehalose	361	40.8432	0.035573	-0.015576	0.000242	0.001002	0.00125	0.02
Fructose	307	25.7157	0.296617	-0.368847	0.004408	0.01308	0.01293	0.01
Raffinose	87	49.0069	0.024816	0.018845	6.15E-05	0.000236	0.00078	0.02
Sedoheptulose anhydride monohydrate	204	27.8032	-0.004646	-0.181400	0.000586	0.002895	0.00121	0.02
Sophorose	319	41.6028	0.000078	0.000116	3.35E-06	4.68E-06	4.6E-06	0.85
Sucrose	361	39.0483	0.174925	-0.074017	0.001662	0.005294	0.00631	0.02
Talose	319	26.5018	-0.112775	0.202728	0.005084	0.000143	0.00025	0.00
ROOTS								
D-glucose	319	26.2202	-0.040919	0.782006	0.005498	0.0101	0.00194	0.24
D-lyxose	307	21.2187	0.000862	0.004194	3.5E-05	0.000128	6.5E-05	0.01
D-(+) trehalose	361	40.8432	0.112013	0.160751	1.56E-05	0.004449	0.00621	0.02
Fructose	307	25.7157	-0.092943	0.551804	0.006759	0.005579	0.00094	0.29
Raffinose	87	49.0069	0.000005	-0.000002	1.71E-07	5.14E-07	0	0.50
Sedoheptulose anhydride monohydrate	204	27.8032	-0.002582	0.030655	0.000125	0.000365	5.7E-05	0.33
Sophorose	319	41.6028	-0.000093	-0.000026	7.05E-06	0	0	0.01
Sucrose	361	39.0483	-0.138201	0.184455	0.009577	0.004935	0.00036	0.11
Talose	319	26.5018	-0.000183	-0.000256	2.59E-05	5.96E-06	4.8E-06	0.12

Table 1e. Polyols

	m/z	RT	Loadings		Control Average	2,5% PEG Average	5% PEG Average	P
			PC-1 (68%)	PC-2 (14%)				
LEAVES								
D-mannitol	319	26.8178	-0.000811	-0.005078	3.63E-06	7.44E-05	1.4E-05	0.08
D-sorbitol	319	27.0474	0.003639	-0.001708	1.55E-06	8.71E-05	0.00015	0.00
myo-inositol	432	29.7437	0.190297	-0.147333	0.002398	0.006161	0.00817	0.00
D-threitol	217	17.3574	0.001082	0.000871	2.66E-06	8.93E-06	3.8E-05	0.01
ROOTS								
D-mannitol	319	26.8178	0.976739	0.087747	0.003333	0.039295	0.04937	0.02
D-sorbitol	319	27.0474	0.000727	-0.002098	6.14E-06	3.37E-05	8.6E-05	0.00
myo-inositol	432	29.7437	0.002516	0.037217	0.000634	0.000622	0.00087	0.78
D-threitol	217	17.3574	0.007972	-0.002757	8.18E-06	7.62E-05	0.00046	0.06

Table 1f. Phosphates

	m/z	RT	Loadings		Control	2,5% PEG	5% PEG	P
			PC-1 (68%)	PC-2 (14%)	Average	Average	Average	
LEAVES								
D-glucose-6-phosphate	387	33.9694	0.000143	-0.006922	9.18E-05	0.000206	8.6E-05	0.01
Glycerol 1-phosphate	357	23.2557	0.001401	-0.005874	2.65E-05	0.000126	0.00014	0.14
Phosphoric acid	299	11.4449	0.105162	-0.168933	0.001059	0.003924	0.00414	0.03
O-phosphocolamine	299	23.5399	0.001582	-0.001420	1.49E-05	4.82E-05	6.8E-05	0.00
Ribulose-5-phosphate	357	30.4788	-0.000040	0.000041	2.4E-06	1.05E-06	5.2E-07	0.04
ROOTS								
D-glucose-6-phosphate	387	33.9694	-0.000618	0.000475	7E-05	2.74E-05	8.6E-06	0.01
Glycerol 1-phosphate	357	23.2557	0.015894	0.003408	5.65E-05	0.000699	0.00066	0.08
Phosphoric acid	299	11.4449	0.046304	0.103660	0.000852	0.003849	0.00255	0.02
O-phosphocolamine	299	23.5399	n/d	n/d	n/d	n/d	n/d	-
Ribulose-5-phosphate	357	30.4788	n/d	n/d	n/d	n/d	n/d	-

Table 1g. Polyamines

	m/z	RT	Loadings		Control	2,5% PEG	5% PEG	P
			PC-1 (68%)	PC-2 (14%)	Average	Average	Average	
LEAVES								
1,3-diaminopropane	174	20.3561	0.002079	-0.001720	2.73E-05	4.68E-05	9.3E-05	0.01
Putrescine	174	22.3494	-0.004973	-0.000420	0.000378	0.000281	0.0002	0.05
ROOTS								
1,3-diaminopropane	174	20.3561	-0.000106	-0.000029	7.87E-06	0	0	0.00
Putrescine	174	22.3494	-0.010487	-0.000437	0.000813	8.66E-05	2.4E-05	0.00

Table 1h. N-compounds

	m/z	RT	Loadings		Control	2,5% PEG	5% PEG	P
			PC-1 (68%)	PC-2 (14%)	Average	Average	Average	
LEAVES								
Acetyl-L-serine	174	14.4827	0.000224	0.000561	1.84E-06	3.93E-06	8E-06	0.44
Adenine	264	24.9660	0.002173	-0.001493	1.45E-05	4.58E-05	9.6E-05	0.00
Adenosine	236	38.4360	0.004016	-0.006006	4.11E-05	0.00018	0.00016	0.01
b-cyano-L-alanine	141	13.8776	0.009878	-0.014927	2.86E-05	0.000283	0.00037	0.01
1-aminocyclopropane-1-carboxylic acid	147	10.8258	0.000573	-0.000186	4.74E-06	9.97E-06	2.3E-05	0.01
5-aminovaleric acid	174	19.7646	-0.000133	0.000270	2E-05	6.64E-06	5.3E-06	0.15

Carbamic acid ethyl ester (urethane)	146	4.7187	0.005169	-0.001134	2.05E-05	0.000109	0.00019	0.00
D-panthenol	277	26.7673	-0.000016	-0.000497	1.54E-06	1.03E-05	1.6E-06	0.00
Gly-pro	174	27.4069	0.000064	-0.000014	3.14E-08	8.69E-07	2.7E-06	0.00
4-hydroxybenzyl cyanide	190	17.1553	0.000233	-0.000282	1.94E-06	6.9E-06	9.7E-06	0.01
Hypoxanthine	265	23.7548	-0.000017	0.000028	7.56E-07	0	0	0.01
N-acetyl-D-glucosamine	319	30.7496	0.000879	-0.002054	2.4E-05	6.75E-05	5.3E-05	0.07
N-acetyl-D-mannosamine	205	29.6268	n/d	n/d	n/d	n/d	n/d	-
N-methylglutamic acid	260	22.0275	0.003073	-0.001360	9.81E-07	5.69E-05	0.00012	0.00
1-methyl nicotinamide	179	16.2677	0.000151	-0.000134	1.09E-06	4.72E-06	6.1E-06	0.00
L-mimosine	188	9.3364	0.000679	-0.000748	1.33E-05	2.7E-05	2.7E-05	0.38
L-norleucine	158	11.3417	0.004475	-0.001549	1.4E-05	6.89E-05	0.00017	0.00
N,N-dimethylglycine	160	5.2735	0.000290	-0.000106	3.3E-06	8.04E-06	1.2E-05	0.04
Pantothenic acid	291	27.6738	0.001003	-0.001440	6.87E-06	4.78E-05	4E-05	0.01
Uracil	241	13.0040	n/d	n/d	n/d	n/d	n/d	-

ROOTS

Acetyl-L-serine	174	14.4827	-0.000042	-0.000008	3.12E-06	0	0	0.00
Adenine	264	24.9660	0.000232	0.000370	6.18E-06	2.08E-05	1.8E-05	0.01
Adenosine	236	38.4360	-0.000118	0.001161	5.78E-05	4.75E-05	2.3E-05	0.10
b-cyano-L-alanine	141	13.8776	0.000089	0.002606	1.72E-06	3.87E-05	7E-06	0.06
1-aminocyclopropane-1-carboxylic acid	147	10.8258	n/d	n/d	n/d	n/d	n/d	-
5-aminovaleric acid	174	19.7646	-0.000479	0.000061	4.66E-05	1.59E-05	0	0.00
Carbamic acid ethyl ester (urethane)	146	4.7187	n/d	n/d	n/d	n/d	n/d	-
D-panthenol	277	26.7673	n/d	n/d	n/d	n/d	n/d	-
Gly-pro	174	27.4069	0.000009	0.000018	0	1.16E-06	0	0.00
4-hydroxybenzyl cyanide	190	17.1553	0.000028	-0.000101	6.23E-07	1.54E-06	3E-06	0.48
Hypoxanthine	265	23.7548	-0.000013	0.000006	9.91E-07	0	0	0.10
N-acetyl-D-glucosamine	319	30.7496	0.001173	-0.000011	1.71E-05	3.47E-05	8.1E-05	0.02
N-acetyl-D-mannosamine	205	29.6268	0.000186	-0.000164	0	3.19E-05	7.5E-06	0.04
N-methylglutamic acid	260	22.0275	0.000093	0.000102	4.03E-07	4.2E-06	3.9E-06	0.20
1-methyl nicotinamide	179	16.2677	0.000124	0.000140	1.59E-06	7.36E-06	3.3E-06	0.42
L-mimosine	188	9.3364	0.000095	0.000150	1.88E-06	1.32E-05	4E-06	0.02
L-norleucine	158	11.3417	0.000218	0.001968	0.000139	0.000131	8.2E-05	0.71
N,N-dimethylglycine	160	5.2735	n/d	n/d	n/d	n/d	n/d	-
Pantothenic acid	291	27.6738	-0.000140	0.000225	1.23E-05	4.71E-06	0	0.01
Uracil	241	13.0040	0.001171	0.000204	9.61E-06	6.69E-05	7.6E-05	0.00

Table 1i. Other compounds

	m/z	RT	Loadings		Control	2,5% PEG	5% PEG	P
			PC-1 (68%)	PC-2 (14%)	Average	Average	Average	
LEAVES								
Acetol	217	19.9778	-0.000023	0.000037	9.93E-07	0	0	0.00
Arbutin	254	37.9619	0.000814	0.000501	1.87E-05	2.12E-05	4.1E-05	0.15
Galacturonic acid	333	24.5397	0.002229	-0.001768	1.75E-05	7.37E-05	8.5E-05	0.01
Isopropyl b-D-1-thiogalactopyranoside	217	27.2213	-0.001081	-0.010687	0.000246	0.000411	0.00018	0.01
2-keto-L-gulonic acid	204	24.4794	-0.001595	-0.015615	0	0.000219	3.8E-05	0.04
Ribonic acid-g-lactone	147	20.6655	-0.000028	0.000032	8.78E-07	3.71E-07	0	0.07
ROOTS								
Acetol	217	19.9778	0.000025	-0.000057	0	0	2.1E-06	0.21
Arbutin	254	37.9619	-0.000200	0.000158	1.54E-05	1.93E-06	4.7E-07	0.00
Galacturonic acid	333	24.5397	-0.000580	0.001859	3.94E-05	1.89E-05	0	0.05
Isopropyl b-D-1-thiogalactopyranoside	217	27.2213	-0.000354	0.001440	2.27E-05	1.22E-05	0	0.15
2-keto-L-gulonic acid	204	24.4794	n/d	n/d	n/d	n/d	n/d	-
Ribonic acid-g-lactone	147	20.6655	-0.000024	0.000004	1.76E-06	0	0	0.03

For each identified metabolite GC-MS characteristics (fragment mass, RI: Retention Index), loadings in principal components and average response (resulting from 4 biological replications) are presented. P value represents the level of significance (ANOVA)