

### IAA content (Figure S1)

Treatments	Time	repetition	IAA content ( $\mu\text{mol}\cdot\text{g}^{-1}\text{FW}$ )	mean	standard error	significant difference between treatments	significant difference between full data	
CK	24 h	1	0.169	0.16	0.01	b	B	
		2	0.157					
		3	0.144					
L		1	0.166	0.16	0.01	b	B	
		2	0.145					
		3	0.164					
L+S		1	0.185	0.18	0.00	a	A	
		2	0.171					
		3	0.178					
L+S+TIBA	48 h	1	0.134	0.13	0.00	c	C	
		2	0.136					
		3	0.132					
L+S+BBO		1	0.139	0.13	0.00	c	C	
		2	0.130					
		3	0.133					
CK		1	0.120	0.12	0.00	a	C	
		2	0.129					
		3	0.114					
L		1	0.086	0.09	0.00	c	E	
		2	0.099					
		3	0.092					
L+S		1	0.103	0.11	0.00	b	D	
		2	0.111					
		3	0.103					
L+S+TIBA		1	0.073	0.08	0.00	cd	F	
		2	0.080					
		3	0.080					
L+S+BBO		1	0.070	0.07	0.00	d	F	
		2	0.064					
		3	0.078					

### Sucrose content (Figure S2a)

Treatments	Time	repetition	Sucrose content (mg g <sup>-1</sup> FW)	mean	standard error	significant difference between treatments	significant difference between full data	
CK	24 h	1	1.5729	1.57	0.167	c	F	
		2	1.8622					
		3	1.2836					
L		1	2.3301	2.40	0.041	b	DE	
		2	2.4030					
		3	2.4729					
L+S	24 h	1	2.7654	2.79	0.012	a	C	
		2	2.7865					
		3	2.8077					
L+S+TIBA		1	2.5019	2.58	0.043	b	CD	
		2	2.5772					
		3	2.6525					
L+S+BBO		1	2.4636	2.40	0.034	b	DE	
		2	2.3487					
		3	2.3907					
CK	48 h	1	2.2761	2.27	0.008	d	E	
		2	2.2790					
		3	2.2542					
L		1	3.2919	3.08	0.121	b	B	
		2	3.0816					
		3	2.8714					
L+S		1	3.4346	3.40	0.034	a	A	
		2	3.4262					
		3	3.3277					
L+S+TIBA		1	2.6631	2.56	0.054	c	CD	
		2	2.5469					
		3	2.4790					
L+S+BBO		1	2.6815	2.72	0.025	c	C	
		2	2.7681					
		3	2.7248					

### Glucose content (Figure S2b)

Treatments	Time	repetition	Glucose content (mg g <sup>-1</sup> FW)	mean	standard error	significant difference between treatments	significant difference between full data	
CK	24 h	1	0.2781	0.31	0.02	c	D	
		2	0.3138					
		3	0.3494					
L		1	0.4188	0.42	0.01	ab	BC	
		2	0.4141					
		3	0.4329					
L+S		1	0.4768	0.46	0.01	a	B	
		2	0.4581					
		3	0.4394					
L+S+TIBA		1	0.4146	0.46	0.03	a	B	
		2	0.4589					
		3	0.5032					
L+S+BBO		1	0.4018	0.40	0.00	b	C	
		2	0.3932					
		3	0.3929					
CK	48 h	1	0.4069	0.40	0.01	c	C	
		2	0.3758					
		3	0.4058					
L		1	0.5474	0.51	0.02	b	A	
		2	0.4925					
		3	0.5047					
L+S		1	0.5470	0.55	0.00	a	A	
		2	0.5497					
		3	0.5444					
L+S+TIBA		1	0.5383	0.53	0.01	ab	A	
		2	0.5206					
		3	0.5277					
L+S+BBO		1	0.4885	0.51	0.01	b	A	
		2	0.5242					
		3	0.5063					

### Fructose content (Figure S2c)

Treatments	Time	repetitio n	Fructose content (mg g <sup>-1</sup> FW)	mean	standar d error	significant difference between treatments	significant difference between full data	
CK	24 h	1	0.3141	0.32	0.00	e	G	
		2	0.3178					
		3	0.3215					
L		1	0.3572	0.36	0.00	d	F	
		2	0.3686					
		3	0.3636					
L+S		1	0.4175	0.42	0.00	c	E	
		2	0.4306					
		3	0.4203					
L+S+TIBA		1	0.4929	0.49	0.01	a	C	
		2	0.5084					
		3	0.4774					
L+S+BBO		1	0.4679	0.46	0.00	b	D	
		2	0.4600					
		3	0.4549					
CK	48 h	1	0.4596	0.45	0.01	c	D	
		2	0.4343					
		3	0.4469					
L		1	0.5805	0.59	0.00	b	B	
		2	0.5907					
		3	0.5911					
L+S		1	0.6540	0.65	0.00	a	A	
		2	0.6453					
		3	0.6432					
L+S+TIBA		1	0.5996	0.60	0.00	b	B	
		2	0.5901					
		3	0.5956					
L+S+BBO		1	0.5889	0.60	0.00	b	B	
		2	0.5962					
		3	0.6035					

### Sorbitol content (Figure S2d)

Treatments	Time	repetition	Sorbitol content (mg g <sup>-1</sup> FW)	mean	standard error	significant difference between treatments	significant difference between full data	
CK	24 h	1	3.3054	3.43	0.07	d	G	
		2	3.4252					
		3	3.5450					
L		1	4.0592	4.06	0.01	c	EF	
		2	4.0397					
		3	4.0787					
L+S		1	4.6826	4.68	0.01	a	D	
		2	4.6979					
		3	4.6672					
L+S+TIBA		1	3.1003	3.28	0.09	d	G	
		2	3.3427					
		3	3.3834					
L+S+BBO		1	4.2597	4.20	0.04	b	E	
		2	4.2316					
		3	4.1203					
CK	48 h	1	3.8458	3.85	0.01	d	F	
		2	3.8666					
		3	3.8250					
L		1	5.5330	5.50	0.02	b	B	
		2	5.4624					
		3	5.4977					
L+S		1	5.8980	5.90	0.03	a	A	
		2	5.8436					
		3	5.9523					
L+S+TIBA		1	5.4538	5.13	0.19	c	C	
		2	4.8115					
		3	5.1326					
L+S+BBO		1	4.7816	4.97	0.10	c	C	
		2	5.1257					
		3	5.0063					

### MDA content (Figure S3)

Treatments	Time	repetition	MDA content (nmol g <sup>-1</sup> Fw)	mean	standard error	significant difference between treatments	significant difference between full data	
CK	24h	1	31.6994	30.46	0.80	b	D	
		2	30.7320					
		3	28.9522					
L		1	38.4904	39.14	1.08	a	BC	
		2	41.2373					
		3	37.6852					
L+S		1	35.3871	30.21	2.59	b	D	
		2	27.6763					
		3	27.5762					
L+S+TIBA		1	37.3683	36.71	0.38	a	C	
		2	36.7091					
		3	36.0500					
L+S+BBO		1	37.0391	36.99	0.16	a	C	
		2	36.6850					
		3	37.2341					
CK	48h	1	36.4390	35.49	1.52	b	C	
		2	32.5010					
		3	37.5192					
L		1	44.5029	46.22	1.64	a	A	
		2	49.5090					
		3	44.6569					
L+S		1	37.1707	37.23	0.04	b	C	
		2	37.2320					
		3	37.2934					
L+S+TIBA		1	42.6357	42.64	0.92	a	AB	
		2	41.0374					
		3	44.2340					
L+S+BBO		1	44.1082	44.79	0.80	a	A	
		2	46.3848					
		3	43.8679					

### O<sub>2</sub><sup>·-</sup> content (Figure S4a)

Treatments	Time	repetition	O <sub>2</sub> <sup>·-</sup> content ( μg·g <sup>-1</sup> Fw)	mean	standard error	significant difference between treatments	significant difference between full data	
CK	24h	1	9.202	9.38	0.10	c	E	
		2	9.565					
		3	9.384					
L		1	15.046	14.16	0.45	a	B	
		2	13.577					
		3	13.847					
L+S		1	9.755	9.57	0.11	c	E	
		2	9.572					
		3	9.389					
L+S+TIBA		1	14.846	14.31	0.31	a	B	
		2	14.305					
		3	13.764					
L+S+BBO		1	11.194	11.49	0.30	b	D	
		2	11.192					
		3	12.085					
CK	48h	1	9.601	9.60	0.19	d	E	
		2	9.931					
		3	9.272					
L		1	16.716	16.28	0.25	a	A	
		2	16.280					
		3	15.845					
L+S		1	9.990	9.91	0.23	d	E	
		2	10.262					
		3	9.484					
L+S+TIBA		1	12.576	12.29	0.18	b	C	
		2	12.332					
		3	11.953					
L+S+BBO		1	11.624	11.62	0.12	c	CD	
		2	11.412					
		3	11.835					

### H<sub>2</sub>O<sub>2</sub> content (Figure S4c)

Treatments	Time	repetition	H <sub>2</sub> O <sub>2</sub> ( mmol g <sup>-1</sup> Fw)	mean	standard error	significant difference between treatments	significant difference between full data	
CK	24 h	1	11.203	11.26	0.12	d	G	
		2	11.489					
		3	11.084					
L		1	16.480	16.60	0.07	a	A	
		2	16.598					
		3	16.716					
L+S		1	13.447	13.71	0.21	c	E	
		2	14.125					
		3	13.556					
L+S+TIBA		1	15.098	15.11	0.01	b	C	
		2	15.124					
		3	15.111					
L+S+BBO		1	15.148	15.15	0.02	b	C	
		2	15.179					
		3	15.117					
CK	48 h	1	12.993	12.92	0.04	d	F	
		2	12.853					
		3	12.923					
L		1	15.722	15.41	0.18	b	C	
		2	15.098					
		3	15.410					
L+S		1	14.125	14.18	0.08	c	D	
		2	14.087					
		3	14.337					
L+S+TIBA		1	16.442	16.12	0.36	a	B	
		2	15.407					
		3	16.507					
L+S+BBO		1	15.573	15.30	0.16	b	C	
		2	15.029					
		3	15.301					

### SOD activity (Figure S5a)

Treatments	Time	repetition	SOD activity ( $\mu\text{mol g}^{-1} \text{min}^{-1}$ Fw )	mean	standard error	significant difference between treatments	significant difference between full data	
CK	24h	1	5.74	5.99	0.34	c	EF	
		2	5.57					
		3	6.67					
L		1	6.61	7.10	0.27	b	BCD	
		2	7.13					
		3	7.56					
L+S		1	8.99	8.99	0.08	a	A	
		2	9.13					
		3	8.85					
L+S+TIBA		1	7.26	6.89	0.42	b	CD	
		2	7.36					
		3	6.04					
L+S+BBO		1	7.33	7.30	0.02	b	BC	
		2	7.27					
		3	7.30					
CK	48h	1	6.37	6.58	0.12	b	DE	
		2	6.58					
		3	6.78					
L		1	5.31	5.63	0.21	d	F	
		2	5.56					
		3	6.02					
L+S		1	7.73	7.63	0.05	a	B	
		2	7.54					
		3	7.63					
L+S+TIBA		1	6.09	6.08	0.17	cd	EF	
		2	5.78					
		3	6.38					
L+S+BBO		1	6.42	6.20	0.12	bc	EF	
		2	6.20					
		3	5.99					

### POD activity (Figure S5b)

Treatments	Time	repetition	POD activity ( $\mu\text{mol}\cdot\text{g}^{-1}\cdot\text{min}^{-1}$ Fw)	mean	standard error	significant difference between treatments	significant difference between full data	
CK	24h	1	41.04	39.84	0.67	b	DE	
		2	38.73					
		3	39.76					
L		1	38.67	39.30	0.36	b	E	
		2	39.30					
		3	39.93					
L+S		1	43.89	43.51	0.21	a	B	
		2	43.47					
		3	43.17					
L+S+TIBA		1	38.29	39.20	0.46	b	E	
		2	39.61					
		3	39.71					
L+S+BBO		1	35.46	35.75	0.46	c	F	
		2	36.65					
		3	35.13					
CK	48h	1	42.50	41.99	0.63	b	BC	
		2	42.72					
		3	40.74					
L		1	40.33	39.81	0.30	c	DE	
		2	39.81					
		3	39.30					
L+S		1	47.93	47.18	0.38	a	A	
		2	46.86					
		3	46.75					
L+S+TIBA		1	41.75	41.54	0.27	bc	CD	
		2	41.00					
		3	41.88					
L+S+BBO		1	42.15	40.42	1.22	bc	CDE	
		2	41.05					
		3	38.06					

### CAT activity (Figure S5c)

Treatments	Time	repetition	CAT activity ( $\mu\text{mol}\cdot\text{g}^{-1}\cdot\text{min}^{-1}$ Fw)	mean	standard error	significant difference between treatments	significant difference between full data	
CK	24h	1	0.57	0.62	0.03	c	F	
		2	0.64					
		3	0.65					
L		1	0.92	0.92	0.01	a	B	
		2	0.93					
		3	0.91					
L+S		1	0.98	0.97	0.01	a	A	
		2	0.97					
		3	0.96					
L+S+TIBA		1	0.84	0.79	0.03	b	E	
		2	0.77					
		3	0.76					
L+S+BBO		1	0.83	0.83	0.01	b	DE	
		2	0.84					
		3	0.82					
CK	48h	1	0.59	0.58	0.01	d	F	
		2	0.58					
		3	0.57					
L		1	0.91	0.91	0.01	b	BC	
		2	0.93					
		3	0.89					
L+S		1	1.00	1.01	0.03	a	A	
		2	1.06					
		3	0.97					
L+S+TIBA		1	0.84	0.85	0.01	c	DE	
		2	0.85					
		3	0.86					
L+S+BBO		1	0.89	0.87	0.02	bc	CD	
		2	0.88					
		3	0.84					

### APX activity (Figure S6a)

Treatments	Time	repetition	APXactivity ( $\mu\text{mol}\cdot\text{g}^{-1}\cdot\text{min}^{-1}$ Fw)	mean	standard error	significant difference between treatments	significant difference between full data	
CK	24 h	1	15.32	15.11	0.12	b	B	
		2	15.11					
		3	14.90					
L		1	16.14	16.09	0.05	a	A	
		2	15.99					
		3	16.14					
L+S		1	16.02	16.08	0.04	a	A	
		2	16.06					
		3	16.15					
L+S+TIBA		1	14.70	15.07	0.21	b	B	
		2	15.43					
		3	15.07					
L+S+BBO		1	14.37	14.37	0.37	c	C	
		2	15.01					
		3	13.74					
CK	48 h	1	14.73	14.99	0.13	b	BC	
		2	15.08					
		3	15.15					
L		1	15.72	15.99	0.19	a	A	
		2	15.88					
		3	16.36					
L+S		1	16.18	16.18	0.04	a	A	
		2	16.25					
		3	16.10					
L+S+TIBA		1	13.10	13.03	0.29	c	D	
		2	13.49					
		3	12.50					
L+S+BBO		1	10.82	11.19	0.35	d	E	
		2	11.89					
		3	10.87					

### MDHAR activity (Figure S6b)

Treatments	Time	repetition	MDHAR activity (nmol g <sup>-1</sup> min <sup>-1</sup> Fw)	mean	standard error	significant difference between treatments	significant difference between full data	
CK	24h	1	116.30	121.04	3.60	c	D	
		2	128.10					
		3	118.72					
L		1	142.71	134.19	4.26	b	C	
		2	130.10					
		3	129.77					
L+S		1	179.86	176.79	1.69	a	B	
		2	176.46					
		3	174.04					
L+S+TIBA		1	134.19	134.19	2.70	b	C	
		2	129.51					
		3	138.86					
L+S+BBO		1	131.13	136.20	2.57	b	C	
		2	139.49					
		3	137.98					
CK	48h	1	128.60	128.60	0.43	b	C	
		2	129.34					
		3	127.85					
L		1	116.60	117.18	0.33	c	D	
		2	117.73					
		3	117.20					
L+S		1	190.32	195.97	3.35	a	A	
		2	195.67					
		3	201.92					
L+S+TIBA		1	120.84	119.13	0.99	c	D	
		2	117.42					
		3	119.13					
L+S+BBO		1	115.64	118.72	1.78	c	D	
		2	118.72					
		3	121.80					

### DHAR activity (Figure S6c)

Treatments	Time	repetition	DHAR activity ( $\mu\text{mol}\cdot\text{g}^{-1}$ min $^{-1}$ Fw)	mean	standard error	significant difference between treatments	significant difference between full data	
CK	24h	1	0.19	0.18	0.01	b	B	
		2	0.16					
		3	0.19					
L		1	0.15	0.14	0.01	c	D	
		2	0.12					
		3	0.16					
L+S		1	0.22	0.22	0.00	a	A	
		2	0.22					
		3	0.22					
L+S+TIBA		1	0.19	0.19	0.00	b	B	
		2	0.19					
		3	0.19					
L+S+BBO		1	0.16	0.18	0.01	b	B	
		2	0.19					
		3	0.18					
CK	48h	1	0.14	0.15	0.01	b	CD	
		2	0.15					
		3	0.16					
L		1	0.16	0.17	0.01	b	BC	
		2	0.17					
		3	0.19					
L+S		1	0.21	0.22	0.01	a	A	
		2	0.22					
		3	0.23					
L+S+TIBA		1	0.07	0.09	0.01	c	E	
		2	0.11					
		3	0.09					
L+S+BBO		1	0.10	0.11	0.01	c	E	
		2	0.12					
		3	0.12					

### GR activity (Figure S6d)

Treatments	Time	repetition	GR activity (nmol • g <sup>-1</sup> • min <sup>-1</sup> Fw)	mean	standard error	significant difference between treatments	significant difference between full data	
CK	24h	1	102.21	103.24	0.61	b	A	
		2	103.19					
		3	104.33					
L		1	93.39	92.69	0.40	c	B	
		2	92.69					
		3	91.99					
L+S		1	114.54	110.90	2.91	a	A	
		2	113.01					
		3	105.15					
L+S+TIBA		1	74.63	76.96	1.35	d	C	
		2	79.29					
		3	76.96					
L+S+BBO		1	61.27	64.44	1.60	e	D	
		2	66.38					
		3	65.66					
CK	48h	1	87.08	91.88	3.20	b	B	
		2	97.94					
		3	90.63					
L		1	59.43	63.84	4.13	c	D	
		2	60.00					
		3	72.09					
L+S		1	108.34	108.62	0.16	a	A	
		2	108.62					
		3	108.89					
L+S+TIBA		1	55.22	56.39	2.00	c	D	
		2	53.66					
		3	60.28					
L+S+BBO		1	71.14	60.56	5.29	c	D	
		2	54.89					
		3	55.66					

### AsA content (Figure S7a)

Treatments	Time	repetition	AsA content ( mg g-1 Fw)	mean	standard error	significant difference between treatments	significant difference between full data	
CK	24h	1	0.538	0.55	0.02	d	F	
		2	0.592					
		3	0.517					
L		1	0.654	0.64	0.01	c	E	
		2	0.643					
		3	0.631					
L+S		1	0.952	0.92	0.02	a	B	
		2	0.898					
		3	0.906					
L+S+TIBA		1	0.645	0.70	0.04	c	DE	
		2	0.780					
		3	0.680					
L+S+BBO		1	0.775	0.78	0.00	b	CD	
		2	0.777					
		3	0.773					
CK	48h	1	0.768	0.72	0.03	b	DE	
		2	0.716					
		3	0.664					
L		1	0.910	0.86	0.03	b	B	
		2	0.845					
		3	0.821					
L+S		1	1.007	1.01	0.01	a	A	
		2	1.024					
		3	0.984					
L+S+TIBA		1	0.806	0.84	0.03	b	BC	
		2	0.905					
		3	0.816					
L+S+BBO		1	0.867	0.86	0.04	b	B	
		2	0.792					
		3	0.927					

### DHA content (Figure S7b)

Treatments	Time	repetition	DHA content (mg g <sup>-1</sup> Fw)	mean	standard error	significant difference between treatments	significant difference between full data	
CK	24h	1	0.657	0.61	0.03	a	C	
		2	0.546					
		3	0.635					
L		1	0.621	0.66	0.02	a	C	
		2	0.701					
		3	0.661					
L+S		1	0.368	0.42	0.04	b	D	
		2	0.389					
		3	0.507					
L+S+TIBA		1	0.670	0.68	0.02	a	C	
		2	0.649					
		3	0.718					
L+S+BBO		1	0.582	0.64	0.04	a	C	
		2	0.632					
		3	0.712					
CK	48h	1	0.605	0.61	0.00	c	C	
		2	0.614					
		3	0.610					
L		1	0.665	0.71	0.05	bc	BC	
		2	0.655					
		3	0.817					
L+S		1	0.446	0.44	0.00	d	D	
		2	0.433					
		3	0.439					
L+S+TIBA		1	0.885	0.84	0.02	a	A	
		2	0.799					
		3	0.842					
L+S+BBO		1	0.735	0.78	0.04	ab	AB	
		2	0.855					
		3	0.747					

### AsA/DHA (Figure S7c)

Treatments	Time	repetition	AsA/DHA	mean	standard error	significant difference between treatments	significant difference between full data	
CK	24h	1	0.819	0.82	0.00	c	D	
		2	0.817					
		3	0.815					
L		1	1.053	1.03	0.06	b	CD	
		2	0.917					
		3	1.133					
L+S		1	2.588	2.45	0.08	a	A	
		2	2.307					
		3	2.448					
L+S+TIBA		1	0.963	1.04	0.08	b	CD	
		2	1.202					
		3	0.947					
L+S+BBO		1	1.379	1.23	0.08	b	C	
		2	1.229					
		3	1.086					
CK	48h	1	1.537	1.51	0.10	b	B	
		2	1.675					
		3	1.323					
L		1	1.368	1.22	0.11	c	C	
		2	1.289					
		3	1.005					
L+S		1	2.259	2.31	0.03	a	A	
		2	2.365					
		3	2.312					
L+S+TIBA		1	0.911	1.00	0.07	c	CD	
		2	1.133					
		3	0.969					
L+S+BBO		1	1.179	1.12	0.10	c	C	
		2	0.925					
		3	1.242					

### GSH content (Figure S7d)

Treatments	Time	repetition	GSH ( $\mu\text{mol}\cdot\text{g}^{-1}$ Fw)	mean	standard error	significant difference between treatments	significant difference between full data	
CK	24h	1	32.63	31.99	0.37	b	B	
		2	31.34					
		3	31.99					
L		1	32.46	32.46	0.00	b	B	
		2	32.46					
		3	32.46					
L+S		1	42.77	39.76	1.51	a	A	
		2	38.02					
		3	38.50					
L+S+TIBA		1	28.66	31.43	1.40	b	B	
		2	32.45					
		3	33.18					
L+S+BBO		1	25.32	26.15	0.99	c	D	
		2	25.01					
		3	28.11					
CK	48h	1	26.30	30.00	2.38	bc	BC	
		2	29.26					
		3	34.43					
L		1	30.75	31.84	0.63	b	B	
		2	31.84					
		3	32.93					
L+S		1	41.75	41.05	1.20	a	A	
		2	38.72					
		3	42.68					
L+S+TIBA		1	25.89	27.30	0.81	cd	CD	
		2	27.30					
		3	28.70					
L+S+BBO		1	25.02	24.69	0.16	d	D	
		2	24.52					
		3	24.54					

### GSSG content (Figure S7e)

Treatments	Time	repetition	GSSG ( $\mu\text{mol} \cdot \text{g}^{-1} \cdot \text{Fw}$ )	mean	standard error	significant difference between treatments	significant difference between full data	
CK	24h	1	11.66	11.44	0.13	c	E	
		2	11.44					
		3	11.21					
L		1	14.11	14.11	0.10	a	A	
		2	13.93					
		3	14.28					
L+S		1	11.30	11.55	0.14	c	E	
		2	11.80					
		3	11.55					
L+S+TIBA		1	12.09	12.67	0.62	b	CD	
		2	12.01					
		3	13.92					
L+S+BBO		1	12.80	12.80	0.02	b	CD	
		2	12.76					
		3	12.83					
CK	48h	1	12.19	12.21	0.06	c	DE	
		2	12.32					
		3	12.11					
L		1	14.30	14.26	0.03	a	A	
		2	14.26					
		3	14.21					
L+S		1	13.56	13.32	0.17	b	BC	
		2	12.98					
		3	13.41					
L+S+TIBA		1	13.75	13.98	0.22	a	AB	
		2	14.42					
		3	13.77					
L+S+BBO		1	14.11	14.32	0.33	a	A	
		2	13.88					
		3	14.98					

### GSH/GSSG (Figure S7f)

Treatments	Time	repetition	GSH/GSSG	mean	standard error	significan t difference between treatments	significan t difference between full data	
CK	24h	1	2.798	2.84	0.02	b	BC	
		2	2.881					
		3	2.839					
		1	2.166	2.17	0.09	cd	F	
		2	2.330					
		3	2.003					
L+S		1	3.784	3.45	0.17	a	A	
		2	3.223					
		3	3.333					
L+S+TIB A	48h	1	2.370	2.49	0.11	c	DE	
		2	2.702					
		3	2.384					
L+S+BBO		1	2.076	2.08	0.07	d	F	
		2	1.960					
		3	2.190					
CK		1	2.610	2.61	0.14	b	CD	
		2	2.376					
		3	2.844					
L		1	2.151	2.23	0.05	c	EF	
		2	2.234					
		3	2.317					
L+S		1	3.080	3.08	0.06	a	B	
		2	2.983					
		3	3.182					
L+S+TIB A		1	1.883	2.06	0.10	c	F	
		2	2.225					
		3	2.084					
L+S+BBO		1	1.773	1.73	0.04	d	G	
		2	1.767					
		3	1.639					

### G6PDH activity (Figure S8)

Treatments	Time	repetition	G6PDH activity ( $\mu\text{mol g}^{-1} \text{min}^{-1}$ Fw )	mean	standard error	significant difference between treatments	significant difference between full data	
CK	24h	1	85.64	90.21	5.48	d	G	
		2	83.87					
		3	101.11					
L		1	122.83	122.33	1.09	bc	D	
		2	120.24					
		3	123.93					
L+S		1	158.24	151.54	3.46	a	B	
		2	146.69					
		3	149.69					
L+S+TIBA		1	84.42	96.12	5.94	d	FG	
		2	103.76					
		3	100.17					
L+S+BBO		1	116.76	117.23	0.37	c	DE	
		2	117.97					
		3	116.97					
CK	48h	1	81.94	88.00	5.95	d	G	
		2	82.17					
		3	99.89					
L		1	141.07	134.79	3.14	b	C	
		2	131.35					
		3	131.96					
L+S		1	172.76	172.43	0.79	a	A	
		2	173.60					
		3	170.94					
L+S+TIBA		1	100.55	108.03	6.87	c	EF	
		2	101.79					
		3	121.76					
L+S+BBO		1	121.99	116.80	2.60	c	DE	
		2	114.20					
		3	114.20					

### NADPH content (Figure S9a)

Treatments	Time	repetition	NADPH content ( $\mu\text{mol}\cdot\text{g}^{-1}\text{Fw}$ )	mean	standard error	significant difference between treatments	significant difference between full data	
CK	24h	1	0.67	0.65	0.03	bc	CDE	
		2	0.68					
		3	0.59					
L		1	0.69	0.70	0.02	b	BC	
		2	0.67					
		3	0.74					
L+S		1	0.87	0.92	0.03	a	A	
		2	0.92					
		3	0.97					
L+S+TIBA		1	0.60	0.62	0.01	c	E	
		2	0.62					
		3	0.64					
L+S+BBO		1	0.68	0.68	0.00	b	CD	
		2	0.69					
		3	0.68					
CK	48h	1	0.70	0.70	0.00	b	CD	
		2	0.70					
		3	0.69					
L		1	0.71	0.70	0.02	b	BC	
		2	0.67					
		3	0.72					
L+S		1	0.77	0.75	0.01	a	B	
		2	0.75					
		3	0.73					
L+S+TIBA		1	0.63	0.65	0.01	c	CDE	
		2	0.65					
		3	0.67					
L+S+BBO		1	0.64	0.64	0.00	c	DE	
		2	0.65					
		3	0.64					

### NADPH/NADP<sup>+</sup> (Figure S9b)

Treatments	Time	repetition	NADPH/NADP <sup>+</sup>	mean	standard error	significant difference between treatments	significant difference between full data	
CK	24h	1	2.286	2.05	0.12	bc	BC	
		2	1.921					
		3	1.934					
L		1	2.281	2.20	0.04	b	B	
		2	2.130					
		3	2.201					
L+S		1	2.842	2.87	0.10	a	A	
		2	2.714					
		3	3.054					
L+S+TIBA		1	1.809	1.79	0.01	d	CD	
		2	1.785					
		3	1.761					
L+S+BBO		1	2.000	1.89	0.09	cd	CD	
		2	1.951					
		3	1.714					
CK	48h	1	2.372	2.18	0.12	a	B	
		2	1.949					
		3	2.212					
L		1	1.711	1.79	0.11	b	CD	
		2	1.653					
		3	1.995					
L+S		1	2.302	2.25	0.04	a	B	
		2	2.285					
		3	2.176					
L+S+TIBA		1	1.656	1.66	0.07	b	D	
		2	1.773					
		3	1.547					
L+S+BBO		1	1.650	1.65	0.07	b	D	
		2	1.779					
		3	1.525					