

Additional file 2: Relative content of primary metabolites obtained from fruit pulp GC-MS analysis and (\log_{10}) fold change in the autopolyploid lines. Mass spectral searching utilized the algorithm incorporated in the Xcalibur® data system and finally normalized by the internal standard ribitol. MP = main product, BP = by product. Asterisks represent significant changes in metabolite content ($P < 0.01$) according to ANOVA. Fold change represents the (\log_{10}) change in the relative content of each metabolite between the autopolyploid lines and their respective control lines

Metabolites	2n		4n		Significance	(\log_{10}) 4n fold change	3n		6n		Significance	(\log_{10}) 6n fold change
	Average	\pm SE	Average	\pm SE			Average	\pm SE	Average	\pm SE		
Sugars	Glucose MP	3601.49	188.77	3575.94	383.46	0.00	27951.79	1519.40	14429.41	3388.09	*	-0.29
	Fructose MP	1281.93	78.08	1180.14	166.83	0.00	22151.62	1260.36	12547.01	2712.50	*	-0.25
	Fructose BP	776.89	47.46	719.29	99.92	0.00	14795.77	884.84	8230.16	2006.52	*	-0.25
	myo-Inositol	621.10	37.09	582.53	61.82	0.00	19140.14	975.76	18915.80	1580.29		-0.01
	Glucose BP	492.65	26.65	453.92	53.56	-0.01	6323.09	444.76	2688.23	799.67	*	-0.37
	Glucopyranose	12.76	0.59	11.17	1.32	-0.03	126.35	7.47	60.48	14.57	*	-0.32
	Sucrose	9.03	0.86	8.42	1.42	0.00	112.51	16.08	83.07	8.84		-0.13
	Arabinose MP	1.78	0.18	1.68	0.21	0.02	4.75	0.55	2.86	0.67		-0.22
	Arabinose BP	1.09	0.08	0.99	0.09	-0.01	1.30	0.20	1.01	0.28		-0.11
	Galacturonate	0.99	0.06	0.93	0.08	-0.01	1.85	0.16	1.19	0.51		-0.19
Amino acids	Threonate	0.86	0.08	0.70	0.05	-0.07	6.94	0.45	3.63	0.46	*	-0.28
	Alanine	0.03	0.00	0.03	0.00	0.07	0.07	0.02	0.13	0.06		0.26
	Proline	3.39	0.59	3.14	0.71	-0.01	128.40	28.06	429.05	80.07	*	0.52
	Pyroglutamate	1.28	0.09	1.24	0.15	0.01	21.96	1.80	70.66	9.26	*	0.51
	Aspartate	0.38	0.03	0.34	0.05	-0.02	3.95	0.12	6.93	0.23	*	0.24
	Valine	0.12	0.02	0.16	0.03	0.16	0.98	0.04	2.99	0.05	*	0.48
	Tryptophan	1.37	0.05	1.31	0.03	0.02	1.40	0.05	4.92	0.10	*	0.55
	Tryosine	0.08	0.02	0.10	0.02	0.13	0.08	0.00	0.11	0.00	*	0.15
TCA cycle	Serine	0.31	0.01	0.30	0.01	0.02	1.16	0.19	1.90	0.25		0.21
	Malate	67.02	4.63	76.32	8.47	0.08	47.75	3.68	272.67	43.84	*	0.76
	Fumarate	5.61	0.58	6.03	0.88	0.07	1.10	0.13	2.50	0.43	*	0.36
	Citrate	3.39	0.32	4.40	0.66	0.15	706.47	70.47	2733.99	384.40	*	0.59
	cis-Aconitate	0.11	0.01	0.12	0.02	0.11	2.32	0.14	7.00	0.89	*	0.48
Organic acids	Succinate	0.21	0.01	0.19	0.01	0.00	0.08	0.00	0.15	0.04	*	0.30
	Maleic acid	25.01	1.26	26.30	2.20	0.05	44.66	2.21	86.70	13.57	*	0.29
	Phosphoric acid	9.81	1.31	9.23	1.48	-0.01	88.44	8.19	116.97	13.70		0.12
	Gluconic acid	1.05	0.08	1.07	0.11	0.04	1.32	0.08	5.65	1.71	*	0.63
	Itaconic acid	0.17	0.03	0.16	0.02	-0.01	0.68	0.06	1.38	0.14	*	0.31
	Lactic acid	0.16	0.01	0.19	0.03	0.11	0.89	0.07	1.74	0.25	*	0.29
	Nicotinic acid	0.10	0.00	0.12	0.01	0.07	1.05	0.04	1.41	0.13	*	0.13
Saturated fatty acids	Caffeic acid	0.01	0.00	0.01	0.00	0.06	0.38	0.02	0.44	0.09		0.06
	Nonanoic acid (C9:0)	0.29	0.03	0.25	0.03	0.02	1.03	0.04	1.30	0.10		0.08
	Decanoic acid (C10:0)	0.04	0.00	0.03	0.00	-0.03	0.21	0.02	0.20	0.03		0.10
	Hexadecanoic acid (C16:0)	0.50	0.02	0.50	0.03	0.04	2.92	0.10	3.66	0.48		0.10
	Octadecanoic acid (C18:0)	1.23	0.04	1.18	0.05	-0.02	4.64	0.23	5.56	0.48		0.00