

**Table S2.** The complete list of candidate postzygotic single-nucleotide mutations identified by our computational pipeline

<b>Individual</b>	<b>ID</b>	<b>Position</b>	<b>Ref base</b>	<b>Alt base</b>	<b>P(mosaic)</b>	<b>Validated?</b>
ACC1-II-1	B1	15:80878576	C	T	1.00	True
ACC1-II-1	B2	4:165099831	C	T	1.00	True
ACC1-II-1	B3	6:85605629	A	T	1.00	True
ACC1-II-1	B4	10:115352263	C	T	0.96	False
ACC1-II-1	B5	6:154692299	G	C	0.95	True
ACC1-II-1	B6	18:31333885	A	G	0.41	False
ACC1-II-1	B7	15:45732364	G	A	0.21	False
ACC1-II-1	B8	6:76955874	G	T	0.17	False
ACC1-II-1	B9	3:158917393	C	T	0.16	False
ACC1-II-1	B10	8:11688362	A	G	0.11	False
ACC1-II-1	B11	19:4032107	G	A	0.08	False
ACC1-II-1	B12	7:19343958	C	T	0.08	False
ACC1-II-1	B13	18:70512197	T	G	0.07	True
ACC1-II-1	B14	7:147539364	G	A	0.06	False
ACC1-II-1	B15	6:123036854	A	T	0.05	False
DS1-II-2	F1	17:52287119	G	T	1.00	True
DS1-II-2	F2	19:55529705	C	T	1.00	True
DS1-II-2	F3	19:56343191	T	C	1.00	True
DS1-II-2	F4	20:4339054	G	A	1.00	False
DS1-II-2	F5	4:190179317	C	T	1.00	False
DS1-II-2	F6	14:71380614	T	G	0.98	True
DS1-II-2	F7	16:84073033	A	C	0.95	True
DS1-II-2	F8	7:132334892	C	T	0.13	False
DS1-II-2	F9	13:79932615	C	A	0.12	True
DS1-II-2	F10	2:166848782	G	C	0.09	True
DS2-I-1	X1	1:72008400	A	G	1.00	True
DS2-I-1	X2	16:64312186	C	T	1.00	True
DS2-I-1	X3	X:68611473	C	A	1.00	True

DS2-I-1	X4	20:60500224	C	T	1.00	False
DS2-I-1	X5	22:18619901	G	A	0.98	False
DS2-I-1	X6	2:166854673	G	T	0.95	True
DS2-I-1	X7	6:35755658	A	G	0.89	False
DS2-I-1	X8	13:89662020	A	C	0.69	True
DS2-I-1	X9	2:79895035	A	G	0.44	False
DS2-I-1	X10	4:3472325	T	C	0.12	False
DS2-I-1	X11	16:48880245	T	C	0.09	False
DS2-I-1	X12	18:11785347	G	A	0.06	False
DS2-I-1	X13	17:59415198	G	T	0.06	False

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