

Suppl. Table 4. Indel details

MA Lines	Chromosome	Position	Reference	Indels	SSR* motif
Hv1	NC_013967.1	315827	TACG	T	-
Hv7	NC_013967.1	709482	CT	C	-
Hv17	NC_013966.1	468209	CGATGATGG	C	-
Hv18	NC_013967.1	683949	GGC	G	3g
Hv18	NC_013967.1	2302447	CA	C	3a
Hv19	NC_013966.1	451640	AGCCCGGCA	A	-
Hv20	NC_013966.1	112265	AC	A	-
Hv25	NC_013967.1	1924512	CGGCGGT	C	-
Hv33	NC_013967.1	1169790	ACGAACCGC	A	-
Hv33	NC_013966.1	187714	AC	A	-
Hv34	NC_013967.1	2772433	TTCCCC	T	-
Hv35	NC_013967.1	2218278	TCA	T	-
Hv42	NC_013968.1	43853	CA	C	-
Hv44	NC_013967.1	1067846	G	GCCA	4gcc
Hv45	NC_013967.1	1289230	CCCGGCA	C	4c
Hv48	NC_013966.1	163783	GT	G	-
Hv87	NC_013967.1	1453364	C	CGTGGAGGC	-
Hv87	NC_013967.1	1617094	CA	C	-
Hv90	NC_013967.1	2121583	CT	C	-

* SSR is the simple sequence repeat size and motif where the insertion/deletion is found (minimum 3 repeats).