

Table S1 WS220 Hawaiian Variants filtered to assist in CloudMap Hawaiian Variant Mapping with WGS Data plot normalization. This table provides details on the numbers of Hawaiian SNPs subtracted from mapping analysis for purposes of SNP mapping plot normalization. Details provided in text and in Fig.7.

Mutant	Trans-gene	Location of trans-gene (LG)	Location of Mutation (LG)	LG I	LG II	LG III	LG IV	LG V	LG X	< .05 or > .95 ratio positions	< .05 or > .95 ratio positions after transgene & mutation LGs removed
ot219	<i>otls114</i>	I	V	✓	✓	✓	✓		✓	15,873	4,003
ot266	<i>vtls1</i> <i>oxlsl2,</i>	V	X	✓	✓	✓	✓		✓*	13,669	11,033
ot628	<i>vsls33</i> <i>otls138,</i>	V, X	I, V		✓	✓	✓			22,268	3,367
ot641	<i>vls33</i> <i>otls138,</i>	V, X	X		✓	✓	✓			9,616	4,314
ot642	<i>vsl33</i>	V, X	X		✓	✓	✓	✓	✓**	13,589	5,021
ot704	<i>otls341</i>	X	I		✓	✓	✓	✓		14,805	7,197
ot705	<i>otls341</i>	X	III	✓	✓		✓	✓	✓	14,686	7,969

✓ Considered SNPs on this chromosome for filtering

* Excluding 6-13 Mb

** Excluding 0-6 Mb

Total dataset used:

WS220_WormMart HA SNPs: 112,061 Variants

<.05 or >.95 ratio positions in at least 2 samples: 8,715 Variants

<.05 & >.95 ratio positions in all 7 samples: 1,563 Variants

Final curated Hawaiian SNP list (WS220.64): 103,346 Variants