

Chr	Site	Ref	Alt	Region	Gene	Detail	Type	Total-depth	Ref-depth	Alt-depth
Chr5	20359626	-	A	upstream	AT5G50050	43831	-	22	0	22
Chr5	20364893	T	-	intronic	AT5G50070	43831	-	14	0	14
Chr5	20364935	-	A	intronic	AT5G50070	43831	-	11	0	11
Chr5	20444003	C	G	exonic	AT5G50210	c.C862G:p.Q288E	nonsynonymous	31	0	31
Chr5	20921186	-	A	upstream	AT5G51510	43831	-	11	0	11
Chr5	21023424	-	A	downstream	AT5G51750	43831	-	13	1	12
Chr5	21046447	T	A	intergenic	AT5G51795	-	-	14	0	14
Chr5	21175547	-	C	downstream	AT5G52120	43831	-	41	0	41
Chr5	21581896	-	T	upstream	AT5G53200	43831	-	26	0	26
Chr5	21672748	-	A	upstream	AT5G53420	43831	-	7	0	7