

**Table 12. Genotypes of wild barley sampled from Sede Boquer**

	<b>64</b>	<b>120</b>	<b>151</b>	<b>194</b>	<b>249</b>	<b>269</b>	<b>277</b>	<b>286</b>	<b>336</b>	<b>339</b>	<b>351</b>	<b>462</b>	<b>546</b>	<b>549</b>	<b>603</b>	<b>625</b>
Consensus	C	C	G	A	G	T	T	A	G	C	T	C	T	C	G	A
20.1																
20.3					T									G	?	?
20.4					T									G	?	?
20.5					T									G		
20.7					T									G	?	?
20.12																
20.13													C	G		
20.24					T									G	?	?
20.25					T									G	?	?
20.27					T									G		
20.28															?	?
20.30	T/C				G/T	T/C						T/C			C/G	
20.33					T									G		
20.35				T							C				?	?
20.36					T									G	?	?
20.37										C					?	?
20.38										C					?	?
20.39					T									G	?	?
20.40					T									G	?	?
20.45													C			

Heterozygous base positions are indicated by a “/” symbol with the two alternative bases indicated. Where clearly readable, the sequence was not obtained for a SNP site; this is indicated by a “?” symbol.