

Table S1. Results from association analysis of *GABRG2* two-marker haplotypes with suicidal behaviour in our schizophrenia sample of European ancestry.

<u>Haplotype window</u>	<u>Haplotype</u>	<u>Case</u>	<u>Control</u>	<u>Suicide Specifier +/- Standard deviation</u>
<u>rs183294-rs209356</u> - - -	<u>T-A</u>	<u>28</u>	<u>83</u>	<u>0.230 (0.074, 0.386)</u>
	<u>C-G</u>	<u>36</u>	<u>101</u>	
	<u>C-A</u>	<u>30</u>	<u>36</u>	
	<u>P</u>	<u>0.00754</u>		
<u>rs209356-rs209357</u> - -				
	<u>p</u>	<u>0.243</u>		<u>0.209</u>
<u>rs209357-rs211037</u> - -				
	<u>p</u>	<u>0.973</u>		<u>0.896</u>
<u>rs211037-rs365054</u> - -				
	<u>p</u>	<u>0.412</u>		<u>0.385</u>
<u>rs365054-rs2422106</u> - -				
	<u>p</u>	<u>0.128</u>		<u>0.117</u>
<u>rs2422106-rs211013</u> - -				
	<u>p</u>	<u>0.825</u>		<u>0.801</u>
<u>rs211013-rs647625</u> - -				
	<u>p</u>	<u>0.746</u>		<u>0.470</u>

Specific haplotypes are shown only for significant haplotype windows.

p-values correspond to omnibus test for each two-marker haplotype window from UNPHASED.