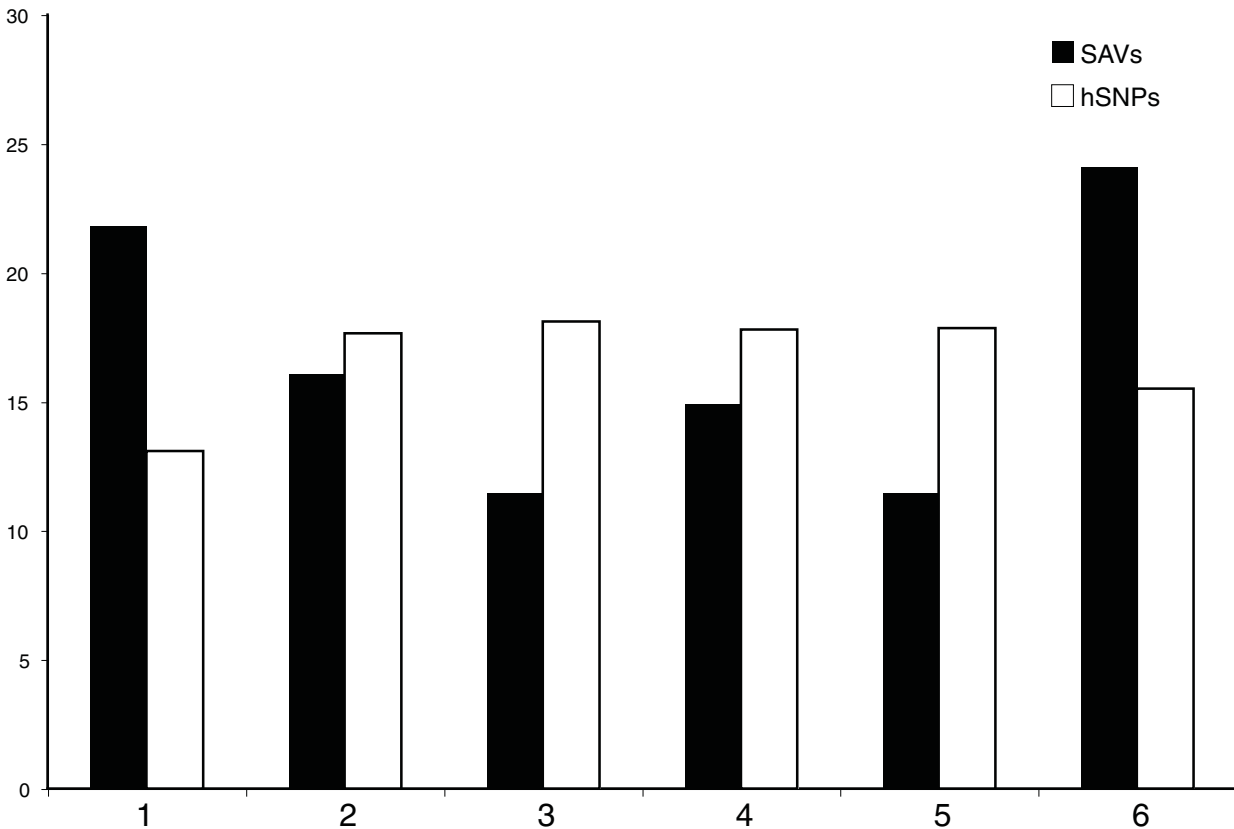


Percentage of variants

■ SAVs
□ hSNPs



← Towards 5' splice site

Exon section

Towards 3' splice site →