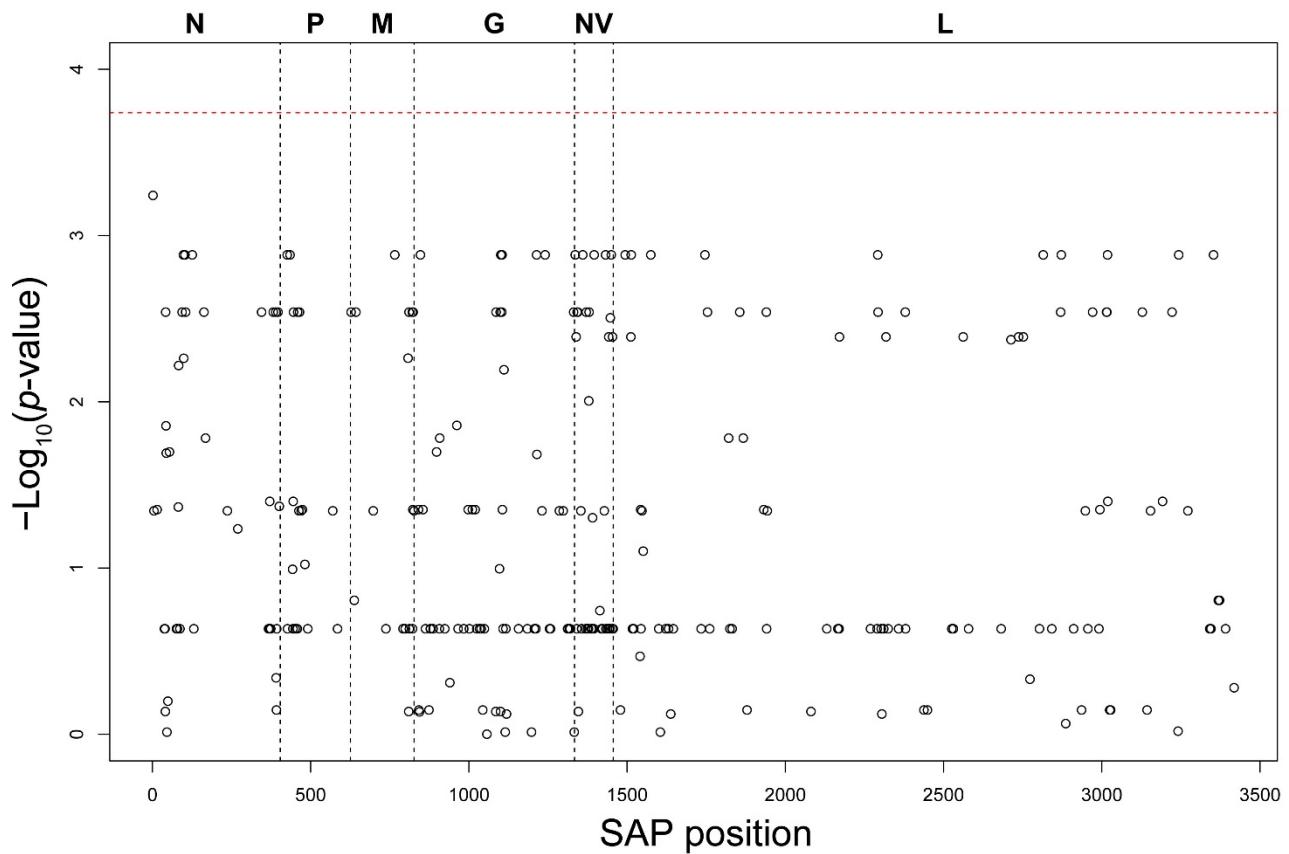


## *Supplementary Material*

### SAPs of low virulent VHSV



**Supplementary Figure 2. Manhattan plot of the association analysis conducted on low virulent isolates SAPs and the trait “virulence”.** The SAP position along the genome coding regions is displayed on the x-axis, while the y-axis reports the negative logarithm of the association  $p$ -value. The red dashed line identifies the association test significance limit. As shown, no SAP was significant associated with virulence in this dataset.