

## S13 File

*Mutation rate, diversity and evenness across the polyprotein*

	<u># Snps per 100bp</u>	<u>Shannon diversity index</u>	<u>Shannon equitability</u>
<b>Early Aegypti</b>	0.397	1.098	0.187
<b>Early Albopictus</b>	0.501	1.22	0.2
<b>Early Human</b>	0.438	1.314	0.22
<b>Late Aegypti</b>	0.536	1.071	0.173
<b>Late Albopictus</b>	0.406	1.152	0.195
<b>Late Human</b>	0.566	1.285	0.206

**S13 File. Mutation rate, diversity and evenness across the polyprotein.** The number of mutations in a 100bp window were calculated across the genome. Diversity and evenness of the nucleotide type change ( $A \leftrightarrow G$ ,  $C \leftrightarrow T$ ,  $A \leftrightarrow C$ ,  $G \leftrightarrow T$ ,  $G \leftrightarrow C$ ,  $A \leftrightarrow T$ ) were calculated using the Shannon diversity index and Shannon equitability measurement respectively