Topology tree for 165 nucleotides of NS5 gene fragment of AeFv from *Ae. aegypti* and *Ae. albopictus.*
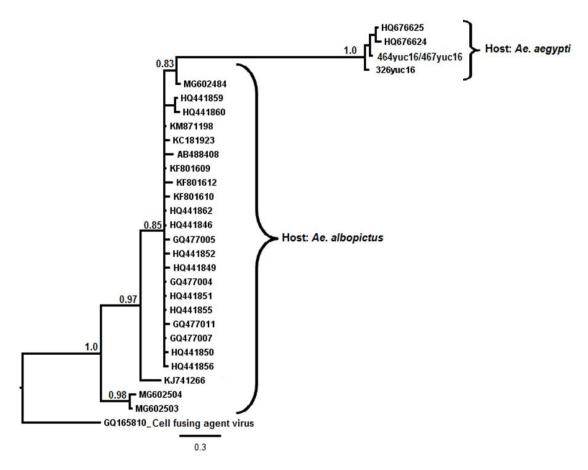


Figure 1. Bayesian Inference (BI) topology tree for 165 nucleotides of NS5 gene fragment of AeFv from *Ae. aegypti* and *Ae. albopictus*. The 464yuc16 and 326yuc16 variants (Yucatan Peninsula, Mexico) were grouped with AeFV from Portugal isolated from *Ae. aegypti*. The numbers on the branches represent posterior probabilities obtained in the BI. The scale bar represents the expected number of nucleotide substitutions per site. Cell fusing agent virus was used as outgroup.