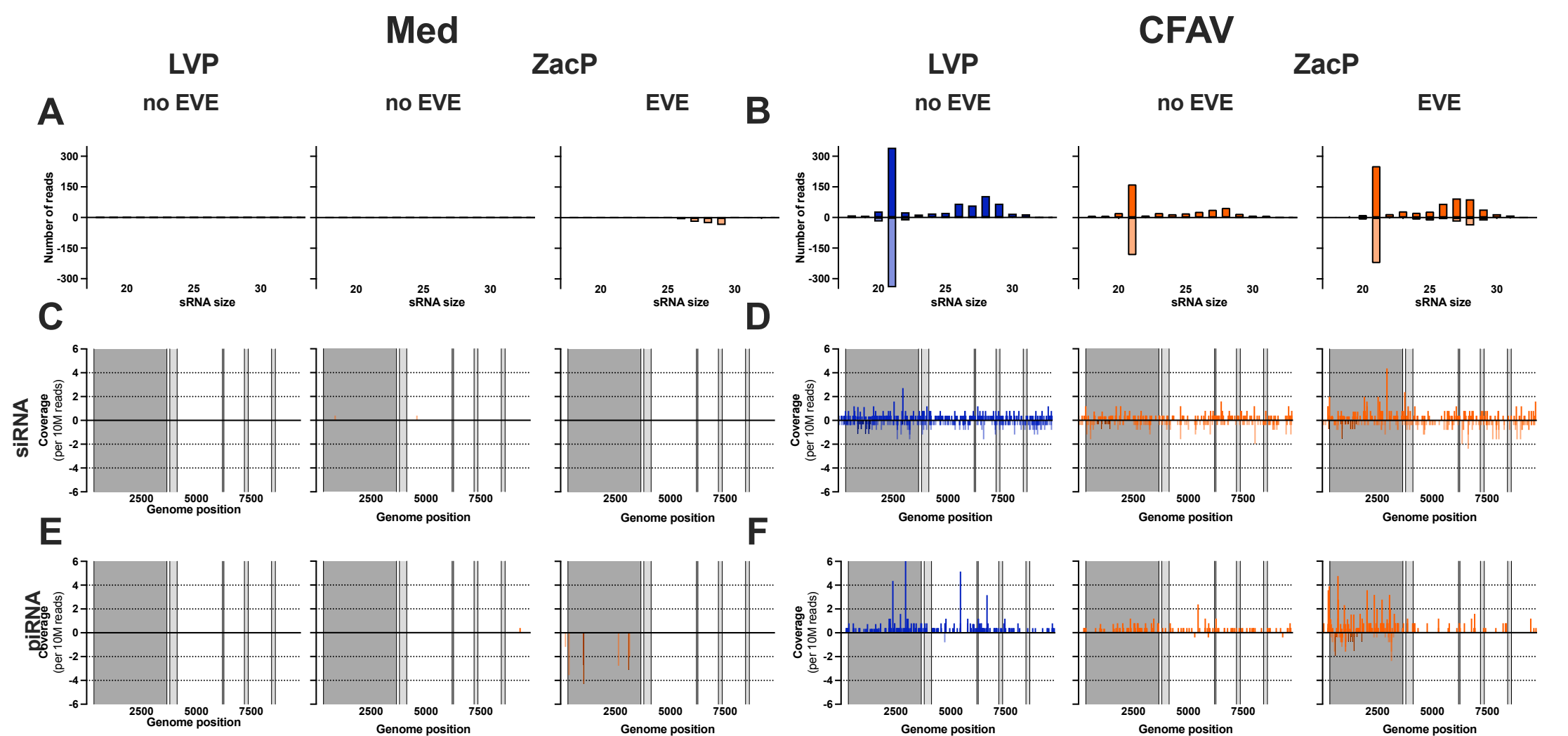




S2 Figure: sRNA profile against CFAV in ovaries of media injected and CFAV infected mosquitoes. In all panels, data from LVP are in blue and from ZacP in orange. **(A)** Size distribution of sRNAs from the ovaries of non-infected LVP and ZacP mosquitoes with or without an EVE mapping to the CFAV genome. **(B)** Size distribution of sRNAs from the ovaries of CFAV infected LVP and ZacP mosquitoes with or without an EVE mapping to the CFAV genome. **(C)** Coverage of 21 nt reads across the CFAV genome from ovaries of non-infected mosquitoes. **(D)** Coverage of 21 nt reads across the CFAV genome from ovaries of CFAV infected mosquitoes. **(E)** Coverage of 26-30 nt reads across the CFAV genome from ovaries of non-infected mosquitoes. **(F)** Coverage of 26-30 nt reads across the CFAV genome from ovaries of CFAV infected mosquitoes. Dark grey and light grey shading represent areas of the CFAV genome with similarity to EVE2 and EVE3, respectively.