

Supplementary figure S6. Identification of putative recombination event involving canine associated cyclovirus CN9E and dragonfly associated cyclovirus 4 isolate US-DFKWGX-2012/Dragonfly/USA/KC512916, as determined by the Recombination Detection Program (RDP v.4.101). A sequence was considered as a recombinant if it was identified by two or more than two methods with a highest acceptable *p-value* of $p < 0.01$ with Bonferroni's correction.

Recombinant: Canine associated cyclovirus CN9E/Dog/KNA

Major parent: Unknown

Minor parent: Dragonfly associated cyclovirus 4 isolate US-DFKWGX-2012/Dragonfly/USA/KC512916

Beginning breakpoint: nt 1303 (without gaps)

Ending breakpoint: nt 1481 (without gaps)



Confirmation table generated by RDP v.4.101 program.

<i>Method</i>	<i>Average p-value</i>
RDP	4.534×10^{-02}
MaxChi	2.027×10^{-02}
Chimaera	1.384×10^{-02}