

Table S3. Number of recombination events calculated by different methods, implemented in the RDP package (Martin et al. 2010): RDP (Martin and Rybicki 2000), GENECONV (Padidam et al. 1999), Maxchi (Maynard Smith 1992), Bootscan (Salminen et al 1995), Chimera (Posada and Crandall 2001), and number of recombination breakpoints calculated in GARD (Kosakovsky Pond et al. 2006) implemented in Datamonkey (Delport et al. 2010)

	No. of sequences	RDP	GENECONV	Maxchi	Bootscan	Chimera	GARD
Entire gradient	57	0	0	1	0	1	0
Northern cluster	14	0	0	0	0	0	0
Southern Cluster	47	0	0	1	0	1	0

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