

Table S8. Comparison of AKMV_VANI10 orthologs to the proteins of AKMV_AKHM13-88 and CPXV_GRI-90 encoded within or flanking the recombination region

AKMV_VANI10 (tentative)	AKMV_AKHM13	CPXV_GRI-90
ORF013	99.53 (AKMV013)	79.32 (C1L)
ORF014	80.34 (AKMV014)	73.60 (C2L)
ORF015*	74.29 (AKMV015)	90.43 (C3L)
ORF016	73.84 (AKMV016)	89.53 (C4L)
ORF017	76.26 (AKMV017)	86.96 (C5R)
ORF018	84.73 (AKMV018)	94.26 (C6L)
OFF019	85.08 (AKMV019)	94.21 (C7R)
ORF020	86.05 (AKMV020)	90.70 (C8L)
ORF021	97.62 (AKMV021)	79.70 (C9L)

Bold number is aa identity; gene names are indicated in parenthesis; *, identity after a 281 aa internal gap was stripped from protein sequence alignment.