# Links between nudivirus-like regions and regions flanking proviral segments

BVs (M. demolitor specifically)

Scaffold	Segment-linked genes Md	Nudivirus-linked genes	Nearest separating distance	Note
				A scaffolds that contains Segment T
Mdem_scaffold_0157	Segment T	HzNVorf93-like	5.4 kb	also has a nudivirus-like gene

BVs

Scaffold	Segment-linked genes Md		Nudivirus-linked genes	Nearest separating distance	Note
		Glyptapanteles Locus 3 (Desjardins et			
		al. 2008): GFP_L6_0020,			
		GFP_L6_0170, and GFP_L6_0160;			Scaffold 938 possesses genes that are similar to those flanking Segments 22
Mdem_scaffold_0938	K425_1716, K425_1718, K425_1728	GIP_L3_0040 and GIP_L3_0210	p47	30.5 kb	and 23 in Glyptapanteles species, and also carries a nudivirus-like gene p47
		Glyptapanteles flavicoxis Locus 7			Scaffold 1213 has a gene similar to a gene flanking Segments 27 and 28 in
Mdem_scaffold_1213	K425_190	(Desjardins et al. 2008): GFP_L5_0170	vlf-1b-1	23.6 kb	Glyptapanteles flavicoxis, and also carries vlf-1b-1
		Glyptapanteles flavicoxis Locus 7			
		(Desjardins et al. 2008): GFP_L5_0170			Scaffold 19 has possesses genes similar to those flanking Segments 27 and 28
Mdem_scaffold_0019	K425_529, K425_534, K425_544	and GFP_L5_0200	int-2	24.2 kb	in Glyptapanteles flavicoxis, and also carries int-2
		Glyptapanteles flavicoxis Locus 7			
		(Desjardins et al. 2008):			
		GFP_L5_0070, GFP_L5_0090,			
		GFP_L5_0190, GFP_L5_0230,			
		GFP_L5_0240; Glyptapanteles			
		indiensis Locus 6, Locus 7:			A conserved uncharacterized gene family member in M. demolitor is present
		GIP_L2_0090, GIP_L2_0100,			in several Glyptapanteles loci containing viral segments (Segment 27 and 28
	K425_820 (conserved	GIP_L7_0040, GIP_L7_0010,	several nudivirus-like genes, closest is		in Gf, Segment 26 in Gi) and is on a Md scaffold that contains several
Mdem_scaffold_0275	uncharacterized protein family 7)	GIP_L7_0080	helicase	58.5 kb	nudivirus-like genes
		Glyptapanteles flavicoxis Locus 7			
		(Desjardins et al. 2008):			
		GFP_L5_0070, GFP_L5_0090,			
		GFP_L5_0190, GFP_L5_0230,			
		GFP_L5_0240; Glyptapanteles			
		indiensis Locus 6, Locus 7:			A conserved uncharacterized gene family member in M. demolitor is present
		GIP_L2_0090, GIP_L2_0100,			in several Glyptapanteles loci containing viral segments (Segment 27 and 28
	K425_412 (conserved	GIP_L7_0040, GIP_L7_0010,			in Gf, Segment 26 in Gi) and is on a Md scaffold that contains a cluster of odv
Mdem_scaffold_1559	uncharacterized protein family 7)	GIP_L7_0080	cluster of odv-e66 genes	1.7 kb	e66 genes
		Glyptapanteles flavicoxis Locus 7			
		(Desjardins et al. 2008):			
		GFP_L5_0070, GFP_L5_0090,			
		GFP_L5_0190, GFP_L5_0230,			
		GFP_L5_0240; Glyptapanteles			
		indiensis Locus 6, Locus 7:			A conserved uncharacterized gene family member in M. demolitor is present
		GIP_L2_0090, GIP_L2_0100,			in several Glyptapanteles loci containing viral segments (Segment 27 and 28
	K425_1388 (conserved	GIP_L7_0040, GIP_L7_0010,	several nudivirus-like genes, closest is		in Gf, Segment 26 in Gi) and is on a Md scaffold that contains a cluster of odv
Mdem_scaffold_0004	uncharacterized protein family 7)	GIP_L7_0080	odv-e66 cluster	197 kb	e66 genes
		Glyptapanteles indiensis Locus 2			A 5' nucleotidase gene is similar to genes flanking segments 9-21 of the
		(Desjardins et al. 2008): GIP_L8_0090			macrolocus in G. indiensis. This gene is located on a scaffold with a cluster of
Mdem scaffold 0040	K425_1276	and GIP_L8_0100	cluster of pif-5 genes	190 kb	pif-5 genes.

## Links between nudivirus-like regions and virulence genes on proviral segments

BVs

Scaffold	Segment-linked genes Md	Segment linked genes other BVs	Nudivirus-linked genes	Nearest separating distance	Note
		hypothetical protein Cotesia			
		conregata bracovirus CcBV_23.4,			
		GIP_L1_00360, and GFP_L7_0430	several nudivirus-like genes, closest is		A gene similar to BV virulence ribonuclease T2 genes is on a scaffold that also
Mdem_scaffold_0004	K425_1471	Ribonuclease T2	odv-e66 cluster	462 kb	contains several nudivirus-like genes
		hypothetical protein Cotesia			A gene similar to a BV virulence gene from CcBV is on a scaffold that also
Mdem_scaffold_0435	K425_1354, K425_1355	conregata bracovirus CcBV_2.3	ac92-like gene	2.2 kb	contains an ac92-like gene

GfVs (Glypta fumiferanae ichnovirus)

Scaffold	Segment-linked genes Md	Segment linked genes other GVs	Nudivirus-linked genes	Nearest separating distance	Note
		aminopeptidase-like Glypta			
		fumiferanae ichnovirus GfV-B16-ORF1			
		protein, Glyptapanteles flavicoxis			
		hypothetical protein, GFP_L5_0170,			A gene similar to Glypta fumiferance IV, Glyptapanteles flavicoxis BV, and
		Chelonus inanitus GfV-B16-ORF1-like	several nudivirus-like genes, closest is		Chelonus inanitus BV is present on a scaffold that also contains several
Mdem_scaffold_0116	K425_142, K425_180	protein	pif-4	15.7 kb	nudivirus-like genes
		aminopeptidase-like Glypta			
		fumiferanae ichnovirus GfV-B16-ORF1			
		protein, Glyptapanteles flavicoxis			
		hypothetical protein, GFP_L5_0170,			
		Chelonus inanitus GfV-B16-ORF1-like			A gene similar to Glypta fumiferance IV, Glyptapanteles flavicoxis BV, and
Mdem_scaffold_0407	K425_1082	protein	lef-4	316 kb	Chelonus inanitus BV is present on a scaffold that also contains lef-4

#### Gene families present in M. demolitor that link nudivirus-like and segment containing scaffolds

		s-like and segment containing  Present on segment-containing	
Gene family name	Present on nudivirus-like scaffolds	scaffolds	
,	Mdem scaffold 0275,		
	Mdem scaffold 0331,		
	Mdem scaffold 0004,		
conserved uncharacterized protein	Mdem scaffold 0040,		
family 3, similar to venom protein 2	Mdem scaffold 0407,		
from Microctonus hyperodae	Mdem scaffold 0919	Mdem scaffold 0157	
· ·	Mdem scaffold 0004,		
cytochrome P450 domain-containing	Mdem scaffold 0275,		
protein family	Mdem scaffold 1154	Mdem scaffold 0025	
	Mdem scaffold 0004,		
	Mdem scaffold 0275,		
	Mdem scaffold 0407,		
	Mdem scaffold 0435,		
EB module-containing protein family	Mdem scaffold 0528	Mdem scaffold 0025	
,		Mdem scaffold 0014,	
ankyrin-repeat containing protein	Mdem scaffold 0004	Mdem scaffold 0025	
Endoribonuclease XendoU domain-			
containing protein family	Mdem scaffold 0407	Mdem scaffold 0025	
Flavin-containing monooxygenase-like			
family	Mdem scaffold 0275	Mdem scaffold 0014	
•		Mdem scaffold 0014,	
protein tyrosine kinase family	Mdem scaffold 0407	Mdem scaffold 0412	
Haloacid dehalogenase-like hydrolase			
domain-containing protein family	Mdem scaffold 0407	Mdem scaffold 0025	
Immunoglobulin domain-containing	Mdem scaffold 0275,		
protein family	Mdem scaffold 1798	Mdem scaffold 0014	
P-type ATPase of unknown pump			
specificity (type V) family	Mdem_scaffold_0067	Mdem scaffold 1462	
Phosphatidylinositol-specific			
phospholipase C, X domain-containing	Mdem_scaffold_0275,		
protein family	Mdem_scaffold_0040	Mdem_scaffold_0157	
acid phosphatase-like protein family	Mdem_scaffold_0040	Mdem_scaffold_1462	
kelch-like protein family	Mdem_scaffold_0275	Mdem_scaffold_0014	
	Mdem_scaffold_0040,		
kinesin-like protein KIF11 family	Mdem scaffold 1154	Mdem scaffold 0014	
scavenger receptor class B member-			
like family	Mdem scaffold 1154	Mdem_scaffold_0025	
ubiquitin thioesterase trabid-like			
protein	Mdem_scaffold_0407	Mdem_scaffold_0025	
transporter, solute:sodium symporter	Mdem_scaffold_0404,	_	
(SSS) family	Mdem scaffold 0407	Mdem scaffold 0025	
lipase domain-containing protein			
family	Mdem scaffold 0240	Mdem scaffold 0025	

## Gene families present in M. demolitor that link segment containing scaffolds

	Present on segment-containing		
Gene family name	scaffolds		
	Mdem_scaffold_0014,		
	Mdem_scaffold_0157,		
	Mdem_scaffold_0446,		
Histone proteins	Mdem_scaffold_1462		

[only includes those not already listed

above]

## Gene families present in M. demolitor that link nudivirus-like scaffolds

Gene family name	Present on nudivirus-like scaffolds
conserved uncharacterized protein	Mdem_scaffold_0919,
family 1	Mdem_scaffold_1213
	Mdem_scaffold_0116,
conserved uncharacterized protein fan	Mdem_scaffold_0407
	Mdem_scaffold_0275,
conserved uncharacterized protein fan	Mdem_scaffold_0919
alpha-tubulin-like family	Mdem_scaffold_0275,
	Mdem_scaffold_0528,
	Mdem_scaffold_0919,
	Mdem_scaffold_0040,
annexin-like family	Mdem_scaffold_0275
	Mdem_scaffold_0004,
another transcription unit-like family	Mdem_scaffold_0331
	Mdem_scaffold_0275,
BTB/POZ domain containing protein fa	Mdem_scaffold_0919,
	Mdem_scaffold_0116,
disintegrin and metalloproteinase dom	Mdem_scaffold_0407
	Mdem_scaffold_0004,
	Mdem_scaffold_0407,
glucose dehydrogenase acceptor-like fa	Mdem_scaffold_1154
	Mdem_scaffold_0004,
lipase 3-like family	Mdem_scaffold_0407
	Mdem_scaffold_0004,
MFS sugar transporter domain-contain	Mdem_scaffold_0938
	Mdem_scaffold_0067,
	Mdem_scaffold_0938,
neprilysin-like family	Mdem_scaffold_1154
	Mdem_scaffold_0004,
clavesin-2-like family	Mdem_scaffold_0407

[only includes those not already listed

above]