Table S1: Identification and distribution of sequences of insect and plant viruses of interest among the various bat specimens and tissue samples analyzed

		Bat species, specimens and tissue samples ^a								
Virus	Sequence detection ^b	Pipistrellus pipistrellus					Hypsugo savii	Myotis nattereri	Eptesicus serotinus	Myotis mystacinus
		b1	b2	b3	b4	b9	b5	b6	b7	b8
		Br Li Lu	Br Li Lu	Br Li Lu	Br Li Lu	Li Lu	Br Li Lu	Br Li Lu	Br Li Lu	Li Lu
Dicistrovirus	HTS	- c (tissue pool)	+c (tissue pool) [ORF1-2]	- (tissue pool)	- (tissue pool)	- (tissue pool)	- (tissue pool)	- (tissue pool)	- (tissue pool)	- (tissue pool)
	PCR/Sanger		+ [ORF1]							
Nodavirus	HTS	- (tissue pool)	- (tissue pool)	- (tissue pool)	- (tissue pool)	- (tissue pool)	- (tissue pool)	- (tissue pool)	+ (tissue pool) [multiple genes]	- (tissue pool)
	PCR/Sanger								+ [ORF1a]	
Luteovirus	HTS	- (tissue pool)	- (tissue pool)	+ (tissue pool) [ORF1-2]	- (tissue pool)	- (tissue pool)	- (tissue pool)	- (tissue pool)	- (tissue pool)	- (tissue pool)
	PCR/Sanger	ND ^d ND ND	ND ND ND	ND ND ND	ND ND ND	ND ND	ND ND ND	ND ND ND	ND ND ND	ND ND
Sobemovirus	HTS	- (tissue pool)	- (tissue pool)	+ (tissue pool) [ORF4]	- (tissue pool)	- (tissue pool)	- (tissue pool)	- (tissue pool)	- (tissue pool)	- (tissue pool)
	PCR/Sanger	ND ND ND	ND ND ND	ND ND ND	ND ND ND	ND ND	ND ND ND	ND ND ND	ND ND ND	ND ND

Legend:

a: Br = brain, Li = liver, Lu = lungs

b: Matching (HTS) or targeting (PCR/Sanger) viral genes (in brackets), with ORF = open reading frame

c: + = positive, - = negative

d: ND = not done