

This form should be used for all taxonomic proposals. Please complete all those modules that are applicable (and then delete the unwanted sections). For guidance, see the notes written in blue and the separate document "Help with completing a taxonomic proposal"

Please try to keep related proposals within a single document; you can copy the modules to create more than one genus within a new family, for example.

MODULE 1: TITLE, AUTHORS, etc

Code assigned:	2015.015	a,bP		(to be con officers)	mpleted by	ICTV
Short title: 56 new and 22 dela (e.g. 6 new species in the genus A Modules attached (modules 1 and 10 are required)	eted species in Z <i>etavirus</i>)	the genus $1 \boxtimes 6 \square$	Begomov 2 🔀 7 🔀	irus 3 🗌 8 🗌	4 🗌 9 🗌	5 🗌 10 🖂

Author(s):

F.M. Zerbini and J. Navas-Castillo, on behalf of the *Geminiviridae* Study Group

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List the ICTV study group(s) that have seen this proposal:

A list of study groups and contacts is provided at <u>http://www.ictvonline.org/subcommittees.asp</u>. If in doubt, contact the appropriate subcommittee chair (fungal, invertebrate, plant, prokaryote or vertebrate viruses)

Geminiviridae

ICTV Study Group comments (if any) and response of the proposer:

Date first submitted to ICTV: Date of this revision (if different to above): 15 June 2015

ICTV-EC comments and response of the proposer:

MODULE 2: NEW SPECIES

creating and naming one or more new species.

If more than one, they should be a group of related species belonging to the same genus. All new species must be placed in a higher taxon. This is usually a genus although it is also permissible for species to be "unassigned" within a subfamily or family. Wherever possible, provide sequence accession number(s) for **one** isolate of each new species proposed.

Code	201	15.015aP (assigned by IC		TV office	ers)	
To crea	ite 56 r	new species within	:			
				Fill in	all that apply.	
G	Genus: Begomovirus		If the higher taxon has yet to be			
Subfa	mily:			 created (in a later module, below) write "(new)" after its proposed name. If no genus is specified, enter 		
Fa	mily:	Geminiviridae				
(Order:			"unassigned" in the genus box.		
Name of new species: Re (or		Representative isol (only 1 per species pl	late: lease)	GenBank sequence accession number(s)		
See Tab	ole 1.					

Table 1. Proposed new species, isolate name, and corresponding GenBank accession numbers. Accession numbers refer to DNA-A and DNA-B (when available) components for bipartite viruses or the single component for monopartite viruses.

Species name	Isolate name	GenBank access #
Abutilon golden mosaic virus	Abutilon golden mosaic virus - [Mexico:Yucatan:2007]	KC430935
Capraria yellow spot virus	Capraria yellow spot virus - [Mexico:Yucatan:Conkal:2007]	KC426927, KC426928
Cassava mosaic Madagascar virus	Cassava mosaic Madagascar virus - [Madgascar:Toliary:2006]	HE617299, HE617300
Catharanthus yellow mosaic virus	Catharanthus yellow mosaic virus - [Pakistan:Lahore:KN4]	HE580234
Chenopodium leaf curl virus	Chenopodium leaf curl virus - [USA:Florida:Citra:2007]	HM626515
Chilli leaf curl India virus	Chilli leaf curl India virus - [India:2008]	FM877858
Chilli leaf curl Kanpur virus	Chilli leaf curl Kanpur virus [India:Kanpur:2008]	HM007106
Chilli leaf curl Vellanad virus	Chilli leaf curl Vellanad virus - [India:Vellanad:2008]	HM007121
Clerodendron yellow mosaic virus	Clerodendron yellow mosaic virus - [India:lari:2006]	EF408037
Clerodendrum golden mosaic China virus	Clerodendrum golden mosaic China virus - [China:Fuzhou 7:2007]	FJ011668, FJ011669
Clerodendrum golden mosaic Jiangsu virus	Clerodendrum golden mosaic Jiangsu virus - [China:Jiangsu XY2:2008]	FN396966
Corchorus yellow vein mosaic virus	Corchorus yellow vein mosaic virus - [India:Maharashtra:2011]	KC196077
Cotton chlorotic spot virus	Cotton chlorotic spot virus - [Brazil:Campina Grande B012:2009]	KF358470
Crassocephalum yellow vein virus	Crassocephalum yellow vein virus - [China:Jinhong:2005]	EF165536
Emilia yellow vein virus	Emilia yellow vein virus - [China:Fuzhou 1:2007]	EU377539
French bean leaf curl virus	French bean leaf curl virus - [India:Kanpur:2011]	JQ866297
Hedyotis uncinella yellow mosaic virus	Hedyotis uncinella yellow mosaic virus - [Vietnam:VN1]	KF429251
Hemidesmus yellow mosaic virus	Hemidesmus yellow mosaic virus - [India:Tirupati:H1:2012]	KC898543
Jatropha leaf curl virus	Jatropha leaf curl virus - [India:New Delhi:2007]	EU798996
Jatropha mosaic Nigeria virus	Jatropha mosaic Nigeria virus - [Nigeria:2:2011]	JX025358
Jatropha mosaic virus	Jatropha mosaic virus - [Jamaica:Spanish Town 1:2004]	KF723258, KF723261
Jatropha yellow mosaic virus	Jatropha yellow mosaic virus - [India:Kathaupahadi:2008]	FJ177030
Malvastrum leaf curl Philippines virus	Malvastrum leaf curl Philippines virus - [Philippines:Mc1:2012]	KC577540
Mesta yellow vein mosaic Bahraich virus	Mesta yellow vein mosaic Bahraich virus - [India:Bahraich:2007]	EU360303
Pepper yellow leaf curl virus	Pepper yellow leaf curl virus - [China:YN65- 1:2010]	KC149938
Pouzolzia golden mosaic virus	Pouzolzia golden mosaic virus - [China:TY01:2012]	JX183732
Pouzolzia mosaic Guangdong virus	Pouzolzia mosaic Guangdong virus - [Taiwan:Miaoli:ML13W1:2013]	KF414123
Premna leaf curl virus	Premna leaf curl virus - [Vietnam:VN7:2011]	JQ793786
Rhynchosia yellow mosaic India virus	Rhynchosia yellow mosaic India virus - [India:Thiruvananthapuram:JRH1:2009]	HM777508, HM777510

Sauropus leaf curl virus	Sauropus leaf curl virus - [Thailand:Kamphaengsaen:AFSP5e:2010]	JN809819
Sida ciliaris golden mosaic virus	Sida ciliaris golden mosaic virus - [Venezuela:Lara:M3:2009]	JX857691
Sida common mosaic virus	Sida common mosaic virus - [Brazil:Coimbra 4:2007]	EU710751
Sida golden mosaic Brazil virus	Sida golden mosaic Brazil virus - [Brazil:Mato Grosso do Sul:2007]	FN436001
Sida golden mosaic Lara virus	Sida golden mosaic Lara virus - [Venezuela:Lara:M1:2009]	JX857693
Sida yellow leaf curl virus	Sida yellow leaf curl virus - [Brazil:Coimbra3:2007]	EU710750
Sidastrum golden leaf spot virus	Sidastrum golden leaf spot virus - [Brazil:DF334:2010]	HM357458
Soybean chlorotic blotch virus	Soybean chlorotic blotch virus - [Nigeria:Soybean 19:2007]	GQ472985, GQ472986
Spinach yellow vein virus	Spinach yellow vein virus - [India:Sikar:AS22]	KF660223
Sunn hemp leaf distortion virus	Sunn hemp leaf distortion virus - [India:Barrackpore 1:2008]	FJ455449
Sweet potato leaf curl Henan virus	Sweet potato leaf curl Henan virus - [China:Henan 10:2012]	KC907406
Sweet potato leaf curl Sichuan virus 1	Sweet potato leaf curl Sichuan virus 1 - [China:Sichuan 15:2012]	KC488316
Sweet potato leaf curl Sichuan virus 2	Sweet potato leaf curl Sichuan virus 2 - [China:Sichuan 14:2012]	KF156759
Tobacco leaf curl Comoros virus	Tobacco leaf curl Comoros virus - [Comoros:Simboussa:2004]	AM701760
Tomato bright yellow mosaic virus	Tomato bright yellow mosaic virus - [Brazil:BA167:2012]	KC791690
Tomato bright yellow mottle virus	Tomato bright yellow mottle virus - [Brazil:TO167:2008]	KC791691
Tomato golden leaf distortion virus	Tomato golden leaf distortion virus - [Brazil:TO45:2007]	HM357456
Tomato interveinal chlorosis virus	Tomato interveinal chlorosis virus - [Brazil:Pernambuco:Mdc2681:2004]	JF803252
Tomato leaf curl Liwa virus	Tomato leaf curl Liwa virus - [Oman:Liwa:LW1:2012]	HF912280
Tomato leaf curl New Delhi virus 2	Tomato leaf curl New Delhi virus 2 - [India:IANDS1:2011]	JQ897969
Tomato leaf curl New Delhi virus 4	Tomato leaf curl New Delhi virus 4 - [India:Junagad:TC306:2011]	KF551592
Tomato leaf curl Palampur virus	Tomato leaf curl Palampur virus - [India:Palampur:2007]	AM884015, AM992534
Tomato leaf curl Patna virus	Tomato leaf curl Patna virus - [India:Patna:2008]	EU862323
Tomato leaf curl Rajasthan virus	Tomato leaf curl Rajasthan virus - [India:Rajasthan:2005]	DQ339117
Tomato leaf curl Sulawesi virus	Tomato leaf curl Sulawesi virus - [Indonesia:Sulawesi:Langowan F101:2006]	FJ237614
Velvet bean severe mosaic virus	Velvel bean severe mosaic virus - [India:Lucknow:2008]	FN543425, FN543426
Vigna yellow mosaic virus	Vigna yellow mosaic virus - [Mexico:Morelos:Yautepec:2007]	KC430936

Reasons to justify the creation and assignment of the new species:

- Explain how the proposed species differ(s) from all existing species.
 - If species demarcation criteria (see module 3) have previously been defined for the genus, **explain how the new species meet these criteria**.
 - If criteria for demarcating species need to be defined (because there will now be more than one species in the genus), please state the proposed criteria.

• Further material in support of this proposal may be presented in the Appendix, Module 9

- 1. All have distinctively begomovirus-like genome organization.
- 2. All are associated with the whitefly vector Bemisia tabaci.
- 3. Phylogenetically, they all group within the *Begomovirus* genus (see Module 10, Figure 1, for phylogenetic trees).
- 4. All taxa proposed as new species meet the current species demarcation threshold of <91% nucleotide sequence identity when considering the DNA-A component or the monopartite genome of the isolate (the DNA-B component of bipartite viruses is not taxonomically informative), when implementing the Muscle alignment method to compute pairwise identity comparisons (one minus Hamming distances of pairwise aligned sequences with pairwise deletion of gaps). This is the approach recently described by the *Geminiviridae* Study Group for the three main genera in the family (Brown *et al.*, 2015; Muhire *et al.*, 2013; Varsani *et al.*, 2014).

New begomovirus species continue to be reported on a regular basis. This is partly due to ease of detection (small circular DNA genomes that can be readily amplified by rolling-circle amplification in a sequence-unbiased fashion) but also reflects (i) the great diversity of this group of viruses, and (ii) the fact that many groups have turned their attention to non-cultivated plants: isolates of 31 out of the 56 proposed new species were obtained from non-cultivated plants. The sequences of all 56 proposed new species were obtained by conventional (Sanger) sequencing of a full-length clone, using a primer-walking strategy. Assembly methods varied with each specific sequence, but included the most commonly used programs such as BioEdit, DNABaser, DNAMan, DNAStar and VectorNTI (Table A1, Annex).

MODULE 7: REMOVE and MOVE

Use this module whenever an existing taxon needs to be removed:

- Either to abolish a taxon entirely (when only part (a) needs to be completed)
- *Either* to abolish a taxon entirely (when only part (a) needs to be completed;
 Or to move a taxon and re-assign it e.g. when a species is moved from one genus to another (when BOTH parts (a) and (b) should be completed)

Part (a) taxon/taxa to be removed or moved

Code	201	5.015bP	(assigned by I	CTV officers)	
To remove the following taxon (or taxa) from their present position:					
Abutilon Ageratur Bhendi y Bitter go Eclipta y Hollyhoo Honeysu Kenaf lea Macropt Malvastr Malvastr Malvastr Okra lea Okra yel Okra yel Sida gola Sida gola Sida gola Sida yell Soybean Tomato	Brazil n leaf ellow urd ye eellow ck leaf ckle ye af curl ilium y rum lee rum ye f curl low m low ve den mo den mo ow vel chlore crinkl leaf cu	l virus curl Cameroon virus vein India virus ellow vein virus crumple virus crumple virus ellow vein Kagoshima vir virus vellow net virus af curl Guangdong virus ilow leaf curl virus llow leaf curl virus curl Guangdong virus vellow net virus ilow vein Changa Manga Cameroon virus ottle Iguala virus cosaic Honduras virus osaic Honduras virus osaic Liguanea virus otic spot virus ce leaf virus rl Kumasi virus	us virus		
The pres	sent ta	axonomic position of the	se taxon/taxa:		
G Subfa Fa C	enus: mily: mily: Order:	Begomovirus Geminiviridae		Fill in all that apply.	
If the taxon/taxa are to be abolished (i.e. not reassigned to another taxon) write "yes" YES In the box on the right YES Reasons to justify the removal: Explain why the taxon (or taxa) should be removed					
Following the reassessment of the species demarcation criteria in the genus <i>Begomovirus</i> (Brown <i>et al.</i> , 2015), the <i>Geminiviridae</i> Study Group is reevaluating the status of all current species. For the above listed species, it was found that one or more isolates have >91% identity, for the monopartite genome or the DNA-A, with isolates of other species (Table 2). Therefore, these species should be merged. The older species is being maintained, and all isolates of the deleted species will be renamed accordingly.					

Table 2. List of species to be deleted from the genus *Begomovirus*. Isolates belonging to these species were found to have >91% nucleotide sequence identity with isolates belonging to other species. The GenBank access numbers of the relevant isolates are presented, as well as the percent identity between their nucleotide sequences (whole genome in the case of monopartite viruses, DNA-A component in the case of bipartite viruses).

Deleted species	Access #	Merges with	Access #	% id.
Abutilon Brazil virus	FN434438	Abutilon mosaic Brazil virus	JF694480	93
Ageratum leaf curl Cameroon virus	FR873229	Tomato leaf curl Ghana virus	HE659517	91
Bhendi yellow vein India virus	GU112039	Bhendi yellow vein mosaic virus	GU112056	94
Bitter gourd yellow vein virus	AM491590	Tomato leaf curl New Delhi virus	AM747291	91
Eclipta yellow vein virus	GQ478343	Hollyhock leaf curl virus	FR772082	91
Hollyhock leaf crumple virus	AY036009	Cotton leaf curl Gezira virus	AF014881	99
Honeysuckle yellow vein Kagoshima virus	AB178949	Honeysuckle yellow vein virus	AB236323	92
Kenaf leaf curl virus	FN806777	Malvastrum yellow vein virus	AJ457824	91
Macroptilium yellow net virus	JN418998	Tomato interveinal chlorosis virus	JF803252	92
Malvastrum leaf curl Guangdong virus	AM236779	Papaya leaf curl Guandong virus	AJ558122	91
Malvastrum yellow leaf curl virus	AJ971524	Tomato yellow leaf curl China virus	AM260703	92
Malvastrum yellow vein Changa Manga virus	FR715681	Mesta yellow vein mosaic virus	FR772081	91
Okra leaf curl Cameroon virus	FR717137	Cotton leaf curl Gezira virus	FM164726	98
Okra yellow mottle Iguala virus	AY751753	Chino del tomate virus	AF101476	91
Okra yellow vein mosaic virus	AJ002451	Bhendi yellow vein mosaic virus	AF241479	95
Sida golden mosaic Honduras virus	Y11097	Sida yellow vein virus	Y11099	91
Sida golden mosaic Liguanea virus	HQ009522	Sida golden yellow vein virus	HQ009519	91
Sida yellow vein Madurai virus	AM259382	Sida leaf curl virus	AM050730	91
Soybean chlorotic spot virus	JX122965	Macroptilium yellow spot virus	JN419013	91
Soybean crinkle leaf virus	AB050781	Ageratum yellow vein virus	JN809816	92
Tomato leaf curl Kumasi virus	EU847739	Tomato leaf curl Ghana virus	EU350585	91
Tomato mosaic leaf curl virus	AY508991	Merremia mosaic virus	AF068636	92

MODULE 10: APPENDIX: supporting material

additional material in support of this proposal

References:

- Brown JK, Zerbini FM, Navas-Castillo J, Moriones E, Ramos-Sobrinho R, Silva JCF, Fiallo-Olive E, Briddon R, Hernández-Zepeda C, Idris A, Malathi VG, Martin DP, Rivera-Bustamante R, Ueda S, Varsani A (2015) Revision of *Begomovirus* taxonomy based on pairwise sequence comparisons. Archives of Virology 160:1593-1619.
- Muhire B, Martin DP, Brown JK, Navas-Castillo J, Moriones E, Zerbini FM, Rivera-Bustamante R, Malathi VG, Briddon RW, Varsani A (2013) A genome-wide pairwiseidentity-based proposal for the classification of viruses in the genus *Mastrevirus* (family *Geminiviridae*). Archives of Virology 158:1411-1424.
- Varsani A, Martin DP, Navas-Castillo J, Moriones E, Hernandez-Zepeda C, Idris A, Zerbini FM, Brown JK (2014) Revisiting the classification of curtoviruses based on genomewide pairwise identity. Archives of Virology 159:1873-1882.

Annex:

Include as much information as necessary to support the proposal, including diagrams comparing the old and new taxonomic orders. The use of Figures and Tables is strongly recommended but direct pasting of content from publications will require permission from the copyright holder together with appropriate acknowledgement as this proposal will be placed on a public web site. For phylogenetic analysis, try to provide a tree where branch length is related to genetic distance.

Α

0.05



Figure A1. Maximum-likelihood phylogenetic trees based on the complete monopartite genome for monopartite viruses and the DNA-A component of bipartite begomoviruses. Bootstrap values (2,000 replications) are represented as filled circles when >90% and as open circles when between 50% and 89%. **A**, New World begomoviruses; **B**, Old World begomoviruses. New species in this proposal are indicated in green.

Species name	GenBank access #	Sequence/Assembly method	Reference
Abutilon golden mosaic virus	KC430935	Sanger/DNAStar	n.a.
Capraria yellow spot virus	KC426927, KC426928	Sanger/DNAStar	n.a.
Cassava mosaic Madagascar virus	HE617299, HE617300	Sanger/DNA Baser	Harimalala et al., Arch Virol 57: 2027, 2012
Catharanthus yellow mosaic virus	HE580234	Sanger/DNAStar	Ilyas et al., Arch. Virol. 158:505, 2013
Chenopodium leaf curl virus	HM626515	Sanger/DNAStar	Ng et al., PLoS ONE 6:E19050, 2011
Chilli leaf curl India virus	FM877858	Sanger/no info	n.a.
Chilli leaf curl Kanpur virus	HM007106	Sanger/no info	n.a.
Chilli leaf curl Vellanad virus	HM007121	Sanger/DNAStar	Sharma et al., Plant Cell Rep 34:1389, 2015
Clerodendron yellow mosaic virus	EF408037	Sanger/no info	Sivalingam et al., Acta Virol 55:357, 2011
Clerodendrum golden mosaic China virus	FJ011668, FJ011669	Sanger/DNAStar	Li & Zhou, Virus Genes 41:250, 2010
Clerodendrum golden mosaic Jiangsu virus	FN396966	Sanger/DNAStar	Li & Zhou, Virus Genes 41:250, 2010
Corchorus yellow vein mosaic virus	KC196077	Sanger/no info	n.a.
Cotton chlorotic spot virus	KF358470	Sanger/Staden	Almeida et al., Genome Announc 1:e00661-13, 2013
Crassocephalum yellow vein virus	EF165536	Sanger/DNAMan	Dong et al., J. Phytopathol. 156:193, 2008
Emilia yellow vein virus	EU377539	Sanger/DNAStar	Yang et al., J Plant Pathol 90:475, 2008
French bean leaf curl virus	JQ866297	Sanger/BioEdit	Kamaal et al., Virus Genes 46:120, 2013
Hedyotis uncinella yellow mosaic virus	KF429251	Sanger/DNAStar	Du et al., Virus Genes 48:557, 2014
Hemidesmus yellow mosaic virus	KC898543	Sanger/BioEdit	Reddy et al., Arch. Virol. 159:1223, 2014
Jatropha leaf curl virus	EU798996	Sanger/no info	n.a.
Jatropha mosaic Nigeria virus	JX025358	Sanger/no info	Kashina et al., Arch. Virol. 158:511, 2013
Jatropha mosaic virus	KF723258, KF723261	Sanger/DNAStar	Simmonds-Gordon et al., Arch Virol 159:2815, 2014
Jatropha yellow mosaic virus	FJ177030	Sanger/no info	Snehi et al., Arch. Virol. 156:2303-2307, 2011
Malvastrum leaf curl Philippines virus	KC577540	Sanger/no info	n.a.
Mesta yellow vein mosaic Bahraich virus	EU360303	Sanger/BioEdit	Das et al., Arch. Virol. 153:1791, 2008
Pepper yellow leaf curl virus	KC149938	Sanger/no info	n.a.
Pouzolzia golden mosaic virus	JX183732	Sanger/DNAStar	Tang et al., Arch. Virol. 158:1617, 2013
Pouzolzia mosaic Guangdong virus	KF414123	Sanger/DNAStar	Tang et al., Arch. Virol. 159:2799, 2014

Table A1. Sequencing and assembly methods used for obtaining the full-length sequence of the proposed new species in the genus *Begomovirus*.

Premna leaf curl virus	JQ793786	Sanger/DNAStar	She et al., Arch. Virol. 158:2425, 2013
Rhynchosia yellow mosaic India virus	HM777508, HM777510	Sanger/no info	Jyothsna et al., Virus Genes 42:407, 2011
Sauropus leaf curl virus	JN809819	Sanger/DNAStar	Shih et al., J Phytopath 161:78, 2013
Sida ciliaris golden mosaic virus	JX857691	Sanger/no info	n.a.
Sida common mosaic virus	EU710751	Sanger/DNA Baser	Castillo-Urquiza et al., Arch Virol 153:1985, 2008
Sida golden mosaic Brazil virus	FN436001	Sanger/BioEdit	Paprotka et al., Virology 404:148, 2010
Sida golden mosaic Lara virus	JX857693	Sanger/no info	n.a.
Sida yellow leaf curl virus	EU710750	Sanger/DNA Baser	Castillo-Urquiza et al., Arch Virol 153:1985, 2008
Sidastrum golden leaf spot virus	HM357458	Sanger/no info	n.a.
Soybean chlorotic blotch virus	GQ472985, GQ472986	Sanger/Vector NTI	Alabi et al., Arch. Virol. 155:643, 2010
Spinach yellow vein virus	KF660223	Sanger/no info	Sahu et al., J Gen Plant Path 81:576, 2015
Sunn hemp leaf distortion virus	FJ455449	Sanger/no info	n.a.
Sweet potato leaf curl Henan virus	KC907406	Sanger/DNAMan	Liu et al., Arch. Virol. 159:1537, 2014
Sweet potato leaf curl Sichuan virus 1	KC488316	Sanger/DNAMan	Liu et al., Virus Genes 47:591, 2013
Sweet potato leaf curl Sichuan virus 2	KF156759	Sanger/no info	n.a.
Tobacco leaf curl Comoros virus	AM701760	Sanger/no info	Lefeuvre et al., J Gen Virol 88:3458, 2007
Tomato bright yellow mosaic virus	KC791690	Sanger/no info	n.a.
Tomato bright yellow mottle virus	KC791691	Sanger/no info	n.a.
Tomato golden leaf distortion virus	HM357456	Sanger/no info	n.a.
Tomato interveinal chlorosis virus	JF803252	Sanger/DNAStar	Albuquerque et al., Arch. Virol. 157:747, 2012
Tomato leaf curl Liwa virus	HF912280	Sanger/DNAStar	Khan et al., Arch. Virol. 159:445, 2014
Tomato leaf curl New Delhi virus 2	JQ897969	Sanger/no info	n.a.
Tomato leaf curl New Delhi virus 4	KF551592	Sanger/no info	n.a.
Tomato leaf curl Palampur virus	AM884015, AM992534	Sanger/no info	Kumar et al., Virus Genes 37:425, 2008
Tomato leaf curl Patna virus	EU862323	Sanger/no info	Kumari et al., Virus Res 152:19, 2010
Tomato leaf curl Rajasthan virus	DQ339117	Sanger/no info	n.a.
Tomato leaf curl Sulawesi virus	FJ237614	Sanger/DNAStar	Tsai et al., Plant Dis. 93:321, 2009
Velvet bean severe mosaic virus	FN543425, FN543426	Sanger/no info	Zaim et al., Virus Genes 43:138, 2011
Vigna yellow mosaic virus	KC430936	Sanger/DNAStar	n.a.