

TVCV OSU	MSIVSYEPKVSDFLNLSSKKEEILPKALTRLKTVISISTKDIISVKESSETLCDIDLLINVPLDKYRYVG	67	<a href="#">Q88921</a>
NZ-587	MSIVSYEPKVSDFLNLSSKKEEILPKALTRLKTVISISTKDIISVKESSETLCDIDLLINVPLDKYRYVG	67	<a href="#">I6NG32</a>
NZ-438	MSIVSYEPKVSDFLNLSSKKEEILPKALTRLKTVISISTKDIISVKESSETLCDIDLLINVPLDKYRYVG	67	<a href="#">I6NH32</a>
Pet-MS	MSIVSYEPKVSDFLNLSSKKEEILPKALTRLKTVISISTKDIISVKESSETLCDIDLLINVPLDKYRYVG	67	<a href="#">B1A0S9</a>
D/A	MSIVSYEPKVSDFLNLSSKKEEILPKALTRLKTVISISTKDIISVKESSETLCDIDLLINVPLDKYRYVG	67	<a href="#">Q88605</a>
YoMV (ORMV)	---MSYEPKVSDFLALTTKKEEILPKALTRLKTVISISTKDVISVKESSESLCDIDLLVNVPDLKYRYVG	64	<a href="#">Q66221</a>
Shanxi	---MSYEPKVSDFLALTTKKEEILPKALTRLKTVISISTKDVISVKESSESLCDIDLLVNVPDLKYRYVG	64	<a href="#">H6X0P4</a>
Br	---MSYEPKVSDFLALTTKKEEILPKALTRLKTVISISTKDVISVKESSESLCDIDLLVNVPDLKYRYVG	64	<a href="#">G9IAP7</a>
R/W	---MSYEPKVSDFLALTTKKEEILPKALTRLKTVISISTKDVISVKESSESLCDIDLLVNVPDLKYRYVG	64	<a href="#">Q7TD01</a>
Shanghai	---MSYEPKVSDFLALTTKKEEILPKALTRLKTVISISTKDVISVKESSESLCDIDLLVNVPDLKYRYVG	64	<a href="#">Q91PA6</a>
722	---MSYEPKVSDFLALTTKKEEILPKALTRLKTVISISTKDVISVKESSESLCDIDLLVNVPDLKYRYVG	64	<a href="#">A0AP61</a>
Wh	---MSYEPKVSDFLALTTKKEEILPKALTRLKTVISISTKDVISVKESSESLCDIDLLVNVPDLKYRYVG	64	<a href="#">B2LS15</a>
RMV (Zhu/Yu)	---MSYEPKVSDFLALTTKKEEILPKALTRLKTVISISTKDVISVKESSESLCDIDLLVNVPDLKYRYVG	64	<a href="#">Q9QD18</a>
Kons.1105	MSMVSYPEKVNDFLFTLNKEKILPKALTRLKTVISISTKDIISVKESSETLCDIDLLIDVPLDKYRYVG	67	<a href="#">F8SRF8</a>
Act.-AC/AD	MSMVSYPEKVNDFLFTLNKEKILPKALTRLKTVISISTKDIISVKESSETLCDIDLLIDVPLDKYRYVG	67	<a href="#">E0WXS9</a>
FSHS	MSMVSYPEKVNDFLFTLNKEKILPKALTRLKTVISISTKDVISVKESSETLCDIDLLIDVPLDKYRYVG	67	<a href="#">I7A7B2</a>
NZ-439	MSMVSYPEKVNDFLFTLNKEKILPKALTRLKTVISISTKDIISVKESSETLCDIDLLIDVPLDKYRYVG	67	<a href="#">F8SRF4</a>
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TVCV OSU	ILGAVFTGEWLVPDFVKGGVTISVIDKRLVNSKECVIGTYRAAAAKSKRFQFKLVPNYFVSTVDADAKR	134
NZ-587	ILGAVFTGEWLVPDFVKGGVTISVIDKRLVNSKECVIGTYRAAAAKSKRFQFKLVPNYFVSTADAKR	134
NZ-438	ILGAVFTGEWLVPDFVKGGVTISVIDKRLVNSKECVIGTYRAAAAKSKRFQFKLVPNYFVSTADAKR	134
Pet-MS	ILGAVFTGEWLVPDFVKGGVTISVIDKRLVNSKECVIGTYRAAAAKSKRFQFKLVPNYFVSTVDADAKR	134
D/A	ILGAVFTGEWLVPDFVKGGVTISVIDKRLANSKECVIGTYRAAAAKSKRFQFKLVPNYFVSTVDADAKR	134
YoMV (ORMV)	VLGVVFTGEWLVPDFVKGGVTVSVIDKRLSNSKECI GITYRAAAKDRRFQFKLVPNYFVSVADAKR	131
Shanxi	VLGVVFTGEWLVPDFVKGGVTVSVIDKRLSNSKECI IGTYRAAAKDRRFQFKLVPNYFVSVADAKR	131
Br	VLGVVFTGEWLVPDFVKGGVTVSVIDKRLSNSRECI IGTYRAAAKDRRFQFKLVPNYFVSVADAKR	131
R/W	VLGVVFTGEWLVPDFVKGGVTVSVIDKRLSNSRECI IGTYRAAAKDRRFQFKLVPNYFVSTADAKR	131
Shanghai	VLGVAFTEGWLPDFVKGGVTVSVIDKRLSNSRESMIGTYRAAAKDRRFQFKLVPNYFVSTADAKR	131
722	VLGVVFTGEWLVPDFVKGGVTVSVIDKRLSNSKECI IGTYRAAAKDRRFQFKLVPNYFVSVADAKR	131
Wh	VLGVVFTGEWLVPDFVKGGVTVSVIDKRLSNSKECI IGTYRAAAKDRRFQFKLVPNYFVSVADAKR	131
RMV (Zhu/Yu)	VLGVAFTEGWLPDFVKGGVTVSVIDKRLSNSRESMIGTYRAAAKDRRFQFKLVPNYFVSTADAKR	131
Kons.1105	ILGAVFTGEWLVPDFVKGGVTISVIDKRLVNSKECVIGTYRAAAAKSKRFQFKLVPNYFVSTADAKR	134
Act.-AC/AD	ILGAVFTGEWLVPDFVKGGVTISVIDKRLVNSKECVIGTYRAAAAKSKRFQFKLVPNYFVSTADAKR	134
FSHS	ILGAVFTGEWLVPDFVKGGVTISVIDKRLVNSKECVIGTYRAAAAKSKRFQFKLVPNYFVSTADAKR	134
NZ-439	ILGAVFTGEWLVPDFVKGGVTISVIDKRLVNSKECVIGTYRAAAAKSKRFQFKLVPNYFVSTADAKR	134

[illegible]

<i>TVCV OSU</i>	FKAVDNF <b>RRKKKK</b> VEERDVV-SKYKYRPEKYAGPDSFNLKEENVLQHYKPESVPVLRSGVGRAHTNA	267
<i>NZ-587</i>	FKAVDNF <b>RRKKKK</b> VEEKGVV-SKYKYRPEKYAGPNSFNFKEENVLQHYEPESVPVFRSGVGRAHTNA	267
<i>NZ-438</i>	FKAVDNF <b>RRKKKK</b> VEEKGVV-RKYKYRPEKYAGPNSFNLKEENVLQDYEPESVPVFRSGVGRAHTNA	267
<i>Pet-MS</i>	FKAVDNF <b>RRKKKK</b> VEERDVV-SKYKYRPEKYAGPDSFNLKEENALQHYKPESVPVLRSGVGRAHTNA	267
<i>D/A</i>	FKAVDNF <b>RRKKKK</b> VEEKGVV-SKYKYRPEKYAGPDSFNLKEENVLQHYKPESVPVFRSGVGRAHSDA	267
<i>YoMV (ORMV)</i>	FKAIDSF <b>RRKKKR</b> IGGRDVNSNKYRYRPERYAGPDSLQYKEENGLQHHELESVPVFRSDVGRAHSDA	265
<i>Shanxi</i>	SKAIDSF <b>RRKKKR</b> IGGKDVNNKKYRYRPERYAGPNSLQYKEGNDLQHHELESVPVFRSDVGRAHGD	265
<i>Br</i>	FKAIDSF <b>RRKKKK</b> IGGKDVNNKKYRYRPERYAGPDSLQYKEENVLQHHELESVPVFRSDVGRAHSDA	265
<i>R/W</i>	FKAIDSF <b>RRKKKK</b> IGGKDVNNKKYRYRPERYAGPDSLQYKEENGLQHHELESVPVFRSDVGRAHSDA	265
<i>Shanghai</i>	FKAIDSF <b>RRKKKK</b> IGGRDVN-NKYRYRPERYAGPDSLQYKEENGLQHHELESVPVFRSDVGRAHSDA	264
<i>722</i>	FKAVDTF <b>RRKKKK</b> IGGKDVNNKKYRYRPERYAGDSLQYKEENVLQHHELESVPVFRSDVGRAHSDA	265
<i>Wh</i>	FKAIDSF <b>RRKKKK</b> IGGKDVSNKKYRYRPETYAGPDSLQYKEGNVLQHHELESVPVFRSNVGRAHSDA	265
<i>RMV (Zhu/Yu)</i>	FKAIDSF <b>RRKKKK</b> IGGRDVNNKKYRYRPERYAGPDSLQYKEENGLQHHELESVPVFRSDVGRAHSDA	265
<i>Kons.1105</i>	FRAVDNF <b>RRKKKR</b> IGEKEAV-SRNKYRPEKHAGPNSLYIKEENVLQHHELESVPVLRSGVGRTHNA	267
<i>Act.-AC/AD</i>	FKAVDNF <b>RRKKKR</b> IGGSEVV-SKNKYRPEKHAGPNSLYIKEENVLQHHELESVPVFRSGVGRAHTNA	267
<i>FSHS</i>	FKAVDNF <b>RRKKKR</b> IGEKEVV-SRNKYRPEKHAGPNSLYIKEENVLQHHELESVPVLRSGVGRTYTNA	267
<i>NZ-439</i>	FKAVDNF <b>RRKKKR</b> IGEKEVV-SRNKYRPEKHAGPNSLYIKEENVLQHHELESVPVLRSGVGRTHNA	267
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**Figure S1. Conservation of MP<sup>TVCV</sup> NLS among all subgroup 3 Tobamoviruses**

Alignment of all sequenced MP proteins from the subgroup 3 Tobamoviruses (TVCV, YoMV and RMV). Conservation of monopartite NLS identified in TVCV in this study (TVCV OSU, top line) as predicted among all available subgroup 3 virus MP sequences is highlighted in yellow (basic residues) with additional residues predicted to be involved in importin- $\alpha$  interactions ([http://nls-mapper.iab.keio.ac.jp/cgi-bin/NLS\\_Mapper\\_form.cgi/](http://nls-mapper.iab.keio.ac.jp/cgi-bin/NLS_Mapper_form.cgi/)) shaded in grey. Accession numbers for each MP sequence is shown at the right of the first line of sequence, as are the residue positions within each protein (every line). Sequence similarity indicated below the aligned sequences. See text for details.

TMGMV (TMV U2)	MAVSLRDTVKISEFINLSKQDEILPAFMTKVKSVRISTVDKIMAVKNDLSLSDVDLLKGVKLVKNGYV	67	P18338
U2	MAVSLRDTVKISEFIDLSKQDEILPAFMTKVKSVRISTVDKIMAVKNDLSLSDVDLLKGVKLVKNGYV	67	A4LAL8
Japanese	MAVSLRDTVKISEFIDLSKQDEILPAFMTKVKSVRISTVDKIMAVKNDLSLSDVDLLKGVKLVKNGYV	67	Q8V701
HP	MAVSLRDTVKISEFIDLSKQDEILPAFMTKVKSVRISTVDKIMAVKNDLSLSDVDLLKGVKLVKNGYV	67	A3QVG8
	*****;*****		
TMGMV (TMV U2)	CLAGLVVSGEWNLPDNCRGGVSVCIVDKRMKRSNEATLGAYHAPACKKNFSFKLIPNYSITSEDAEK	134	
U2	CLAGLVVSGEWNLPDNCRGGVSVCIVDKRMKRSKEATLGAYHAPACKKNFSFKLIPNYSITSEDAEK	134	
Japanese	CLAGLVVSGEWNLPDNCRGGVSVCIVDKRMKRSKEATLGAYHAPACKKNFSFKLIPNYSITSEDAEK	134	
HP	CLAGLVVSGEWNLPDNCRGGVSVCIVDKRMKRSKEATLGAYHAPACKKNFSFKLIPNYSITSEDAEK	134	
	*****;*****		
TMGMV (TMV U2)	NPWQVLVNIKGVAMEEGYCPLSLEFVSICVVHKNNVRKGLRERILRVTDGSPIELTEKVVEEFVDEV	201	
U2	HPWQVLVNIKGVAMEEGYCPLSLEFVSICVVHKNNVRKGLRERILSVTDGSPIELTEKVVEEFVDEV	201	
Japanese	HPWQVLVNIKGVAMEEGYCPLSLEFVSICVVHKNNVRKGLRERILRVTDGLPIELTEKVVEEFVDEV	201	
HP	HPWQVLVNIKGVAMEEGYCPLSLEFVSICVVHKNNVRKGLRERILRVTDGLLIELTEKVVEEFVDEV	201	
	*****;*****		
TMGMV (TMV U2)	PMAVKLERFRKTKKGGKKRKKKKRVVGNVNNKKINNSGKKGLKVEEIEDNVSDDESIIASSSTF	266	
U2	PMAVKLERFRKTK-----KKVVGNNVNNKKINNSGKKGFKIEEIEDNVSDDESIIASSSTF	256	
Japanese	PMAVKLERFRKTK-----KRVVGNVNNKKINNSGKKGLKVEEIEDSVSDDESIIASSSTF	256	
HP	PMAVKLERFRKTK-----KRVVGNVNNKKINNSGKKGLKVEEIEDSVSDDESIIASSSTF	256	
	*****;*****;*****;*****;*****;*****		

**Figure S2. A putative basic monopartite NLS is not conserved among TMGMV isolates.** Alignment of MP protein sequences from the four sequenced isolates of *Tobacco mild green mosaic virus* (TMGMV). The monopartite NLS predicted to be present in one isolate, and to correspond in sequence and position to the MP<sup>TVCV</sup> NLS, is highlighted in yellow with additional residues predicted to be involved in importin- $\alpha$  interactions ([http://nls-mapper.iab.keio.ac.jp/cgi-bin/NLS\\_Mapper\\_form.cgi/](http://nls-mapper.iab.keio.ac.jp/cgi-bin/NLS_Mapper_form.cgi/)) shaded in grey. Accession numbers for each MP sequence are shown at the right of the first line of sequence, as are the residue positions within each protein (every line). Sequence similarity indicated below the aligned sequences. See text for details.