

Virus	Genus	aa position	eee	hhhhhhh	e	eeeeeeee
TSWV	<i>Tospovirus</i>	111	--ANKV	KICP---	TWD---	SRKQY-----
RCNMV	<i>Dianthovirus</i>	33	--GVSE	APLIPA---	SMMSKITDYA---	KTTAK-----
SBWMV	<i>Furovirus</i>	44	--RYVE	KALTOP---	GVVTQMQDAW---	TNWTKTNKEEGTPYN-----
CMoV	<i>Umbravirus</i>	42	--TLHT	PLYP---	RRQLTLSRFF---	SGRWR-----
CNSV	<i>Nepovirus B</i>	432	-----	VLNRP	-----	YNALNKLA-----
RSV	<i>Tenuivirus</i>	88	-----	ATFNILV	-----	DPYWFHQ-----
ACLSV	<i>Tricovirus</i>	62	-----	RLASIPVIP	-----	SSEVQAVL-----
SDV	<i>Sadwavirus</i>	53	-----	QQQDIAIPL	-----	SSRF-----
PYMoV	<i>Badnavirus</i>	100	-----	NQVDRFTIQ	-----	EESHKLV-----
CLBV	<i>Citriovirus</i>	55	-----	IKLANILT	-----	ADRLQSI-----
PYFV	<i>Sequivirus</i>	115	-----	RAEITPTA	-----	ASKLDKTL-----
RBDV	<i>Idaeovirus</i>	138	-----	GKNEFLV	-----	ASVFDKLI-----
RRV	<i>Emaravirus</i>	63	-----	KTQLLPISV	-----	YNEMYKFY-----
TMV	<i>Tobamovirus</i>	36	CSKVDK	KIMVHENESLSEVNLLKGVK	-----	YNEFYK-----
CPsV	<i>Ophiovirus</i>	84	-----	HKKKLKLGT	-----	LKSIDTKL-----
CPMV	<i>Comovirus</i>	204	-----	TVLDIPVTK	-----	DKTAMAS-----
CiLV-C	<i>Cilevirus</i>	47	-----	RIGVGLFTP	-----	NDMISRLVGF-----
PrLbAV		244	-----	PVYIPLVP	-----	ALA-SDIE-----
CRLV	<i>Cheravirus</i>	109	-----	DPYVAVYTPLP	-----	HITEEQARKLM-----
GFLV	<i>Nepovirus A</i>	118	-----	NTEILYNFV	-----	SEEMEYR-----
OuMV	<i>Ourmiavirus</i>	57	-----	VEDPIALIP	-----	HGIWSIFKSK-----
PoLV	<i>Aureovirus</i>	30	-----	TKAILPLAP	-----	ISQFSKWKI-----
TBSV	<i>Tombusvirus</i>	32	-----	EDQEVPLLP	-----	SNFLTQVVG-----
ToTV	<i>Torradovirus</i>	66	-----	KKAMELFD	-----	PEDVHNITSLWSSPKKFTTSS-----
CaMV	<i>Caulimovirus</i>	97	-----	GKVVPLIIT	-----	KEEINKRLSS-----
CMV	<i>Cucumovirus</i>	52	-----	AISVRPLVP	-----	EVTHGRIASF-----
AMV	<i>Alfamovirus</i>	60	-----	APVVLVLSVS	-----	KEKKSILNRM-----
LNyV	<i>Cytohabdovirus</i>	53	-----	IMRQVPLFD	-----	KEDLDAMK-----
AcVB	<i>Vitivirus</i>	78	-----	VDQNILDVP	-----	EKTIDGLN-----
CaLCuV	<i>Begomovirus</i>	38	-----	LTEILQF	-----	PSTAAQFT-----

Virus	Genus	aa position	eeeeee	ee eeeee	eeee
TSWV	<i>Tospovirus</i>	IPN	-----	PTGKLVVALVDPNMP	-----
RCNMV	<i>Dianthovirus</i>	IGV	-----	ATPGHVTVLIPNPNVEG	-----
SBWMV	<i>Furovirus</i>	VPM	-----	GYAGTVEVSLDLSGLSP	-----
CMoV	<i>Umbravirus</i>	VPD	-----	TSPQAVIVVHDTALPG	-----
CNSV	<i>Nepovirus B</i>	LPO	-----	QMNAYAFCVIMWGHSSD	-----
RSV	<i>Tenuivirus</i>	-----	-----	RASGITTLRIIDKSYVN	-----
ACLSV	<i>Tricovirus</i>	FRK	-----	NAGVSGWCYVYDNRWET	-----
SDV	<i>Sadwavirus</i>	VLT	-----	GVDTAMAIMVSDASHDR	-----
PYMoV	<i>Badnavirus</i>	HRK	-----	EEGTLALVFRDNRWKG	-----
CLBV	<i>Citriovirus</i>	LPS	-----	GNEQIWRGRCALVDTSRTR	-----
PYFV	<i>Sequivirus</i>	MHQ	-----	GDSRECIIMYLCDRRFKD	-----
RBDV	<i>Idaeovirus</i>	VGK	-----	GTPGTARIKIRIDARLDD	-----
RRV	<i>Emaravirus</i>	SK	-----	RNEMATLLIMDERFRDDG	-----
TMV	<i>Tobamovirus</i>	LPD	-----	NCRGVSVCVLDKRMER	-----
CPsV	<i>Ophiovirus</i>	FSRV	-----	DGNGEITVSLIDDGKEA	-----
CPMV	<i>Comovirus</i>	ASP	-----	ESDLMGGFLLVDSLHTD	-----
CiLV-C	<i>Cilevirus</i>	IA	-----	LPAELSLKLVDSAN	-----
PrLbAV		LAE	-----	SSGMRRTSFSVEDASQ	-----
CRLV	<i>Cheravirus</i>	VGO	-----	GTPLLAMCGIMDSRTDD	-----
GFLV	<i>Nepovirus A</i>	GMP	-----	GNPDTLVVATYGHEDR	-----
OuMV	<i>Ourmiavirus</i>	VAT	-----	GLARGQIAVVDTRVNHTS	-----
PoLV	<i>Aureovirus</i>	ISE	-----	RAQVRGVKLVDSRDL	-----
TBSV	<i>Tombusvirus</i>	VSR	-----	KAGVSGKLYLRDISDT	-----
ToTV	<i>Torradovirus</i>	VHG	-----	GEGSVKISLCSNSDPTN	-----
CaMV	<i>Caulimovirus</i>	FRN	-----	GIDTPIKIALIDDRINS	-----
CMV	<i>Cucumovirus</i>	VST	-----	DAEGSLRIYLADLGDKE	-----
AMV	<i>Alfamovirus</i>	ILK	-----	SSSGSITLKLFEATGE	-----
LNyV	<i>Cytohabdovirus</i>	LHQRYMKNFGKT	-----	IAGNCAIDSTFRK	-----
AcVB	<i>Vitivirus</i>	GR	-----	AMKGTQIKVEDQRFKE	-----
CaLCuV	<i>Begomovirus</i>	VPI	-----	NATGVSIVEIHDKRTMD	-----

Virus	Genus	aa position	eee	eeeeeee	eeeeee
TSWV	<i>Tospovirus</i>	-----	WSIPKMNNTP	-----	ENCCQLHLMCSQ
RCNMV	<i>Dianthovirus</i>	-----	HQLNSDH	-----	EPRVRITNTG
SBWMV	<i>Furovirus</i>	-----	YSIPLNDKGR	-----	AVKLAFKIDAEM
CMoV	<i>Umbravirus</i>	-----	LCEIPLSDSAMN	-----	APRCFSLVTLQEGVRLEQG
CNSV	<i>Nepovirus B</i>	-----	LYEYVGHNLQDF	-----	EAYKRSVLVSTVPEFSGIADGKAMFGITSIEF
RSV	<i>Tenuivirus</i>	-----	FLALEDK	-----	HNLQVSVVDDSSVQNC
ACLSV	<i>Tricovirus</i>	-----	FPVSLDDPGL	-----	SNSISVAVMFENLNFKFE
SDV	<i>Sadwavirus</i>	-----	ITVFPDPLV	-----	DRFLRLSVLTGGSPMADG
PYMoV	<i>Badnavirus</i>	-----	IMMVTNDFY	-----	HNTHISVLTTRGVEGWQHEANLLVTRGM
CLBV	<i>Citriovirus</i>	-----	NCKVSVG	-----	SIQVLELHGVDLREE
PYFV	<i>Sequivirus</i>	-----	EATAAD	-----	RLQVLYVKGAKLERTONT
RBDV	<i>Idaeovirus</i>	-----	YSVPKSEI	-----	DYKINIDFAGVPIKDG
RRV	<i>Emaravirus</i>	-----	QFVTGSLDFATDKDI	-----	NKIKFYISFPDTRM
TMV	<i>Tobamovirus</i>	-----	YAITTQDAM	-----	KNVQVVLVNIIRNVKMSAG
CPsV	<i>Ophiovirus</i>	-----	FFVEKDM	-----	DFIGIHVSAENVVQD
CPMV	<i>Comovirus</i>	-----	TLAPVSCDL	-----	NNRFKLCISLPNCDIVQG
CiLV-C	<i>Cilevirus</i>	-----	HISIPNEDRTHVNG	-----	SEVHRRLLGIYQVDCDNISGR
PrLbAV		-----	NDSDHKSNS	-----	TPQPELSFIFGLPKVS
CRLV	<i>Cheravirus</i>	-----	INFLPNKEDFD	-----	DYMRGLYLCTMFHNVRGFQNNKALCSYSA
GFLV	<i>Nepovirus A</i>	-----	LQYSQEP	-----	ESIIRLVASTNATVTD
OuMV	<i>Ourmiavirus</i>	-----	YCLFHPKKEGGDKSLESQNP	-----	IRGVIIYDTSRY
PoLV	<i>Aureovirus</i>	-----	FCLPVG	-----	DYPLQFVTVLQSQFRET
TBSV	<i>Tombusvirus</i>	-----	FSVSTRSD	-----	VPIVFGFEELVSPFLEG
ToTV	<i>Torradovirus</i>	-----	ITLFPVKRG	-----	PMFYIMECLGTRA
CaMV	<i>Caulimovirus</i>	-----	FGISLNTQRL	-----	NQTLSLIHDFENKLNMGKDVMTITYV
CMV	<i>Cucumovirus</i>	-----	YDCPMETVGN	-----	RRKCFVAVIERHGYIQT
AMV	<i>Alfamovirus</i>	-----	RSILAKDAK	-----	GHDKLVVHAVASTNAN
LNyV	<i>Cytohabdovirus</i>	-----	HCLSLDPMI	-----	QKRLSVLLGIKIGDVEPG
AcVB	<i>Vitivirus</i>	-----	YFVSTTDVKN	-----	GYALNKLKVAEGMCMVEG
CaLCuV	<i>Begomovirus</i>	-----	FFSLKDP	-----	IPKLYRVSVDTNVHQ

Figure S3. Multiple sequence alignment of movement proteins from PrLbAV and plant viruses of the 30K superfamily. Sequences are from representative members of the virus genera in

the 30K superfamily. PrLBaV is underlined. The GenBank identifier, virus genus and the distance in aminoacids from the N-terminus of the protein are shown on the right side of each sequence. Consensus secondary structure elements, predicted by PSIREN within PROMALS3D program, are reported on the top with strands and helices indicated with e and h, respectively. The nearly invariant aspartic acid residue (the D motif) is in bold.