

**Table 1**

Comparisons of amino acid sequence identity of the viral proteins or motif domains between BdMoV and related virus sequences available in the GenBank database.

Virus <sup>a</sup>	RNA1 (replicase, 249 kDa protein)				RNA2						RefSeq/Accession no.
	MTR	HEL	Pro	RdRp	CP	RTD	TGB1	TGB2	TGB3	CRP	
BNYVV	62 <sup>b</sup>	67	44	81	39	24	48	48	38 <sup>8</sup>	nh <sup>c</sup>	NC_003514, NC_003515
BSBMV	58	67	42	80	35	31	46	39	34	nh	NC_003506, NC_003503
RNSV	44	66	33	76	31	nh	42	43	nh	nh	EU099844, EU099845
NVMV*	ud <sup>c</sup>	ud	ud?	ud	nh	–?	32	38	nh	–?	D00906
CAV*	ud	30	– <sup>c</sup>	35	nh	–	–	–	–	–	JF824737
HEV	nh	nh	nh	30	nh	–	–	–	–	–	M74506
DpTV	nh	25	–?	25	nh	–	–	–	–	–	NC_005898, NC_005899

<sup>a</sup> Unassigned viruses are indicated by their abbreviation with asterisks (see [Supplementary Table S2](#)).

<sup>b</sup> The percent of amino acid sequence identity in the sequence comparison analysis is shown. Each scores are calculated by the BLASTp with the expect value significance cut-off below 10 (the default parameter setting).

<sup>c</sup> nh: no hit against the corresponding target; ud: undetermined ; –: not present.