

Table S1. Record of reference isolates, their host species and geographic origin, used to generate probes through specific RT-PCR. Virus acronyms are as follows: GarVA, -B, -C, and -X = *garlic virus A, -B, -C, and -X*, GCLV = *garlic common latent virus*, SLV = *shallot latent virus*, OYDV = *onion yellow dwarf virus*, SYSV = *shallot yellow stripe virus* and LYSV = *leek yellow stripe virus*.

Isolate number	Host species	Country of Origin	Virus specific probes
3185	<i>Allium sativum</i> , garlic	Gatton, Australia	GarVA, GarVX, GCLV, OYDV
3188	<i>Allium sativum</i> , garlic	Gatton, Australia	GarVC
3191	<i>Allium sativum</i> , garlic	Gatton, Australia	GarVB, SLV
Q4608	<i>Allium cepa</i> , shallot	Cirebon, Indonesia	SLV, SYSV
Q4609	<i>Allium cepa</i> , shallot	Indonesia (IVEGRI)	SLV
PC-0451 DSMZ	<i>Allium ampeloprasum</i> var. <i>porrum</i> , leek	Indonesia	LYSV
PV-0622	<i>Allium ascalonicum</i> , shallot	Slovenia	ShVX

Table S2 Primer sequence information for amplification of degenerate PCR products for the allexi-, carla-, and potyviruses.

Primer	Sequence (5'-3')	Target	Product size w/ Poty1 (bp)	Reference	Use
Poty1	GGATCCCGGGTTTTTTTTTTTTTTT TV	Poly A tail	na	Gibbs and MacKenzie, 1997	cDNA and reverse
AlcarF	TGCTGCYTTTGATACYTTCGAT	Carlavirus es	715	This study	Forward
pGV3t	TGGNCNTGCTACCACAANGG	Allexivirus es	950	Chen et al, 2004	Forward
U341	CCGGAATTCATGRTITGGTGYATIG AIAAYGG ¹	Potyviruse s	600-850	Langeveld et al., 1991	Forward

Table S3 Primer sequence information for PCR labelling of species-specific probes

Primer name	Sequence (5'-3')	Genus	Product size (bp)
GarVA_F	TCGTCGAAAACCTATTGCACCT	Allexi	507
GarVA_R	GAAGTGCCTTGAAATGGCTTA		
GarVB_F	GGGCAAACAGCAGAATAAGC	Allexi	610
GarVB_R	ATAGCTTAGCGGGTCCTTCA		
GarVC_F	GCTGAACTCAAAGACTTAGT	Allexi	509
GarVC_R	TCTTACGTCCTAATTTCA		
GarVX_F	CACCTTGTGGCATGCCTC	Allexi	500
GarVX_R	GCTTTACGTCCTAATTTTC		
ShVX_F	GTGAGTGTCTTTGGTCACGT	Allexi	500
ShVX_R	ACTAGAGCCGACCTCCTAGG		
GCLV_F	GGCAAACCAAAGGTTACACT	Carla	572
GCLV_R	GTAAGATACCTAATCGTCCC		
SLV_F	GCTTTCGCAACACAAGAA	Carla	500
SLV_R	TCGTTCCCACTTATACATTA		
LYSV_F	AGCAGGCTTATATGCCAAGG	Poty	550
LYSV_R	AAAACCGTACTCCGGGTAGA		
OYDV_F	TCAGGGAATTGGACTATGATG	Poty	535
OYDV_R	TCAGATTAAACGGAAATGAAGA		
SYSV_F	TTGGACGATGATGGATGGCG	Poty	500
SYSV_R	CTATGCCACAAACGGTGTT		
IYSV_F	GTTTAGAAGACTCACCAATG	Tospo	837
IYSV_R	ACCAGAGGAAGCCCGCAG		

Table S4 Primer sequence information for amplicon sequencing of the coat protein (CP) of nine garlic virus species

Primer name	Sequence (5'-3')	Genus	Product size (bp)
GVA_CP_F	ACTGGAAGGGTGAATTAGATGAGT	Allexi	854
GVA_CP_R	TGGCAAARTGYAGGCAGCAA	Allexi	
GVB_CP_F	TCAGGGATGATGGTTTGTGTT	Allexi	878
GVB_CP_R	GGGAGRTTGGGTTGAGCAAA	Allexi	
GVC_CP_F	TCGTGAATGCGGCCATGACT	Allexi	916
GVC_CP_R1	GTGCCCTGGAAGGGCTTATT	Allexi	
GVC_CP_R2	GTGAAATGAAGGCAGCAGAG	Allexi	772
GVX_CP_F	TGCAACCATTGAGCGACTCT	Allexi	910
GVX_CP_R	TGAAGACAGGCGAGCARGTT	Allexi	
GCLV_CP_F	ACGCTTGAGGTAGCCAAACA	Carla	1073
GCLV_CP_R	TAGGCACGCCACTAYCACAC	Carla	
SLV_CP_F	ACAGCGCTCTAAATTGATATTATGACT	Carla	935
SLV_CP_R	ACGGTAAGTTCTYTGTTGATCA	Carla	
LYSV_CP_F	GATGCACCTCCGGATGAAA	Poty	938
LYSV_CP_R	TTTAAGTGGAAGTCYATAACCGRT	Poty	
OYDV_CP_F	GCGGGGGAAGGAGAAGATGCAG	Poty	730
OYDV_CP_R	CCGCAGCTGTGTGTCTTTCCG	Poty	
SYSV_CP_F	AATCCTGGATGATCGTATACCG	Poty	923
SYSV_CP_R	ACTATGCCACAAACGYTGGT	Poty	

Table S5 Distance matrix showing percent nucleotide identities of coat protein sequences of potyviruses. Australian isolates from this study (Eastern Australia) are shown in orange and from Western Australia in light blue (GenBank accessions).

4	Y11826	MH686302	MH686301	MH686300	KY347900	KR025485	KF862691	KF632715	KF632714	KF623541	KF623540	JX433020	JX433019	JX429964	JN127342	HQ918256	HQ873798	HQ258894	GQ475410	F1765739	DQ925455	AY390253	AJ292224	KF597285	JX429967	JN127340	GQ475418	DQ402056	DQ293982	DQ925458	AB693978	AB669435
Y11826 OYDV		82.13	82.04	81.97	96.29	98.07	82.64	81.45	82.94	87.63	95.19	98.98	96.06	81.67	82.25	84.31	82.82	82.28	82.07	82.96	82.49	83.98	82.67	62.54	61.79	62.28	61.9	63.79	59.88	74.37	74.45	74.23
MH686302 OYDV	82.13		86.9	99.7	81.53	81.38	85.44	86.34	86.49	81.56	82.13	81.98	81.98	86.41	98.8	81.61	86.96	87.24	86.81	80.06	93.39	81.03	80.06	63.1	62.52	63.17	63.27	63.32	60.16	72.13	71.85	72.15
MH686301 OYDV	82.04	86.9		86.6	81.58	81.43	86.91	86.3	87.22	80.4	81.89	82.04	81.28	98.25	87.52	84.13	86.47	99.7	85.71	84.04	86.61	85.3	84.65	62.16	62.31	62.52	62.78	62.39	60.34	75.38	73.8	74.78
MH686300 OYDV	81.97	99.7	86.6		81.36	81.21	85.76	86.67	86.52	81.39	81.97	81.82	81.82	86.14	98.79	81.52	86.99	86.82	86.84	79.88	93.33	80.85	79.88	62.78	62.19	62.84	62.94	63.29	60.16	72.33	72.35	72.35
KY347900 OYDV	96.29	81.53	81.58	81.36		95.85	81.6	81.16	81.31	87.56	94.66	96.44	94.96	81.01	81.75	82.66	82.37	81.75	81.63	82.07	82.64	83.36	82.22	62.37	61.79	62.28	62.1	63.89	60.34	73.92	74.45	73.94
KR025485 OYDV	98.07	81.38	81.43	81.21	95.85		82.94	81.16	82.05	87.26	95.25	98.22	95.25	80.86	81.31	83.26	82.52	81.6	81.78	82.37	81.75	83.67	82.07	62.23	62.08	61.98	61.96	63.89	60.34	74.37	74.74	74.09
KF862691 OYDV	82.64	85.44	86.91	85.76	81.6	82.94		91.1	86.94	81.93	82.79	82.34	81.6	86.28	85.61	81.91	90.96	87.24	90.22	82.52	85.31	83.05	83.26	62.81	62.23	63.17	62.25	64.6	60.95	74.37	73.42	73.5
KF632715 OYDV	81.45	86.34	86.3	86.67	81.16	81.16	91.1		85.76	80.59	81.6	81.45	80.71	85.98	86.2	81.31	97.48	86.8	96.74	83.11	85.46	83.67	83.56	62.37	60.76	62.13	61.1	61.15	60.34	73.92	73.86	72.91
KF632714 OYDV	82.94	86.49	87.22	86.52	81.31	82.05	86.94	85.76		81.2	80.91	82.14	81.55	86.61	87.28	86.56	85.93	87.43	85.19	82.67	85.46	83.83	82.96	64.91	62.08	63.17	63.3	64.35	60.51	73.32	73.42	74.38
KF623541 OYDV	87.63	81.56	80.4	81.39	87.56	87.26	81.93	80.59	81.2		88.57	87.86	87.43	80.61	81.68	81.93	81.66	80.97	80.92	80.33	81.63	81.52	80.62	62.1	60.91	62.43	61.34	61.94	58.51	72.43	73.17	73.21
KF623540 OYDV	95.19	82.13	81.89	81.97	94.66	95.25	82.79	81.6	80.91	88.57		95.72	96.58	81.61	82.1	82.36	82.67	82.13	81.93	82.22	82.05	83.52	82.22	62.39	62.23	62.13	62.14	63.57	58.73	73.77	73.42	73.06
JX433020 OYDV	98.98	81.98	82.04	81.82	96.44	98.22	82.34	81.45	82.14	87.86	95.72		96.53	81.75	82.1	83.56	82.82	82.28	82.07	82.37	82.34	83.67	82.37	62.25	61.64	62.28	61.67	63.48	59.73	74.37	73.86	74.23
JX433019 OYDV	96.06	81.98	81.28	81.82	94.96	95.25	81.6	80.71	81.55	87.43	96.58	96.53		80.88	81.95	82.81	81.78	81.39	81.04	81.63	82.34	83.36	81.93	62.39	62.23	61.54	61.43	62.67	58.46	73.47	72.98	73.5
JX429964 OYDV	81.67	86.41	98.25	86.14	81.01	80.86	86.28	85.98	86.61	80.61	81.61	81.75	80.88		87.06	83.78	86	98.15	85.56	83.41	85.68	84.45	84	61.95	61.64	62.5	62.7	62.61	60.09	74.67	73.42	74.52
JN127342 OYDV	82.25	98.8	87.52	98.79	81.75	81.31	85.61	86.2	87.28	81.68	82.1	82.1	81.95	87.06		81.91	86.96	87.87	86.52	80.59	93.18	81.49	80.89	63.45	62.81	63.17	63.65	63.27	59.63	71.98	71.95	71.89
HQ918256 OYDV	84.31	81.61	84.13	81.52	82.66	83.26	81.91	81.31	86.56	81.93	82.36	83.56	82.81	83.78	81.91		81.76	84.16	81.91	83.26	81.91	85.15	83.26	62.96	60.62	62.28	62.34	63.99	61.49	73.32	72.85	73.44
HQ873798 OYDV	82.82	86.96	86.47	86.99	82.37	82.52	90.96	97.48	85.93	81.66	82.67	82.82	81.78	86	86.96	81.76		86.96	99.26	83.11	86.67	83.67	83.41	62.37	60.76	62.43	61.24	62.16	60.95	74.52	74.6	73.65
HQ258894 OYDV	82.28	87.24	99.7	86.82	81.75	81.6	87.24	86.8	87.43	80.97	82.13	82.28	81.39	98.15	87.87	84.16	86.96		86.22	84.15	86.35	85.38	84.74	62.48	62.37	62.87	62.98	62.75	60.55	74.96	73.72	74.38
GQ475410 OYDV	82.07	86.81	85.71	86.84	81.63	81.78	90.22	96.74	85.19	80.92	81.93	82.07	81.04	85.56	86.52	81.91	99.26	86.22		82.37	86.22	83.2	82.67	62.23	60.62	62.28	61.1	61.58	60.95	74.67	74.45	73.5
F1765739 OYDV	82.96	80.06	84.04	79.88	82.07	82.37	82.52	83.11	82.67	80.33	82.22	82.37	81.63	83.41	80.59	83.26	83.11	84.15	82.37		80.74	97.51	97.63	61.64	61.2	61.98	60.81	62.88	59.72	73.77	72.1	73.61
DQ925455 OYDV	82.49	93.39	86.61	93.33	82.64	81.75	85.31	85.46	85.46	81.63	82.05	82.34	82.34	85.68	93.18	81.91	86.67	86.35	86.22	80.74		81.8	80.59	62.52	62.37	62.87	62.97	63.02	60.18	72.28	72.54	73.35
AY390253 OYDV	83.98	81.03	85.3	80.85	83.36	83.67	83.05	83.67	83.83	81.52	83.52	83.67	83.36	84.45	81.49	85.15	83.67	85.38	83.2	97.51	81.8		97.36	63.05	62.6	63.36	62.14	64.79	61.73	75	74.38	75.62
AJ292224 OYDV	82.67	80.06	84.65	79.88	82.22	82.07	83.26	83.56	82.96	80.62	82.22	82.37	81.93	84	80.89	83.26	83.41	84.74	82.67	97.63	80.59	97.36		62.67	62.37	61.98	61.82	62.73	59.57	74.07	72.54	74.34
KF597285 LYSV	62.54	63.1	62.16	62.78	62.37	62.23	62.81	62.37	64.91	62.1	62.39	62.25	62.39	61.95	63.45	62.96	62.37	62.48	62.23	61.64	62.52	63.05	62.67		89.91	82.01	92.63	79.09	80.79	61.79	61.02	62.28
JX429967 LYSV	61.79	62.52	62.31	62.19	61.79	62.08	62.23	60.76	62.08	60.91	62.23	61.64	62.23	61.64	62.81	60.62	60.76	62.37	60.62	61.2	62.37	62.6	62.37	89.91		80.81	93.78	79.88	81.42	62.37	60.88	61.99

JN127340 LYSV WA	62.28	63.17	62.52	62.84	62.28	61.98	63.17	62.13	63.17	62.43	62.13	62.28	61.54	62.5	63.17	62.28	62.43	62.87	62.28	61.98	62.87	63.36	61.98	82.01	80.81		81.14	79.67	82.8	61.69	62.48	62.13
GQ475418 LYSV	61.9	63.27	62.78	62.94	62.1	61.96	62.25	61.1	63.3	61.34	62.14	61.67	61.43	62.7	63.65	62.34	61.24	62.98	61.1	60.81	62.97	62.14	61.82	92.63	93.78	81.14		79.49	79.16	62.81	61.24	62.16
DQ402056 LYSV	63.79	63.32	62.39	63.29	63.89	63.89	64.6	61.15	64.35	61.94	63.57	63.48	62.67	62.61	63.27	63.99	62.16	62.75	61.58	62.88	63.02	64.79	62.73	79.09	79.88	79.67	79.49		82.1	60.97	61.53	60.63
DQ299382 LYSV	59.88	60.16	60.34	60.16	60.34	60.34	60.95	60.34	60.51	58.51	58.73	59.73	58.46	60.09	59.63	61.49	60.95	60.55	60.95	59.72	60.18	61.73	59.57	80.79	81.42	82.8	79.16	82.1		60.13	60.64	60.86
DQ925458 SYSV	74.37	72.13	75.38	72.33	73.92	74.37	74.37	73.92	73.32	72.43	73.77	74.37	73.47	74.67	71.98	73.32	74.52	74.96	74.67	73.77	72.28	75	74.07	61.79	62.37	61.69	62.81	60.97	60.13		90.05	90.18
A8693978 SYSV	74.45	71.85	73.8	72.35	74.45	74.74	73.42	73.86	73.42	73.17	73.42	73.86	72.98	73.42	71.95	72.85	74.6	73.72	74.45	72.1	72.54	74.38	72.54	61.02	60.88	62.48	61.24	61.53	60.64	90.05		88.87
A8669435 SYSV	74.23	72.15	74.78	72.35	73.94	74.09	73.5	72.91	74.38	73.21	73.06	74.23	73.5	74.52	71.89	73.44	73.65	74.38	73.5	73.61	73.35	75.62	74.34	62.28	61.99	62.13	62.16	60.63	60.86	90.18	88.87	

Table S6 Distance matrix showing percent nucleotide identities of coat protein sequences of carlaviruses. Australian isolates from this study (Eastern Australia) are shown in orange and from Western Australia in light blue (GenBank accessions).

	MH686306	MH686305	LC27526	JX429968	JX429966	JQ899443	JF320811	HQ258896	GU355922	GU120175	DQ520093	AJ409116	AB048002	AB046886	AB046886	AB04567	KP208802	KJ801307	KF862699	JQ899445	JQ818259	JF320810	HQ873863	HQ873861	MH686304	MH686303	DQ520092	AF228416	AB048004	AB04566
MH686306 SLV		92.58	78.11	89.22	91.44	86.47	89.21	92.34	97.48	88.62	89.33	92.58	87.65	94.01	94.01	89.09	55.75	54.82	55.05	54.36	55.05	54.01	54.94	54.82	55.52	55.75	55.17	54.82	54.01	55.75
MH686305 SLV	92.58		78.20	89.93	96.52	85.95	89.10	98.83	92.73	88.76	89.45	98.94	88.04	92.85	92.85	89.22	55.73	55.52	55.06	54.49	55.06	54.73	54.72	54.61	55.29	55.52	55.29	55.29	54.72	55.75
LC279526 SLV	78.11	78.20		79.39	79.87	78.51	78.92	78.10	78.92	78.83	79.63	78.31	80.56	79.51	79.51	79.39	54.02	54.28	53.62	52.85	53.84	53.46	52.96	52.85	53.51	53.51	54.06	54.28	53.51	53.29
JX429968 SLV	89.22	89.93	79.39		89.49	88.19	90.07	89.97	90.04	88.29	89.10	90.28	88.98	89.92	89.92	89.57	56.38	55.70	55.48	54.61	55.15	55.51	54.72	54.61	55.81	56.03	55.92	55.81	55.15	54.93
JX429966 SLV	91.44	96.52	79.87	89.49		87.89	92.17	96.81	91.61	89.95	91.10	96.65	89.81	91.87	91.87	90.71	56.97	55.96	55.46	54.61	54.84	54.41	55.21	54.96	54.89	55.00	55.46	55.58	54.81	55.54
JQ899443 SLV	86.47	85.95	78.51	88.19	87.89		90.16	86.32	87.10	89.46	89.80	86.89	89.33	86.17	86.17	90.39	56.76	56.36	56.59	55.57	55.91	56.21	55.11	55.00	56.02	56.25	56.71	55.34	55.46	55.57
JF320811 SLV	89.21	89.10	78.92	90.07	92.17	90.16		89.50	89.67	92.03	93.90	89.45	91.78	89.32	89.32	94.37	56.27	55.81	55.59	55.04	54.72	55.73	54.93	54.72	55.81	56.03	55.92	55.04	55.15	55.59
HQ258896 SLV	92.34	98.83	78.10	89.97	96.81	86.32	89.50		92.73	89.34	89.92	99.06	88.63	92.97	92.97	89.68	55.56	54.46	53.92	53.00	54.14	53.78	53.71	53.60	54.08	54.18	54.35	54.46	53.28	54.02
GU355922 GVL	97.48	92.73	78.92	90.04	91.61	87.10	89.67	92.73		88.75	89.32	92.97	88.03	94.25	94.25	89.32	55.49	54.84	55.06	54.04	55.18	54.28	55.06	54.95	55.52	55.75	55.29	54.61	54.15	55.86
GU120175 SLV	88.62	88.76	78.83	88.29	89.95	89.46	92.03	89.34	88.75		92.15	89.34	91.33	88.28	88.28	92.26	56.22	55.97	55.29	55.40	55.86	55.19	54.04	53.93	55.75	55.97	55.75	55.06	54.84	56.43
DQ520093 GVL	89.33	89.45	79.63	89.10	91.10	89.80	93.90	89.92	89.32	92.15		90.15	90.85	89.32	89.32	98.12	56.46	56.31	56.77	55.29	55.75	55.64	55.52	55.40	56.66	56.88	56.54	55.75	55.29	55.75
AJ409116 GVL	92.58	98.94	78.31	90.28	96.65	86.89	89.45	99.06	92.97	89.34	90.15		88.28	93.08	93.08	89.92	55.98	55.40	55.40	54.61	55.18	54.85	54.95	54.84	55.18	55.40	55.18	55.18	54.84	55.86
AB048002 SLVJ	87.65	88.04	80.56	88.98	89.81	89.33	91.78	88.63	88.03	91.33	90.85	88.28		87.91	87.91	91.43	56.71	56.88	56.09	56.31	55.86	56.21	55.18	55.06	56.77	57.00	56.77	55.97	56.31	55.63
AB046886 GVL	94.01	92.85	79.51	89.92	91.87	86.17	89.32	92.97	94.25	88.28	89.32	93.08	87.91		100.00	89.09	55.61	54.61	54.72	53.58	54.61	53.82	54.49	54.38	54.72	54.95	54.49	54.04	54.49	55.18
AB046886 GLV	94.01	92.85	79.51	89.92	91.87	86.17	89.32	92.97	94.25	88.28	89.32	93.08	87.91	100.00		89.09	55.61	54.61	54.72	53.58	54.61	53.82	54.49	54.38	54.72	54.95	54.49	54.04	54.49	55.18
AB04567 GLV	89.09	89.22	79.39	89.57	90.71	90.39	94.37	89.68	89.32	92.26	98.12	89.92	91.43	89.09	89.09		56.83	55.86	56.54	55.40	55.52	55.30	55.06	54.95	55.97	56.20	56.20	55.29	55.29	55.86
KP208802 GCLV	55.75	55.73	54.02	56.38	56.97	56.76	56.27	55.56	55.49	56.22	56.46	55.98	56.71	55.61	55.61	56.83		90.45	93.27	88.77	89.26	89.02	93.28	93.40	89.47	89.72	89.99	89.36	90.45	90.09
KJ801307 GCLV	54.82	55.52	54.28	55.70	55.96	56.36	55.81	54.46	54.84	55.97	56.31	55.40	56.88	54.61	54.61	55.86	90.45		89.07	93.46	95.85	95.35	89.29	89.40	96.57	96.88	96.57	95.32	94.70	89.81
KF862699 GCLV	55.05	55.06	53.62	55.48	55.46	56.59	55.59	53.92	55.06	55.29	56.77	55.40	56.09	54.72	54.72	56.54	93.27	89.07		87.94	87.73	87.79	91.88	91.88	88.24	88.45	88.15	88.03	89.29	87.62
JQ899445 GCLV	54.36	54.49	52.85	54.61	54.61	55.57	55.04	53.00	54.04	55.40	55.29	54.61	56.31	53.58	53.58	55.40	88.77	93.46	87.94		92.52	92.82	88.06	88.16	92.47	92.12	93.45	92.63	93.90	88.53
JQ818259 GCLV	55.05	55.06	53.84	55.15	54.84	55.91	54.72	54.14	55.18	55.86	55.75	55.18	55.86	54.61	54.61	55.52	89.26	95.85	87.73	92.52		93.95	88.45	88.55	95.33	95.64	96.35	94.17	93.36	89.41
JF320810 GCLV	54.01	54.73	53.46	55.51	54.41	56.21	55.73	53.78	54.28	55.19	55.64	54.85	56.21	53.82	53.82	55.30	89.02	95.35	87.79	92.82	93.95		88.27	88.39	93.92	94.05	95.00	93.50	93.83	90.51
HQ873863 GCLV	54.94	54.72	52.96	54.72	55.21	55.11	54.93	53.71	55.06	54.04	55.52	54.95	55.18	54.49	54.49	55.06	93.28	89.29	91.88	88.06	88.45	88.27		99.58	88.05	88.36	88.66	88.65	89.10	88.88
HQ873861 GCLV	54.82	54.61	52.85	54.61	54.96	55.00	54.72	53.60	54.95	53.93	55.40	54.84	55.06	54.38	54.38	54.95	93.40	89.40	91.88	88.16	88.55	88.39	99.58		88.15	88.46	88.76	88.75	89.20	88.88
MH686304 GCLV	55.52	55.29	53.51	55.81	54.89	56.02	55.81	54.08	55.52	55.75	56.66	55.18	56.77	54.72	54.72	55.97	89.47	96.57	88.24	92.47	95.33	93.92	88.05	88.15		99.30	96.47	94.80	93.97	88.93
MH686303 GCLV	55.75	55.52	53.51	56.03	55.00	56.25	56.03	54.18	55.75	55.97	56.88	55.40	57.00	54.95	54.95	56.20	89.72	96.88	88.45	92.12	95.64	94.05	88.36	88.46	99.30		96.78	94.91	93.68	88.50
DQ520092 GCLV	55.17	55.29	54.06	55.92	55.46	56.71	55.92	54.35	55.29	55.75	56.54	55.18	56.77	54.49	54.49	56.20	89.99	96.57	88.15	93.45	96.35	95.00	88.66	88.76	96.47	96.78		95.32	94.09	89.20
AF228416 GCLV	54.82	55.29	54.28	55.81	55.58	55.34	55.04	54.46	54.61	55.06	55.75	55.18	55.97	54.04	54.04	55.29	89.36	95.32	88.03	92.63	94.17	93.50	88.65	88.75	94.80	94.91	95.32		93.15	88.57
AB048004 GCLV	54.01	54.72	53.51	55.15	54.81	55.46	55.15	53.28	54.15	54.84	55.29	54.84	56.31	54.49	54.49	55.29	90.45	94.70	89.29	93.90	93.36	93.83	89.10	89.20	93.97	93.68	94.09	93.15		90.47
AB004566 GCLV	55.75	55.75	53.29	54.93	55.54	55.57	55.59	54.02	55.86	56.43	55.75	55.86	55.63	55.18	55.18	55.86	90.09	89.81	87.62	88.53	89.41	90.51	88.88	88.88	88.93	88.50	89.20	88.57	90.47	

Table S7 Distance matrix showing percent nucleotide identities of coat protein sequences of allexiviruses. Australian isolates from this study (Eastern Australia) are shown in orange and from Western Australia in light blue (GenBank accessions).

	KU933691	JX97952	JX97951	JX488633	JX429970	JX99446	JX99444	JN019812	MH686308	MH686307	AF478197	AB010300	KP55793	KM379144	JX488620	JN019813	MH686310	MH686309	AF543829	AB010301	KX034780	KP55566	JQ899447	JN019814	HM777004	KF555653	KF446210	JX488630	JN019815	LN67527	KP71369	KF51028	JX429971	JX429969	JQ807994	GM7542	MH686311	MH686312	MH686313	AJ32229
KU933691	83	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	
JX97952	83	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	
JX97951	99	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	
JX488633	99	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	
JX429970	99	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	
JX99446	99	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	
JX99444	99	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	
JN019812	99	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	
MH686308	99	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	
MH686307	99	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	
AF478197	99	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	
AB010300	99	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	
KP55793	99	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	
KM379144	99	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	
JX488620	99	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	
JN019813	99	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	
MH686310	99	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	
MH686309	99	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	
AF543829	99	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	
AB010301	99	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	
KX034780	99	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	
KP55566	99	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	
JQ899447	99	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	
JN019814	99	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	
HM777004	99	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	
KF555653	99	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	
KF446210	99	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	
JX488630	99	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	
JN019815	99	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	
LN67527	99	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	
KP71369	99	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	
KF51028	99	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	
JX429971	99	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	
JX429969	99	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	
JQ807994	99	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	
GM7542	99	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	
MH686311	99	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	
MH686312	99	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	
MH686313	99	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	
AJ32229	99	99	78	97	85	97	74	85	97	97	97	97	61	62	61	62	61	61	61	61	60	59	62	62	59	71	71	68	68	64	64	64	65	63	64	63	63	64	62	

MH686311	64.	65.	62.	65.	67.	64.	60.	63.	64.	64.	63.	64.	76.	76.	77.	76.	76.	75.	74.	74.	74.	65.	64.	65.	65.	66.	72.	61.	62.	61.	62.	99.	91.	95.	91.	95.	94.	94.	94.	94.	94.
GarVX	83.	11.	04.	09.	09.	11.	96.	36.	42.	08.	57.	15.	93.	72.	21.	79.	04.	64.	90.	94.	23.	60.	55.	26.	72.	04.	06.	24.	65.	32.	27.	01.	81.	21.	60.	13.	67.	51.	94.	95.	
AJ292229	62.	64.	61.	62.	66.	61.	60.	62.	62.	62.	61.	62.	75.	75.	75.	75.	74.	74.	73.	74.	63.	63.	64.	64.	63.	60.	61.	60.	61.	94.	91.	95.	91.	94.	94.	93.	94.	94.	94.		
GarVX	99.	42.	91.	99.	30.	92.	68.	30.	61.	12.	62.	18.	31.	91.	58.	98.	57.	08.	11.	15.	97.	72.	86.	82.	83.	50.	67.	84.	99.	81.	68.	05.	76.	77.	41.	99.	60.	27.	65.		