

Supplementary Fig. 2: Alignment of the minus-strand priming site of *Agapanthus tungro virus* (AgTV) with whole genome sequences of *Rice tungro bacilliform virus* (RTBV)

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MT501685	AgTV	1	19-3001	TGGTATCAGAGC-ATG
MT501686	AgTV	1	19-3004	TGGTATCAGAGC-ATG
MT501687	AgTV	2	19-3005	TGGTATCAGAGC-ATG
MT579870	AgTV	1	19-3006	TGGTATCAGAGC-ATG
MT501688	AgTV	1	19-3007	TGGTATCAGAGC-ATG
MT501689	AgTV	2	19-3010	TGGTATCAGAGC-ATG
MT501691	AgTV	1	19-3013	TGGTATCAGAGC-ATG
MT501690	AgTV	2	19-3013	TGGTATCAGAGC-ATG
AF076470	RTBV		Serdang	TGGTATCAGAGCGATG
AF220561	RTBV		Chainat	TGGTATCAGAGCGATG
AF113830	RTBV		G1	TGGTATCAGAGCGATG
AF113832	RTBV		Ic	TGGTATCAGAGCGATG
FN377814	RTBV		Chinsura West Bengal	TGGTATCAGAGCGATG
JX255736	RTBV		Punjab	TGGTATCAGAGCAATG
HQ385226	RTBV		Kanyakumari	TGGTATCAGAGCGATG
AJ314596	RTBV		West Bengal	TGGTATCAGAGCGATG
MK552377	RTBV		Seberang Perai	TGGTATCAGAGCGATG
NC001914	RTBV			TGGTATCAGAGCGATG
AJ292232	RTBV			TGGTATCAGAGCAATG