

Table S3: Oligonucleotides used in this work

	Name	Sequence (5' to 3')	Specificity	
Plasmid and strain construction	egfp_r	ACTTCAGGGTCAGCTTGC	<i>egfp</i> gene	
	PcvelA_f	TCGGTCGACATGGCCAACAGA CCATCTC	<i>PcvelA</i> gene	
ChIP-PCR	qPCR_NC1_f	TTCTTCCGCAATCAAGCTCA	chr1:5375234-5375253	
	qPCR_NC1_r	GAAAAATTGCCGCTGGACTC	chr1:5375364-5375383	
	qPCR_NC2_f	GGTCGTTGATTCCCTTGAGC	chr2:7621179-7621198	
	qPCR_NC2_r	GGATCGGATTATTCGGGTGA	chr2:7621294-7621313	
	qPCR_Pc21g02240_f	CGAGAGAGAGGAACCCGGGA	chr2:5277047-5277066	
	qPCR_Pc21g02240_r	TTCCCGTACCAGGCTGTCG	chr2:5277176-5277195	
	qPCR_Pc22g17530_f	AGGCACCGAAACCGTGAAGA	chr1:2788884-2788903	
	qPCR_Pc22g17530_r	ACGCCAGGCCAGAGTTCAAT	chr1:2788799-2788818	
	qPCR_Pc20g02880_f	CGTGAAATTCGAAGTTCCCGA	chr2:2995008-2995029	
	qPCR_Pc20g02880_r	AGAAATTAAGCCGAAAACCCA GA	chr2:2994880-2994903	
	qPCR_Pc20g14090_f	GTGGAAATTTCCGATGGGGTA GC	chr2:378180-378202	
	qPCR_Pc20g14090_r	GATGCCCTGGTATCGGCAAAA	chr2:378053-378073	
	qRT-PCR	qRT_Pc21g02240_f	ACAAGGAAATCGGTGCATC	chr2:5275851-5275870
		qRT_Pc21g02240_r	GCCCTCTCCATATGCTCCTG	chr2:5275747-5275766
qRT_Pc18g01840_f		TTCGGCAAGGACATGACATC	chr1:5930185-5930204	
qRT_Pc18g01840_r		TGGTACCGACCAAGCTCCTT	chr1:5930361-5930380	
qRT_Pc21g12700_f		GGGTTTGTGATACCCAGGA	chr2:7743588-7743607	
qRT_Pc21g12700_r		CCGCAGTCCACTGATGGTAA	chr2:7743677-7743696	
qRT_Pc18g04780_f		TGATGATGCGAAGACCATCC	chr1:6636202-6636221	
qRT_Pc18g04780_r		TGAACCAATGCACCAGCTCT	chr1:6636094-6636113	
qRT_Pc18g06010_f		CCCCGATACCAACGCATACT	chr1:6920825-6920844	
qRT_Pc18g06010_r		CGTGATCTTCAACCCAGCAG	chr1:6920718-6920737	
qRT_Pc13g15570_f		GGACCCGAACTCTGTTGCTC	chr4:2949445-2949464	
qRT_Pc13g15570_r		GCATCCACCACCTTCTCAA	chr4:2949330-2949349	
qRT_Pc22g01170_f		TCGCTCGCTTCTTGTATGA	chr3:5122377-5122396	
qRT_Pc22g01170_r		CAGGACTCGCAGACCAACAG	chr3:5122469-5122488	
EMSA ^a	SSU1	ATCCAAGGAAGGCAGCAGGC	SSUr RNA	
	SSU2	TGGAGCTGGAATTACCGCG	SSUr RNA	
	PcLImA_2	TAGCGTCATTTATTTTTTCTTC CAAGGTTTTCCCTCTTCTTCG GAGT	chr2:5277074-5277122	
	PcLImA_2_m	TAGCGTCATTTATTTTTTCTTC AgaGTTTTCCCTCTTCTTCGGA GT	chr2:5277074-5277122 (with mutations)	
EMSA ^a	PcLImA_4	TCCGACAGCCTGGTACGGGAA ACCTTGAACCCATTCCAAATC GGTCTG	chr2:5277174-5277222	
	PcLImA_4_m	TCCGACAGCCTGGTACGGGAA ACtTGaAgCCCATTCCAAATCG GTCTG	chr2:5277174-5277222 (with mutations)	

^a in case of double-stranded oligonucleotides used for EMSAs, only the sense sequences are given.