

Locus ID	CHIP (5'-3')	RT-qPCR (5'-3')
PSF113_1559 (<i>fleQ</i>)	Fw: AGGCGTCTACAGGCATGACAA Rv: TTAAAGATCACCGCCAAGTCG	
PSF113_4752 (<i>algD</i>)	Fw: TGCACTATGACCGCTGAACAC Rv: GCGACATCTACGCCAACTACG	
PSF113_4470 (<i>amrZ</i>)	Fw: CATTCCGTTATCCGCCAGAC Rv: TCATGCCTCAAAATTGCCAGA	
PSF113_0714 (<i>yfiR</i>)	Fw: GGGTTGATGCTGGAGTTGGAG Rv: CCAGCGAGACAGCGTGTATG	
PSF113_4360	Fw: ATGGTGATGCTCGTCGGAGT Rv: CTGTCGTGGTCATGGCAATCT	Fw: GCGCATCGTGGGTTATGAAG Rv: CCAGTTCCAATTCGCTGAGG
PSF113_5392	Fw: CGATGGCGTTTGTTCAGTCAG Rv: GAGGGGACGGTTTTCTTAGGG	Fw: CGACGCTGGTGTGTTGCT Rv: CCAGGAGGAAGTAGGCGATGT
PSF113_1274	Fw: GATAGGCTCCGCTGATTCGTC Rv: GCAAGAAACCGGCAAGTAAGG	Fw: CCACCTGAAAAAGCCCTGGA Rv: GCGCTGATGGTCACTTGTG
PSF113_1322	Fw: TACCGAAGGCGGCAATCTC Rv: ATCATGGCGAACATCCTGCTT	Fw: GGCGAAGGCAAAGACAAGG Rv: TAGACGATGTCGGCGTAGTGC
PSF113_1749 (<i>pvdS</i>)	Fw: TCGCACCTGCTGTGGATACT Rv: CTAATTTGGCGCTCGATGTGT	Fw: CGCCAGAAACCTCCCATATCA Rv: GCGGTACATCTCGAAGGCATAG
PSF113_1750 (<i>pvdL</i>)	Fw: TCGCACCTGCTGTGGATACT Rv: CTAATTTGGCGCTCGATGTGT	Fw: AGGCAGGAAACCCCATGAC Rv: CGCGATAACTCAACACCACAC
PSF113_4569	Fw: TTCAAGGAACCGGACTCAATGT Rv: GCGCATTTACGAGTGTCAAACC	Fw: GAGATGCTTGAGTGCACCTG Rv: CGATGTGCTCTTCTCGGATT
PSF113_0750-0751 (<i>flhDC</i>)	Fw: GGAACATCCGCCAAGTACGA Rv: TCACATACTCGACCCTGCTCAA	Fw: TGGAACAGGTCGGTCATCAAG Rv: GCGTGAGAGTGAGCAACGAA
PSF113_3554	Fw: ATAGTTTCGCGGCCATCACTT Rv: TTGCCTGTCTTTCACCTTCA	Fw: GTGTGGAGCCTGCGTTTATTG Rv: CCTCACCAATCCTTCATCAG
PSF113_4205	Fw: ACTGGCTACGGATCAATGTGG Rv: AAGATGCTGGCGTCGGAATAA	Fw: GGATTTGTCTGGTGCTGCT Rv: GAGGGCCGACATCTGTTTCA
PSF113_4460 (<i>flgZ</i>)	Fw: GCCAGTCGATCCTGATGCAA Rv: ATCATCCGCGTTTGAGGCA	Fw: CCGCGAGACCAACTCCATT Rv: GTCAGTGGACCCTTGCCATC
PSF113_0118	Fw: TGCCATCAGTAAAAGCGGTATG Rv: CCAGGTCACGAATGAACGAAG	Fw: TGCTTCGTTTCATTCGTGACCT Rv: GGCATTTCCAATCGCACA
PSF113_5133 (<i>ladS</i>)	Fw: TGACACTTATTGGGGCTGCTTC Rv: GTTGCCTTGGTGAGGTCGTAA	Fw: CTCTCGATCAGCGTTCTGGTT Rv: CGCATCGGGTAGGTACAAGTC
PSF113_5263	Fw: ACAGTCCAGAGCGGTGTCAGT Rv: GTTTGGTCGCAAGTCCATGTC	Fw: ATGGACTTGCGACCAAACACC Rv: GCAGCCCTTCAACGACAC
PSF113_5333	Fw: CTTTCATCGCCAAGACTCCACT Rv: ATGCTGGTCGGTCAGGTCTC	Fw: AAGCCATGTTCTCGGTGAGT Rv: CCCAGCAAGAGCATTTCATC
PSF113_5334 (<i>vfr</i>)	Fw: CTTTCATCGCCAAGACTCCACT Rv: ATGCTGGTCGGTCAGGTCTC	Fw: GCCATCCAGCCAAACACAAC Rv: ACCGTCGTCATCTTCTATCAGG
PSF113_2409 (<i>vgrG</i>)	Fw: CCAGAATGGAAACGCCAATAA Rv: CGGTGAGTGATAGCGTGGTGT	Fw: GACAGCAACACCACGCTATCA Rv: CGTTGAGGATGTGCGAACC
<i>dif</i> region	Fw: TTGAAGCCAATCCCATCCTCT Rv: TTCGTTCTTCTTGTGTCG	
PSF113_4631		Fw: TGTTGTTGACCACTGTCTGCT Rv: CGGAGATCAGGCTTGATTG
PSF113_5797 (<i>impA</i>)		Fw: GTGGATGTGCCTTGTGCTC Rv: CATGCTACGTTCCGGCTGAC
PSF113_1837 (<i>pvdD</i>)		Fw: GACAAGGAGGGCGTGGAGTT

Locus ID	ChIP (5'-3')	RT-qPCR (5'-3')
PSF113_0889		Rv: CAGGTTGTAAGCACCGCTGTC Fw: CGACCGTGTCTACCTGCTGA
PSF113_2589		Rv: ACTGCGCCTTGAAGCTCTTGT Fw: TGTTTCAGAAATGGGCATCCAG
PSF113_3220		Rv: TCCTCGCTCAGCTCATACTCG Fw: TCCGCCAACGATCTGTCTCTA
PSF113_4045		Rv: ACGTATCCTGGGCAAGGTCTG Fw: CTTCGGTTTGGTCCTGAGCAT
PSF113_4845		Rv: GGGTTTCTGGCTACGCTTGAG Fw: GAAGAGGAACGGGCAATCATC
PSF113_0933		Rv: TGACCGTCGCCAATGAAATAC Fw: CACTGCGCGACCTCGAACAAC
PSF113_1554 (<i>fliC</i>)		Rv: TACTCATCGGGAGATCCTTGC Fw: TTCCGATGCTCTGTCCACTTC
PSF113_0740 (<i>fliC2</i>)		Rv: GATACCGTCGTTGGCGTTCTT Fw: GACCAGACCATCACCATCCAG Rv: AGCGTTGTGTCATAGCCGTTTC

AmrZ fusions (5'-3')^a

HAamrZF	ATGTCT <u>ATCCATACGATGTTCCAGATTATGCTCGCCCATTGAAACAGGCA</u>
amrZextR	TCAGGTCGCGTCTGCGG
amrZextF	ATGCGCCCATTGAAACAGG
amrZHAR	TTA <u>AGCATAATCTGGAACATCGTATGGATAGGTCGCGTCTGCGGCCA</u>

^aHA sequence is underlined