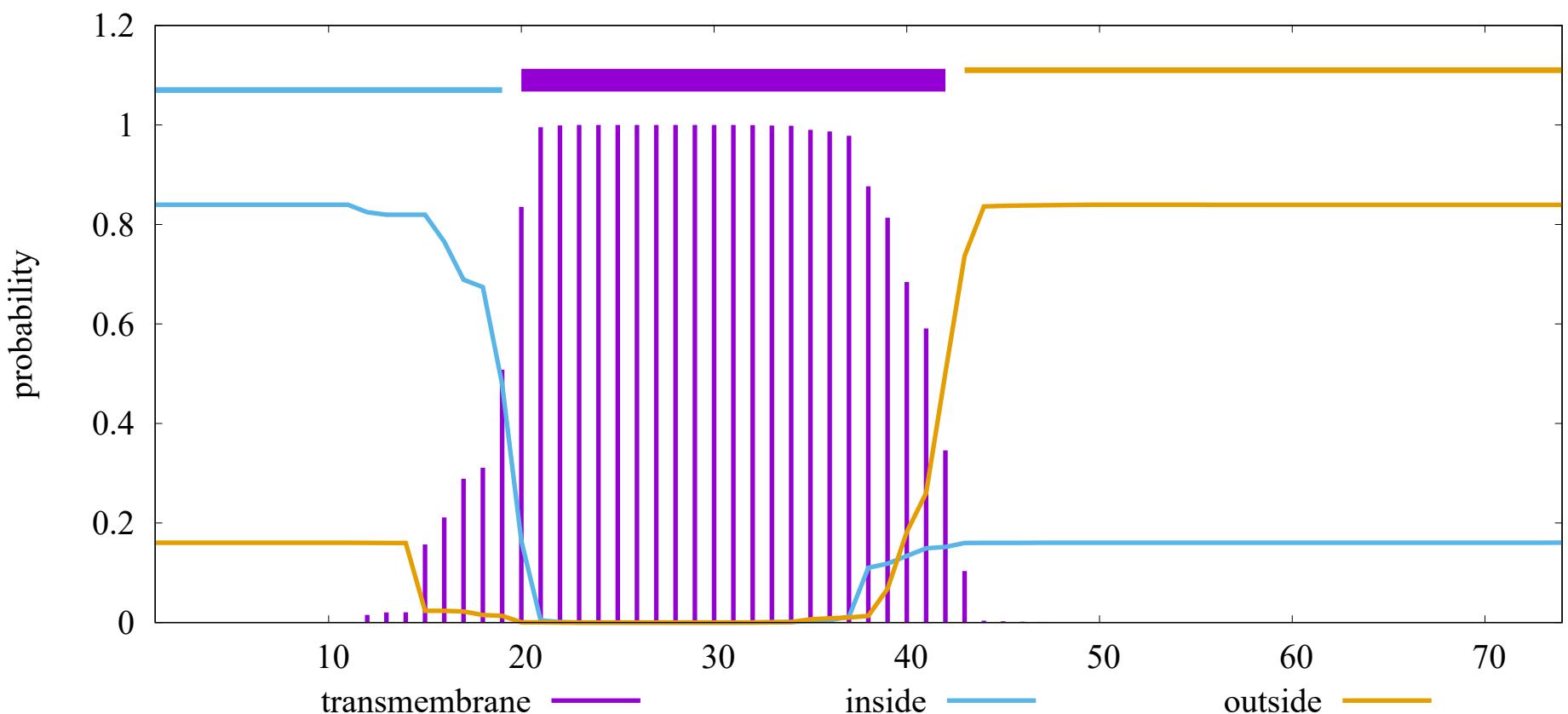


Huangshi Niviventer confucianus morbilli-like virus (Huangshi10)



Shiyan Crocidura tanakae henipavirus (Shiyan201)

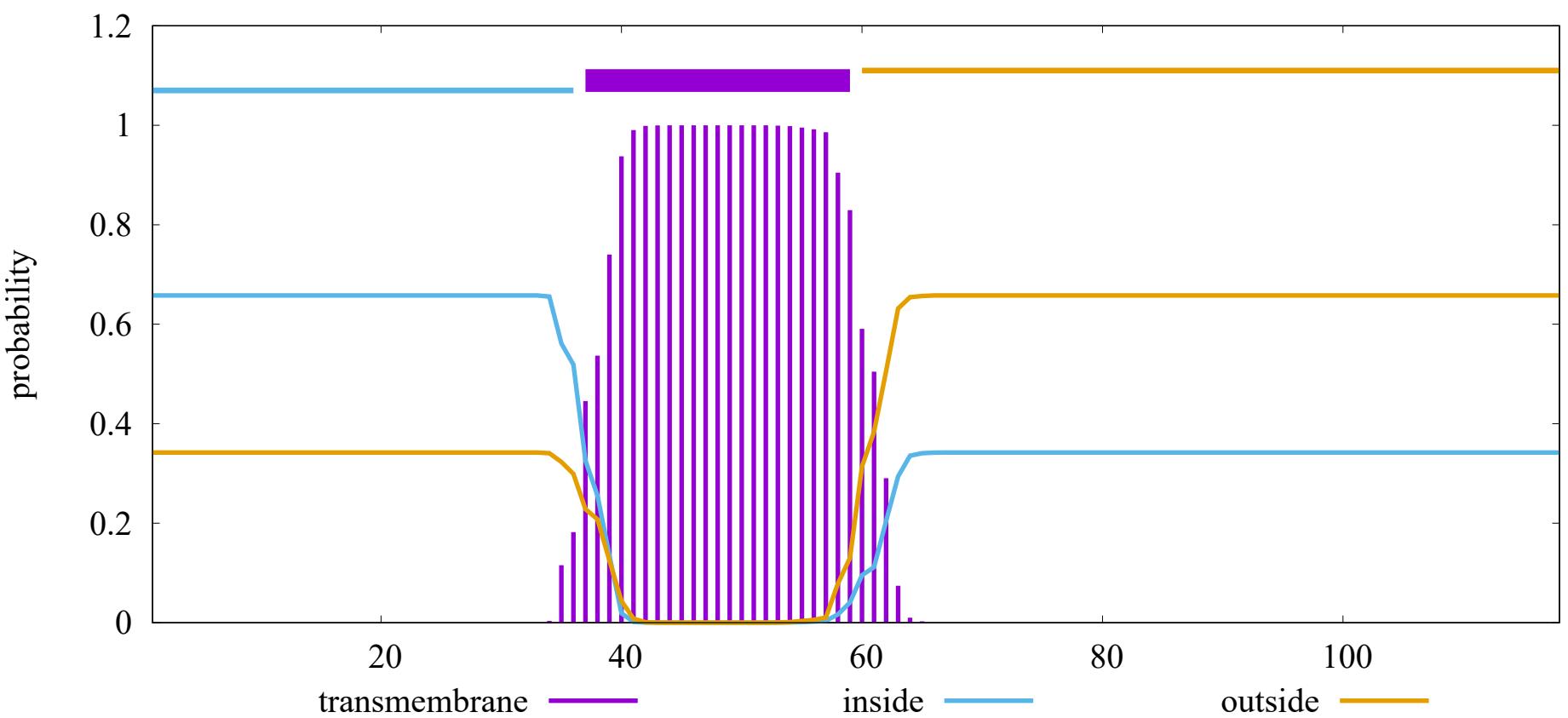


Figure S1: TMHMM posterior probabilities of ORF X Y and ORF Z in Huangshi Niviventer confucianus morbilli-like virus and Shiyan Crocidura tanakae henipavirus. The x-axis represents the amino acid length and the y-axis represents the posterior probability.

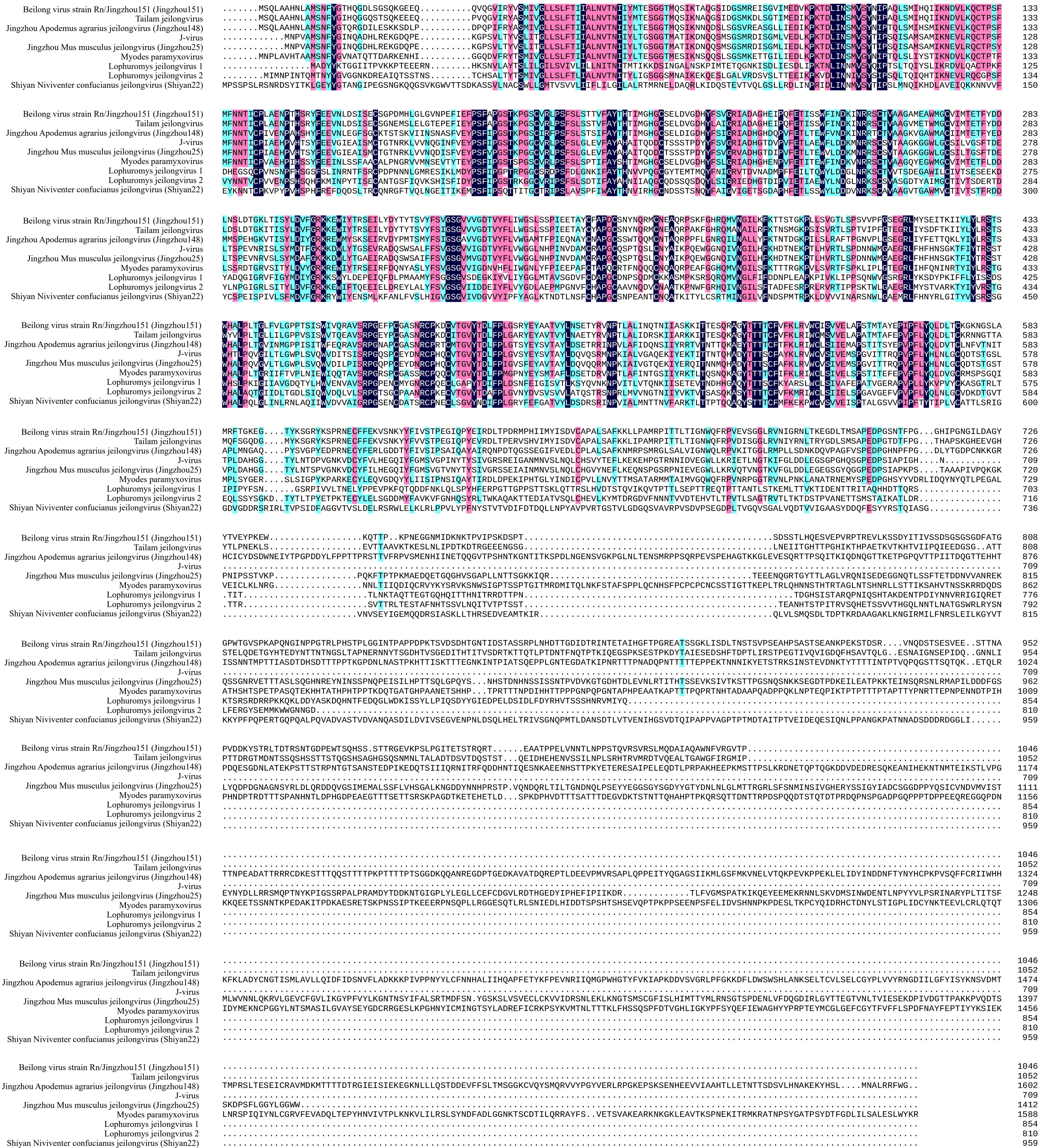


Figure S2: Multiple alignment of amino acid sequences of G proteins of Jeilongviruses.

Table S1. Primers used for nested RT-PCR to obtain complete virus genome sequences.

Sample	First round		Second round	
	Primer name	Primer sequence (5'-3')	Primer name	Primer sequence (5'-3')
Huangshi10	1F	GCCTACCAACCATGTCAACA	N-1F	ATCAAACCACCTGATCCTTCTG
	1R	CTGACTACTCACTGAATCCTAAG	N-1R	GTTCCAAC TGCCATGAATCTC
	2F	AAGTCCTTAGATGGCTTGAT	N-2F	GAACACAACCAGGTATTACTG
	2R	TCTCTGACTTCCCYCATATG	N-2R	CTTGTRGCATTAGTCTGYTGTG
	2F	GGACGACCATATAGCTCTGT	N-2F	CGATCTAACCTATTCTTGG
	2R	CCARTGAACCTGAGCATTKG	N-2R	ACTAGATAATAGATCAGGAGCATG
	3F	GTCTACTGTGGAGGAACCAG	N-3F	CATCAGCTCCGAAAGAGTT
	3R	GCTCTGATTGATCTAGCAGAAC	N-3R	GTTGAGTCTATCCATAGCATGC
	3F	GCATGCTATGGATAGACTCAAC	N-3F	GTTCTGCTAGATCAATCAGAGC
	3R	GCCACGAGCAATTAGCTG	N-3R	AAGGGTCTTCTCTCCCATC
	4F	GATGGGAGAAGAAGACCCCT	N-4F	CAGCTAAAATTGCTCGTGG
	4R	TGTTATYCCACAATCYTGYTGG	N-4R	GAAGAGCCYTCTARTTTGTAG
	4F	GACATAGGAAGAATACCCATTCA	N-4F	GATTGCTACATTATCACAGTATGGA
	4R	AATCTYCCWACRCCGTTAGATAT	N-4R	CATGCTCTCATYTRTAGGTAC
	5F	ACTATCAATCCAACATGACAGC	N-5F	AGACCTATACTTGACAATCCACA
	5R	GCATAGATAACCTGCTGCTTC	N-5R	TTTCTGGAATTGAACGAAGGC
	5F	TCGYGACAARAGCACTCAAGT	N-5F	GGGAGRMGAGTRGACACYAAC
	5R	ATGGATTGAACCTCGCTGAC	N-5R	ACCTAAGTAATGTTGATAGACTGC
	6F	TYGGTRTMTCACTCAACAATTACG	N-6F	ACTCAACTGCWTGTTAYAARGCT
	6R	CAAAATTGCCATAGAAACTTACT	N-6R	TCAGGGACTGGAAGTGGAC
Shiyan201	1F	ACGTAACTCTTAAGAACCAAAGAT	N-1F	GTTCAAGGGAGTATCAAGCAAAC
	1R	TGCCATCATYTGACCRA GTCTG	N-1R	CCATRGCRARCTCCA WAGYAA
	2F	CAGTGAGAAAGTTCATGGTCGA	N-2F	GAAGTCAGAACAGAGCGGAGC
	2R	CYAAACATCAGWG CYTTGACTCC	N-2R	CCTCAGGTACAGAGTTGATGAAC
	2F	GAGGC GATAAGT GAAAGGCAT	N-3F	CATGACGACAAGCTCAAACAAAT
	2R	TTAGTCATRTCAAGTGGCTCA	N-3R	AGGK CCTYTCATTCWGATGACA
	3F	GATACAAGGAACAGAGTAGATGC	N-4F	CGTAGTGTAGGGAAATTAGGTCT
	3R	CWACWGCTCCYAAKGAGAAC	N-4R	CCTATATGAAGCATRAAYGTGG
	3F	GATTTCAGAGAACACAATGCA	N-5F	CAATCTGCTTGTCTCTTGCT
	3R	AKAYCATYAAWCCTCTRATCARTC	N-5R	TRCAACTRTCTGCATYTGWT
	4F	GATCAACCCGGAACAAATACCT	N-6F	TGTCAAGATAAGCACCTACCTTG
	4R	GGTGTWGCA TARTCGTTG TAGC	N-6R	TCAAGAGAACGTATCGAGGGAC
	4F	CACTGGCAATATAATAGGTGTGG	N-7F	GCCCTGGAGATAACATTTCTA
	4R	TGRCTKGGTATGGTKACACTRAC	N-7R	TTCGCCTTRACAATTGRTCTA
	5F	TTGCTGATGGGGTTACTAAT	N-8F	GGCTAACATGCTCTAATAATCA
	5R	AAGCACTTAGTTAGTTGAYCC	N-8R	CTGRTCTATRACCTCATTYGGT
	5F	GGTAYGGWCARACAGARAGGA	N-9F	GYARTGGWGGTMGWATATGYCC
	5R	GTTRCAYTCKGGRTAYGGTACATG	N-9R	GACCYAA TTCCCATTYTGTATT
	6F	TCRGATGTGYTKTATCCWGAATG	N-10F	TRGAYAGT CCTATWGTWGCWGG
	6R	ACTCTGGATCATTCTTATARTCACC	N-10R	TTCTGAACARATAGGTGATAATGC
	6F	GRGCWCATTCTAATGGYGAAGC	N-11F	CATGARATGTGTGCWACYAAYTGG
	6R	CCTARWGCWGCYARATTDTCYCT	N-11R	GWG CYCKYTCWGC ACTRATTCT
	7F	GCCMTTRATACARAATCARTCATGG	N-12F	GCATGTCCTTACCTGCACC
	7R	GRTGWAGATCMGWTATAGTCCATT	N-12R	GRTCACTTCKATCAAWGGRTTRC
	7F	TTTCARCAR GTKATGCTCTAGG	N-13F	CACCCAAGGARAATATCATAATGC
	7R	CATTGGYTCYTTMCMATKCCAC	N-13R	CCAKGTTTTCWGGCTSC
	8F	GGATTTACTCCCCATACATACTCC	N-14F	GACAGGGTATTAGGTCCAAGC
	8R	AACTCACTGTCTGACTAGATC	N-14R	TACTTGGTTTGGGACTGGC