Primers with 5' flaps improve real-time PCR

Irina Afonina, Irina Ankoudinova, Alan Mills, Sergey Lokhov, Phan Huynh, and Walt Mahoney

Nanogen, Bothell, WA, USA

BioTechniques 43:770-774 (December 2007) doi 10.2144/000112631

Primers that contain portions noncomplementary to the target region are usually used to add to the PCR product a utility sequence such as a restriction site or a universal probe binding site. We have demonstrated that primers with short 5'AT-rich overhangs increase real-time PCR fluorescent signal. The improvement is particularly significant for difficult to amplify templates, such as highly variable viral sequences or bisulfite-treated DNA.

Primers that contain portions noncomplementary to the target (often called overhang, flap, or tail) are usually used to add to the PCR product a utility sequence such as a restriction site (1) or a universal detection site (2,3). Stemloop primers make PCR amplification of short templates such as microRNA possible (4). Flap primers are used to reduce the number of sequencing errors in short PCR products (5). We have noticed that primers with short 5' AT-rich flaps increase realtime PCR fluorescent signal, and this improvement is particularly significant for sequences that are difficult to amplify, such as bisulfite-treated DNA or highly variable viral sequences. The real-time PCR experimental data presented here (see Table 1 for oligo-

nucleotide sequences) was generated by 5' minor groove binder (MGB) fluorescent hybridization probes (6), although we have observed similar effects for other platforms such as TaqMan probes (BioSearch Technologies, Novato, CA, USA) (data not shown). An example of the typical benefits of primers containing 5' AT-rich flaps is shown in Figure 1, in which they are used to amplify sodium bisulfite-treated DNA. The bisulfite treatment leads to a conversion of all unmethylated cytosines to uracils, leaving methylated cytosines unchanged (7), which constrains primer design. Human genomic DNA was treated with sodium bisulfite. and a portion of the differentially methylated region of the H19 gene was amplified using primers with or without 12-mer 5' AT-rich flaps (Table 1). Fluorescent signal is increased, and PCR yield is higher when bisulfite-treated DNA is amplified with flap primers and fluorescence is generated by a hybridization probe (Figure 1A). This increase in signal intensity is also observed when fluorescence is generated by a free intercalating dye such as SYBR Green (Figure 1B), and the bands on ethidium bromide-stained agarose gel are also more pronounced (Figure 1C).

We have determined that a 12-bp 5' AT-rich flap sequence is optimal. Addition of flap sequences longer than 12 bp did not show any significant improvement, while shorter sequences had a lesser impact on the fluorescent signal (see Supplementary Table S1, available online at www.BioTechniques. com). The addition of a flap to either one or the other primer provides a boost to the fluorescent signal, but is not as beneficial as having both primers with flaps. 5' GC-rich flaps overall proved not as useful as 5' AT-rich flaps, because they are more prone to form stable secondary structures and negatively impact the PCR (data not shown). The positive effect of the 12bp 5' AT-rich flap is more pronounced for shorter primers (Supplementary Table S2). To study the effect of flap addition to primers of decreasing length on cycle threshold (C_T) and

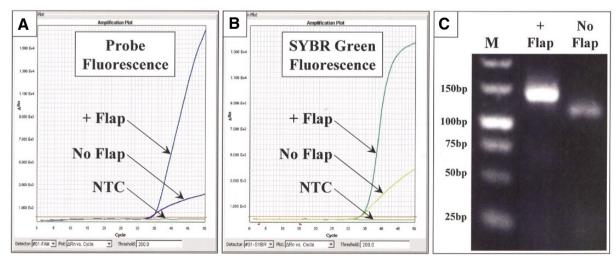


Figure 1. Primers with 5' flaps improve real-time PCR. Bisulfite-treated genomic DNA with 50% methylated H19 differentially methylated region (DMR) was amplified with flap and non-flap primers using (A) JumpStart Taq ReadyMix and (B) SYBR Green JumpStart Taq ReadyMix (both from Sigma-Aldrich, St. Louis, MO, USA) using the following cycling parameters: 2 min at 95°C, followed by 50 cycles of 15 s at 95°C and 1 min at 60°C. (C) The JumpStart Taq ReadyMix PCR products were run on a 4% agarose gel.

Benchmarks

Table 1. Primers and Probes Used for Real-Time PCR

Primer or Probe	Sequence ^a	Final Concentration
	VZV Reagents	Concentration
VZV-forward primer	5'- AATAAATCATAA GAACAATCACGAACCG-3'	0.25 μΜ
VZV-reverse primer	5'-AATAAATCATAACGGGCATGCTTACGGACGT-3'	1.0 μΜ
VZV-probe	5'-MGB-FAM-CAGGACGTCAGATCACGT-NFQ-3'	0.2 μΜ
VZV DNA	VZV, strain Webster; ATCC VR-916	105 copies/rxn
	Enterovirus Reagents	
ENV-forward primer	5'- AATAAATCATAA GAAGAGYCZ*ATTGAGCTA-3'	1.0 μΜ
ENV-reverse primer	5'-AATAAATCATAAGGA*TTRGCCGCA*TTC-3'	1.0 μΜ
ENV-probe	5'-MGB-FAM-TCCGGCCCCTGAATGC-NFQ-3'	0.2 μΜ
ENV RNA	Coxsackievirus isolate A6; ATCC VR-165	
	H19 Differentially Methylated Region Reagents	
DMR-forward primerb	5'- AATAAATCATAA TTGG A*TGGTACGGA*ATTGG-3'	1.0 μΜ
DMR-reverse primer	5'-AATAAATCATAATCCACGAACGAACCC-3'	0.25 ^b μM
DMR-probe	5'-MGB-FAM-ACGA*ACT*CGA*ACT*ATAAT-NFQ-3'	0.2 μΜ
Bisulfite-treated DNA template	Human genomic DNA (DMR 50% methylated) was a generous gift of Alain Carrié [Institut Fédératif de Recherche des NeuroSciences (IFRNS), France]	1 ng/rxn

^aBolded sequences are noncomplementary flaps. Primers without bold type lack flaps.

fluorescence, we chose a primer pair optimized for the cycling conditions recommended by the PCR Master Mix manufacturer (8) and designed several shorter primer pairs. As expected, shorter primers without flaps had higher C_T values and lower fluorescence gain. The addition of a 12-bp 5' AT-rich flap nullified the difference in performance. Fourteen- to fifteen-mer primers with these flaps had approximately the same C_T value and fluorescence as the original 22- to 25-mer primers without flaps (Supplementary Table S2). This is useful information if primer design is constrained to short stretches of sequence for any reason.

Figure 2 demonstrates the effect of 5' flap primers in real-time PCR with varicella-zoster virus (VZV) DNA template and one-step reverse transcription PCR (RT-PCR) with enterovirus RNA template. The region chosen for the VZV design [open reading frame (ORF) 38 gene] is conservative and does not impose any restrictions on primer design. An increase in total fluorescence gain of approximately 50% due to 5' primer flaps (Figure 2A), and no significant shift in C_T is typical for such unconstrained primer designs. In contrast to

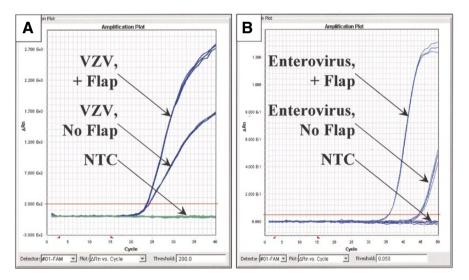


Figure 2. Real-time PCR amplification of viral targets with 5' flap primers. (A) Varicella-zoster virus (VZV) DNA template was amplified with the LightCycler FastStart DNA Master HybProbe master mixture (Roche Applied Science, Indianapolis, IN, USA) using the following cycling parameters: 10 min at 95°C, followed by 40 cycles of 5 s at 95°C, 20 s at 56°C, and 20 s at 76°C. (B) Enterovirus RNA (coxsackievirus A6) template was amplified by one-step reverse transcription PCR (RT-PCR) with the QuantiTect Probe RT-PCR master mixture (Qiagen, Venlo, The Netherlands) using the following cycling parameters: 30 min at 50°C (RT), 15 min at 95°C, followed by 50 cycles of 15 s at 95°C, 30 s at 56°C, and 30 s at 76°C. In both cases detection was done during the annealing stage of PCR.

VZV, the sequences of the enterovirus genus (Taxonomy ID 12059; www. ncbi.nlm.nih.gov/Taxonomy) are highly variable. Therefore, to enable detection of all known isolates, the primer design is constrained to a short

conservative region in the 5' untranslated region (UTR). The primers also include degenerate and modified bases to account for unavoidable single nucleotide polymorphisms (SNPs) and to boost primer stability (9). As a result,

^bRaised to 1 μM for SYBR Green.

^{*}Indicates modified base with increased stability.

VZV, varicella-zoster virus; MGB, minor groove binder; NFQ, nonfluorescent quencher; ATCC, American Type Culture Collection, Manassas, VA, USA; rxn, reaction; ENV, enterovirus; DMR, differentially methylated region

Benchmarks

the PCR primers perform suboptimally, and PCR efficiency is compromised. When the flap is added to the suboptimal (non-flap) enterovirus primers, there is a significant increase in performance for both fluorescence and C_T values (Figure 2B).

It is not clear why PCR primers containing the 5' AT-rich flaps are better, but the benefits are obvious. This innovation could be of benefit to every real-time PCR laboratory, especially when sequence choice for primer design is constrained.

COMPETING INTERESTS STATEMENT

The authors declare no competing interests.

REFERENCES

- 1. Espelund, M. and K.S. Jacobsen. 1992. Cloning and direct sequencing of plant promoters using primer-adapter mediated PCR on DNA coupled to a magnetic solid phase. BioTechniques *13*:74-81.
- Li, X., Y. Huang, Y. Guan, M. Zhao, and Y. Li. 2006. Universal molecular beacon-based tracer system for real-time polymerase chain reaction. Anal. Chem. 78:7886-7890.
- 3. Xu, S.X., K. Lan, Y.L. Shan, H. Wang, J.Y. Luo, F. Cui, Y.S. Huang, J.P. Meng, et al. 2007. Real-time quantitative assay of HCV RNA using the duplex scorpion primer. Arch. Virol. *152*:431-440.
- 4. Chen, C., D.A. Ridzon, A.J. Broomer, Z. Zhou, D.H. Lee, J.T. Nguyen, M. Barbisin, N.L. Xu, et al. 2005. Real-time quantification of microRNAs by stem-loop RT-PCR. Nucleic Acids Res. 33:e179.
- 5. Binladen, J., M.T. Gilbert, P.F. Campos, and E. Willerslev. 2007. 5'-tailed sequencing primers improve sequencing quality of PCR products. BioTechniques 42:174-176.
- 6. Lukhtanov, E.A., S.G. Lokhov, V.V. Gorn, M.A. Podyminogin, and W. Mahoney. 2007. Novel DNA probes with low background and high hybridization-triggered fluorescence. Nucleic Acids Res. 35:e30.
- 7. Shen, L., Y. Guo, X. Chen, S. Ahmed, and J.-P. Issa. 2007. Optimizing annealing temperature overcomes bias in bisulfite PCR methylation analysis. BioTechniques 42:48-
- 8.MGB Eclipse™ PCR Reagent Kit.
 Technical Bulletin. Product Code M 4568.
 Sigma-Aldrich, St. Louis, MO.
- 9. Afonina, I.A., A. Mills, S. Sanders, A. Kulchenko, R. Dempcy, S. Lokhov, N.M. Vermeulen, and W. Mahoney. 2006. Improved biplex quantitative real-time polymerase chain reaction with modified primers for gene ex-

pression analysis. Oligonucleotides *16*:395-403.

Received 25 July 2007; accepted 1 October 2007.

Address correspondence to Irina Afonina, Nanogen, Inc., 21720 23rd Drive SE, #150, Bothell, WA 98021, USA. e-mail: iafonina@nanogen.com

To purchase reprints of this article, contact: Reprints@BioTechniques.com