

## SimpleChIP<sup>®</sup> Human MS4A7 Promoter Primers

500 μl (250 PCR reactions)



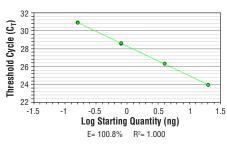
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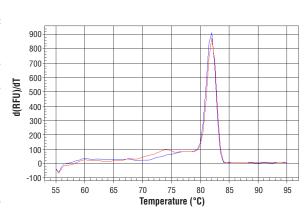
## For Research Use Only. Not For Use In Diagnostic Procedures.

Applications	Species Cross-Reactivity	Primer Anneal/Extension	PCR Product Length
ChIP	н	65°C	120 bp

**Description:** SimpleChIP<sup>®</sup> Human MS4A7 Promoter Primers contain a mix of forward and reverse PCR primers that are specific to a region of the human membrane-spanning 4-domains subfamily A member 7 promoter. These primers can be used to amplify DNA that has been isolated using chromatin immunoprecipitation (ChIP). Primers have been optimized for use in SYBR<sup>®</sup> Green quantitative real-time PCR and have been tested in conjunction with SimpleChIP<sup>®</sup> Enzymatic Chromatin IP Kits #9002 and #9003, and ChIP-validated antibodies from Cell Signaling Technology<sup>®</sup>. MS4A7 is expressed in the monocytic cellular lineage and is involved in mature cellular function and proliferation. IRF-8 plays a role in the regulation of MS4A7 in differentiated THP-1 cells.



SimpleChIP® Human MS4A7 Promoter Primers were tested on DNA isolated from cross-linked cells using the SimpleChIP® Enzymatic Chromatin IP Kit (Magnetic Beads) #9003. Real-time PCR was performed in duplicate on a serial dilution of 2% total input DNA (20 ng, 4 ng, 0.8 ng, and 0.16 ng) using a real-time PCR detection system and SYBR® Green reaction mix. The PCR amplification efficiency (E) and correlation coefficient (R<sup>e</sup>) were calculated based on the corresponding threshold cycle (C<sub>2</sub>) of each dilution sample during 40 cycles of real-time PCR (95°C denaturation for 15 sec, 65°C anneal/extension for 60 sec).



PCR product melting curves were obtained for real-time PCR reactions performed using SimpleChIP® Human MS4A7 Promoter Primers. Data is shown for both duplicate PCR reactions using 20 ng of total DNA. The melt curve consists of 80 melt cycles, starting at 55°C with increments of 0.5°C per cycle. Each peak is formed from the degradation of a single PCR product.

## Entrez-Gene ID #58475 Swiss-Prot Acc. #Q9GZW8

Storage: Supplied in nuclease-free water at a concentration of 5  $\mu$ M (each primer is at a final concentration of 5  $\mu$ M). Store at -20°C.

## Directions for Use:

1. Label the appropriate number of PCR tubes or PCR plates compatible with the model of real-time PCR machine to be used. PCR reactions should be performed in duplicate and should include a tube with no DNA to control for contamination, and a serial dilution of a 2% total input chromatin DNA (undiluted, 1:5, 1:25, 1:125), which is used to create a standard curve and determine amplification efficiency.

2. Add 2  $\mu l$  of the appropriate ChIP DNA sample to each tube or well of the PCR plate.

3. Prepare a master PCR reaction mix as described below. Add enough reagents for two extra reactions to account for loss of volume. Add 18  $\mu$ l of the master PCR reaction mix to each PCR reaction tube or well of the PCR plate.

Reagent	Volume for 1 PCR Reaction (20 $\mu l)$
Nuclease-free H <sub>2</sub> O	6 µl
5 µM SimpleChIP®	Primers 2 µl
2X SYBR <sup>®</sup> Green Reaction Mix	

4. Start the following PCR reaction program:

- a. Initial Denaturation: 95°C for 3 min
- b. Denaturation: 95°C for 15 sec
- c. Anneal and Extension: Primer-specific temp. for 60 sec
- d. Repeat steps b and c for a total of 40 cycles.

5. Analyze quantitative PCR results using software provided with the real-time PCR machine.

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| pchemistry **ChIP**—Chromatin Immunoprecipitation **IF**—Immunofluorescence **F**—Flow cytometry **E-P**—ELISA-Peotide

Applications Key: W-Western IP-Immunoprecipitation IHC-Immunohistochemistry ChIP-Chromatin Immunoprecipitation IF-Immunofluorescence F-Flow cytometry E-P-ELISA-P Species Cross-Reactivity Key: H-human M-mouse R-rat Hm-hamster Mk-monkey Mi-mink C-chicken Dm-D. melanogaster X-Xenopus Z-zebratish B-bovine

Dg—dog Pg—pig Sc—S. cerevisiae Ce—C. elegans Hr—horse

All-all species expected

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Species enclosed in parentheses are predicted to react based on 100% homology.