

SimpleChIP® Mouse RPL30 Intron 2 Primers



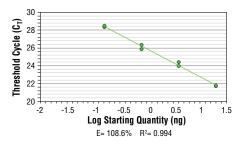
Orders	877-616-CELL (2355)
	orders@cellsignal.com
Support	877-678-TECH (8324)
	info@cellsignal.com
Web	www.cellsignal.com

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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	М	60°C	158 bp

Description: SimpleChIP® Mouse RPL30 Intron 2 Primers contain a mix of forward and reverse PCR primers that are specific to intron 2 of the mouse RPL30 gene. These primers can be used to amplify DNA that has been isolated using chromatin immunoprecipitation (ChIP). Primers have been optimized for use in quantitative real-time PCR using SimpleChIP® Universal qPCR Master Mix #88989. Primers have been tested in conjunction with SimpleChIP® Enzymatic Chromatin IP Kits #9002 and #9003 and ChIPvalidated antibodies from Cell Signaling Technology®. The RPL30 gene is actively transcribed in all cell types and its promoter is highly enriched for histone modifications associated with active transcription, such as histone H3 Lys4 tri-methylation and general histone acetylation. This gene promoter shows very low levels of histone modifications associated with heterochromatin, such as histone H3 Lys9 and Lys27 tri-methylation.



SimpleChIP® Mouse RPL30 Intron 2 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 SimpleChIP® Universal qPCR Master Mix #88989. The PCR amplification efficiency (E) and correlation coefficient (R²) were calculated based on the corresponding threshold cycle (C₁) of each dilution sample during 40 cycles of real-time PCR (95°C denaturation for 15 sec, 60°C anneal/ extension for 60 sec).

Storage: Supplied in nuclease-free water at a concentration of 5 μM (each primer is at a final concentration of 5 μM). 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 μl 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 μ l of the master PCR reaction mix to each PCR reaction tube or well of the PCR plate.

Reagent	Volume for 1 PCR Reaction (2	20 µl)
Nuclease-free H ₂ O		6 µl
5 µM SimpleChIP [®] Primers		2 µl
2X SimpleChIP® Un	iversal qPCR Master Mix #88989	10 µl
4. Ctart the following	a DCD reaction program.	

- 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.

1100 1000 900-800 700d(RFU)/dT 600 -500-400 300-200-100 -0. -100 = 55 60 65 70 75 80 85 90 95 Temperature (°C)

PCR product melting curves were obtained for real-time PCR reactions performed using SimpleChIP[®] Mouse RPL30 Intron 2 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.

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Applications Key: W—Western IP—Immunoprecipitation IHC—Immunohistochemistry ChIP—Chromatin Immunoprecipitation IF—Immunofluorescence F—Flow cytometry E-P—ELISA-Peptide

Species Cross-Reactivity Key: H—human M—mouse R—rat Dg—dog Pg—pig Sc—S. cerevisiae Ce—C. elegans Hr—horse

All-all species expected

Hm—hamster

tad Spacias anclosed in

Mk—monkey Mi—mink C—chicken Dm—D. melanogaster X—Xenopus Z—zebrafish B—bovine

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