

## G-protein coupled receptor 13 (GPR113) / CXCR3

<b>Alternate names:</b>	Fractalkine Receptor, CMK-BRL-1, CMKBLR1, CMKDR1, GPR13, GPRV28, V28, C-X3-C CKR-1, CX3C Chemokine Fractalkine Receptor 1, CX3C Chemokine Receptor 1, G-protein Coupled Receptor 13, Beta Chemokine Receptor-Like 1
<b>Catalog No.:</b>	SP6326CP
<b>Quantity:</b>	50 µg
<b>Background:</b>	The signal transducer and activator of transcription (STAT) family of transcription factors is composed of seven family members (STATs 1, 2, 3, 4, 5A, 5B and 6) that are involved in many cell processes (apoptosis, anti-apoptosis, cell differentiation and proliferation) depending upon the cell type. STATs are activated by ligand binding of mostly cytokine receptors, that receptor associating with a Janus kinase (JAK) family member that phosphorylates a STAT protein, causing it to translocate to the nucleus, where it can bind to DNA and regulate the transcription of particular genes. Phosphorylation on serine 727 by ERK1&2, JNK, or MSK1, together with phosphorylation of tyrosine 705 by JAK1&2, is required for maximum transcriptional activity.
<b>Immunogen:</b>	Chemically synthesized phosphopeptide derived from the region of human STAT3 that contains serine 727. <b>Remarks:</b> This region is conserved between human, mouse and rat.
<b>Format:</b>	<b>State:</b> Liquid Ig fraction <b>Purification:</b> Epitope-specific affinity chromatography. The antibody has been negatively pre-adsorbed using a non-phosphopeptide corresponding to the site of phosphorylation to remove antibody that is reactive with non-phosphorylated STAT3. The final product is generated by affinity chromatography using a STAT3-derived peptide that is phosphorylated at serine 727. <b>Buffer System:</b> Dulbecco's phosphate buffered saline (without Mg <sup>2+</sup> and Ca <sup>2+</sup> ), pH 7.3 (+/- 0.1), 50% glycerol, with 1.0 mg/mL BSA (IgG, protease free) as a carrier, containing 0.05 % sodium azide
<b>Applications:</b>	Western blot (1:1000). Positive Control Used: 3T3-L1 adipocytes +/- Leukemia Inhibitory Factor (LIF). Other applications not tested. Optimal dilutions are dependent on conditions and should be determined by the user.
<b>Specificity:</b>	This antibody detects STAT3. <b>Species:</b> Human, Mouse, Rat. Others not tested.
<b>General References:</b>	Visconti, R., et al. (2000) Importance of the MKK6/p38 pathway for interleukin-12-induced STAT4 serine phosphorylation and transcriptional activity. <i>Blood</i> 96(5):1844-1852. Ortmann, R.A., et al. (2000) Janus kinases and signal transducers and activators of transcription: their roles in cytokine signaling, development and immunoregulation.

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Arthritis Res. 2(1):16-32.

Ihle, J.N. (1996) STATs: signal transducers and activators of transcription. Cell 84(3):331-334.

Wen, Z. (1995) Maximal activation of transcription by STAT1 and STAT3 requires both tyrosine and serine phosphorylation. Cell 82(2):241-250.

#### Protocols:

##### Western Blotting Procedure

1. Lyse approximately 10e7 cells in 0.5 mL of ice cold Cell Lysis Buffer (formulation provided below). This buffer, a modified RIPA buffer, is suitable for recovery of most proteins, including membrane receptors, cytoskeletal-associated proteins, and soluble proteins. Other cell lysis buffer formulations, such as Laemmli sample buffer and Triton-X 100 buffer, are also compatible with this procedure. Additional optimization of the cell stimulation protocol and cell lysis procedure may be required for each specific application.
2. Remove the cellular debris by centrifuging the lysates at 14,000 x g for 10 minutes. Alternatively, lysates may be ultracentrifuged at 100,000 x g for 30 minutes for greater clarification.
3. Carefully decant the clarified cell lysates into clean tubes and determine the protein concentration using a suitable method, such as the Bradford assay. Polypropylene tubes are recommended for storing cell lysates.
4. React an aliquot of the lysate with an equal volume of 2x Laemmli Sample Buffer (125 mM Tris, pH 6.8, 10% glycerol, 10% SDS, 0.006% bromophenol blue, and 130 mM dithiothreitol [DTT]) and boil the mixture for 90 seconds at 100°C.
5. Load 10-30 µg of the cell lysate into the wells of an appropriate single percentage or gradient minigel and resolve the proteins by SDS-PAGE.
6. In preparation for the Western transfer, cut a piece of PVDF membrane slightly larger than the gel. Soak the membrane in methanol for 1 minute, then rinse with ddH<sub>2</sub>O for 5 minutes. Alternatively, nitrocellulose may be used.
7. Soak the membrane, 2 pieces of Whatman paper, and Western apparatus sponges in transfer buffer (formulation provided below) for 2 minutes.
8. Assemble the gel and membrane into the sandwich apparatus.
9. Transfer the proteins at 140 mA for 60-90 minutes at room temperature.
10. Following the transfer, rinse the membrane with Tris buffered saline for 2 minutes.
11. Block the membrane with blocking buffer (formulation provided below) for one hour at room temperature or overnight at 4°C.
12. Incubate the blocked blot with primary antibody at a 1:1000 dilution in Tris buffered saline supplemented with 3% BSA and 0.1% Tween 20 overnight at 4°C or for one hour at room temperature.
13. Wash the blot with several changes of Tris buffered saline supplemented with 0.1% Tween 20.
14. Detect the antibody band using an appropriate secondary antibody, such as goat F(ab)<sub>2</sub> anti-rabbit IgG alkaline phosphatase conjugate or goat F(ab)<sub>2</sub> anti-rabbit IgG horseradish peroxidase conjugate in conjunction with your chemiluminescence reagents and instrumentation.

##### Cell Lysis Buffer Formulation:

10 mM Tris, pH 7.4  
100 mM NaCl  
1 mM EDTA  
1 mM EGTA  
1 mM NaF  
20 mM Na<sub>4</sub>P<sub>2</sub>O<sub>7</sub>  
2 mM Na<sub>3</sub>VO<sub>4</sub>

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0.1% SDS  
0.5% sodium deoxycholate  
1% Triton-X 100  
10% glycerol  
1 mM PMSF (made from a 0.3 M stock in DMSO)  
or 1 mM AEBSF (water soluble version of PMSF)  
60 µg/mL aprotinin  
10 µg/mL leupeptin  
1 µg/mL pepstatin  
(alternatively, protease inhibitor cocktail such as Sigma Cat. # P2714 may be used)

**Transfer Buffer Formulation:**

2.4 gm Tris base  
14.2 gm glycine  
200 mL methanol  
Q.S. to 1 liter, then add 1 mL 10% SDS.  
Cool to 4°C prior to use.

**Tris Buffered Saline Formulation:**

20 mM Tris-HCl, pH 7.4  
0.9% NaCl

**Blocking Buffer Formulation:**

100 mL Tris buffered saline  
5 gm BSA  
0.1 mL Tween 20

**Peptide Competition Experiment**

The specificity of a Phosphorylation Site Specific Antibody (PSSA) in each experimental system can be confirmed through peptide competition. In this technique, aliquots of antibody are pre-incubated with peptide containing the sequence of the phosphopeptide immunogen used to raise the PSSA and the corresponding non-phosphopeptide. Following preincubation with the peptide, each antibody preparation is then used as a probe in antibody-based detection methods, such as Western blotting, immunocytochemistry, flow cytometry, or ELISA. With a PSSA specific for the phosphorylated target protein, pre-incubation with an excess of peptide containing the sequence of the phosphopeptide immunogen will block all antigen binding sites, while pre-incubation with the corresponding non-phosphopeptide will not affect the antibody.

In performing the Peptide Competition Experiment, it is important to note that the optimal dilutions of both antibody and peptide should be determined empirically for each specific application. The optimal dilution of antibody in these procedures is below saturating, as determined by previous experiments in your system.

The optimal dilution of peptide used in these procedures will depend on the overall affinity or avidity of the antibody, as well as the quantity of the target antigen. A 50-150 fold molar excess of peptide to antibody is found to be effective for most peptide competition experiments.

In the example presented below, the PSSA is used as a dilution of 1:1000 and the peptides are used at a concentration of 333 nM. The total volume of the phosphopeptide and nonphosphopeptide pre-incubated antibody preparations is 2 mL, sufficient for probing

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Western blot strips, as well as for use in other antibody-based detection methods. Under these conditions, the molar excess of peptide to antibody is  $> / = 50$ .

**Procedure:**

1. Prepare three identical test samples, such as identical PVDF or nitrocellulose strips to which the protein of interest has been transferred. The test samples should be blocked using a blocking buffer, such as Tris buffered saline supplemented with 0.1% Tween 20, and either 5% BSA or 5% non-fat dried milk.
2. Prepare 6.5 mL of working antibody stock solution (1:1000 in this example) by adding 6.5  $\mu$ L of antibody stock solution to 6.5 mL of buffer containing blocking protein, such as TBS supplemented with 0.1% Tween 20, and either 3% BSA or 3% non-fat dried milk.
3. Apportion the unused PSSA into working aliquots and store at  $-20^{\circ}\text{C}$  for future use (the stock PSSA contains 50% glycerol and will not freeze at this temperature).
4. Allow the lyophilized control peptides to reach room temperature, ideally under desiccation.
5. Reconstitute each of the control peptides to a concentration of 66.7  $\mu\text{M}$  with nanopure water. (i.e. for a peptide with a molecular mass of 1500, reconstitution with 1 mL water yields a solution with a concentration of 66.7  $\mu\text{M}$ ).
6. Apportion the unused reconstituted peptide solutions into working aliquots and store at  $-20^{\circ}\text{C}$  for future use.
7. Label 3 test tubes as follows:
  - tube 1: water only no peptide control
  - tube 2: phosphopeptide
  - tube 3: non-phosphopeptide
8. Into each tube, pipette the following components
  - tube 1: 2 mL diluted PSSA solution plus 10  $\mu$ L nanopure water
  - tube 2: 2 mL diluted PSSA solution plus 10  $\mu$ L phosphopeptide
  - tube 3: 2 mL diluted PSSA solution plus 10  $\mu$ L non-phosphopeptide
9. Incubate the three tubes for 30 minutes at room temperature with gentle rocking. During this incubation, the peptides have the chance to bind to the combining site of the antibody.
10. At the end of the incubation step, transfer the contents of each of the three tubes to clean reaction vessels containing one of the three identical test samples.

**For Western blotting strips:**

Incubate the strips with the pre-incubated antibody preparations for 1 hour at room temperature or overnight at  $4^{\circ}\text{C}$ .

Wash each strip four times, five minutes each, to remove unbound antibody. Transfer each strip to a new solution containing a labeled secondary antibody [e.g., goat F(ab)<sub>2</sub> anti-rabbit IgG alkaline phosphatase conjugate or goat F(ab)<sub>2</sub> anti-rabbit IgG horseradish peroxidase conjugate].

Remove unbound secondary antibody by thorough washing, and develop the signal using your chemiluminescent reagents and instrumentation.

The signal obtained with antibody incubated with the "Water Only, No Peptide Control" (Tube 1), represents the maximum signal in the assay. This signal should be eliminated by preincubation with the "Phosphopeptide" (Tube 2), while pre-incubation with the "Non-Phosphopeptide" (Tube 3) should not impact the signal. If the "Phosphopeptide" only partially eliminates the signal, repeat the procedure using twice the volume of water or peptide solutions listed in Step 8. If partial competition is seen following pre-incubation with the "Non-Phosphopeptide", repeat the procedure using half the volumes of water or peptide solutions listed in Step 8.

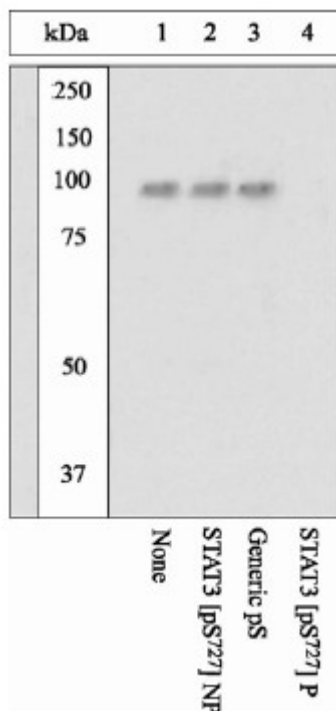
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**Pictures:**



**Peptide Competition** Extracts prepared from 3T3L1 cells stimulated with LIF were resolved by SDS-PAGE on a 10% polyacrylamide gel and transferred to PVDF. Membranes were blocked with a 5% BSA-TBST buffer overnight at 4°C, then were incubated with STAT3 [pS727] antibody for two hours at room temperature in a 1% BSA-TBST buffer, following prior incubation with: no peptide (1), the non-phosphopeptide corresponding to the immunogen (2), a generic phosphoserine-containing peptide (3), or, the phosphopeptide immunogen (4). After washing, membranes were incubated with goat F(ab')<sub>2</sub> anti-rabbit IgG HRP-conjugate and bands were detected using the Pierce SuperSignal™ method. The data show that only the peptide corresponding to STAT3 [pS727] blocks the antibody signal, demonstrating the specificity of the antibody.

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