

DESCRIPTION

Source Mouse myeloma cell line, NS0-derived mouse Mast Cell Protease-11/Prss34 protein
Met20-Ser318, with a C-terminal 10-His tag
Accession # Q80UR4

N-terminal Sequence Analysis Met20

Structure / Form Pro form

Predicted Molecular Mass 34 kDa

SPECIFICATIONS

SDS-PAGE 48 kDa doublet, reducing conditions

Activity Measured by its ability to cleave a colorimetric peptide substrate, N-carbobenzyloxy-Arg-ThioBenzyl ester (Z-R-SBzl), in the presence of 5,5'Dithio-bis (2-nitrobenzoic acid) (DTNB). Edwards, K.M. *et al.* (1999) J. Biol. Chem. **274**:30468.
The specific activity is >20,000 pmol/min/μg, as measured under the described conditions.

Endotoxin Level <1.0 EU per 1 μg of the protein by the LAL method.

Purity >95%, by SDS-PAGE under reducing conditions and visualized by silver stain.

Formulation Supplied as a 0.2 μm filtered solution in MES and NaCl. See Certificate of Analysis for details.

Activity Assay Protocol

- Materials**
- Activation Buffer: 50 mM Tris, 0.15 M NaCl, 10 mM CaCl₂, pH 7.5 (TCN)
 - Assay Buffer: 50 mM Tris, pH 8.0
 - Recombinant Mouse Mast Cell Protease-11/Prss34 (rmMCP-11) (Catalog # 2857-SE)
 - Bacterial Thermolysin (Thermolysin) (Catalog # 3097-ZN)
 - 1,10 Phenanthroline (Sigma, Catalog # 320056), 0.6 M stock in DMSO
 - Substrate: Z-Arg-SBzl (SM Biochemicals, Catalog # SMSB01), 10 mM in DMSO
 - 5,5'Dithio-bis(2-nitrobenzoic acid) (DTNB) (Sigma, Catalog # D-8130)
 - 96-well Clear Plate (Costar, Catalog # 92592)
 - Plate reader (Model: SpectraMax Plus by Molecular Devices) or equivalent

- Assay**
1. Dilute rmMCP-11 to 200 μg/mL with Activation Buffer.
 2. Dilute Thermolysin to 0.4 μg/mL with Activation Buffer.
 3. Mix equal volumes of 200 μg/mL rmMCP-11 and 0.4 μg/mL Thermolysin for final concentrations of 50 μg/mL and 0.2 μg/mL, respectively.
 4. Incubate at 37 °C for 30 minutes.
 5. Stop the reaction with 10 mM 1,10 Phenanthroline.
 6. Dilute activated rmMCP-11 to 0.1 ng/μL in Assay Buffer.
 7. Dilute substrate to 200 μM in Assay Buffer with 200 μM of DTNB.
 8. Load 50 μL of the 0.1 ng/μL rmMCP-11 into plate, and start the reaction by adding 50 μL of the substrate/DTNB mixture to wells. Include a Substrate Blank containing 50 μL Assay Buffer and 50 μL substrate mix without any rmMCP-11.
 9. Read in kinetic mode for 5 minutes at an absorbance of 405 nm.
 10. Calculate specific activity:

$$\text{Specific Activity (pmol/min/}\mu\text{g)} = \frac{\text{Adjusted } V_{\text{max}}^* (\text{OD/min}) \times \text{well volume (L)} \times 10^{12} \text{ pmol/mol}}{\text{ext. coeff}^{**} (\text{M}^{-1}\text{cm}^{-1}) \times \text{path corr.}^{***} (\text{cm}) \times \text{amount of enzyme } (\mu\text{g})}$$

*Adjusted for Substrate Blank

**Using the extinction coefficient 13260 M⁻¹cm⁻¹

***Using the path correction 0.32 cm

Note: the output of many spectrophotometers is in mOD

Final Assay Conditions Per Well:

- rmMCP-11: 0.005 μg
- DTNB: 100 μM
- Substrate: 100 μM

PREPARATION AND STORAGE

Shipping The product is shipped with dry ice or equivalent. Upon receipt, store it immediately at the temperature recommended below.

Stability & Storage Use a manual defrost freezer and avoid repeated freeze-thaw cycles.

- 6 months from date of receipt, -20 to -70 °C as supplied.
- 3 months, -20 to -70 °C under sterile conditions after opening.

BACKGROUND

Mast Cell Protease-11 (MCP-11) is encoded by Prss34, one of 13 genes on mouse chromosome 17A3.3 that correspond to functional trypsin-like serine proteases (1). The deduced amino acid sequence of mouse MCP-11 consists of 318 residues with a signal peptide (residues 1 to 19), a pro region (residue 20 to 34), and a catalytic domain (35 to 318). The mRNA is preferentially expressed in spleen and bone marrow. The mouse MCP-11 (residues 20 to 318) was expressed in the NS0 cells with a foreign signal peptide. After being treated with thermolysin, the purified enzyme is active against a peptide substrate described in the Activity Assay Protocol. Apparently, the human gene corresponding to Prss34 encodes a protein that is not enzymatically active due to a mutation that leads to a premature translation termination codon.

References:

1. Wong, G.W. *et al.* (2004) *J. Biol. Chem.* **279**:2438.