

**DESCRIPTION**

**Source** *E. coli*-derived  
Met1-Asp187, with a C-terminal 6-His tag  
Accession # P00374

**N-terminal Sequence Analysis** Met1 & Val2

**Predicted Molecular Mass** 22 kDa

**SPECIFICATIONS**

**SDS-PAGE** 23 kDa, reducing conditions

**Activity** Measured by the reduction of dihydrofolic acid (DHF).  
The specific activity is >5,500 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 Colloidal Coomassie® Blue stain at 5 μg per lane.

**Formulation** Supplied as a 0.2 μm filtered solution in Tris, NaCl, Glycerol, Brij-35 and DTT. See Certificate of Analysis for details.

**Activity Assay Protocol**

- Materials**
- Assay Buffer: 50 mM MES, 25 mM Tris, 100 mM NaCl, 25 mM Ethanolamine, 2 mM DTT
  - Recombinant Dihydrofolate Reductase/DHFR (rhDHFR) (Catalog # 8456-DR)
  - β-Nicotinamide adenine dinucleotide phosphate reduced, tetrasodium salt (β-NADPH) (Sigma, Catalog # N7505), 10 mM stock in deionized water
  - Dihydrofolic acid (DHF) (Sigma, Catalog # D7006), 10 mM stock in Assay Buffer + 4.5 mM NaOH
  - 96-well Clear Plate (Catalog # DY990)
  - Plate Reader (Model: SpectraMax Plus by Molecular Devices) or equivalent

- Assay**
1. Dilute rhDHFR to 1 μg/mL in Assay Buffer.
  2. Prepare a Substrate Mixture containing 0.2 mM DHF and 0.25 mM β-NADPH in Assay Buffer.
  3. Load 50 μL of 1 μg/mL rhDHFR into a plate, and start the reaction by adding 50 μL of Substrate Mixture. Include a Substrate Blank containing 50 μL of Assay Buffer and 50 μL of Substrate Mixture.
  4. Read at an absorbance of 339 nm in kinetic mode for 5 minutes.
  5. Calculate specific activity:

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

\*Adjusted for Substrate Blank

\*\*Using the extinction coefficient 6220 M<sup>-1</sup>cm<sup>-1</sup>

\*\*\*Using the path correction 0.32 cm

Note: the output of many spectrophotometers is in mOD.

- Final Assay Conditions**
- Per Well:
- rhDHFR: 0.05 μg
  - DHF: 0.1 mM
  - β-NADPH: 0.125 mM

**PREPARATION AND STORAGE**

**Shipping** The product is shipped with polar packs. 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**

Dihydrofolate Reductase (DHFR) is an approximately 21 kDa enzyme that catalyzes the NADPH-dependent reduction of dihydrofolate to tetrahydrofolate, which is crucial for the synthesis of purines, thymidylate, and certain amino acids (1-4). Structurally, DHFR consists of an eight-stranded  $\beta$ -sheet and four  $\alpha$ -helices (2). Human DHFR shares 90% amino acid identity with mouse and rat DHFR. In addition to acting as a cofactor, NADPH protects DHFR from degradation *in vitro* and in a cellular context (5-8). DHFR protein binds to its own mRNA to self-regulate translation (9-12). The enzymatic activity and subcellular localization of DHFR are at least partially regulated by post-translational modification. Mono-ubiquitination of DHFR by MDM2 reduces the activity of DHFR (13). DHFR is primarily cytoplasmic, but it has been shown to translocate to the nucleus in a SUMO-dependent manner (14). Due to its role in nucleotide biosynthesis, DHFR has long been a therapeutic target for the treatment of various cancers and infectious diseases (15, 16).

**References:**

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