

## **Product datasheet for TP720732M**

## OriGene Technologies, Inc.

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## HMGB2 (NM 002129) Human Recombinant Protein

**Product data:** 

**Product Type:** Recombinant Proteins

**Description:** Purified recombinant protein of Human high mobility group box 2 (HMGB2), transcript

variant 1

Species: Human Expression Host: HEK293

Expression cDNA Clone

or AA Sequence:

Gly2-Glu209

Tag: C-His

Predicted MW: 25.07 kDa

Concentration: lot specific

**Purity:** >95% as determined by SDS-PAGE and Coomassie blue staining

**Buffer:** Provided lyophilized from a 0.2 μm filtered solution of 20 mM Tris-HCl, 150 mM NaCl

Endotoxin: Endotoxin level is < 0.1 ng/μg of protein (< 1 EU/μg)

**Reconstitution Method:** Always centrifuge tubes before opening. Do not mix by vortex or pipetting. Dissolve the

lyophilized protein in ddH2O. It is not recommended to reconstitute a concentration less than 100  $\mu$ g/ml. Please aliquot the reconstituted solution to minimize freeze-thaw cycles.

Storage: Store at -80°C.

Stability: Stable for at least 6 months from date of receipt under proper storage and handling

conditions.

**RefSeq:** <u>NP 002120</u>

 Locus ID:
 3148

 UniProt ID:
 P26583

RefSeq Size: 1527 Cytogenetics: 4q34.1

RefSeg ORF: 627

Synonyms: HMG2





## HMGB2 (NM\_002129) Human Recombinant Protein - TP720732M

**Summary:** 

This gene encodes a member of the non-histone chromosomal high mobility group protein family. The proteins of this family are chromatin-associated and ubiquitously distributed in the nucleus of higher eukaryotic cells. In vitro studies have demonstrated that this protein is able to efficiently bend DNA and form DNA circles. These studies suggest a role in facilitating cooperative interactions between cis-acting proteins by promoting DNA flexibility. This protein was also reported to be involved in the final ligation step in DNA end-joining processes of DNA double-strand breaks repair and V(D)J recombination. [provided by RefSeq, Jul 2008]

**Protein Families:** 

Druggable Genome, Transcription Factors