

## 32-4150: Recombinant Human Macrophage Erythroblast Attacher

**Alternative Name :** Macrophage Erythroblast Attacher, GID Complex Subunit 9 FYV10 Homolog, Lung Cancer-Related Protein 10, Proliferation-Inducing Gene 5, Human Lung Cancer Oncogene 10 Protein, Cell Proliferation-Inducing Gene 5 Protein, Erythroblast Macrophage Protein,

### Description

Source : Escherichia Coli. MAEA Human Recombinant (isoform 1) produced in E.coli is a single, non-glycosylated polypeptide chain containing 419 amino acids (1-396) and having a molecular mass of 47.7kDa. MAEA is fused to a 23 amino acid His-tag at N-terminus & purified by proprietary chromatographic techniques. MAEA is a 396aa widely expressed adhesion protein which has 5 alternatively spliced isoforms. MAEA holds one CTLH domain and one LisH domain. MAEA forms a complex with F-actin, which takes part in regulating actin distribution in erythroblasts and macrophages and is known to assist in cell division and nuclear architecture. MAEA is confined with condensed chromatin at prophase, nuclear spindle poles at metaphase and in the contractile ring throughout cytokinesis phase and telophase.

### Product Info

**Amount :** 10 µg  
**Purification :** Greater than 85% as determined by SDS-PAGE.  
**Content :** The MAEA solution contains 20mM Tris-HCl buffer (pH 8.0), 1M Urea and 10% glycerol.  
**Storage condition :** Store at 4°C if entire vial will be used within 2-4 weeks. Store, frozen at -20°C for longer periods of time. For long term storage it is recommended to add a carrier protein (0.1% HSA or BSA). Avoid multiple freeze-thaw cycles.  
**Amino Acid :** MGSSHHHHHH SSGLVPRGSH MGSMVQESA AQLSMTLVQ EYPTLVKPYE TLNKRFRRAQ  
 KNIDRETSV TMVVAELEKT LSGCPAVDSV VSLLDGVVEK LSVLKRKAVE SIQAEDSAK LCKRRIEHLK  
 EHSSDQAAAA SVWKRKRMDR MMVEHLLRCG YYNTAVKLAR QSGIEDLVNI EMFLTAKVE ESLETERAT  
 CLAWCHDNKS RLRKMKSCLE FSLRIQEFIE LIRQNKRLDA VRHARKHFSQ AEGSQLDEV R QAMGMLAFPP  
 DTHISPYKDL LDPARWRMLI QQFRYDNYRL HQLGNNSVFT LTLQAGLSAI KTPQCYKEDG SSKSPDCPVC  
 SRSLNKLAP LPMAHCANR LVCKISGDVM NENPPMMLP NGVYGYNSL LSIRQDDKVV CPRTKEVFHF  
 SQAQKYYIM

