



# hnRNP U

## Mouse monoclonal Antibody

### #53118

**Catalog Number:** 53118

**Amount:** 100µg/100µl

**Swiss-Prot No. :**Q00839

**Gene name:**hnrnpu

**Gene id:**3192

**Clone Number:** 1B11-H2-B7

**Form of Antibody:**Purified mouse monoclonal in buffer containing 0.1M Tris-Glycine (pH 7.4, 150 mM NaCl) with 0.2% sodium azide, 50%.glycerol

**Storage/Stability:** Store at -20°C/1 year

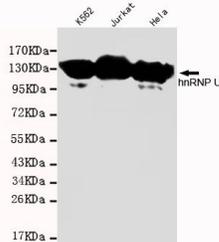
**Immunogen:** Purified recombinant human hnRNP U protein fragments expressed in E.coli

**Purification:** affinity-chromatography

**Specificity/Sensitivity:**This antibody detects endogenous levels of hnRNP U and does not cross-react with related proteins

**Reactivity:** Human,

**Applications:** Predicted MW: 110kd WB: 1:1000



Western blot detection of hnRNP U in K562, Jurkat and HeLa cell lysates and using hnRNP U mouse mAb(1:1000diluted). Predicted band size: 110KDa. Observed band size: 110KDa.

**Background:**This gene belongs to the subfamily of ubiquitously expressed heterogeneous nuclear ribonucleoproteins (hnRNPs). The hnRNPs are RNA binding proteins and they form complexes with heterogeneous nuclear RNA (hnRNA). These proteins are associated with pre-mRNAs in the nucleus and appear to influence pre-mRNA processing and other aspects of mRNA metabolism and transport. While all of the hnRNPs are present in the nucleus, some seem to shuttle between the nucleus and the cytoplasm. The hnRNP proteins have distinct nucleic acid binding properties. The protein encoded by this gene contains a RNA binding domain and scaffold-associated region (SAR)-specific bipartite DNA-binding domain. This protein is also thought to be involved in the packaging of hnRNA into large ribonucleoprotein complexes. During apoptosis, this protein is cleaved in a caspase-dependent way. Cleavage occurs at the SALD site, resulting in a loss of DNA-binding activity and a concomitant detachment of this protein from nuclear structural sites. But this cleavage does not affect the function of the encoded protein in RNA metabolism. At least two alternatively spliced transcript variants have been identified for this gene.