

Product datasheet for TP724138

OriGene Technologies, Inc.

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Human CD117 Protein, His Tag

Product data:

Product Type: Recombinant Proteins

Description: Human CD117 Protein, His Tag

Expression Host: HEK293
Tag: C-6×His

Predicted MW: The protein has a predicted molecular mass of 56.5 kDa after removal of the signal

peptide. The apparent molecular mass of CD117-His is approximately 70-100 kDa due to

glycosylation.

Purity: The purity of the protein is greater than 95% as determined by SDS-PAGE and Coomassie

blue staining.

Reconstitution Method: Lyophilized from sterile PBS, pH 7.4. Normally 5 % - 8% trehalose is added as protectants

before lyophilization.

Storage: Store at -20°C to -80°C for 12 months in lyophilized form. After reconstitution, if not intended

for use within a month, aliquot and store at -80°C (Avoid repeated freezing and thawing).

Lyophilized proteins are shipped at ambient temperature.

Stability: 12 months from date of despatch
Synonyms: C-Kit, CD117, MASTC, PBT, SCFR, KIT

Summary: This gene encodes a receptor tyrosine kinase. This gene was initially identified as a homolog

of the feline sarcoma viral oncogene v-kit and is often referred to as proto-oncogene c-Kit.

The canonical form of this glycosylated transmembrane protein has an N-terminal

extracellular region with five immunoglobulin-like domains, a transmembrane region, and an intracellular tyrosine kinase domain at the C-terminus. Upon activation by its cytokine ligand, stem cell factor (SCF), this protein phosphorylates multiple intracellular proteins that play a role in in the proliferation, differentiation, migration and apoptosis of many cell types and thereby plays an important role in hematopoiesis, stem cell maintenance, gametogenesis, melanogenesis, and in mast cell development, migration and function. This protein can be a

membrane-bound or soluble protein. Mutations in this gene are associated with gastrointestinal stromal tumors, mast cell disease, acute myelogenous leukemia, and

piebaldism. Multiple transcript variants encoding different isoforms have been found for this

gene. [provided by RefSeq, May 2020]

