

Product Data Sheet

EB3/MAPRE3 Protein, Human (His)

Cat. No.:	HY-P75275
Synonyms:	Microtubule-associated protein RP/EB family member 3; EBF3; EB3; RP3; MAPRE3
Species:	Human
Source:	E. coli
Accession:	Q9UPY8-1 (M1-Y281)
Gene ID:	22924
Molecular Weight:	Approximately 34 kDa

PROPERTIES	
AA Sequence	MAVNVYSTSVTSENLSRHDMLAWVNDSLHLNYTKIEQLCSGAAYCQFMDMLFPGCVHLRKVKFQAKLEHEYIHNFKVLQAAFKKMGVDKIIPVEKLVKGKFQDNFEFIQWFKKFFDANYDGKDYNPLLARQGQDVAPPPNPGDQIFNKSKKLIGTAVPQRTSPTGPKNMQTSGRLSNVAPPCILRKNPPSARNGGHETDAQILELNQQLVDLKLTVDGLEKERDFYFSKLRDIELICQEHESENSPVISGIIGILYATEEGFAPPEDDEIEEHQQEDQDEYYYYY
Appearance	Lyophilized powder
Formulation	Lyophilized from a 0.2 μm filtered solution of PBS, pH 7.4.
Endotoxin Level	<1 EU/µg, determined by LAL method.
Reconsititution	It is not recommended to reconstitute to a concentration less than 100 μg/mL in ddH ₂ O. For long term storage it is recommended to add a carrier protein (0.1% BSA, 5% HSA, 10% FBS or 5% Trehalose).
Storage & Stability	Stored at -20°C for 2 years. After reconstitution, it is stable at 4°C for 1 week or -20°C for longer (with carrier protein). It is recommended to freeze aliquots at -20°C or -80°C for extended storage.
Shipping	Room temperature in continental US; may vary elsewhere.

DESCRIPTION	
Background	EB3, a plus-end tracking protein (+TIP), intricately modulates the dynamics of the microtubule cytoskeleton by binding to the plus-end of microtubules. Functionally, EB3 serves as a promoter of microtubule growth and potentially contributes to spindle function by stabilizing microtubules and anchoring them at centrosomes. Additionally, it acts as a crucial regulate of minus-end microtubule organization, participating in the recruitment of CAMSAP2 to the Golgi apparatus through

interaction with the complex formed by AKAP9 and PDE4DIP. This interaction is pivotal for tethering non-centrosomal minus-end microtubules to the Golgi, a process essential for polarized cell movement. EB3 forms homodimers and heterodimers with MAPRE1, binding both monomeric and polymerized GTP-bound tubulin. Its intricate network of interactions includes APC2, DCTN1, SRCIN1, CLIP1, SLAIN2, SLAIN1, AKAP9, and PDE4DIP, underscoring its multifaceted role in microtubule dynamics and cellular organization.

Caution: Product has not been fully validated for medical applications. For research use only.

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