Elabscience®

Recombinant Human F9 Protein(His Tag)

Catalog Number: PDMH100254

Note: Centrifuge before opening to ensure complete recovery of vial contents.

Description		
Species	Human	
Source	Mammalian-derived Human F9 protein Met1-Thr461, with an C-terminal His	
Calculated MW	50.6 kDa	
Observed MW	60-80 kDa	
Accession	P00740	
Bio-activity	Not validated for activity	
Properties		
Purity	> 90% as determined by reducing SDS-PAGE.	
Endotoxin	< 1.0 EU/mg of the protein as determined by the LAL method	
Storage	Generally, lyophilized proteins are stable for up to 12 months when stored at -20 to - $^{\circ}$ C. Reconstituted protein solution can be stored at 4-8°C for 2-7 days. Aliquots of reconstituted samples are stable at < -20°C for 3 months.	
Shipping	This product is provided as lyophilized powder which is shipped with ice packs.	
Formulation	Lyophilized from a 0.2 μ m filtered solution in PBS with 5% Trehalose and 5%	
	Mannitol.	
Reconstitution	It is recommended that sterile water be added to the vial to prepare a stock solution of	
	0.5 mg/mL. Concentration is measured by UV-Vis.	

Data

	М	R
kDa		
80	-	818
60	-	***
40	-	
30	-	
20	-	
12		

SDS-PAGE analysis of Human F9 proteins, 2 µg/lane of Recombinant Human F9 proteins was resolved with an SDS-PAGE under reducing conditions, showing bands at 50.6 KD

Background

This gene encodes vitamin K-dependent coagulation factor IX that circulates in the blood as an inactive zymogen. This factor is converted to an active form by factor XIa, which excises the activation peptide and thus generates a heavy chain and a light chain held together by one or more disulfide bonds. The role of this activated factor IX in the blood coagulation cascade is to activate factor X to its active form through interactions with an Ca+2 ions, membrane phospholipids, and factor VIII. Alterations of this gene, including point mutations, insertions and deletions, cause factor IX deficiency, which is a recessive X-linked disorder, also called hemophilia B or Christmas disease. Alternative splicing results in multiple transcript variants encoding different isoforms that may undergo similar proteolytic processing.

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