for research use only Certificate of Analysis

Product Name	Recombinant Human Cytochrome b-245 heavy chain (CYBB), partial				
Catalog Number	AAA18493				
Expression host	E.coli				
Tag Info	N-terminal 6xHis-tagged				
Buffer	10 mM Tris-HCl, 1 mM EDTA, pH 8.0, 50% glycerol				
Batch Number	YA04543b1g5				
Nature	Human CYBB-(AA 283-570)- P04839 -Partial Protein				
Purification	Affinity purified using IMAC				
Recommended Storage	Short term	2 to 8 ℃, c	of receipt		
	Long term	-20 to -80 °C, six months from the date of receipt			
Form	Liquid				
Date of detection	2024.09.20				
Test Items	Specifications			Results	
Appearance	Clear Solution			pass	
Concentration	0.1-5 mg/ml, by the Bradford Method.			0.5 mg/ml	
Purity	≥90%,by SDS-PAGE quandensitometry by Coomass Staining.		kDa M 116.0 66.2 45.0 35.0	90%	
Molecular Weight	Predicted band size: 37.3 kDa		25.0 18.4 14.4	Observed band size: 37 kDa	

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Electrophoretic	(Tris-Glycine gel) Discontinuou SDS-PAGE (reduced) with 5% enrichment gel and 15% separation
parameters	gel.
Aseptic	
Processing	Not done
Endotoxin Level	Untreated
Activity	Not tested
Conclusion	pass

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Datasheet

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Catalog Number	AAA18493		
Expression host	E.coli		
Tag Info	N-terminal 6xHis-tagged		
Buffer	10 mM Tris-HCl, 1 mM EDTA, pH 8.0, 50% glycerol		
Storage	Store at -20°C, for extended storage, conserve at -20°C or -80°C.		
Notes	Repeated freezing and thawing is not recommended. Store working aliquots at 4°C for up to one week.		
Relevance	Critical component of the membrane-bound oxidase of phagocytes that generates superoxide. It is the terminal component of a respiratory chain that transfers single electrons from cytoplasmic NADPH across the plasma membrane to molecular oxygen on the exterior. Also functions as a voltage-gated proton channel that mediates the H(+) currents of resting phagocytes. It participates in the regulation of cellular pH and is blocked by zinc.		
AA sequence	ERLVRFWRSQQKVVITKVVTHPFKTIELQMKKKGFKMEVGQYIFVKCPKVSKLEWHPFTLTSAPEEDFFSIHIRIV GDWTEGLFNACGCDKQEFQDAWKLPKIAVDGPFGTASEDVFSYEVVMLVGAGIGVTPFASILKSVWYKYCN NATNLKLKKIYFYWLCRDTHAFEWFADLLQLLESQMQERNNAGFLSYNIYLTGWDESQANHFAVHHDEEKD VITGLKQKTLYGRPNWDNEFKTIASQHPNTRIGVFLCGPEALAETLSKQSISNSESGPRGVHFIFNKENF		