

Recombinant Enzyme Product Specification Sheet

Cat. No.:	PRO-E0355	add this product to cart
LOT:	2012-0355-1	view other feruloyl esterases
Activity:	Feruloyl esterase	
Synonyms:	Ferulic acid esterase; hydroxycinnamoyl esterase; FAE-III; cinnamoyl ester hydrolase; FAEA; cinnAE; FAE-I; FAE-II; 4-hydroxy-3-methoxycinnamoyl-sugar hydrolase	
Nomenclature:	CAZy [CE1, carbohydrate esterase family 1]	
Source organism:	<i>Clostridium thermocellum</i> DSM 1313	
Enzyme Commission No.:	3.1.1.73	
Activity:	1.501 U/mL	} (37°C; pH 6.5; 187 µM methyl ferulate)
Specific activity:	0.453 U/mg	
Purity:	>95% as judged by SDS-PAGE	
Form and storage:	Supplied in 3.2 M ammonium sulphate. Store at 4°C (shipped at room temperature)	
pH optimum:	-	
Temperature optimum:	≥ 37°C	
[Protein]:	3.317 mg/mL	
Sequence length:	261 amino acids (view sequence)	
Accession No.:	E6UT93	
Molecular weight:	32092.0 Da	(theoretical)
	~ 30000 Da	(observed by SDS-PAGE)
	-	(observed by mass spectrometry)
Biological function:	Likely to hydrolyse natural esters such as FAXX, FAX3 and PAX3. Feruloyl-polysaccharide + H ₂ O = ferulate + polysaccharide	
Potential application(s):	Biomass conversion , carbohydrate research	
Comments:	This esterase is the N-terminal CE1 catalytic module of modular enzyme also containing a C-terminal CBM6 module and dockerin sequences. All microbial ferulate esterases are secreted into the culture medium. They are sometimes called hemicellulase accessory enzymes, since they help xylanases and pectinases to break down plant cell wall hemicellulose	

Usage: Agitate vial sufficiently to fully homogenise enzyme precipitate before use

Assay: One unit is defined as the amount of enzyme required to release 1 μmol of ferulic acid per minute from 0.187 mM methyl ferulate in 50 mM sodium phosphate buffer, pH 6.5, at 37°C, and at 335 nm, and using a molar extinction coefficient of $14000 \text{ M}^{-1} \text{ cm}^{-1}$.

Primary sequence:

AASLPTMPPSGYDQVRGGIQRGQVVNISYYSTATNGTRPAKVYLP PGYSTSKRYSVLYLLHGIGGSEGDWFADWG
GRASIIADNLI AEGKIKPLIIVT PNTNAAGPGIGDGYENFTKDLINCLIPYIESRYSVYTDREHRAIAGLSMGGG
QSFNIGLTNLDKFAYIGPISSAPNTYPNNR LFPDGGAAARQK LKLLFIACGTNDSLIGFGQRVHEFCVANNINHI
YWLIQGGGH DYNVWKAGLWNFLQLAEQAGLTDYNAP

Literature: -