

Recombinant Enzyme Product Specification Sheet

Cat. No.:	PRO-E0063
LOT:	2009-0063
Activity:	α -Galactosidase
Synonyms:	Melibiose; α -D-galactosidase; α -galactosidase A; α -galactoside galactohydrolase; α -D-galactoside galactohydrolase; alpha-galactosidase; alpha-D-galactosidase; alpha-galactosidase A; alpha-galactoside galactohydrolase; alpha-D-galactoside galactohydrolase
Nomenclature:	CAZy [GH27, glycoside hydrolase family 27 , member of clan GH-D], Gal27A, gal27A, Ccel_1237
Source organism:	<i>Clostridium cellulolyticum</i> H10
Enzyme Commission No.:	3.2.1.22
Activity:	804.1 U/mL
Specific activity:	264.2 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:	6.0
Temperature optimum:	-
[Protein]:	3.04 mg/mL
Sequence length:	577 amino acids (view sequence)
Accession No.:	Q0PRN2 , DQ778333 , ACL75593.1 , ABG76969.1
Molecular weight:	64179.5 Da (theoretical) ~ 65000 Da (observed by SDS-PAGE) - (observed by mass spectrometry)
Biological function:	Hydrolysis of terminal, non-reducing α -D-galactose residues in α -D-galactosides, including galactose oligosaccharides and galactomannans
Potential application(s):	Biomass conversion , carbohydrate research
Comments:	No detectable activity against p NP- β -D-glucopyranoside or p NP- β -D-xylopyranoside. This enzyme exhibits strong inhibition by Tris/HCl buffer

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

Assay: One unit is defined as the amount of enzyme required to release 1 μmol of *p*NP per minute from *p*NP- α -D-galactopyranoside (1 mM) in 50 mM sodium phosphate buffer, pH 6.0, at 40°C, and using an extinction coefficient of 18000 $\text{M}^{-1} \text{cm}^{-1}$

Primary sequence:

WDNGLAKTPPMGWNSWNI FHGDINETKIKQIADTMVSSGMKEAGYVYLNLDNWMANPARDSNGNLRADPTRFPS
GIRALADYVHAKGLKLG IYGCRGTMTCMNI PQSGSKGYEDKDAKTFASWGI DYLYKYNDCNI PNGSDMKTDYQKMQ
TALANCGRP I VFS ICAWGYQSWMPATGNLWRTTGDIADKWDNGNEWFKGI INAIDGNAQYTSSAAPGAWNDPDML
EIGNGGCTTEEYRTQMSMWSMMASPLIAGNDIRTMSQTTKDILLNKEVIAIDQDPAGVQGKRVKSANGLEI WVKP
LGTNGTTKAVALLNRNSATSNI TVNWSDIGVSGSVTVRDLWAKSDKGSFTG SYTASVPSHGTVLIKISTEPPAPV
DATKQIEAESYSNQSGIQ TETCSEGGEDVGFIE NGDYTVYSNVDFGDGVGGFQARVASATSGGNIEIRLDS PAGT
LIGTCPVAGTGDWQTYT DVKCTVSGATGKH DVYLVFKGDSGYLFNLNWF TFTP GSVNTGTLGDLNSDGQVDAIDL
QLLKKYILGLGAIENTKLADLDANGDINAIDFSLKQFLLGIRTSFPGQGA

Literature: 1. [Blouzard et al. \(2007\) J. Bacteriol. 189, 2300-2309](#)