

Product name

ABATACEPT Labelled standard

ORF90261

Catalog number Product description

Labelled Abatacept is a recombinant, stable isotope labelled, human fusion protein consisting of extracellular domain of human cytotoxic T-lymphocyte-associated antigen 4 (CTLA-4) linked to the Fc portion of human IgG1. Labelled Abatacept is designed for use as an internal standard for quantitative analysis of Abatacept in plasma samples.

Abatacept is a selective T cell costimulation modulator indicated for treatment of rheumatoid arthrisis.

Similar products: Labelled Rituximab, Labelled Tocilizumab-like

Fc-fusion protein sequence

MHVAQPAVVLASSRGIASFVCEYASPGKATEVRVTVLRQADSQVTEVCAATYMMGNELTFLDDSICTGTSSGNQVNLTIQGLRAMDTGLYI CKVELMYPPPYYLGIGNGTQIYVIDPEPCPDSDQEPKSSDKTHTSPPSPAPELLGGSSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVKF NWYVDGVEVHNAKTKPREEQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDELTKNQVSLTC LVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK

Product features and protocols

Key features

1	Purity	>90%
		as determined by SDS-PAGE

2	Labelling	Arg- ¹³ C ₆ , ¹⁵ N ₄ Lys- ¹³ C ₆ , ¹⁵ N ₂
Z	Labelling	$Aig^{}C_6,^{}N_4 \mid Lys^{}C_6,^{}N_2$

3	Isotopic	>98%
3	incorporation	as determined by LC-MS/MS
		analysis of digested SIL-protein

Other features

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Expression System	Mammalian cells
Predicted MW	39.4 kDa
Protein content	Quantitation is carried out by UV Absorbance at 280 nm
Formulation	10 mM sodium phosphate pH7.2 50 mM sodium chloride 5% sucrose 0.01% Polysorbate 80

Product preparation

For product preparation we recommend the following steps:

- 1. Thaw in ice
- 2. Vortex gently
- 3. Centrifuge briefly before opening

Product storage

The product is liquid and shipped with dry ice.

Store at -80 °C upon receipt.

Once thawed, the protein can be preserved at 4°C for a few weeks.

Avoid multiple freeze-thaw cycles



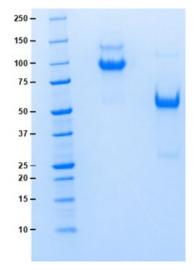
How to use our product



SIL proteins are used as internal standards by addition at the start of a sample preparation workflow. Both Labelled and target proteins will be processed throughout the entire analytical procedure controlling for variability caused by pre-analytical treatments or incomplete enzymatic digestion (1).



Supporting information



References

1. G.Picard, D. Lebert, et al. PSAQ standards for accurate MS-based quantification of proteins: from the concept to biomedical applications, J. Mass Spectrom. 2012, 47, 1353-1363

SDS-PAGE gel analysis of ABATACEPT labelled protein in Reducing/Heated conditions (right lane) and in non-reducing/ non heated conditions (left lane) Stained with Coomassie blue



The product is intended for research use only. Not for diagnostic or therapeutic use.

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