Secukinumab ELISA Kit

Catalogue No.:abx395150

Secukinumab ELISA Kit is a quantitative ELISA kit for detection of Secukinumab.

Secukinumab (Cosentyx) is a fully human IgG1/κ-class monoclonal antibody (mAb) composed of 215 light chain amino acids and 457 heavy chain amino acids. It was discovered and developed by Novartis with the developmental name AIN457, for the treatment of uveitis, rheumatoid arthritis, ankylosing spondylitis, and psoriasis. It is produced in Chinese Hamster Ovary (CHO) cells using recombinant DNA technology.

Target:	Secukinumab
Research Area:	Biosimilars
Reactivity:	Human
Tested Applications:	ELISA
Recommended dilutions:	Optimal dilutions/concentrations should be determined by the end user.
Storage:	Shipped at 4°C. Upon receipt, store the kit according to the storage instruction in the kit's manual.
Validity:	The validity for this kit is 6 months.
Stability:	The stability of the kit is determined by the rate of activity loss. The loss rate is less than 5% within the expiration date under appropriate storage conditions. To minimize performance fluctuations, operation procedures and lab conditions should be strictly controlled. It is also strongly suggested that the whole assay is performed by the same user throughout.
Test Range:	937 ng/ml - 15000 ng/ml
Sensitivity:	< 716 ng/ml
Standard Form:	Lyophilized
Detection Method:	Colorimetric
Assay Type:	Indirect
Assay Data:	Quantitative
Sample Type:	Serum and plasma.
CAS Number:	1229022-83-6

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Note:

THIS PRODUCT IS FOR RESEARCH USE ONLY. NOT FOR USE IN DIAGNOSTIC OR THERAPEUTIC PROCEDURES.

The range and sensitivity is subject to change. Please contact us for the latest product information. For accurate results, sample concentrations must be diluted to mid-range of the kit. If you require a specific range, please contact us in advance or write your request in your order comments.

Please note that our kits are optimised for detection of native samples, rather than recombinant proteins. We are unable to guarantee detection of recombinant proteins, as they may have different sequences or tertiary structures to the native protein.

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