Datasheet

Version: 4.0.0 Revision date: 12 Mar 2025



Ubiquitin Carboxyl Terminal Hydrolase L1 (UCHL1) CLIA Kit

Catalogue No.:abx495338

Ubiquitin Carboxyl Terminal Hydrolase L1 (UCHL1) Chemiluminescent Immunoassay (CLIA) Kit is a Sandwich Chemiluminescent Immunoassay (CLIA) Kit for use with Tissue homogenates, cell lysates, cell culture supernates and other biological fluids.

Target: Ubiquitin Carboxyl Terminal Hydrolase L1 (UCHL1)

Reactivity: Human, Mouse, Rat

Tested Applications: CLIA

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 at least 6 months. Up to 12 months validity can be provided on request.

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: 0.781 ng/ml - 50 ng/ml

Sensitivity: < 0.28 ng/ml

Standard Form: Lyophilized

Detection Method: Chemiluminescent

Assay Type: Sandwich

Assay Data: Quantitative

Sample Type: Tissue homogenates, cell lysates, cell culture supernatants and other biological fluids.

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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.

