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## DATASHEET

## Human Apolipoprotein B mRNA Editing Enzyme Catalytic Subunit 3C (APOBEC3C) ELISA Kit

Catalogue No.:abx385797



Human Apolipoprotein B mRNA Editing Enzyme Catalytic Polypeptide 3C (APOBEC3C) ELISA Kit is an ELISA kit against Apolipoprotein B mRNA Editing Enzyme Catalytic Polypeptide 3C (APOBEC3C).

| Target:               | Apolipoprotein B mRNA Editing Enzyme Catalytic Polypeptide 3C (APOBEC3C)  |
|-----------------------|---|
| Reactivity:           | Human   |
| Tested Applications:  | ELISA   |
| Recommended dilutions | : Optimal dilutions/concentrations should be determined by the end user.  |
| Test Range:           | 0.313 ng/ml - 20 ng/ml  |
| Sensitivity:          | < 0.19 ng/ml  |
| Validity:             | The validity for this kit is 6 months.  |
| Storage:              | Store at 2°C to 8°C upon receipt.   |
| 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. |
| Swiss Prot:           | Q9NRW3  |
| GenelD:               | 27350   |
| Gene Symbol:          | APOBEC3C  |
| OMIM:                 | <u>607750</u>   |

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| HGNC:            | 17353  |
|------------------|--|
| Ensembl:         | ENSG0000244509   |
| Standard Form:   | Lyophilized  |
| ELISA Detection: | Colorimetric   |
| ELISA Type:      | Sandwich   |
| ELISA Data:      | Quantitative   |
| Note:            | This product is for research use only.<br>The range and sensitivity is subject to change. Please contact us for the latest product information.<br>For accurate results, sample concentrations must be diluted to mid-range of the kit. If you require a<br>specific range, please contact us in advance or write your request in your order comments.<br>Please note that our ELISA and CLIA kits are optimised for detection of native samples, rather than<br>recombinant proteins. We are unable to guarantee detection of recombinant proteins, as they may<br>have different sequences or tertiary structures to the native protein. |