

Abbexa Ltd, Innovation Centre, Cambridge Science Park, Cambridge, CB4 0EY, UK
Telephone: +44 (0) 1223 755950 - Fax: +44 (0) 1223 755951 - E-Mail: info@abbexa.com

Human Polymeric Immunoglobulin Receptor/Membrane Secretory Component (PIGR/SC) CLIA Kit

Catalogue No.: abx197522



Human Polymeric Immunoglobulin Receptor/Membrane Secretory Component (PIGR/SC) CLIA Kit is a CLIA kit against Polymeric Immunoglobulin Receptor/Membrane Secretory Component (PIGR/SC).

Target:	Polymeric Immunoglobulin Receptor/Membrane Secretory Component (PIGR/SC)
Reactivity:	Human
Tested Applications:	CLIA
Recommended dilutions:	Optimal dilutions/concentrations should be determined by the end user.
Test Range:	15.6 pg/ml - 1000 pg/ml
Sensitivity:	9.38 pg/ml
Validity:	The validity for this kit is 6 months.
Storage:	Upon receipt, store at 4 °C.
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.
Standard Form:	Lyophilized
ELISA Detection:	Fluorometric
ELISA Data:	Quantitative
Sample Type:	Serum, plasma and other biological fluids.

Abbexa Ltd, Innovation Centre, Cambridge Science Park, Cambridge, CB4 0EY, UK
Telephone: +44 (0) 1223 755950 - Fax: +44 (0) 1223 755951 - E-Mail: info@abbexa.com

Note:

This product is for research use only.

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 ELISA and CLIA 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.