

DESCRIPTION

Species Reactivity	Human/Rat
Specificity	Detects human PIWIL2 in direct ELISAs and human and rat PIWIL2 in Western blots. In direct ELISAs, less than 4% cross-reactivity with recombinant human PIWIL4 is observed.
Source	Polyclonal Sheep IgG
Purification	Antigen Affinity-purified
Immunogen	<i>E. coli</i> -derived recombinant human PIWIL2 Lys216-Asp379 Accession # Q8TC59
Conjugate	Alexa Fluor 350 Excitation Wavelength: 346 nm Emission Wavelength: 442 nm
Formulation	Supplied 0.2mg/ml in 1X PBS with RDF1 and 0.09% Sodium Azide *Contains <0.1% Sodium Azide, which is not hazardous at this concentration according to GHS classifications. Refer to the Safety Data Sheet (SDS) for additional information and handling instructions.

APPLICATIONS

Please Note: Optimal dilutions should be determined by each laboratory for each application. [General Protocols](#) are available in the Technical Information section on our website.

Western Blot Optimal dilution of this antibody should be experimentally determined.

PREPARATION AND STORAGE

Shipping The product is shipped with polar packs. Upon receipt, store it immediately at the temperature recommended below.

Stability & Storage Protect from light. Do not freeze. 12 months from date of receipt, 2 to 8 °C as supplied

BACKGROUND

PIWIL2 (PIWI [P-element-induced wimpy testis]-like 2; also HILI) is a 100-110 kDa cytosolic member of the PIWI subfamily, Argonaute family of proteins. It is expressed in the germ line, particularly in premeiotic and early meiotic germ cells, plus mesenchymal stem cells, and in select cancer stem cells such as CD44⁺CD24⁻ breast cancer cells. PIWIL2 likely silences mobile genetic elements in germ cells, and appears to depress stem cell proliferation by inhibiting the expression of cell cycle genes. Human PIWIL2 is 973 amino acids (aa) in length. It contains one PAZ domain (aa 384-496) and a C-terminal PIWI domain (aa 668-959) that interact with RNA. There are multiple methylation sites on arginines. Two potential splice forms are reported, one that shows a deletion of aa 887-922, and a second that contains a 12 aa substitution for aa 486-801. Over aa 216-379, human PIWIL2 shares 96% aa identity with rat PIWIL2.

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