

SL3202R

Product Name:	phospho-IRS1 (Ser789)
Chinese Name:	磷酸化胰岛素受体底物-1抗体
Alias:	p-IRS1; IRS1(phospho-Ser789); IRS1 (phospho S789); p-IRS1 (phospho S789); HIRS 1; HIRS1; Insulin Receptor Substrate 1; IRS-1; IRS 1; IRS1; OTTHUMP00000164234; IRS1_HUMAN.
Organism Species:	Rabbit
Clonality:	Polyclonal
React Species:	Human,Mouse,Rat,
Applications:	ELISA=1:500-1000IHC-P=1:400-800IHC-F=1:400-800Flow-Cyt=1µg /testIF=1:100- 500 (Paraffin sections need antigen repair) not yet tested in other applications. optimal dilutions/concentrations should be determined by the end user.
Molecular weight:	132kDa
Cellular localization:	The nucleuscytoplasmicThe cell membrane
Form:	Lyophilized or Liquid
<b>Concentration:</b>	1mg/ml
immunogen:	KLH conjugated synthesised phosphopeptide derived from human IRS1 around the phosphorylation site of Ser789:SS(p-S)SG
Lsotype:	IgG
Purification:	affinity purified by Protein A
Storage Buffer:	0.01M TBS(pH7.4) with 1% BSA, 0.03% Proclin300 and 50% Glycerol.
Storage:	Store at -20 °C for one year. Avoid repeated freeze/thaw cycles. The lyophilized antibody is stable at room temperature for at least one month and for greater than a year when kept at -20°C. When reconstituted in sterile pH 7.4 0.01M PBS or diluent of antibody the antibody is stable for at least two weeks at 2-4 °C.
PubMed:	PubMed
Product Detail:	Insulin receptor substrates (IRS) are responsible for several insulin related activities, such as glucose homeostasis, cell growth, cell transformation, apoptosis and insulin signal transduction. Serine/threonine phosphorylation of IRS1 has been demonstrated to



be a negative regulator of insulin signaling and is responsible for its degradation, although IRS1 degradation pathways are not well understood. IRS1 has also been shown to be constitutively activated in cancers such as breast cancer, Wilm's tumors, and adrenal cortical carcinomas, thus making IRS1 phosphorylation and subsequent degradation an attractive therapeutic target. To date there have been four subtypes identified: IRS1, 2, 3 and 4, with IRS1 being widely expressed.

### Function:

May mediate the control of various cellular processes by insulin. When phosphorylated by the insulin receptor binds specifically to various cellular proteins containing SH2 domains such as phosphatidylinositol 3-kinase p85 subunit or GRB2. Activates phosphatidylinositol 3-kinase when bound to the regulatory p85 subunit.

## Subunit:

Interacts with UBTF and PIK3CA. Interacts (via phosphorylated YXXM motifs) with PIK3R1. Interacts with ROCK1 and FER. Interacts (via PH domain) with PHIP. Interacts with GRB2. Interacts with SOCS7. Interacts (via IRS-type PTB domain) with IGF1R and INSR (via the tyrosine-phosphorylated NPXY motif). Interacts with ALK.

## Subcellular Location:

Membrane; Single-pass type I membrane protein.

# **Tissue Specificity:**

Isoform Long and isoform Short are predominantly expressed in tissue targets of insulin metabolic effects: liver, adipose tissue and skeletal muscle but are also expressed in the peripheral nerve, kidney, pulmonary alveoli, pancreatic acini, placenta vascular endothelium, fibroblasts, monocytes, granulocytes, erythrocytes and skin. Isoform Short is preferentially expressed in fetal cells such as fetal fibroblasts, muscle, liver and kidney. Found as a hybrid receptor with IGF1R in muscle, heart, kidney, adipose tissue, skeletal muscle, hepatoma, fibroblasts, spleen and placenta (at protein level). Overexpressed in several tumors, including breast, colon, lung, ovary, and thyroid carcinomas.

# **Post-translational modifications:**

Serine phosphorylation of IRS1 is a mechanism for insulin resistance. Ser-312 phosphorylation inhibits insulin action through disruption of IRS1 interaction with the insulin receptor.

Phosphorylation of Tyr-896 is required for GRB2-binding. Phosphorylated by ALK. Phosphorylated at Ser-270, Ser-307, Ser-636 and Ser-1101 by RPS6KB1; phosphorylation induces accelerated degradation of IRS1.

### **DISEASE:**

Polymorphisms in IRS1 may be involved in the etiology of non-insulin-dependent diabetes mellitus (NIDDM) [MIM:125853].

# Similarity:

	Contains 1 IRS-type PTB domain.
	Contains 1 PH domain.
	SWISS: P35568
	<b>Gene ID:</b> 3667
	Database links:
	Entrez Gene: 3667Human
	Entrez Gene: 16367 Mouse
	Entrez Gene: 25467Rat
	Omim: 147545Human
	SwissProt: P35568Human
	Entrez Gene: 16367Mouse Entrez Gene: 25467Rat Omim: 147545Human SwissProt: P35568Human SwissProt: P35569Mouse SwissProt: P35570Rat Unigene: 471508Human Unigene: 4952Mouse
	SwissProt: P35570Rat
	Unigene: 471508Human
	Unigene: 4952Mouse
	Unigene: 10476Rat
	Important Note:
	This product as supplied is intended for research use only, not for use in human, therapeutic or diagnostic applications.
4	胰岛素受体底物-1是细胞中分子量为165-
	185kDa的磷蛋白,胰岛素受体的内源性底物。经胰岛素刺激后其酪氨酸残基被磷酸
	化。



