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SYEP Rabbit pAb

CatalogNo: YN2956

Key Features

Host Species Rabbit 	Reactivity Human,Mouse 	Applications WB,ELISA
MW • 166kD (Observed)	Isotype • IgG	

Recommended Dilution Ratios

WB 1:500-2000 ELISA 1:5000-20000

Storage

Storage*	-15°C to -25°C/1 year(Do not lower than -25°C)
Formulation	Liquid in PBS containing 50% glycerol, 0.5% BSA and 0.02% sodium azide.

Basic Information

Clonality Polyclonal

Immunogen Information

ImmunogenSynthesized peptide derived from part region of human protein

Specificity SYEP Polyclonal Antibody detects endogenous levels of protein.

Target Information

Gene name EPRS GLNS PARS QARS QPRS PIG32

Protein Name

Bifunctional glutamate/proline--tRNA ligase (Bifunctional aminoacyl-tRNA synthetase) (Cell proliferation-inducing gene 32 protein) [Includes: Glutamate--tRNA ligase (Glutamyl-tRNA synthetase) (GluRS); Proline--tRNA ligase (Prolyl-tRNA synthetase)]

	Organism	Gene ID	UniProt ID
	Human	<u>2058;</u>	<u>P07814;</u>
	Mouse		<u>Q8CGC7;</u>
Cellular Localization	Cytoplasm, cytosol . Membrane ; Peripheral membrane protein . Translocates from cytosol to membranes upon phosphorylation at Ser-999		
Tissue specificity	Bone marrow,Brain,Cervix carcinoma,Duodenum,Epithelium,Eye,Lung,Placenta,		
Function	Catalytic activity:ATP + L-glutamate + tRNA(Glu) = AMP + diphosphate + L-glutamyl- tRNA(Glu).,Catalytic activity:ATP + L-proline + tRNA(Pro) = AMP + diphosphate + L-prolyl- tRNA(Pro).,Caution:Was originally thought to be a glutaminyl-tRNA synthetase.,Domain:The WHEP-TRS domain is involved in RNA binding.,Function:Catalyzes the attachment of the cognate amino acid to the corresponding tRNA in a two-step reaction: the amino acid is first activated by ATP to form a covalent intermediate with AMP and is then transferred to the		

acceptor end of the cognate tRNA., sequence Caution: Contaminating sequence. Potential poly-A sequence., sequence Caution: Sequencing errors., similarity: Belongs to the class-I aminoacyl-tRNA synthetase family., similarity: Contains 3 WHEP-TRS domains., similarity: In the C-terminal section; belongs to the class-II aminoacyl-tRNA synthetase family., similarity: In the N-terminal section; belongs to the class-I aminoacyI-tRNA synthetase family., subunit: Component of the multisynthetase complex which is comprised of a bifunctional glutamyl-prolyl-tRNA synthetase, the monospecific isoleucyl, leucyl, glutaminyl, methionyl, lysyl, arginyl, and aspartyl-tRNA synthetases as well as three auxiliary proteins, p18, p48 and p43. Interacts with DUS2L.,

Validation Data

Contact information

order.cn@immunoway.com
support.cn@immunoway.com
400-8787-807(China)
http://www.immunoway.com.cn
2200 Ringwood Ave San Jose, CA 95131 USA



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