Applications

WB,IF,ELISA



Osteocalcin Rabbit pAb

CatalogNo: YN2113

Key Features

Host Species Reactivity

Rabbit
 Human, Mouse, Rat

MW Isotype
• 11kD (Observed) • IgG

Recommended Dilution Ratios

WB 1:500-2000 IF 1:100-300

ELISA 1:5000-20000

Not yet tested in other applications.

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

Immunogen Synthesized peptide derived from part region of human protein AA range: 31-80

Specificity OSTCN Polyclonal Antibody detects endogenous levels of protein.

| Target Information

Gene name BGLAP

Protein Name Osteocalcin (Bone Gla protein) (BGP) (Gamma-carboxyglutamic acid-containing protein)

Organism	Gene ID	UniProt ID	
Human	<u>632</u> ;	<u>P02818;</u>	
Mouse		<u>P86546;</u>	
Rat		<u>P04640;</u>	

Cellular Localization Secreted.

Tissue specificity Brain, Cerebellum, Colon, Lung carcinoma, Skin, Testis, Uterus,

Function Function:Constitutes 1-2% of the total bone protein. It binds strongly to apatite and

calcium.,Function:Part of the MIS12 complex which is required for normal chromosome alignment and segregation and kinetochore formation during mitosis. May act as a

cotranscription partner of NFE2L2 involved in regulation of polyamine-induced transcription of SSAT.,induction:By polyamine analogs in analog-sensitive H157 cells.,online information:Osteocalcin entry,PTM:Gamma-carboxyglutamate residues are formed by

vitamin K dependent carboxylation. These residues are essential for the binding of calcium.,similarity:Belongs to the osteocalcin/matrix Gla protein family.,similarity:Contains 1 Gla (gamma-carboxy-glutamate) domain.,subcellular location:Associated with the kinetochore.,subunit:Component of the MIS12 complex composed of MIS12, DSN1, NSL1 and PMF1. Interacts with COPS7A. Interacts via its coiled-coil domain with the leucine-zipper domain of NFE2L2. The interaction with NFE2L2 is required for the transcriptional regulation

of SSAT., tissue specificity: Highest levels of expression in heart and skeletal muscle, with significant levels expressed in kidney and liver..

I Validation Data

Contact information

Orders: order@immunoway.com
Support: tech@immunoway.com

Telephone: 877-594-3616 (Toll Free), 408-747-0185

Website: http://www.immunoway.com

Address: 2200 Ringwood Ave San Jose, CA 95131 USA



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