



Polyclonal Anti-Connexin 26 (Sepharose Bead Conjugate)

Catalogue No. PA1025-S	Immunogen
	A peptide mapping at the middle region of rat Connexin 26,
Lot No. 03A01	different from the relative sequence of human by three amino
	acids.
Ig type: rabbit	
	Purification
IgG Size: 100μg/vial	Immunogen affinity purified.
Specificity	Formulation
Human, mouse, rat. No cross reactivity with other	50% slurry in PBS pH 7.2 with 0.01mg NaN $_3a_3$ preservative.
proteins.	
	Storage
Recommended application	Store at 4°C for frequent use.
Immunoprecipitation(IP)	
	Description:
	This Antagene antibody is immobilized via covalent binding of
	primary amino groups to N-hydroxysuccinimide
	(NHS)-activated sepharose beads. It is useful for

immunoprecipitation assays

BACKGROUND

Connexin26(CX26), also known as GAP junction protein, beta2, GJB2. Gap junctions were first characterized by electron microscopy as regionally specialized structures on plasma membranes of contacting adherent cells. These structures were shown to consist of cell-to-cell channels. Proteins, called connexins, purified from fractions of enriched gap junctions from different tissues differ. The 3-prime untranslated region of the CX26 transcript contains a putative mRNA instability sequence. The deduced 226-amino acid protein has a calculated molecular mass of about 26 kD. CX26 shares 92.5% identity with rat Cx26. connexin 26 (GJB2) is assigned to human chromosome 13q11-q12. Connexin 26 regulates epidermal barrier and wound remodeling and promotes psoriasiform response. Connexin 26 gene (GJB2) mutation modulates the severity of hearing loss associated with the 1555A-G mitochondrial mutation.

REFERENCE

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2. Djalilian, A. R.; McGaughey, D.; Patel, S.; Seo, E. Y.; Yang, C.; Cheng, J.; Tomic, M.; Sinha, S.; Ishida-Yamamoto, A.; Segre, J. A. : Connexin 26 regulates epidermal barrier and wound remodeling and promotes psoriasiform response. J. Clin. Invest. 116: 1243-1253, 2006. 3. Abe, S.; Kelley, P. M.; Kimberling, W. J.; Usami, S. : Connexin 26 gene (GJB2) mutation modulates the severity of hearing loss associated with the 1555A-G mitochondrial mutation. Am. J. Med. Genet. 103: 334-338, 2001.