## RPPB4313

## Product Information Protein Information

## Product SKU:

RPPB4313

## Host:

Escherichia Coli.

## Protein description:

Recombinant Protein-L produced in E.Coli is a single non-glycosylated polypeptide chain fused with a $6 \times$ His tag at N -terminus and a Cys on C-terminus. Protein-L is comprised of 5 IgG -binding regions of protein L ( $\mathrm{B} 1-\mathrm{B} 2-\mathrm{B} 3-\mathrm{B} 4-\mathrm{B} 5$ ) containing 373 amino acids in total and having a molecular mass of 41.6 kDa , however, it migrates with an apparent molecular mass of 46 kDa on SDS-PAGE. Cell wall binding region, cell membrane binding region and albumin binding region have been eliminated from the recombinant Protein-L to guarantee the maximum specific $\operatorname{lgG}$ binding.

## Appearance:

Sterile Filtered White lyophilized (freeze-dried) powder.

## Formulation:

Protein-L was lyophilized without any additives.

## Purity:

Greater than $96.0 \%$ as determined by SDS-PAGE.

## Solubility:

It is recommended to reconstitute the lyophilized Protein-L in sterile $18 \mathrm{M}-\mathrm{cm} \mathrm{H} 2 \mathrm{O}$ not less than $0.1 \mathrm{mg} / \mathrm{ml}$, which can then be further diluted to other aqueous solutions.

## Stability:

Lyophilized Protein-L although stable at room temperature for 3 weeks, should be stored desiccated below $-18^{\circ} \mathrm{C}$. Upon reconstitution Protein-L should be stored at $4^{\circ} \mathrm{C}$ between 2-7 days and for future use below $-18^{\circ} \mathrm{C}$. For long term storage it is recommended to add a carrier protein ( $0.1 \% \mathrm{HSA}$ or BSA). Please prevent freeze-thaw cycles.

## Amino Acid Sequence:

MHHHHHHKEE TPETPETDSE EEVTIKANLI FANGSTQTAE FKGTFEKATS EAYAYADTLK KDNGEYTVDV ADKGYTLNIK FAGKEKTPEE PKEEVTIKAN LIYADGKTQT AEFKGTFEEA TAEAYRYADA LKKDNGEYTV DVADKGYTLN IKFAGKEKTP EEPKEEVTIK ANLIYADGKT QTAEFKGTFE EATAEAYRYA DLLAKENGKY TVDVADKGYT LNIKFAGKEK TPEEPKEEVT IKANLIYADG KTQTAEFKGT FAEATAEAYR YADLLAKENG KYTADLEDGG YTINIRFAGK KVDEKPEEKE QVTIKENIYF EDGTVQTATF KGTFAEATAE AYRYADLLSK EHGKYTADLE DGGYTINIRF AGC.

