

**bs-5010R**

**[ Primary Antibody ]**

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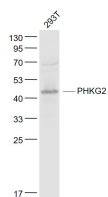
400-901-9800

## PHKG2 Rabbit pAb

### — DATASHEET —

<b>Host:</b> Rabbit	<b>Isotype:</b> IgG	<b>Applications:</b> WB (1:500-2000)
<b>Clonality:</b> Polyclonal		
<b>GeneID:</b> 5261		
<b>Target:</b> PHKG2		
<b>Immunogen:</b> KLH conjugated synthetic peptide derived from human PHKG2: 81-180/406.		
<b>Purification:</b> affinity purified by Protein A		
<b>Concentration:</b> 1mg/ml		
<b>Storage:</b> 0.01M TBS (pH7.4) with 1% BSA, 0.02% Proclin300 and 50% Glycerol. Shipped at 4°C. Store at -20°C for one year. Avoid repeated freeze/thaw cycles.		<b>Reactivity:</b> Human (predicted: Mouse, Rat, Rabbit, Pig, Cow, Dog, Horse)
<b>Background:</b> Phosphorylase kinase is a polymer of 16 subunits, four each of alpha, beta, gamma and delta. The alpha subunit includes the skeletal muscle and hepatic isoforms, encoded by two different genes. The beta subunit is the same in both the muscle and hepatic isoforms, and encoded by one gene. The gamma subunit also includes the skeletal muscle and hepatic isoforms, and the hepatic isoform is encoded by this gene. The delta subunit is a calmodulin and can be encoded by three different genes. The gamma subunits contain the active site of the enzyme, whereas the alpha and beta subunits have regulatory functions controlled by phosphorylation. The delta subunit mediates the dependence of the enzyme on calcium concentration. Mutations in this gene cause glycogen storage disease type 9C, also known as autosomal liver glycogenosis. Alternatively spliced transcript variants encoding different isoforms have been identified in this gene.		<b>Predicted MW.:</b> 46 kDa
		<b>Subcellular Location:</b> Cytoplasm

### — VALIDATION IMAGES —



Sample: 293T(Human) Cell Lysate at 30 ug  
Primary: Anti- PHKG2 (bs-5010R) at 1/1000  
dilution Secondary: IRDye800CW Goat Anti-  
Rabbit IgG at 1/20000 dilution Predicted band  
size: 46 kD Observed band size: 44 kD

### — SELECTED CITATIONS —

- **[IF=2.2]** Zhou Hua. et al. A ferroptosis-related signature predicts the clinical diagnosis and prognosis, and associates with the immune microenvironment of lung cancer. Discover Oncology. 2024 Dec;15(1):1-19 IHC ;Human. 38743344