

**AKT1 antibody - N-terminal region**  
**Rabbit Polyclonal Antibody**  
**Catalog # AI16186****Specification**

---

**AKT1 antibody - N-terminal region - Product Information**

Application	WB
Primary Accession	<a href="#">P31749</a>
Other Accession	<a href="#">NM_005163</a> , <a href="#">NP_005154</a>
Reactivity	Human, Mouse, Rat, Pig, Sheep, Bovine, Neisseria Gonorrhoeae, Dog
Predicted	Human, Mouse, Rat, Pig, Chicken, Sheep, Bovine, Neisseria Gonorrhoeae, Dog
Host	Rabbit
Clonality	Polyclonal
Calculated MW	56kDa KDa

**AKT1 antibody - N-terminal region - Additional Information****Gene ID 207**

Alias Symbol **AKT, PKB, RAC, PRKBA, PKB-ALPHA, RAC-ALPHA**

**Other Names**

RAC-alpha serine/threonine-protein kinase, 2.7.11.1, Protein kinase B, PKB, Protein kinase B alpha, PKB alpha, Proto-oncogene c-Akt, RAC-PK-alpha, AKT1, PKB, RAC

**Format**

Liquid. Purified antibody supplied in 1x PBS buffer with 0.09% (w/v) sodium azide and 2% sucrose.

**Reconstitution & Storage**

Add 50 ul of distilled water. Final anti-AKT1 antibody concentration is 1 mg/ml in PBS buffer with 2% sucrose. For longer periods of storage, store at 20°C. Avoid repeat freeze-thaw cycles.

**Precautions**

AKT1 antibody - N-terminal region is for research use only and not for use in diagnostic or therapeutic procedures.

**AKT1 antibody - N-terminal region - Protein Information**

**Name** AKT1

**Synonyms** PKB, RAC

**Function**

AKT1 is one of 3 closely related serine/threonine-protein kinases (AKT1, AKT2 and AKT3) called the AKT kinase, and which regulate many processes including metabolism, proliferation, cell survival, growth and angiogenesis (PubMed:<a href="http://www.uniprot.org/citations/11882383">

target="\_blank">11882383</a>, PubMed:<a href="http://www.uniprot.org/citations/15526160" target="\_blank">15526160</a>, PubMed:<a href="http://www.uniprot.org/citations/15861136" target="\_blank">15861136</a>, PubMed:<a href="http://www.uniprot.org/citations/21432781" target="\_blank">21432781</a>, PubMed:<a href="http://www.uniprot.org/citations/21620960" target="\_blank">21620960</a>, PubMed:<a href="http://www.uniprot.org/citations/31204173" target="\_blank">31204173</a>). This is mediated through serine and/or threonine phosphorylation of a range of downstream substrates (PubMed:<a href="http://www.uniprot.org/citations/11882383" target="\_blank">11882383</a>, PubMed:<a href="http://www.uniprot.org/citations/15526160" target="\_blank">15526160</a>, PubMed:<a href="http://www.uniprot.org/citations/21432781" target="\_blank">21432781</a>, PubMed:<a href="http://www.uniprot.org/citations/21620960" target="\_blank">21620960</a>, PubMed:<a href="http://www.uniprot.org/citations/31204173" target="\_blank">31204173</a>). Over 100 substrate candidates have been reported so far, but for most of them, no isoform specificity has been reported (PubMed:<a href="http://www.uniprot.org/citations/11882383" target="\_blank">11882383</a>, PubMed:<a href="http://www.uniprot.org/citations/15526160" target="\_blank">15526160</a>, PubMed:<a href="http://www.uniprot.org/citations/21432781" target="\_blank">21432781</a>, PubMed:<a href="http://www.uniprot.org/citations/21620960" target="\_blank">21620960</a>). AKT is responsible of the regulation of glucose uptake by mediating insulin-induced translocation of the SLC2A4/GLUT4 glucose transporter to the cell surface (By similarity). Phosphorylation of PTPN1 at 'Ser-50' negatively modulates its phosphatase activity preventing dephosphorylation of the insulin receptor and the attenuation of insulin signaling (By similarity). Phosphorylation of TBC1D4 triggers the binding of this effector to inhibitory 14-3-3 proteins, which is required for insulin-stimulated glucose transport (PubMed:<a href="http://www.uniprot.org/citations/11994271" target="\_blank">11994271</a>). AKT regulates also the storage of glucose in the form of glycogen by phosphorylating GSK3A at 'Ser-21' and GSK3B at 'Ser-9', resulting in inhibition of its kinase activity (By similarity). Phosphorylation of GSK3 isoforms by AKT is also thought to be one mechanism by which cell proliferation is driven (By similarity). AKT regulates also cell survival via the phosphorylation of MAP3K5 (apoptosis signal-related kinase) (PubMed:<a href="http://www.uniprot.org/citations/11154276" target="\_blank">11154276</a>). Phosphorylation of 'Ser-83' decreases MAP3K5 kinase activity stimulated by oxidative stress and thereby prevents apoptosis (PubMed:<a href="http://www.uniprot.org/citations/11154276" target="\_blank">11154276</a>). AKT mediates insulin-stimulated protein synthesis by phosphorylating TSC2 at 'Ser-939' and 'Thr-1462', thereby activating the mTORC1 signaling pathway, and leading to both phosphorylation of 4E-BP1 and in activation of RPS6KB1 (PubMed:<a href="http://www.uniprot.org/citations/12150915" target="\_blank">12150915</a>, PubMed:<a href="http://www.uniprot.org/citations/12172553" target="\_blank">12172553</a>). Also regulates the mTORC1 signaling pathway by catalyzing phosphorylation of CASTOR1 and DEPDC5 (PubMed:<a href="http://www.uniprot.org/citations/31548394" target="\_blank">31548394</a>, PubMed:<a href="http://www.uniprot.org/citations/33594058" target="\_blank">33594058</a>). AKT is involved in the phosphorylation of members of the FOXO factors (Forkhead family of transcription factors), leading to binding of 14-3-3 proteins and cytoplasmic localization (PubMed:<a href="http://www.uniprot.org/citations/10358075" target="\_blank">10358075</a>). In particular, FOXO1 is phosphorylated at 'Thr-24', 'Ser-256' and 'Ser-319' (PubMed:<a href="http://www.uniprot.org/citations/10358075" target="\_blank">10358075</a>). FOXO3 and FOXO4 are phosphorylated on equivalent sites (PubMed:<a href="http://www.uniprot.org/citations/10358075" target="\_blank">10358075</a>). AKT has an important role in the regulation of NF- kappa-B-dependent gene transcription and positively regulates the activity of CREB1 (cyclic AMP (cAMP)-response element binding protein) (PubMed:<a href="http://www.uniprot.org/citations/9829964" target="\_blank">9829964</a>). The phosphorylation of CREB1 induces the binding of accessory proteins that are necessary for the transcription of pro- survival genes such as BCL2 and MCL1 (PubMed:<a href="http://www.uniprot.org/citations/9829964" target="\_blank">9829964</a>). AKT phosphorylates 'Ser-454' on ATP citrate lyase (ACLY), thereby potentially regulating ACLY activity and fatty acid synthesis (By similarity). Activates the 3B isoform of cyclic nucleotide phosphodiesterase (PDE3B) via phosphorylation of 'Ser-273', resulting in reduced cyclic AMP levels and inhibition of lipolysis (By similarity). Phosphorylates PIKFYVE on 'Ser-318', which results in

increased PI(3)P-5 activity (By similarity). The Rho GTPase-activating protein DLC1 is another substrate and its phosphorylation is implicated in the regulation cell proliferation and cell growth (By similarity). AKT plays a role as key modulator of the AKT-mTOR signaling pathway controlling the tempo of the process of newborn neurons integration during adult neurogenesis, including correct neuron positioning, dendritic development and synapse formation (By similarity). Signals downstream of phosphatidylinositol 3-kinase (PI(3)K) to mediate the effects of various growth factors such as platelet-derived growth factor (PDGF), epidermal growth factor (EGF), insulin and insulin-like growth factor 1 (IGF1) (PubMed:<a href="http://www.uniprot.org/citations/12176338" target="\_blank">12176338</a>, PubMed:<a href="http://www.uniprot.org/citations/12964941" target="\_blank">12964941</a>). AKT mediates the antiapoptotic effects of IGF1 (By similarity). Essential for the SPATA13-mediated regulation of cell migration and adhesion assembly and disassembly (PubMed:<a href="http://www.uniprot.org/citations/19934221" target="\_blank">19934221</a>). May be involved in the regulation of the placental development (By similarity). Phosphorylates STK4/MST1 at 'Thr-120' and 'Thr-387' leading to inhibition of its: kinase activity, nuclear translocation, autophosphorylation and ability to phosphorylate FOXO3 (PubMed:<a href="http://www.uniprot.org/citations/17726016" target="\_blank">17726016</a>). Phosphorylates STK3/MST2 at 'Thr-117' and 'Thr-384' leading to inhibition of its: cleavage, kinase activity, autophosphorylation at Thr-180, binding to RASSF1 and nuclear translocation (PubMed:<a href="http://www.uniprot.org/citations/20086174" target="\_blank">20086174</a>). Phosphorylates SRPK2 and enhances its kinase activity towards SRSF2 and ACIN1 and promotes its nuclear translocation (PubMed:<a href="http://www.uniprot.org/citations/19592491" target="\_blank">19592491</a>). Phosphorylates RAF1 at 'Ser-259' and negatively regulates its activity (PubMed:<a href="http://www.uniprot.org/citations/10576742" target="\_blank">10576742</a>). Phosphorylation of BAD stimulates its pro-apoptotic activity (PubMed:<a href="http://www.uniprot.org/citations/10926925" target="\_blank">10926925</a>). Phosphorylates KAT6A at 'Thr-369' and this phosphorylation inhibits the interaction of KAT6A with PML and negatively regulates its acetylation activity towards p53/TP53 (PubMed:<a href="http://www.uniprot.org/citations/23431171" target="\_blank">23431171</a>). Phosphorylates palladin (PALLD), modulating cytoskeletal organization and cell motility (PubMed:<a href="http://www.uniprot.org/citations/20471940" target="\_blank">20471940</a>). Phosphorylates prohibitin (PHB), playing an important role in cell metabolism and proliferation (PubMed:<a href="http://www.uniprot.org/citations/18507042" target="\_blank">18507042</a>). Phosphorylates CDKN1A, for which phosphorylation at 'Thr-145' induces its release from CDK2 and cytoplasmic relocation (PubMed:<a href="http://www.uniprot.org/citations/16982699" target="\_blank">16982699</a>). These recent findings indicate that the AKT1 isoform has a more specific role in cell motility and proliferation (PubMed:<a href="http://www.uniprot.org/citations/16139227" target="\_blank">16139227</a>). Phosphorylates CLK2 thereby controlling cell survival to ionizing radiation (PubMed:<a href="http://www.uniprot.org/citations/20682768" target="\_blank">20682768</a>). Phosphorylates PCK1 at 'Ser-90', reducing the binding affinity of PCK1 to oxaloacetate and changing PCK1 into an atypical protein kinase activity using GTP as donor (PubMed:<a href="http://www.uniprot.org/citations/32322062" target="\_blank">32322062</a>). Also acts as an activator of TMEM175 potassium channel activity in response to growth factors: forms the lysoK(GF) complex together with TMEM175 and acts by promoting TMEM175 channel activation, independently of its protein kinase activity (PubMed:<a href="http://www.uniprot.org/citations/32228865" target="\_blank">32228865</a>). Acts as a regulator of mitochondrial calcium uptake by mediating phosphorylation of MICU1 in the mitochondrial intermembrane space, impairing MICU1 maturation (PubMed:<a href="http://www.uniprot.org/citations/30504268" target="\_blank">30504268</a>). Acts as an inhibitor of tRNA methylation by mediating phosphorylation of the N-terminus of METTL1, thereby inhibiting METTL1 methyltransferase activity (PubMed:<a href="http://www.uniprot.org/citations/15861136" target="\_blank">15861136</a>). In response to LPAR1 receptor pathway activation, phosphorylates Rabin8/RAB3IP which alters its activity and phosphorylates WDR44 which induces WDR44 binding to Rab11, thereby switching Rab11 vesicular function from preciliary trafficking to endocytic recycling (PubMed:<a href="http://www.uniprot.org/citations/31204173" target="\_blank">31204173</a>).

### Cellular Location

Cytoplasm {ECO:0000250|UniProtKB:P31750}. Nucleus. Cell membrane. Mitochondrion intermembrane space {ECO:0000250|UniProtKB:P31750}. Note=Nucleus after activation by integrin-linked protein kinase 1 (ILK1). Nuclear translocation is enhanced by interaction with TCL1A. Phosphorylation on Tyr-176 by TNK2 results in its localization to the cell membrane where it is targeted for further phosphorylations on Thr-308 and Ser-473 leading to its activation and the activated form translocates to the nucleus Colocalizes with WDFY2 in intracellular vesicles (PubMed:16792529) Also localizes to mitochondrial intermembrane space in response to rapamycin treatment (By similarity). {ECO:0000250|UniProtKB:P31750, ECO:0000269|PubMed:16792529}

### Tissue Location

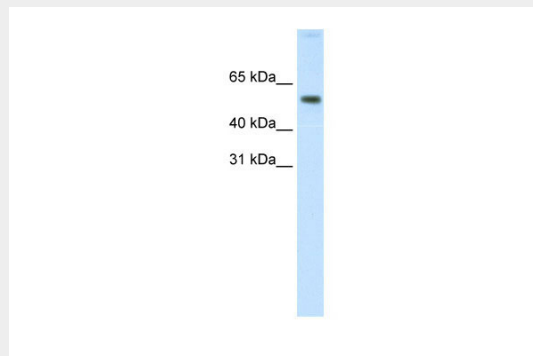
Expressed in prostate cancer and levels increase from the normal to the malignant state (at protein level). Expressed in all human cell types so far analyzed. The Tyr-176 phosphorylated form shows a significant increase in expression in breast cancers during the progressive stages i.e. normal to hyperplasia (ADH), ductal carcinoma in situ (DCIS), invasive ductal carcinoma (IDC) and lymph node metastatic (LNMM) stages.

### AKT1 antibody - N-terminal region - Protocols

Provided below are standard protocols that you may find useful for product applications.

- [Western Blot](#)
- [Blocking Peptides](#)
- [Dot Blot](#)
- [Immunohistochemistry](#)
- [Immunofluorescence](#)
- [Immunoprecipitation](#)
- [Flow Cytometry](#)
- [Cell Culture](#)

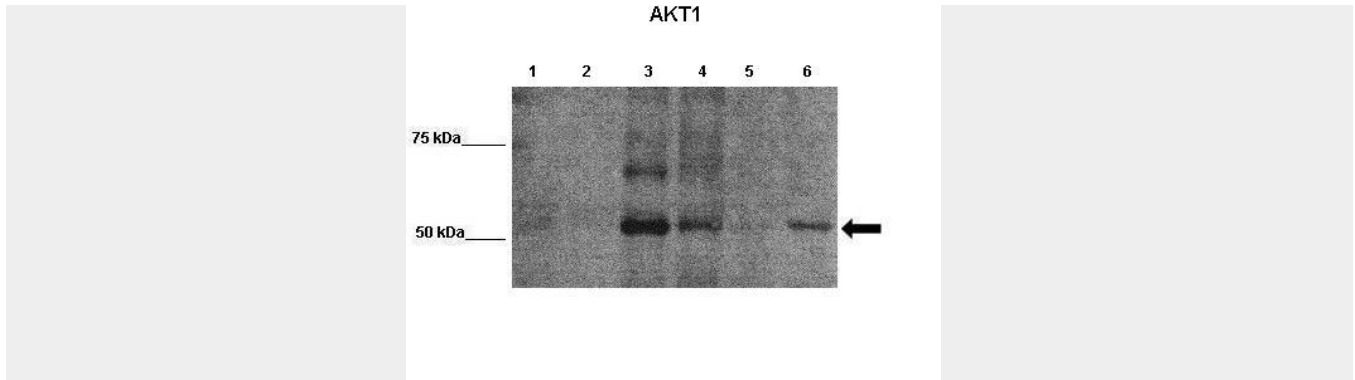
### AKT1 antibody - N-terminal region - Images



WB Suggested Anti-AKT1 Antibody Titration: 0.2-1 µg/ml

ELISA Titer: 1:312500

Positive Control: Human Small Intestine



#### WB Suggested Anti-AKT1 Antibody

Positive Control: Lane 1: 50ug preterm baboon muscle homogenate Lane 2: 50ug preterm baboon muscle homogenate Lane 3: 50ug term baboon muscle homogenate Lane 4: 50ug term baboon muscle homogenate Lane 5: 50ug adult baboon muscle homogenate Lane 6: 50ug adult baboon muscle homogenate

Primary Antibody Dilution : 1:1666

Secondary Antibody : Anti-rabbit-HRP

Secondary Antibody Dilution : 1:1200

Submitted by: Cynthia Blanco, University of Texas Health Science Center

#### AKT1 antibody - N-terminal region - Background

AKT1 is one of 3 closely related serine/threonine- protein kinases (AKT1, AKT2 and AKT3) called the AKT kinase, and which regulate many processes including metabolism, proliferation, cell survival, growth and angiogenesis. This is mediated through serine and/or threonine phosphorylation of a range of downstream substrates. Over 100 substrate candidates have been reported so far, but for most of them, no isoform specificity has been reported. AKT is responsible of the regulation of glucose uptake by mediating insulin-induced translocation of the SLC2A4/GLUT4 glucose transporter to the cell surface. Phosphorylation of PTPN1 at 'Ser-50' negatively modulates its phosphatase activity preventing dephosphorylation of the insulin receptor and the attenuation of insulin signaling. Phosphorylation of TBC1D4 triggers the binding of this effector to inhibitory 14-3-3 proteins, which is required for insulin-stimulated glucose transport. AKT regulates also the storage of glucose in the form of glycogen by phosphorylating GSK3A at 'Ser-21' and GSK3B at 'Ser-9', resulting in inhibition of its kinase activity. Phosphorylation of GSK3 isoforms by AKT is also thought to be one mechanism by which cell proliferation is driven. AKT regulates also cell survival via the phosphorylation of MAP3K5 (apoptosis signal-related kinase). Phosphorylation of 'Ser-83' decreases MAP3K5 kinase activity stimulated by oxidative stress and thereby prevents apoptosis. AKT mediates insulin-stimulated protein synthesis by phosphorylating TSC2 at 'Ser-939' and 'Thr-1462', thereby activating mTORC1 signaling and leading to both phosphorylation of 4E-BP1 and in activation of RPS6KB1. AKT is involved in the phosphorylation of members of the FOXO factors (Forkhead family of transcription factors), leading to binding of 14-3-3 proteins and cytoplasmic localization. In particular, FOXO1 is phosphorylated at 'Thr-24', 'Ser-256' and 'Ser-319'. FOXO3 and FOXO4 are phosphorylated on equivalent sites. AKT has an important role in the regulation of NF-kappa-B-dependent gene transcription and positively regulates the activity of CREB1 (cyclic AMP (cAMP)- response element binding protein). The phosphorylation of CREB1 induces the binding of accessory proteins that are necessary for the transcription of pro-survival genes such as BCL2 and MCL1. AKT phosphorylates 'Ser-454' on ATP citrate lyase (ACLY), thereby potentially regulating ACLY activity and fatty acid synthesis. Activates the 3B isoform of cyclic nucleotide phosphodiesterase (PDE3B) via phosphorylation of 'Ser-273', resulting in reduced cyclic AMP levels and inhibition of lipolysis. Phosphorylates PIKFYVE on 'Ser-318', which results in increased PI(3)P-5 activity. The Rho GTPase-activating protein DLC1 is another substrate and its phosphorylation is implicated in the regulation cell proliferation and cell growth. AKT plays a role as key modulator of the AKT-mTOR signaling pathway controlling the tempo of the process of newborn neurons integration during adult neurogenesis, including correct neuron positioning, dendritic development

and synapse formation. Signals downstream of phosphatidylinositol 3-kinase (PI(3)K) to mediate the effects of various growth factors such as platelet-derived growth factor (PDGF), epidermal growth factor (EGF), insulin and insulin-like growth factor I (IGF-I). AKT mediates the antiapoptotic effects of IGF-I. Essential for the SPATA13-mediated regulation of cell migration and adhesion assembly and disassembly. May be involved in the regulation of the placental development. Phosphorylates STK4/MST1 at 'Thr-120' and 'Thr-387' leading to inhibition of its: kinase activity, nuclear translocation, autophosphorylation and ability to phosphorylate FOXO3. Phosphorylates STK3/MST2 at 'Thr-117' and 'Thr-384' leading to inhibition of its: cleavage, kinase activity, autophosphorylation at Thr-180, binding to RASSF1 and nuclear translocation. Phosphorylates SRPK2 and enhances its kinase activity towards SRSF2 and ACIN1 and promotes its nuclear translocation. Phosphorylates RAF1 at 'Ser-259' and negatively regulates its activity. Phosphorylation of BAD stimulates its proapoptotic activity. Phosphorylates KAT6A at 'Thr-369' and this phosphorylation inhibits the interaction of KAT6A with PML and negatively regulates its acetylation activity towards p53/TP53.

### **AKT1 antibody - N-terminal region - References**

- Jones P.F., et al. Proc. Natl. Acad. Sci. U.S.A. 88:4171-4175(1991).  
Matsubara A., et al. Diabetologia 44:910-913(2001).  
Ota T., et al. Nat. Genet. 36:40-45(2004).  
Goshima N., et al. Nat. Methods 5:1011-1017(2008).  
Heilig R., et al. Nature 421:601-607(2003).