

Rabbit Anti-phospho-Androgen Receptor (Ser94)/AP Conjugated antibody

SL12473R-AP

Product Name	Anti-phospho-Androgen Receptor (Ser94)/AP
Chinese Name	碱性磷酸酶 (AP) 标记的磷酸化雄激素受体抗体
Alias	Androgen Receptor (phospho S94); Androgen Receptor (Phospho-Ser94); Androgen Receptor (phospho Ser94); p-Androgen Receptor (Ser94); ANDR_HUMAN; HYSP1; AIS; Androgen receptor (dihydrotestosterone receptor; testicular feminization; spinal and bulbar muscular atrophy; Kennedy disease); AR; DHTR; Dihydro Testosterone Receptor; Dihydrotestosterone receptor; HUMARA; Nuclear receptor subfamily 3 group C member 4; SBMA; SMAX1; Spinal and bulbar muscular atrophy; TFM.
Product Type	Phosphorylated anti
Research Area	Tumour Cell biology Developmental biology Chromatin and nuclear signals Signal transduction transcriptional regulatory factor Kinases and Phosphatases Epigenetics
Immunogen Species	Rabbit
Clonality	Polyclonal
React Species	Human
Applications	IHC-P=1:100-500,IHC-F=1:100-500 not yet tested in other applications. optimal dilutions/concentrations should be determined by the end user.
Molecular weight	99kDa
Form	Lyophilized or Liquid
Concentration	1mg/ml
immunogen	KLH conjugated synthesised phosphopeptide derived from human Androgen Receptor around the phosphorylation site of Ser94
Lsotype	IgG
Purification	affinity purified by Protein A
Storage Buffer	1M TBS(pH7.4) with 1% BSA, 3% Proclin300 and 50% Glycerol.
Storage	Store at -20 °C for one year. Avoid repeated freeze/thaw cycles. The lyophilized antibody is stable at room temperature for at least one month and

for greater than a year when kept at -20°C . When reconstituted in sterile pH 7.4 1M PBS or diluent of antibody the antibody is stable for at least two weeks at $2-4^{\circ}\text{C}$.

background:

The androgen receptor gene is more than 90 kb long and codes for a protein that has 3 major functional domains: the N-terminal domain, DNA-binding domain, and androgen-binding domain. The protein functions as a steroid-hormone activated transcription factor. Upon binding the hormone ligand, the receptor dissociates from accessory proteins, translocates into the nucleus, dimerizes, and then stimulates transcription of androgen responsive genes. This gene contains 2 polymorphic trinucleotide repeat segments that encode polyglutamine and polyglycine tracts in the N-terminal transactivation domain of its protein. Expansion of the polyglutamine tract causes spinal bulbar muscular atrophy (Kennedy disease). Mutations in this gene are also associated with complete androgen insensitivity (CAIS). Two alternatively spliced variants encoding distinct isoforms have been described. [provided by RefSeq, Jul 2008]

Function:

Steroid hormone receptors are ligand-activated transcription factors that regulate eukaryotic gene expression and affect cellular proliferation and differentiation in target tissues. Transcription factor activity is modulated by bound coactivator and corepressor proteins. Transcription activation is down-regulated by NR0B2. Activated, but not phosphorylated, by HIPK3.

Product Detail

Subunit:

Binds DNA as a homodimer. Part of a ternary complex containing AR, EFCAB6/DJBP and PARK7. Interacts with HIPK3 and NR0B2 in the presence of androgen. The ligand binding domain interacts with KAT7/HBO1 in the presence of dihydrotestosterone. Interacts with EFCAB6/DJBP, PELP1, PQBP1, RANBP9, RBAK, SPDEF, SRA1, TGFB1I1, ZNF318 and RREB1. Interacts with ZMIZ1/ZIMP10 and ZMIZ2/ZMIP7 which both enhance its transactivation activity. Interacts with SLC30A9 and RAD54L2/ARIP4. Interacts via the ligand-binding domain with LXXLL and FXXLF motifs from NCOA1, NCOA2, NCOA3, NCOA4 and MAGEA11. The AR N-terminal poly-Gln region binds Ran resulting in enhancement of AR-mediated transactivation. Ran-binding decreases as the poly-Gln length increases. Interacts with HIP1 (via coiled coil domain). Interacts (via ligand-binding domain) with TRIM68. Interacts with TNK2. Interacts with USP26. Interacts with RNF6. Interacts (regulated by RNF6 probably through polyubiquitination) with RNF14; regulates AR transcriptional activity. Interacts with PRMT2 and TRIM24. Interacts with GNB2L1/RACK1. Interacts with RANBP10; this interaction enhances dihydrotestosterone-induced AR transcriptional activity. Interacts with PRPF6

in a hormone-independent way; this interaction enhances dihydrotestosterone-induced AR transcriptional activity. Interacts with STK4/MST1. Interacts with ZIPK/DAPK3. Interacts with LPXN. Interacts with MAK. Part of a complex containing AR, MAK and NCOA3.

Subcellular Location:

Nucleus. Cytoplasm. Predominantly cytoplasmic in unliganded form but translocates to the nucleus upon ligand-binding. Can also translocate to the nucleus in unliganded form in the presence of GNB2L1.

Tissue Specificity:

Isoform 2 is mainly expressed in heart and skeletal muscle.

Post-translational modifications:

Sumoylated on Lys-386 (major) and Lys-520. Ubiquitinated. Deubiquitinated by USP26. 'Lys-6' and 'Lys-27'-linked polyubiquitination by RNF6 modulates AR transcriptional activity and specificity.

Phosphorylated in prostate cancer cells in response to several growth factors including EGF. Phosphorylation is induced by c-Src kinase (CSK). Tyr-534 is one of the major phosphorylation sites and an increase in phosphorylation and Src kinase activity is associated with prostate cancer progression.

Phosphorylation by TNK2 enhances the DNA-binding and transcriptional activity and may be responsible for androgen-independent progression of prostate cancer.

DISEASE:

Defects in AR are the cause of androgen insensitivity syndrome (AIS) [MIM:300068]; previously known as testicular feminization syndrome (TFM). AIS is an X-linked recessive form of pseudohermaphroditism due end-organ resistance to androgen. Affected males have female external genitalia, female breast development, blind vagina, absent uterus and female adnexa, and abdominal or inguinal testes, despite a normal 46,XY karyotype. Defects in AR are the cause of spinal and bulbar muscular atrophy X-linked type 1 (SMA1) [MIM:313200]; also known as Kennedy disease. SMA1 is an X-linked recessive form of spinal muscular atrophy. Spinal muscular atrophy refers to a group of neuromuscular disorders characterized by degeneration of the anterior horn cells of the spinal cord, leading to symmetrical muscle weakness and atrophy. SMA1 occurs only in men. Age at onset is usually in the third to fifth decade of life, but earlier involvement has been reported. It is characterized by slowly progressive limb and bulbar muscle weakness with fasciculations, muscle atrophy, and gynecomastia. The disorder is clinically similar to classic forms of autosomal spinal muscular atrophy. Note=Caused by trinucleotide CAG repeat expansion. In SMA1 patients the number of Gln ranges from 38 to 62. Longer expansions result in

earlier onset and more severe clinical manifestations of the disease.

Note=Defects in AR may play a role in metastatic prostate cancer. The mutated receptor stimulates prostate growth and metastases development despite of androgen ablation. This treatment can reduce primary and metastatic lesions probably by inducing apoptosis of tumor cells when they express the wild-type receptor.

Defects in AR are the cause of androgen insensitivity syndrome partial (PAIS) [MIM:312300]; also known as Reifenstein syndrome. PAIS is characterized by hypospadias, hypogonadism, gynecomastia, genital ambiguity, normal XY karyotype, and a pedigree pattern consistent with X-linked recessive inheritance. Some patients present azoospermia or severe oligospermia without other clinical manifestations.

Similarity:

Belongs to the nuclear hormone receptor family.

NR3 subfamily.

Contains 1 nuclear receptor DNA-binding domain.

Database links:

[Entrez Gene: 367](#) Human

[Entrez Gene: 11835](#) Mouse

[Entrez Gene: 24208](#) Rat

[Omim: 313700](#) Human

[SwissProt: P10275](#) Human

[SwissProt: P19091](#) Mouse

[SwissProt: P15207](#) Rat

[Unigene: 496240](#) Human

[Unigene: 39005](#) Mouse

[Unigene: 394224](#) Mouse

[Unigene: 439657](#) Mouse

[Unigene: 9813](#) Rat



SunLong Biotech Co.,LTD

Tel: 0086-571-56623320 Fax:0086-571-56623318

E-mail:sales@sunlongbiotech.com

www.sunlongbiotech.com

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