

Datasheet for ABIN6136020

anti-ATR antibody (pSer428)



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| Quantity: | 100 μL | | | |
|--|--|--|--|--|
| Target: | ATR | | | |
| Binding Specificity: | pSer428 | | | |
| Reactivity: | Human | | | |
| Host: | Rabbit | | | |
| Clonality: | Polyclonal | | | |
| Conjugate: | This ATR antibody is un-conjugated | | | |
| Application: | Immunohistochemistry (IHC), Western Blotting (WB), Immunofluorescence (IF) | | | |
| Product Details | | | | |
| Immunogen: | A synthetic phosphorylated peptide around S428 of human ATR (NP_001175.2). | | | |
| Sequence: | GISPK | | | |
| | | | | |
| Isotype: | IgG | | | |
| Isotype: Cross-Reactivity: | | | | |
| | IgG | | | |
| Cross-Reactivity: | IgG Human, Mouse, Rat | | | |
| Cross-Reactivity: Characteristics: | IgG Human, Mouse, Rat Phosphorylated Antibodies | | | |
| Cross-Reactivity: Characteristics: Purification: | IgG Human, Mouse, Rat Phosphorylated Antibodies | | | |

Target Details

| Background: | The protein encoded by this gene belongs the PI3/PI4-kinase family, and is most closely related | | |
|---------------------|---|--|--|
| | to ATM, a protein kinase encoded by the gene mutated in ataxia telangiectasia. This protein and | | |
| | ATM share similarity with Schizosaccharomyces pombe rad3, a cell cycle checkpoint gene | | |
| | required for cell cycle arrest and DNA damage repair in response to DNA damage. This kinase | | |
| | has been shown to phosphorylate checkpoint kinase CHK1, checkpoint proteins RAD17, and | | |
| | RAD9, as well as tumor suppressor protein BRCA1. Mutations of this gene are associated with | | |
| | Seckel syndrome. An alternatively spliced transcript variant of this gene has been reported, | | |
| | however, its full length nature is not known. Transcript variants utilizing alternative polyA sites | | |
| | exist.,ATR,FCTCS,FRP1,MEC1,SCKL,SCKL1,Epigenetics & Nuclear Signaling,Chromatin | | |
| | Modifying Enzymes,Phosphorylation,DNA Damage & Repair,Cancer,Tumor suppressors,Signal | | |
| | Transduction,Kinase,Serine/threonine kinases,Cell Biology & Developmental | | |
| | Biology,Apoptosis,Mitochondrial Control of Apoptosis,Cell Cycle,G1/S Checkpoint,G2/M DNA | | |
| | Damage Checkpoint,Protein phosphorylation,ATR | | |
| Molecular Weight: | 294 kDa/297 kDa/301 kDa | | |
| Gene ID: | 545 | | |
| UniProt: | Q13535 | | |
| Pathways: | Positive Regulation of Response to DNA Damage Stimulus | | |
| Application Details | | | |
| Application Notes: | WB,1:500 - 1:2000,IHC,1:50 - 1:200,IF,1:50 - 1:200 | | |
| Comment: | HIGH QUALITY | | |
| Restrictions: | For Research Use only | | |
| Handling | | | |
| Format: | Liquid | | |
| Buffer: | PBS with 0.02 % sodium azide,50 % glycerol, pH 7.3. | | |
| Preservative: | Sodium azide | | |
| Precaution of Use: | This product contains Sodium azide: a POISONOUS AND HAZARDOUS SUBSTANCE which | | |
| | should be handled by trained staff only. | | |
| Storage: | -20 °C | | |
| Storage Comment: | Store at -20°C. Avoid freeze / thaw cycles. | | |