

Certificate of Analysis

Aurora B, active

(Recombinant enzyme expressed in Sf21 insect cells)

Item # 14-835, 14-835-K, 14-835M

Parent Lot # D8PN002N

The data presented in this document apply to the parent lot shown above and to all pack sizes derived from subsequent vialling runs of this parent lot. An alphabetical suffix after the parent lot number is used to denote each vialling run.

Product Description: N-terminal 6His-tagged, recombinant, human Aurora B, full length and N-terminal GST-tagged, recombinant, human INCENP, amino acids 821-end. Co-expressed by baculovirus in Sf21 insect cells. Purified using glutathione agarose. Combined purity 84% by SDS-PAGE and Coomassie blue staining. Aurora B MW = 40kDa, INCENP MW = 38kDa

Formulation: 0.93mg/ml of 50mM Tris/HCl pH7.5, 300mM NaCl, 0.1mM EGTA, 0.03% Brij-35, 270mM sucrose, 1mM benzamidine, 0.2mM PMSF, 0.1% 2-mercaptoethanol. Frozen solution.

Storage and Stability: On receipt of material store at -70°C. Unopened reagent is stable for a minimum of 1 year from date of shipment when stored at recommended storage temperature. Avoid repeat freeze/thaw cycles. For maximum recovery of product, centrifuge original vial prior to removing the cap.

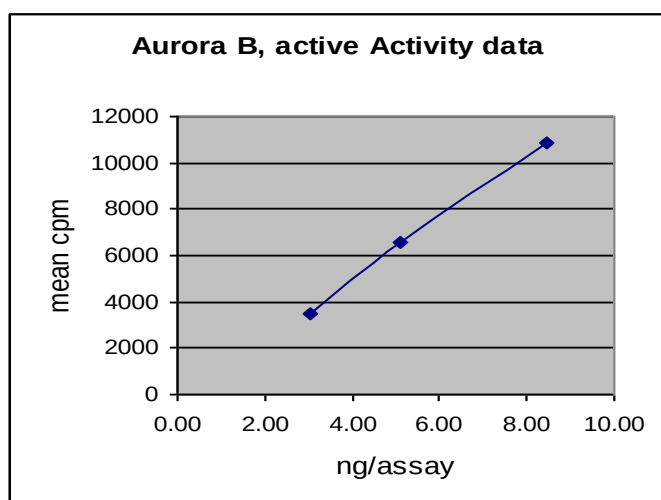
Specific Activity (Parent lot # D8PN002N): 734U/mg, where one unit of Aurora B, active activity is defined as 1nmol phosphate incorporated into 30µM (AKRRRLSSLRA) per minute at 30°C with a final ATP concentration of 100µM.

Handling Recommendations: Rapidly thaw the vial under cold water and immediately place on ice. Aliquot unused material into pre-chilled micro-centrifuge tubes and immediately snap-freeze the vials in liquid nitrogen prior to re-storage at -70°C.

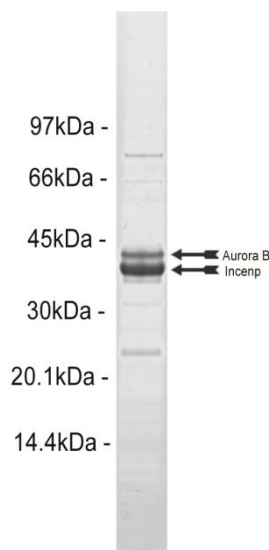
**FOR IN VITRO RESEARCH USE ONLY
NOT FOR USE IN HUMANS OR ANIMALS**

Quality Control Testing

Kinase Assay: 3.06–8.44ng of this lot of enzyme phosphorylated 30µM (AKRRRLSSLRA) in the assay described on page two. Assay background was subtracted from the actual counts to yield the results shown below.



MS Tryptic Fingerprint: Confirmed identity as Aurora B with the translated sequence listed on page three. Confirmed identity as INCENP with the translated sequence listed on page four.



SDS-PAGE and Coomassie Stain: Purity was assessed by SDS-PAGE and Coomassie blue staining using 3µg of Aurora B, active.

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Kinase Assay Protocol

Stock Solutions:

1. **5 x Reaction Buffer:** 40mM MOPS/NaOH pH7.0, 1mM EDTA, other components.
2. **(AKRRRLSSLR):** Use at a final assay concentration of 30µM. Prepare a 0.3mM stock and add 2.5µl of stock per assay point.
3. **Aurora B, active:** Dilute with 20mM MOPS/NaOH pH7.0, 1mM EDTA, 0.01% Brij-35, 5% glycerol, 0.1% 2-mercaptoethanol, 1mg/ml BSA. Use 3.06–8.44ng per assay point.
4. **[γ-³³P]ATP:** 2.5 x MgAc/[γ-³³P]ATP cocktail: 25mM MgAc and 0.25mM ATP to which is added [γ-³³P]ATP (specific activity approximately 500 - 800cpm/pmol as required).

Assay Procedure (96 well plate format):

1. Add 5µl of 5 x reaction buffer per assay to wells.
2. Add 2.5µl of **(AKRRRLSSLRA)**.
3. Add **2.5µl (3.06–8.44ng) Aurora B**.
4. Add 5µl of dH₂O.
5. Add 10µl of diluted [γ-³³P]ATP mixture.
6. Incubate for 10 minutes at 30°C.
7. Stop the reaction by adding 5µl of 3% phosphoric acid.
8. Transfer a 10µl aliquot onto the appropriate area of a **P30 Filtermat**.
9. Wash the filtermat three times for 5 minutes with 75mM phosphoric acid.
10. Wash the filtermat once for 2 minutes with methanol.
11. Transfer the filtermat to a sealable plastic bag and add 4ml of scintillation cocktail.
12. Read in a scintillation counter. Compare cpm of enzyme samples with cpm of control samples that contain all assay components plus 1µl of 30% phosphoric acid.

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Aurora B Sequence Information

<u>Protein</u>	human Aurora B
<u>Tags</u>	N-terminal 6His
<u>Native sequence</u>	M8 of the recombinant protein is equivalent to M1 of human Aurora B
<u>Accession number</u>	GenBank NM_004217

Recombinant Aurora B amino acid sequence:

```

1  MHHHHHHMAQ KENSYPWPYG RQTAPSGLST LPQRVLRKEP VTPSALVLMS RSNVQPTAAP
61  GQKVMENSSG TPDILTRHFT IDDFEIGRPL GKGKFGNVYL AREKKSHFIV ALKVLFKSQI
121 EKEGVEHQLR REIEIQAHLH HPNILRLYNY FYDRRIYLI LEYAPRGELY KELQKSCTFD
181 EQRTATIMEE LADALMYCHG KKVIHRDIKP ENLLLGLKGE LKIADFGWSV HAPSLRRKTM
241 CGTLDYLPPE MIEGRMHNEK VDLWCIGVLC YELLVGNNPF ESASHNETYR RIVKVDLKFP
301 ASVPTGAQDL ISKLLRHNP S ERLPLAQVSA HPWVRANSRR VLPPSALQSV A

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Recombinant Aurora B nucleotide sequence:

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1  atgcaccatc accatcacca tatggcccag aaggagaact cctacccctg gccctacggc
61  cgacagacgg ctccatctgg cctgagcacc ctgccccagc gaggctctcg gaaagagcct
121 gtcaccccat ctgcacttgt cctcatgagc cgctccaatg tccagccac agctgcccct
181 ggccagaagg tgatggagaa tagcagtggg acaccgcaca tcttaacgag gcacttcaca
241 attgatgact ttgagattgg gcgtcctctg ggcaaaggca agtttggaag cgtgtacttg
301 gctcgggaga agaaaagcca ttctatcgtg gcgtcgaagg tcctcttcaa gtcccagata
361 gagaaggagg gcgtggagca tcagctgcgc agagagatcg aaatccaggc ccacctgcac
421 catcccaaca tcctgctgtc ctacaactat ttttatgacc ggaggaggat ctacttgatt
481 ctagagtatg cccccgcggg ggagctctac aaggagctgc agaagagctg cacatttgac
541 gagcagcgaa cagccacgat catggaggag ttggcagatg ctctaattga ctgccatggg
601 aagaagggtga ttacacagaga cataaagcca gaaaatctgc tcttagggct caagggagag
661 ctgaagattg ctgacttcgg ctggtctgtg catgcgccct ccctgaggag gaagacaatg
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781 gtggatctgt ggtgcattgg agtgctttgc tatgagctgc tgggtgggaa cccacccttt
841 gagagtgcac cacacaacga gacctatcgc cgcacgtcga aggtggacct aaagtccccc
901 gcttctgtgc ccacgggagc ccaggacctc atctccaaac tgctcaggca taaccctctg
961 gaacggctgc ccctggccca ggtctcagcc cacccttggg tccgggcca ctctcggagg
1021 gtgctgcctc cctctgcctc tcaatctgtc gctctga

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INCENP Sequence Information

<u>Protein</u>	human INCENP
<u>Tags</u>	N-terminal GST
<u>Native sequence</u>	D231 of the recombinant protein is equivalent to D821 of human INCENP
<u>Accession number</u>	GenBank NM_020238

Recombinant INCENP amino acid sequence:

```

1  MSPILGYWKI  KGLVQPTRLL  LEYLEEKYEE  HLYERDEGDK  WRNKKFELGL  EFPNLPYYID
61  GDVKLTQSMa  IIRYIADKHN  MLGGCPKERA  EISMLEGAVL  DIRYGVSRIA  YSKDFETLKV
121  DFLSKLP EML  KMFEDRLCHK  TYLNGDHVTH  PDFMLYDALD  VVLYMDPMCL  DAFPKLVCFK
181  KRIEAI PQID  KYLKSSKYIA  WPLQGWQATF  GGGDHPPKSD  LEVLFQGPEF  DLNSDDSTDD
241  EAHPRKPIPT  WARGTPLSQA  IIHQYYHPPN  LLELFGTILP  LDLEDIFKKS  KPRYHKRTSS
301  AVWNSPPLQG  ARVPSSLAYS  LKKH

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Recombinant INCENP nucleotide sequence:

```

1  atgtccccta  tactaggtta  ttggaaaatt  aagggccttg  tgcaaccac  tgcatttctt
61  ttggaatatc  ttgaagaaaa  atatgaagag  catttgatg  agcgcgatga  aggtgataaa
121  tggcgaaaca  aaaagtttga  attgggtttg  gagtttccca  atcttcctta  ttatatattg
181  ggtgatgta  aattaacaca  gtctatggcc  atcatagctt  atatagtga  caagcacaac
241  atgttgggtg  gttgtccaaa  agagcgtgca  gagatttcaa  tgcttgaagg  agcgggtttg
301  gatattagat  acggtgtttc  gagaattgca  tatagtaaa  actttgaaac  tctcaaagtt
361  gattttctta  gcaagctacc  tgaaatgctg  aaaatgttcg  aagatcgttt  atgtcataaa
421  acatatatta  atggtgatca  tgtaacccat  cctgacttca  tgttgatga  cgctcttgat
481  gttgttttat  acatggaccc  aatgtgcctg  gatgcgttcc  caaaattagt  ttgttttaaa
541  aaacgtattg  aagctatccc  acaaattgat  aagtacttga  aatccagcaa  gtatatagca
601  tggcctttgc  agggctggca  agccacgttt  ggtggtggcg  accatcctcc  aaaatcggat
661  ctggaagtgc  tggtccagg  gcccgaaatt  gatctgaata  ggcacgactc  caccgatgat
721  gaggccatc  cccggaagcc  catccccacc  tggggccgag  gcaccccgct  cagccaggct
781  atcattcacc  agtactacca  cccaccgaac  cttctggagc  tctttggaac  cattctccca
841  ctggacttgg  aggatatctt  caagaagagc  aagccccgct  atcacaagcg  caccagctct
901  gctgtctgga  actcaccgcc  cctgcagggc  gccagggctc  ccagcagcct  ggcctacagc
961  ctgaagaagc  actga

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Reviewed and approved by site quality representative.

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