

# IκB alpha ProtéGene™ Set

Catalog# 11000  
 Lot# On label

## Materials Provided:

1. pMEV-IκBα-WT (I1000a): 20 μg in 40 μl TE (pH7.5), 0.5 mg/ml.
2. pMEV-IκBα-DN (I1000b): 20 μg in 40 μl TE (pH7.5), 0.5 mg/ml.
3. Product Information Sheets.

**Note:** Individual plasmids can be ordered separately. Some plasmids are shipped as lyophilized pellet.

## Receiving and Storage:

If received in lyophilized form, add 40μl sterile DI water to the vial, dissolve thoroughly by vortex and then collect the contents by centrifuging the vials briefly in a microcentrifuge.

If received in liquid form, spin the vials briefly in a microcentrifuge to collect the contents. Store the products at 2-8°C if used immediately or at -20°C for extended storage.

## Expression Vector:

pMEV-2HA (a): Cat# P1001a.

## Affinity Tag:

N-terminal 2 x HA, a 9-aa peptide derived from influenza virus (MGYPYDVPDYAYPYDVPDYAGS...).

## Prokaryotic Selection:

The kanamycin-resistance gene (aminoglycoside 3' phospho-transferase) expression cassette in the plasmids confers Kanamycin resistance to bacteria cells. Bacterial cells transformed with the plasmids should be maintained and grown in media containing 25-50μg/ml Kanamycin (e.g. #LK-1100, Prepared LB Agar plates, Biomyx, San Diego, California).

## Eukaryotic Selection:

The neomycin resistance gene, driven by SV40 early promoter, confers G418 resistance to eukaryotic cells. Stable mammalian cell lines can be selected with G418 after transfection.

## Description of IκBα and Mutants

I kappa B proteins (IκBs) bind to and inhibit the NF-κB complex by trapping it in the cytoplasm (OMIM #164008). IκBs could be phosphorylated by IKKs and subsequently polyubiquitinated for degradation to allow translocation of NF-κB into the nucleus (reviewed in 1 and 2). Serines 32 and 36 in the N terminal regulatory domain of IκBα are the conserved phosphorylation sites (3). Substitution with alanines (S32A and S36A) of the phosphoacceptor sites would block the phosphorylation and degradation of IκBα by IKKs (4, 5).

## Molecular Features of the Inserts:

**Gene:** *Homo sapiens* I kappa B alpha (IκBα)  
**GenBank Reference Sequence:** NM\_020529  
**5'-Cloning Site:** Bam HI  
**5'-Junction Sequence:** 5'-...tac gct gga tcc ATG TTC CAG-...3'  
**3'-Cloning Site:** Xba I  
**3'-Junction Sequence:** 5'-... gtcgactctaga TCA TAA CGT-...3'

## hIKBA Nucleotide and Protein Sequences

(954 bps encoding 317 amino acid residues. S32, S36 and respective encoding nucleotides are in bold and underlined)

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1 ATGTTCCAGG CGGCCGAGCG CCCCAGGAG TGGGCCATGG AGGGCCCCCG CGACGGGCTG
  m f q a a e r p q e w a m e g p r d g l
61 AAGAAGGAGC GGCTACTGGA CGACGCCAC GACAGCGGCC TGGATCCAT GAAAGCAGG
  k k e r l l d d r h d s g l d s m k d e
121 GAGTACGAGC AGATGGTCAA GGAGCTGCAG GAGATCCGCC TCGAGCCGCA GGAGGTCCCG
  e y e q m v k e l q e i r l e p q e v p
181 CGCGGCTCGG AGCCCTGGAA GCAGCAGCTC ACCGAGGACG GGGACTCGTT CCGTCACTTG
  r g s e p w k q q l t e d g d s f l h l
241 GCCATCATCC ATGAAGAAAA GGCAGTGACC ATGGAAGTGA TCCGCCAGGT GAAGGGAGAC
  a i i h e e k a l t m e v i r q v k g d
301 CTGGCTTTCC TCAACTTCCA GAACAACCTG CAGCAGACTC CACTCCACTT GGCTGTGATC
  l a f l n f q n n l q q t p l h l a v i
361 ACCAACCAGC CAGAAATTGC TGAGGCACCT CTGGGAGCTG GCTGTGATCC TGAGTCCGA
  t n q p e i a e a l l g a g c d p e l r
421 GACTTTCGAG GAAATACCCC CCTACACCTT GCCTGTGAGC AGGGCTGCCT GGCCAGCGTG
  d f r g n t p l h l a c e q g c l a s v
481 GGAGTCTGTA CTCAGTCTTG CACCACCCCG CACTCCACTC CCATCCTGAA GGCTACCAAC
  g v l t q s c t t p h l h s i l k a t n
541 TACAATGGCC ACACGTGTCT ACACCTAGCC TCTATCCATG GCTACTCGGG CATCGTGGAG
  y n g h t c l h l a s i h g y l g i v e
601 CTTTTGGTGT CCTTGGGTGC TGATGTCAAT GCTCAGGAGC CTTGTAATGG CCGGACTGCC
  l l v s l g a d v n a q e p c n g r t a
661 CTTCACTCGC CAGTGGACCT GCAAAATCCT GACTGGTGT CACTCGTGT GAAAGTGTGG
  l h l a v d l q n p d l v s l l l k c g
721 GCTGATGTCA ACAGAGTTAC CTACCAGGCG TATTCTCCCT ACCAGCTCAC CTGGGGCCCG
  a d v n r v t y q g y s p y q l t w g r
781 CCAAGCACCC GGATACAGCA GCAGCTGGCC CAGCTGACAC TAGAAAACCT TCAGATGCTG
  p s t r i q q q l g q l t l e n l q m l
841 CCAGAGAGTG AGGATGAGGA GAGCTATGAC ACAGAGTCAG AGTTCACGGA GTTCACAGAG
  p e s e d e e s y d t e s e f t e f t e
901 GACGAGCTGC CCTATGATGA CTGTGTGTTT GGAGGCCAGC GTCTGACGTT ATGA
  d e l p y d d c v f g g q r l t l -
  
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## Mutations:

pMEV-IKBA-WT (I1000a): No mutation  
 pMEV-IKBA-DN (I1000b): S32A; S36A

## References:

1. Baldwin, A. S. (1996) *Annu. Rev. Immunol.* 14: 649-681 The NF-κappa B and I kappa B proteins: new discoveries and insights
2. Verma, I. M., Stevenson, J. K., Schwarz, E. M., et al., (1995) *Genes Dev.* 9, 2723-2735. Rel/NF-κappa B/I kappa B family: intimate tales of association and dissociation
3. Karin M. (1999) *J Biol Chem.* 274(39): 27339-42 The Beginning of the End: IκB Kinase (IKK) and NF-κB Activation
4. J DiDonato, F Mercurio, C Rosette et al., (1996) *Mol. Cell. Biol.*, 16(4): 1295-1304 Mapping of the inducible IκappaB phosphorylation sites that signal its ubiquitination and degradation
5. Jaspers I., Samet, JM, and Reed W., (1999). Arsetite exposure of cultured airway epithelial cells activates κB-dependent interleukin-8 gene expression in the absence of nuclear factor-κB nuclear translocation. *J. Biol. Chem.*, 274: 31025-33