## 32-4179: Recombinant Human Methyl-CpG Binding Domain Protein 3

Alternative Name : Methyl-CpG Binding Domain Protein 3,Methyl-CpG-Binding Domain Protein 3,Methyl-CpG-Binding Protein

## Description

Source : E.coli. MBD3 Human Recombinant produced in E. coli is a single polypeptide chain containing 314 amino acids (1-291) and having a molecular mass of 35.2 kDa . MBD3 is fused to a 23 amino acid His-tag at N -terminus \& purified by proprietary chromatographic techniques. Methyl-CpG-binding domain protein 3 (MBD3), belongs to the MBD family of transcriptional repressors. Human proteins MECP2, MBD1, MBD2, MBD3, and MBD4 includes a family of nuclear proteins related by the presence of a methyl-CpG binding domain (MBD) in each protein. Nevertheless, not like the other family members, MBD3 is not capable of binding to methylated DNA. MBD3 is a subunit of the NuRD, a multisubunit complex having nucleosome remodeling and histone deacetylase activities. The predicted MBD3 protein shares $71 \%$ and $94 \%$ identity with MBD2 (isoform 1) and mouse Mbd3. MBD3 mediates the association of metastasis-associated protein 2 (MTA2) with the core histone deacetylase complex.

## Product Info

| Amount : | $10 \mu \mathrm{~g}$ |
| :---: | :---: |
| Purification : | Greater than $85 \%$ as determined by SDS-PAGE. |
| Content : | The MBD3 solution ( $0.25 \mathrm{mg} / 1 \mathrm{ml}$ ) contains 20 mM Tris- HCl buffer ( pH 8.0 ), $0.2 \mathrm{M} \mathrm{NaCl}, 1 \mathrm{mM}$ DTT, 1 mM EDTA and $40 \%$ glycerol. |
| Storage condition : | Store at $4^{\circ} \mathrm{C}$ if entire vial will be used within $2-4$ weeks. Store, frozen at $-20^{\circ} \mathrm{C}$ for longer periods of time. For long term storage it is recommended to add a carrier protein ( $0.1 \%$ HSA or BSA).Avoid multiple freeze-thaw cycles. |
| Amino Acid : | MGSSHHHHHH SSGLVPRGSH MGSMERKRWE CPALPQGWER EEVPRRSGLS AGHRDVFYYS |
|  | PSGKKFRSKP QLARYLGGSM DLSTFDFRTG KMLMSKMNKS RQRVRYDSSN QVKGKPDLNT |
|  | ALPVRQTASI FKQPVTKITN HPSNKVKSDP QKAVDQPRQL FWEKKLSGLN AFDIAEELVK |
|  | TMDLPKGLQG VGPGCTDETL LSAIASALHT STMPITGQLS AAVEKNPGVW LNTTQPLCKA |
|  | FMVTDEDIRK QEELVQQVRK RLEEALMADM LAHVEELARD GEAPLDKACA EDDDEEDEEE |
|  | EEEEPDPDPE MEHV. |



