

TFEB Rabbit mAb

Catalog # AP76734

Product Information

Application	WB, IHC-P, IHC-F, FC, IP
Primary Accession	P19484
Reactivity	Human
Host	Rabbit
Clonality	Monoclonal Antibody
Isotype	IgG
Conjugate	Unconjugated
Purification	Affinity Purified
Calculated MW	52865

Additional Information

Gene ID	7942
Other Names	TFEB
Dilution	WB~~1:1000 IHC-P~~N/A IHC-F~~N/A FC~~1:10~50 IP~~N/A
Format	Liquid in 50mM Tris-Glycine(pH 7.4), 0.15M NaCl, 40%Glycerol, 0.01% sodium azide and 0.05% BSA.
Storage	Store at 4°C short term. Aliquot and store at -20°C long term. Avoid freeze/thaw cycles.

Protein Information

Name	TFEB {ECO:0000303 PubMed:2115126, ECO:0000312 HGNC:HGNC:11753}
Function	Transcription factor that acts as a master regulator of lysosomal biogenesis, autophagy, lysosomal exocytosis, lipid catabolism, energy metabolism and immune response (PubMed:21617040, PubMed:22343943, PubMed:22576015, PubMed:22692423, PubMed:25720963, PubMed:30120233, PubMed:31672913, PubMed:32612235, PubMed:32753672, PubMed:35662396, PubMed:36697823, PubMed:36749723, PubMed:37079666). Specifically recognizes and binds E-box sequences (5'-CANNTG-3'); efficient DNA-binding requires dimerization with itself or with another MiT/TFE family member such as TFE3 or MITF (PubMed:1748288, PubMed:19556463, PubMed:29146937). Involved in the cellular response to amino acid availability by acting downstream of MTOR: in the presence of nutrients, TFEB phosphorylation by MTOR promotes its cytosolic retention and subsequent inactivation (PubMed:21617040, PubMed:22343943, PubMed:22576015, PubMed:22692423, PubMed:25720963, PubMed:32612235, PubMed:32753672,

PubMed:[35662396](#), PubMed:[36697823](#)). Upon starvation or lysosomal stress, inhibition of MTOR induces TFEB dephosphorylation, resulting in nuclear localization and transcription factor activity (PubMed:[22343943](#), PubMed:[22576015](#), PubMed:[22692423](#), PubMed:[25720963](#), PubMed:[32612235](#), PubMed:[32753672](#), PubMed:[35662396](#), PubMed:[36697823](#)). Specifically recognizes and binds the CLEAR-box sequence (5'-GTCACGTGAC-3') present in the regulatory region of many lysosomal genes, leading to activate their expression, thereby playing a central role in expression of lysosomal genes (PubMed:[19556463](#), PubMed:[22692423](#)). Regulates lysosomal positioning in response to nutrient deprivation by promoting the expression of PIP4P1 (PubMed:[29146937](#)). Acts as a positive regulator of autophagy by promoting expression of genes involved in autophagy (PubMed:[21617040](#), PubMed:[22576015](#), PubMed:[23434374](#), PubMed:[27278822](#)). In association with TFE3, activates the expression of CD40L in T-cells, thereby playing a role in T-cell-dependent antibody responses in activated CD4(+) T-cells and thymus-dependent humoral immunity (By similarity). Specifically recognizes the gamma-E3 box, a subset of E-boxes, present in the heavy- chain immunoglobulin enhancer (PubMed:[2115126](#)). Plays a role in the signal transduction processes required for normal vascularization of the placenta (By similarity). Involved in the immune response to infection by the bacteria *S.aureus*, *S.typhimurium* or *S.enterica*: infection promotes itaconate production, leading to alkylation, resulting in nuclear localization and transcription factor activity (PubMed:[35662396](#)). Itaconate-mediated alkylation activates TFEB- dependent lysosomal biogenesis, facilitating the bacteria clearance during the antibacterial innate immune response (PubMed:[35662396](#)). In association with ACS2, promotes the expression of genes involved in lysosome biogenesis and both autophagy upon glucose deprivation (PubMed:[28552616](#)).

Cellular Location

Nucleus. Cytoplasm, cytosol. Lysosome membrane. Note=Mainly present in the cytoplasm (PubMed:[23434374](#), PubMed:[33691586](#), PubMed:[35662396](#)). When nutrients are present, recruited to the lysosomal membrane via association with GDP-bound RagC/RRAGC (or RagD/RRAGD): it is then phosphorylated by MTOR (PubMed:[23401004](#), PubMed:[32612235](#), PubMed:[36697823](#)). Phosphorylation by MTOR prevents nuclear translocation and activity by promoting interaction with 14-3-3 proteins, such as YWHAZ (PubMed:[22343943](#), PubMed:[22692423](#), PubMed:[23401004](#), PubMed:[25720963](#), PubMed:[32612235](#), PubMed:[32753672](#), PubMed:[35662396](#), PubMed:[36697823](#), PubMed:[37079666](#)). Under aberrant lysosomal storage conditions, it translocates from the cytoplasm to the nucleus (PubMed:[21617040](#), PubMed:[22576015](#), PubMed:[23434374](#), PubMed:[25720963](#), PubMed:[32753672](#)). The translocation to the nucleus is regulated by ATP13A2 (PubMed:[23434374](#), PubMed:[27278822](#)). Conversely, inhibition of mTORC1, starvation and lysosomal disruption, promotes dephosphorylation and translocation to the nucleus (PubMed:[22343943](#), PubMed:[22692423](#), PubMed:[37079666](#)). Exported from the nucleus in response to nutrient availability (PubMed:[30120233](#)). In macrophages, translocates into the nucleus upon live *S.enterica* infection (PubMed:[27184844](#)).

Background

Transcription factor that acts as a master regulator of lysosomal biogenesis, autophagy, lysosomal exocytosis, lipid catabolism, energy metabolism and immune response (PubMed:[21617040](#), PubMed:[22576015](#), PubMed:[22343943](#), PubMed:[22692423](#), PubMed:[30120233](#), PubMed:[31672913](#))