## CPTC-HRAS-1 (CAB080330)

Uniprot ID: P01112

Protein name: RASH\_HUMAN Full name: GTPase HRas

Tissue specificity: Widely expressed.

Function: Involved in the activation of Ras protein signal transduction (PubMed:22821884). Ras proteins bind GDP/GTP and possess intrinsic GTPase activity

(PubMed:12740440, PubMed:14500341, PubMed:9020151).

Unnamed:

Cell membrane (Topo: Lipid-anchor; Orientation: Cytoplasmic side)

Golgi apparatus

Golgi apparatus membrane (Topo: Lipid-anchor)

NOTE: The active GTP-bound form is localized most strongly to membranes than the inactive GDP-bound form (By similarity). Shuttles between the plasma membrane

and the Golgi apparatus.

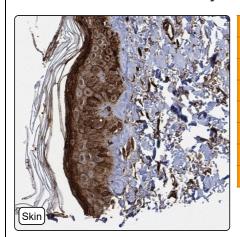
Isoform 2: Nucleus Cytoplasm

Cytoplasm > Perinuclear region

NOTE: Colocalizes with RACK1 to the perinuclear region. Protein existence: Experimental evidence at protein level

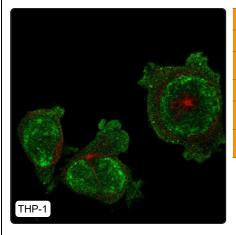
Comment:

## **Immunohistochemistry**



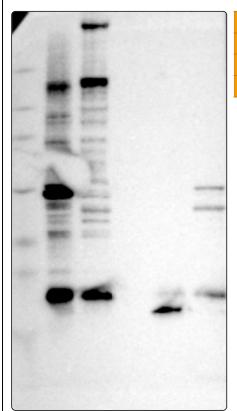
	LUED 110 BU 6 1 400	
IHC protocol:	HIER pH6, Dilution 1:400	
IHC test staining:	Cytoplasmic positivity in most tissues.	
Literature conformance:	Consistent with extensive gene/protein characterization data	
Literature significance:		
RNA similarity:	Medium consistency between antibody staining and RNA expression data	
RNA tissue specificity:	Low tissue specificity	
RNA tissue distribution:	Detected in all	
IHC Sibling similarity:	Other antibody shows similar IHC staining pattern	

## Immunofluorescence



IF Overlay:	antibody (green), anti-tubulin (red) and DAPI (blue)
IF main location:	Nuclear speckles - 3: <b>Supportive</b> (auto)
IF additional location:	Plasma membrane - 3: <b>Supportive</b> (auto) Vesicles - 5: <b>Approved</b> (auto)
IF approved for publication on HPA:	No
IF in THP-1:	Nuc speckles Plasma membrane (Edge)
IF in U2OS:	Nuc speckles Vesicles

## Western blot



WB Size markers (kDa):	250, 130, 100, 70, 55, 35, 25, 15, 10	
WB Lanes:	Marker (1), RT-4 (2), U-251MG (3), Plasma (4), Liver (5), Tonsil (6)	
WB Target weight (kDa):	19, 19, 21, 21, 21	
WB Validation:	Supported (Band of predicted size in kDa (+/-20%) with additional bands present.)	