# **Anti-Senataxin** [OY11]

Catalogue number: 151842 **Sub-type:** Primary antibody

Images:

### Contributor

**Inventor:** Stephen West

Institute: Cancer Research UK, London Research Institute: Clare Hall Laboratories

Images:

### **Tool details**

#### \*FOR RESEARCH USE ONLY

Name: Anti-Senataxin [OY11]

Alternate name:

Class: Polyclonal

Conjugate: Unconjugated

Cancer Tools.org **Description:** Defects in Senataxin are the cause of neurodegenerative diseases AOA-2 and ALS4. Senataxin play vital roles in DNA repair and transcription termination. The staining pattern of the antibody is located in the nucleus and chromatin.

Purpose: Parental cell: Organism: Tissue: Model: Gender: Isotype:

Reactivity: Human

Selectivity: **Host:** Rabbit

**Immunogen:** A mixture of four peptides corresponding to the following amino acid ranges of human

Senataxin: aa8-30, aa884-895, aa1173-1192, aa2654-2677

**Immunogen UNIPROT ID:** 

Sequence:

**Growth properties: Production details:** 

Formulation:

Recommended controls: Whole-cell or Chromatin extracts of HeLa or HEK293 cells

**Bacterial resistance:** 

# Selectable markers: Additional notes:

### **Target details**

Target: Senataxin

#### **Target alternate names:**

**Target background:** Defects in Senataxin are the cause of neurodegenerative diseases AOA-2 and ALS4. Senataxin play vital roles in DNA repair and transcription termination. The staining pattern of the antibody is located in the nucleus and chromatin.

Cancer Tools.org

Molecular weight: 303 kDa

Ic50:

## **Applications**

**Application:** IP; WB **Application notes:** 

## **Handling**

Format: Liquid

Concentration: 1 mg/ml

Passage number:
Growth medium:
Temperature:
Atmosphere:
Volume:

Storage medium:

Storage buffer: 0.1 M Tris-Glycine (pH 7.4), 150 mM NaCl.

**Storage conditions:** -15° C to -25° C **Shipping conditions:** Shipping at 4° C

### **Related tools**

Related tools:

### References

**References:** Frye et al. 2010. Cancer Lett. 289(1):71-80. PMID: 19740597. ; Genomic gain of 5p15 leads to over-expression of Misu (NSUN2) in breast cancer. ; Hussain et al. 2009. J Cell Biol. 186(1):27-40. PMID: 19596847. ; The nucleolar RNA methyltransferase Misu (NSun2) is required for mitotic spindle stability. ; Frye et al. 2006. Curr Biol. 16(10):971-81. PMID: 16713953. ; The RNA methyltransferase Misu (NSun2) mediates Myc-induced proliferation and is upregulated in tumors.

