

EIF4A2 (eukaryotic translation initiation factor 4A, isoform 2)

Identity

| | |
|-------------|------------------------|
| Other names | DDX2B |
| Hugo | EIF4A2 |
| Location | 3q28 |

DNA/RNA

Transcription 1.9 kb mRNA; 1223 bp coding sequence

Protein

| | |
|-------------|---|
| Description | 407 amino acids; 4.6 kDa; NH2-term, ATP binding site, DEAD box, COOH-term |
| Expression | wide; also expressed in the embryo |
| Function | ADP/ATP binding; RNA helicase; binds mRNA to the ribosome: role in initiation of protein synthesis; also -binds single strand DNA |
| Homology | DEAD box proteins, in particular other EIF4A |

Implicated in

| | |
|---------------------|--|
| Entity | t(3;18)(q27;p11.2) --> EIF4A2/ BCL6 |
| Disease | non-Hodgkin lymphoma |
| Prognosis | yet unknown (only 1case available) |
| Hybrid/Mutated Gene | 5' EIF4A2 - 3' BCL6; EIF4A2 fuses with the second exon of BCL6 |
| Abnormal Protein | no fusion protein, but promoter exchange |
| Oncogenesis | BCL6 is a transcription repressor; it is supposed that substitution of the promoter of BCL6 may be responsible for BCL6 deregulation |

External links

| Nomenclature | |
|-----------------------------|--|
| Hugo | EIF4A2 |
| GDB | EIF4A2 |
| Entrez_Gene | EIF4A2_1974 eukaryotic translation initiation factor 4A, isoform 2 |

| Cards | |
|-----------------------|-----------------------------|
| Atlas | EIF4A2ID262 |

[GeneCards](#) [EIF4A2](#)
[Ensembl](#) [EIF4A2](#)
[Genatlas](#) [EIF4A2](#)
[GeneLynx](#) [EIF4A2](#)
[eGenome](#) [EIF4A2](#)
[euGene](#) [1974](#)

Genomic and cartography

[GoldenPath](#) [EIF4A2](#) - [3q28](#) [chr3:187984055-187990377](#) + [3q27.3](#) (hg18-Mar_2006)
[Ensembl](#) [EIF4A2 - 3q27.3 \[CytoView\]](#)
[NCBI](#) [Genes Cyto](#) [Gene Seg](#) [Map View - NCBI]
[OMIM](#) [Disease map \[OMIM\]](#)
[HomoloGene](#) [EIF4A2](#)

Gene and transcription

[Genbank](#) [AB209021](#) [ENTREZ]
[Genbank](#) [AF208852](#) [ENTREZ]
[Genbank](#) [AK130087](#) [ENTREZ]
[Genbank](#) [AL117412](#) [ENTREZ]
[Genbank](#) [AL137681](#) [ENTREZ]
[RefSeq](#) [NM_001967](#) [SRS] [NM_001967](#) [ENTREZ]
[AceView](#) [EIF4A2](#) AceView - NCBI
[TRASER](#) [EIF4A2](#) Traser - Stanford
[Unigene](#) [Hs.478553](#) [SRS] [Hs.478553](#) [NCBI] [HS478553](#) [spliceNest]

Protein : pattern, domain, 3D structure

[SwissProt](#) [Q14240](#) [SRS] [Q14240](#) [EXPASY] [Q14240](#) [INTERPRO]
[Prosite](#) [PS00039 DEAD ATP HELICASE](#) [SRS] [PS00039 DEAD ATP HELICASE](#) [Expasy]
[Interpro](#) [IPR001410 DEAD](#) [SRS] [IPR001410 DEAD](#) [EBI]
[Interpro](#) [IPR011545 DEAD/DEAH_N](#) [SRS] [IPR011545 DEAD/DEAH_N](#) [EBI]
[Interpro](#) [IPR000629 DEAD_box](#) [SRS] [IPR000629 DEAD_box](#) [EBI]
[Interpro](#) [IPR001650 Helicase_C](#) [SRS] [IPR001650 Helicase_C](#) [EBI]
[CluSTr](#) [Q14240](#)
[Pfam](#) [PF00270 DEAD](#) [SRS] [PF00270 DEAD](#) [Sanger] [pfam00270](#) [NCBI-CDD]
[Pfam](#) [PF00271 Helicase_C](#) [SRS] [PF00271 Helicase_C](#) [Sanger]
[Smart](#) [SM00487 DEXDc](#) [EMBL]
[Smart](#) [SM00490 HELICc](#) [EMBL]
[Blocks](#) [Q14240](#)

Protein Interaction databases

[DIP](#) [Q14240](#)

[IntAct](#) [Q14240](#)

Polymorphism : SNP, mutations, diseases

[OMIM](#) [601102](#) [[map](#)]

[GENECLINICS](#) [601102](#)

[SNP](#) [EIF4A2](#) [dbSNP-NCBI]

[SNP](#) [NM_001967](#) [SNP-NCI]

[SNP](#) [EIF4A2](#) [GeneSNPs - Utah] [EIF4A2](#) [HGBASE - SRS] [EIF4A2](#) [SNP - HAPMAP]

General knowledge

[Family Browser](#) [EIF4A2](#) [UCSC Family Browser]

[SOURCE](#) [NM_001967](#)

[SMD](#) [Hs.478553](#)

[SAGE](#) [Hs.478553](#)

[Enzyme](#) [3.6.1.-](#) [Enzyme-SRS] [3.6.1.-](#) [Brenda-SRS] [3.6.1.-](#) [KEGG] [3.6.1.-](#) [WIT]

[Amigo](#) [nucleotide binding](#)

[Amigo](#) [nucleic acid binding](#)

[Amigo](#) [DNA binding](#)

[Amigo](#) [RNA binding](#)

[Amigo](#) [translation initiation factor activity](#)

[Amigo](#) [protein binding](#)

[Amigo](#) [ATP binding](#)

[Amigo](#) [protein biosynthesis](#)

[Amigo](#) [regulation of translational initiation](#)

[Amigo](#) [ATP-dependent helicase activity](#)

[Amigo](#) [eukaryotic translation initiation factor 4F complex](#)

[Amigo](#) [hydrolase activity](#)

[BIOCARTA](#) [Regulation of eIF4e and p70 S6 Kinase](#) [[Genes](#)]

[BIOCARTA](#) [Eukaryotic protein translation](#) [[Genes](#)]

[PubGene](#) [EIF4A2](#)

Other databases

Probes

[Probe](#) [EIF4A2 Related clones \(RZPD - Berlin\)](#)

PubMed

[PubMed](#) [5 Pubmed reference\(s\) in LocusLink](#)

Bibliography

Isolation and mapping of the human EIF4A2 gene homologous to the murine protein synthesis initiation factor 4A-II gene Eif4a2.

Sudo K, Takahashi E, Nakamura Y.
Cytogenet Cell Genet 1995; 71: 385-388.
Medline [96103476](#)

The DEAD box protein eIF4A. 2. A cycle of nucleotide and RNA-dependent conformational changes.

Lorsch JR, Herschlag D.
Biochemistry 1998; 37: 2194-2206.
Medline [98153098](#)

Eukaryotic translation initiation factor 4AIII (eIF4AIII) is functionally distinct from eIF4AI and eIF4AII.

Li Q, Imataka H, Morino S, Rogers GW Jr, Richter-Cook NJ, Merrick WC, Sonenberg N.
Mol Cell Biol 1999; 19: 7336-7346.
Medline [99454990](#)

Identification of heterologous translocation partner genes fused to the BCL6 gene in diffuse large B-cell lymphomas: 5'-RACE and LA - PCR analyses of biopsy samples.

Yoshida S, Kaneita Y, Aoki Y, Seto M, Mori S, Moriyama M.
Oncogene 1999; 18: 7994-7999.
Medline [20105774](#)

[REVIEW articles](#) *automatic search in PubMed*

[Last year publications](#) *automatic search in PubMed*

[BiblioGene - INIST](#)

Contributor(s)

Written 07-2000 Jean-Loup Huret

Citation

This paper should be referenced as such :

Huret JL . EIF4A2 (eukaryotic translation initiation factor 4A, isoform 2). Atlas Genet Cytogenet Oncol Haematol. July 2000 .

URL : <http://AtlasGeneticsOncology.org/Genes/EIF4A2ID262.html>

© *Atlas of Genetics and Cytogenetics in Oncology and Haematology*
