

Gene Section

Review

GZMA (granzyme A (granzyme 1, cytotoxic T-lymphocyte-associated serine esterase 3))

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Identity

Other names: CTLA3, HFSP

HGNC (Hugo): GZMA

Location: 5q11.2

Local order: Size: 7607 bases. Coordinates: 54398473.

DNA/RNA

Description

The GZMA gene, with 7607 bases in length, consists of 5 exons and 4 introns. GZMA gene is located in a gene

cluster together with granzyme K (figure 1) (Grossman et al., 2003).

Transcription

There are at least two transcripts of human GZMA whose expression is differentially regulated by glucocorticoid (Ruike et al., 2007). These transcripts generate two isoforms, GZMA α and GZMA β , which have respective first exons: exon 1a and exon 1b (figure 1):

GZMA α (exon 1a): canonical sequence,

GZMA β (exon 1b): lack aa 1-17; aa 18-23 LLLIPE --> MTKGLR.

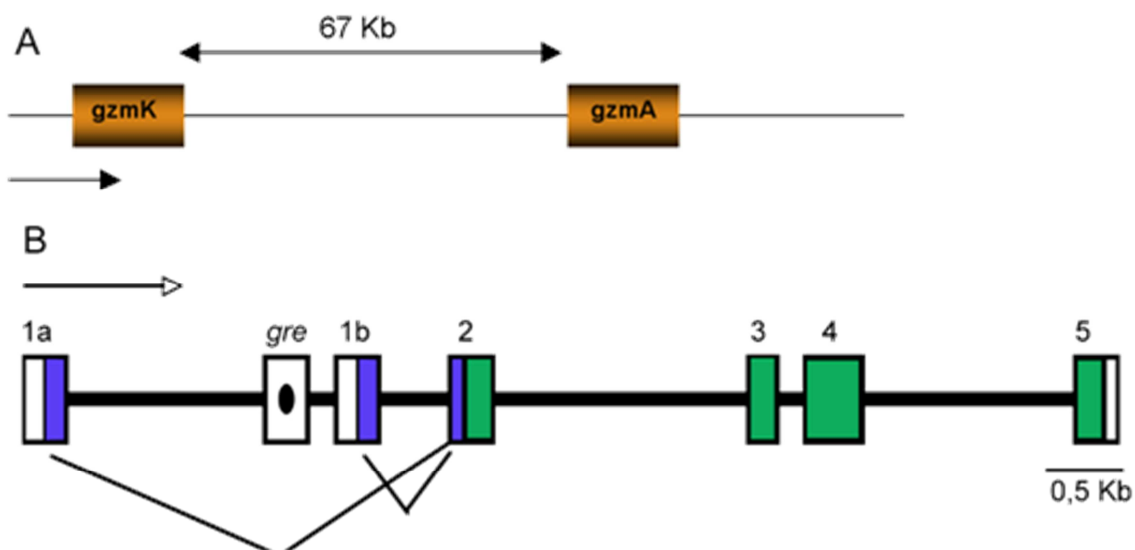


Figure 1. Genomic organization of human GZMA. A, human GZMA cluster. Arrow indicate the direction of transcription. B, representation of the GZMA genetic locus. White: untranslated regions; Blue: leader sequence; Green: mature enzyme. Solid lanes: splicing between the first and second exons. gre: glucocorticoid response element (adapted from Ruike et al., 2007).

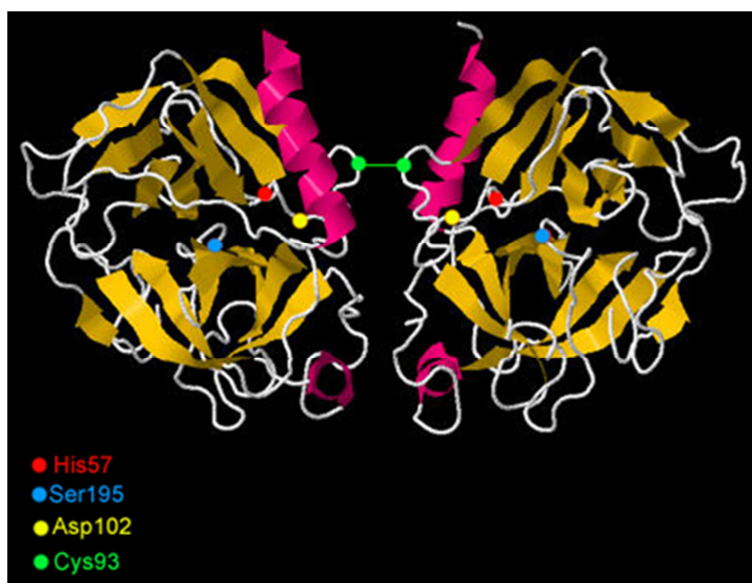


Figure 2. Diagram of the crystal structure of human granzyme A dimer (Bell et al., 2003; Hink-Schauer et al., 2003). The cysteine groups involved in disulphide bond-mediated dimer (green) and the three aminoacids forming the catalytic triad (red, blue and yellow) are shown. Representation from PDB (accession code 1OP8) deposited by Hink-Schauer C, Estébanez-Perpiñá E, Kurschus FC, Bode W, Jenne DE. *Nat Struct Biol.* 2003 Jul;10(7):535-40.

Protein

Description

Granzyme A is a tryptase (cleave proteins after Lys or Arg residues) expressed mainly in cytotoxic cells (cytotoxic T and Natural Killer cells) (Masson et al., 1986; Simon et al., 1986; Young et al., 1986). Protein is expressed as a proenzyme (Jenne et al., 1988) containing a signal sequence that mediates targeting of the nascent enzyme to the ER. Cleavage of the signal peptide produces an inactive proenzyme that contains an N-terminal dipeptide that needs to be cleaved to produce an active protease. In the Golgi, a mannose-6-phosphate tag is added for transporting the proenzyme to cytotoxic granules. Within the cytotoxic granule, the N-terminal dipeptide is removed by cathepsin C (dipeptidyl peptidase I) (Pham et al., 1999), producing the active enzyme that is kept inactive at low pH. Native granzyme A is expressed as a dimer (Bell et al., 2003; Hink-Schauer et al., 2003).

Expression

Cytotoxic CD8+ T cells, Natural Killer cells, CD4+ T cells, gamma-delta T cells, type II pneumocytes, alveolar macrophages, bronchiolar epithelial cells.

Localisation

Cytotoxic granules.

Function

Granzyme A is delivered from CTL or NK cytotoxic granules to the cytoplasm of target cell by a mechanism dependent on perforin (Baran et al., 2009; Praper et al., 2011; Thiery et al., 2011).

There are some controversial findings about the physiological function of *gzmA*.

It has been reported that human *GzmA* induces perforin-mediated caspase-independent cell death in some tumors cell lines (Hayes et al., 1989; Shi et al., 1992; Beresford et al., 1999; Shresta et al., 1999; Pardo et al., 2004). *GzmA* translocates to the nucleus and mitochondria where key substrates such as mitochondrial complex I protein, NADH dehydrogenase Fe-S protein 3 (NDUFS3) is cleaved, inducing the production of Radical Oxygen Species (ROS). ROS production induces the activation of the SET complex that translocates into the nucleus in order to repair DNA damage induced by ROS. Once there, granzyme A cleaves components of the endoplasmic reticulum-associated SET complex, releasing the endonuclease NM23H1 that induces single strand nicks in the DNA and ultimately cell death (Lieberman, 2011).

Other authors have reported that the cytotoxic potential of granzyme A is low, but induce expression of pro-inflammatory cytokines in monocytes-like cells by a caspase-1 dependent mechanism (Metkar et al., 2008). Granzyme A is able to cleave several extracellular substrates like thrombin receptor, fibronectin, collagen IV, proteinase-activated receptor-2, Pro-urokinase plasminogen activator and myelin basic protein (Kramer et al., 1987; Buzza et al., 2006; Hendel et al., 2011).

Granzyme and granzyme B double deficient mice are more susceptible than granzyme B deficient mice to transplanted tumors suggesting a contribution of granzyme A to tumor control in vivo (Pardo et al., 2002; Cao et al., 2007).

Homology

Mouse granzyme A; Rat granzyme A; Chicken granzyme A; Fish granzyme A (Common Carp, Atlantic cod, Channel catfish) (Praveen et al., 2006; Praveen et al., 2006; Wernersson et al., 2006).

Mutations

Note: Not known.

Implicated in

Septis (Froelich et al., 2009; Hendel et al., 2011)

Disease

Several findings suggest that *gzmA* contributes to septic shock. Native and recombinant human granzyme A as well as a human NK cell line expressing *gzmA* induces human adherent peripheral blood mononuclear cells to express proinflammatory cytokines including interleukin-1beta, interleukin-6, interleukin-8 and TNF-alpha (Sower et al., 1996; Metkar et al., 2008). Granzyme A deficient mice are more resistant than wild type mice to septic shock induced by LPS (Metkar et al., 2008).

Rheumatoid arthritis

Prognosis

Granzyme A levels are higher in serum and synovial fluid of patients with rheumatoid arthritis (Griffiths et al., 1992; Nordstrom et al., 1992; Kummer et al., 1994; Tak et al., 1994; Muller-Ladner et al., 1995; Spaeny-Dekking et al., 1998; Tak et al., 1999).

Chronic obstructive pulmonary disease

Prognosis

Granzyme A is expressed in type II pneumocytes of patients with severe chronic obstructive pulmonary disease (Vernooy et al., 2007).

Hypersensitivity pneumonitis

Prognosis

Granzyme A is elevated in bronchoalveolar lavage fluid from patients with hypersensitivity pneumonitis (Tremblay et al., 2000).

Sjögren's syndrome

Prognosis

Granzyme A is expressed in salivary glands from patients with Sjögren's syndrome (Alpert et al., 1994).

Poxvirus infection

Disease

Granzyme A deficient mice are more susceptible than wild type mice to mousepox virus (ectromelia) (Mullbacher et al., 1996).

Herpes virus infection

Disease

Granzyme A deficient mice are more susceptible than wild type mice to herpes simplex virus type 1 (HSV-1)

(Pereira et al., 2000) and mouse cytomegalovirus (CMV) infection (Riera et al., 2000).

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