

Cn β 1 shifts cardiac metabolism

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Cardiovascular diseases have the highest mortality rate worldwide, and their incidence increases with aging. Among them, heart failure after myocardial infarction or maladaptive hypertrophy represents a major health challenge, especially among the elderly. The calcium-regulated phosphatase calcineurin plays a major role in the development of pathological cardiac hypertrophy and heart failure through the activation of the transcription factor NFAT. Calcineurin is composed of a catalytic and a regulatory subunit, CnA and CnB respectively [1]. CnA has 4 major domains: catalytic, CnB-binding, calmodulin-binding and autoinhibitory. In the absence of Ca²⁺, the autoinhibitory domain blocks the catalytic domain, preventing access of the substrate and inhibiting CnA's activity [2]. Following Ca²⁺ increase in the cytoplasm, the autoinhibitory domain in CnA is removed, the substrate reaches the catalytic domain and is dephosphorylated.

Three different genes encode CnA: CnA α and CnA β are ubiquitously expressed, whereas CnA γ is mainly re-

stricted to brain and testis. Notably, the CnA β gene expresses an alternative isoform regulated by differential alternative polyadenylation of Exon12 called CnA β 1 (Figure 1A). Contrary to all other CnAs, CnA β 1 lacks the classical autoinhibitory domain and instead contains a unique C-terminal region not present in any other protein. This alternative sequence contains two different α -helices, comprising an LXVP inhibitory motif and a new Golgi localization signal (Figure 1B) [3, 4]. Unlike other calcineurin isoforms, CnA β 1 promotes Akt phosphorylation by mTORC2, rather than NFAT dephosphorylation. Akt activation depends on the localization of CnA β 1 in the Golgi apparatus, which is regulated by its interaction with the Golgi transmembrane protein Cog8 (Figure 1C). The interaction between CnA β 1 and mTORC2 occurs through its alternative C-terminal region and the Golgi localization of CnA β 1 is necessary for the relocalization of mTORC2 from the cytoplasm to the membranes of the cell, and the subsequent phosphorylation of Akt (Figure 1C).

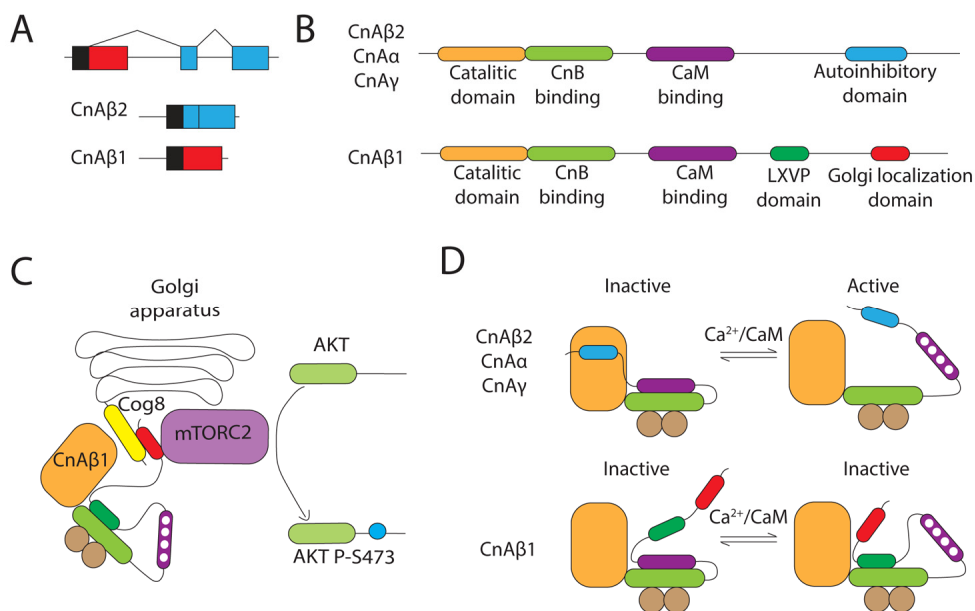


Figure 1. Cn β 1 alternative signalling promotes activation of the mTORC2/Akt pathway. (A) Cn β 1 is the result of an alternative polyadenylation of Exon 12 in the CnA β gene. **(B)** Cn β 1's alternative C-terminal region includes an LXVP motif and a Golgi localization sequence. **(C)** Cn β 1 is localized in the Golgi apparatus through its interaction with Cog8 and modulates mTORC2 phosphorylation of Akt. **(D)** The LXVP inhibitory motif blocks Cn β 1's catalytic domain even in the presence of Ca²⁺ and calmodulin. The schematic is based on [3].

The recent identification of an LXVP motif within the alternative C-terminal region of CnAβ1 has provided a better characterization of its biochemistry [3]. The LXVP peptide was previously found to be a potent inhibitor of CnA activity [2]. The incorporation of an LXVP motif provides CnAβ1's C-terminal region with a similar function, reducing its phosphatase activity even in the presence of Ca²⁺ and calmodulin (Figure 1D). This is in agreement with previous results showing that a CnAβ1 catalytic-dead mutant had a similar capacity to activate Akt, suggesting that CnAβ1 works as an adaptor protein rather than as a phosphatase [5]. Moreover, the LXVP motif has an unprecedented importance in the context of Ca²⁺ oscillations in the Golgi apparatus.

Unlike all other CnA isoforms, which strongly promote maladaptive cardiac hypertrophy, CnAβ1 reduces hypertrophy by inducing genes involved in the serine and one-carbon metabolic pathway. Activation of this pathway in cardiomyocytes results in reduced protein oxidation in the mitochondria and preserved ATP production, which in turn improves systolic function and prevents adverse ventricular remodelling [6]. Activation of the Akt signalling pathway by CnAβ1 also improves cardiac function after myocardial infarction and promotes skeletal muscle regeneration [5, 7, 8]. The development of strategies to increase CnAβ1 expression and/or activation of the serine and one-carbon pathway in the heart will increase the quality of patients suffering from maladaptive cardiac hypertrophy or myocardial infarction, and reduce the burden of heart failure, especially among the elderly.

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