



**Katedry biochémie a genetiky**  
Prírodovedeckej fakulty Univerzity Komenského

Vás pozývajú na **39.** prednášku v rámci Kuželových seminárov:

## **Dr. Carolyn Suzuki**

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*New Insights into the Lon Protease and  
DnaJ-like Chaperone of Human Mitochondria:  
Roles in mtDNA metabolism and cell signaling*

ktorá sa uskutoční

**30.10. 2003** (štvrtok)  
o **15:00** v miestnosti B1-501 PriF UK

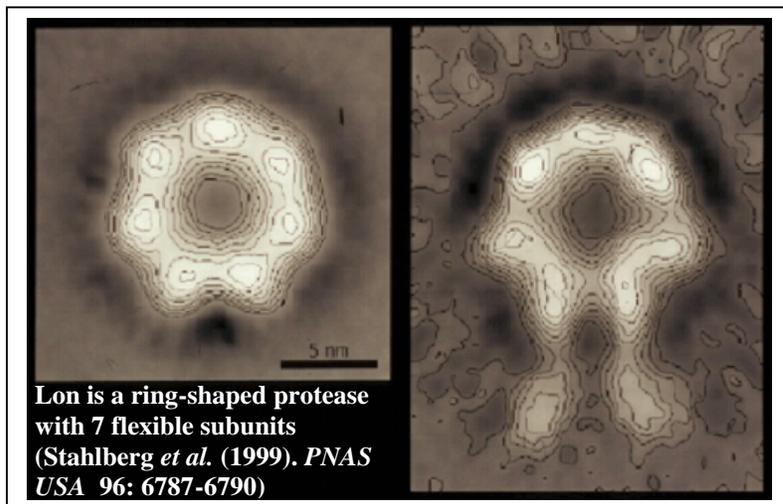
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## Dr. Carolyn Suzuki

[http://njms.umdnj.edu/biochemistry/fb\\_Suzuki\\_Carolyn\\_2002.htm](http://njms.umdnj.edu/biochemistry/fb_Suzuki_Carolyn_2002.htm)

ATP-dependent proteases are highly conserved enzymes present in humans as well as in *Archaea* and *Eubacteria*. In eukaryotes, regulated degradation by the ubiquitin-proteasome system is understood in great detail. Far less is known however, about the regulatory function of ATP-dependent proteases localized to mitochondria. Recent work has shown that the human disease Hereditary Spastic Paraplegia is caused by mutations in a gene that has high sequence similarity to ATP-dependent proteases localized to the mitochondrial inner membrane of yeast. Studies performed in the budding yeast *Saccharomyces cerevisiae* have demonstrated the crucial role of mitochondrial proteases and chaperones in organelle biogenesis,

respiration and mitochondrial DNA stability and gene expression.



Human Tid1-Long and -Short (hTid1-L and hTid1-S respectively), are mitochondrial matrix proteins that show a high degree of sequence similarity to the tumor suppressor Tid56 of *Drosophila melanogaster* and belong to the DnaJ family of molecular chaperones. The absence of Tid56 results in defective differentiation and morphogenesis giving rise to

tumorous imaginal discs that proliferate to form lethal tumors, which maintain their neoplastic characteristics when transplanted into wild-type flies. hTid1-L and -S are produced by alternative splicing and are 98% identical, differing only at their carboxyl-terminal tails. Work by Karl Munger's lab strikingly showed that hTid1-L and a dominant negative mutant of the Short isoform augment cell death induced by an apoptotic signal, whereas hTid1-S and a dominant negative mutant of the Long isoform inhibit cell death. Although these proteins are principally localized to the mitochondrial matrix, recent data suggest that hTid1-L and -S also function within the cytosol or nucleus by interacting with the human papillomavirus oncoprotein- E7, the Ras GTPase activating protein- Ras-GAP and the Janus kinase- Jak2.

### Recent publications:

Suzuki, C.K., K. Suda, N. Wang and G. Schatz (1994). Requirement of the yeast gene LON in intramitochondrial proteolysis and maintenance of respiration. *Science* **264**: 273-276 (1994).

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Suzuki, C.K., M. Rep, J. M. van Dijl, K. Suda, L.A. Grivell and G. Schatz (1997). ATP-dependent proteases that also chaperone protein biogenesis. *Trends Biochem. Sci.* **22**: 118-123.

van Dijl, J.M., E. Kutejova, K. Suda, D. Perecko, G. Schatz and C.K. Suzuki (1998). The ATPase and protease domains of yeast mitochondrial Lon: Roles in proteolysis and respiration-dependent growth. *Proc. Natl. Acad. Sci. USA* **18**: 10584 – 10589.

Beilharz, T., C. K. Suzuki and T. Lithgow (1998). A toxic fusion protein accumulating between the mitochondrial membranes inhibits protein assembly *in vivo*. *J. Biol. Chem.* **273**: 35268-35272.

Stahlberg, H., E. Kutejova, K. Suda, B. Wolpensinger, A. Lustig, G. Schatz, A. Engel, and C.K. Suzuki (1999). Mitochondrial Lon of *Saccharomyces cerevisiae* is a ring-shaped protease with seven flexible subunits. *Proc. Natl. Acad. Sci. USA* **96**: 6787-6790.

Lu, B., T. Liu, J.A. Crosby, J. Thomas-Wohlever, J., I. Lee and C.K. Suzuki (2003). The ATP-dependent Lon Protease of *Mus musculus* is a DNA-binding Protein that is Functionally Conserved between Yeast and Mammals. *Gene* **306**: 45-55.