Functional Roles of O-GlcNAc Modification on Nucleocytoplasmic Proteins

Jin Won Cho

Department of Biology, School of Biological Sciences, Yonsei University, Seoul, 120-749, Korea

E-mail: chojw311@yonsei.ac.kr, Tel:82-2-2123-4083, Fax: 82-2-363-4083

O-GlcNAc modification on nucleocytoplasmic proteins is found in all eukaryotes. Like phosphorylation, this modification is ubiquitous and very dynamic. The O-GlcNAc is modified at serine and threonine residues of the targeted proteins and its dynamic modification is caused by the action of O-GlcNAc transferase (OGT) and O-GlcNAcase. Because O-GlcNAc modification sites are shared with phosphorylation, this modification appears to have a 'yin-yang' relationship with phosphorylation with many regulatory proteins and to play an important role in regulation of protein function. Recently we found function and attachment sites of O-GlcNAc modification in p53 and NF-kB. Research of O-GlcNAc modification is rapidly growing field and especially O-GlcNAc modification is very important in diabetes, neural disease and development, and many cancers.

Recent Publications

- 1. Won Ho Yang, Ji Eun Kim, Hyung Wook Nam, Jung Won Ju, Hoe Suk Kim, Yu Sam Kim, and Jin Won Cho (2006) Modification of p53 with O-linked N-acetylglucosamine regulates p53 activity and stability. *Nature Cell Biol* 8:1073-1083.
- 2. Hoe Suk Kim, Eun Mi Kim, Jiae Lee, Won Ho Yang, Tae Yun Park, Young Min Kim and Jin Won Cho (2006) Heat shock protein 60 modified with

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- O-linked N-acetylglucosamine is involved in pancreatic β-cell death under hyperglycemic conditions. FEBS Letters 580: 2311-2316.
- 3. Hong-Kyung Kim, Jeong-Gu Kang, Shigehiko Yumura, Charles J. Walsh, Jin Won Cho, and Joo Hun Lee (2005) *De novo* Formation of Basal bodies in *Naegleria gruberi*: Regulation by Phosphorylation. *J Cell Biol* **169:**719-724.