Selective Gene Express Profiles in Rat Uterus during Estrus Cycle

<u>Do Rim Kim</u>, Seong Jin Yu, Jee Yun Kim, Mi Young Youm, Chae Kwan Lee*, Sung Goo Kang

Department of Biotechnology & Biomedical Science, Inje University, Kimhae 621-749. Korea, *Institute of Industrial Medicine, Inje University

The uterus undergoes dynamic changes during the cycle and displays many features typical of developmental process. In order to be prepared for implantation, endometrium undergoes predictable, sequential phases of proliferation and secretory changes. The uterus during estrus cycle synthesize a complex of signaling molecules with specific spatial and temporal modes of expression and which are critical for cell proliferlation and differentiation. The purpose of this investigation was to use cDNA microarrays to evaluate the expression of genes of rat uterus in estrus cycle. Animals were sacrificed on proestrus, estrus, metestrus, diestrus. Differential gene expression profiles were revealed(growthrelated c-myc reponsive protein RCL, heat shock 47-kDa protein (HSP47), cytochrome c oxidase polypeptide VIc2 (COX6C2), calreticulin (CALR)). Reverse transcription polymerase chain reaction (RT-PCR) was used to validate the relative expression pattern. Using this approach, we found several genes whose expression in rat uterus was altered with estrus cycle. Our long-term goal is to determine the role of these differentially expressed genes during estrus cycle. This study was supported by through the Biohealth Products Research Center(BPRC), Inje-University

Key words) endometrium, estrus cycle, cDNA array