

neurons. A 200 bp patch of the first intron may act in pharyngeal neurons and another 300 bp patch may act in mechanosensory neurons. We expect that each motif confers binding site for transcription activator(s) in different subsets of neuron cells, which may differ in cell lineage and function.

**D126**

**Interaction of Rbm, a Male Infertility Protein, with hnRNP K Suggests Its Function in mRNA Processing during Spermatogenesis.**

**Euisu Kim<sup>1\*</sup>, Sung Key Jang<sup>2</sup>, Kyungjin Kim<sup>1</sup> and Kunsoo Rhee<sup>1</sup>**

School of Biological Sciences, Seoul National University, Seoul 151-742<sup>1</sup> and Department of Life Science, Pohang University of Science and Technology, Pohang, Kyungbuk 790-784<sup>2</sup>

Microdeletions in the *AZFb* are strongly associated with male infertility. Multiple copies of the *Rbm* genes are located in the *AZFb* region on the Y chromosome. *Rbm* expression is restricted to the testis, especially the nuclei of the male germ cells. To understand biological functions of Rbm, we tried to identify proteins that interact with the Rbm protein. When we carried out yeast two-hybrid screening with a Rbm protein as a bait, we were able to observe a positive interaction with hnRNP K. Interacting domains of Rbm and hnRNP K were defined in yeast using several truncated mutants of both genes. Since the hnRNP K protein has multi-functions as a docking platform for controlling gene expression at the post-transcriptional level, we propose that Rbm plays an important role in spermatogenesis by controlling gene expression at the post-transcriptional level.

**D127**

**Subcellular Localization of Nek2 Suggests Its Dual Functions as a Cell Cycle Regulator**

**Jun Youl Choi<sup>1</sup>, Yongha Kim, Yeon-Tae Jeong and Kunsoo Rhee**

Seoul National University, School of Biological Sciences, Seoul 151-742

Nek2 is a mammalian protein kinase, structurally homologous to *Aspergillus* NIMA. Since NIMA functions as a cell cycle regulator by controlling chromatin condensation, it was proposed that Nek2 may also play a similar role as a cell cycle regulator in association with chromatin. On the contrary, it was reported that Nek2 can associate with centrosome, playing a key role in centrosomal behaviors during mitosis. To define biological functions of Nek2, we determined expression pattern and subcellular localization of endogenous Nek2 in the ovarian follicular cells as well as of the exogenous Nek2 mutants in NIH3T3 cells. The results revealed that Nek2 was expressed in a cell cycle-specific manner, in that the Nek2 protein was present abundantly in S/G2 phase of the cell cycle. The cell-cycle dependence of Nek2 expression may be regulated, at least in part, with proteolytic mechanisms. The Nek2 protein was localized both in the nucleus and in the cytoplasm. In the nucleus, Nek2 appeared to associate with mitotic chromatin. These results suggest that Nek2 may have dual functions related to the chromatin condensation and the centrosome cycle.

**D128**

**Postnatal Changes of the Steroidogenic Acute Regulatory Protein mRNA Expression in the Rat Brain**

**김현준, 박창환, 강상수\*, 최완성**

Department of Anatomy, College of Medicine,

Gyeongsang National University, 92 Chilam-dong,  
Chinju 660-751, Korea

Neurosteroids have been known to be synthesized in the central and peripheral nervous systems. In past years, many investigators participated in elucidation of the regulatory mechanism involved in postnatal brain development, especially, concerning developmental and regional specific expression of steroidogenic enzymes. Now, it is well accepted that the steroidogenic acute regulatory protein (StAR) plays essential roles and consists major rate limiting step in steroidogenesis. However, there is yet no evidence about StAR mRNA expression in developing brain. Thus, in this study, we firstly revealed changes of the expression pattern of StAR mRNA in several brain areas where other steroidogenic enzymes mainly expressed. As a result, the pattern of StAR mRNA expression was mainly changed in hypothalamus, hippocampus and cerebellum. We also detected variations of StAR expression according to their own developmental stages in the peripheral steroidogenic organs, adrenal glands and gonads. These results implicated that StAR might have a role in the neuronal cell growth and differentiation in the rat brain development likewise other steroidogenic enzymes.

#### D129

##### Effects of Ethanol on the Onset of Female Rat Puberty

정명희, 노구섭, 김현준, 김진현,  
강상수, 최완성\*

Department of Anatomy and Neurobiology,  
College of Medicine, Gyeongsang National  
University, Chinju, Korea

The present study was undertaken to examine the effects of ethanol on the hypothalamus-pituitary-gonad reproductive

neuroendocrine axis during prepubertal and onset of puberty in the immature female rat. From day 25, each rat began receiving either a control saline or ethanol. Animals were sacrificed on day 27 and 32, and their ovaries and blood were collected. In the present results, ethanol treatment significantly decreased serum luteinizing hormone contents at both time points. Uterine weights of ethanol-treated group were significantly lighter than control group at early time point, while there was no noticeable discrepancy at late time point. Viginal openings, a marker of onset of puberty, also clearly delayed in ethanol-treated group. Using an in situ hybridization histochemistry, we determined the expression of mRNAs encoding StAR. Ovaries from ethanol-treated rats showed a suppressed expression of StAR mRNA. These results demonstrate that ethanol affect the reproductive activity at the level of brain thereby disturb the prepubertal ovarian function and onset of puberty, at least in part, through the inhibition of ovarian StAR gene expression.

#### D 201

##### Control of Self-incompatibility by CO<sub>2</sub> Gas Treatment in *Brassica campestris*: Structural Alteration of Papillae Surface and Differential Gene Expression upon CO<sub>2</sub> Gas Treatment

Sang-Hee Lee<sup>1</sup>, Moon-Young Hong<sup>1</sup>,  
Shinjo Kim<sup>2</sup>, Byung-Dong Kim<sup>2</sup>,  
Byung-Hoon Min<sup>3</sup>, Nam-Kwon Baek<sup>4</sup>, and  
Yong-Yoon Chung<sup>1,2</sup>

Department of Biology, Korea University, Seoul 136-701<sup>1</sup>; Center for Plant Molecular Genetics & Breeding Research, Seoul National University, Suwon 441-744<sup>2</sup>; Department of Horticulture, Paichai University, Taejon<sup>3</sup>; Osan breeding institute, Choong Ang Seed Co., LTD, Osan 445-810<sup>4</sup>