

第10回 埼玉大学脳科学セミナー

主催：埼玉大学脳科学融合研究センター

**Molecular Basis of Normal and Disordered
Brain Development: mouse cerebellar
development transcriptome project,
CAPS2-mediated BDNF secretion, and
susceptibility to developmental disorders**

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CREST

日時： 2009年 7月10日（金曜日）
16:00 ~ 17:00

場所： 大学会館 2階 小集会室

本セミナー修了後に簡単な懇親会を予定しております（参加費300円）

問い合わせ先 坂井貴文（内線 4308）

脳科学融合研究センターは定期的に脳科学セミナーを開催する予定です。誰でも自由に参加出来るセミナーですので、奮ってご参加下さい。

セミナー要旨 Abstract

In the post-genomic sequencing era, genome-scale and genome-based approaches are important for comprehensive understanding the molecular mechanisms of brain development, function, and diseases. Our laboratory aims to elucidate the molecular mechanisms underlying normal brain development and its disorders by systematic identification and functional analyses of brain development genes. We focus on the genetic design for mouse cerebellar circuit development as a model system and have characterized the transcriptomic basis of mouse cerebellar circuit development by generating the Cerebellar Development Transcriptome Database (CDT-DB, <http://www.cdtdb.brain.riken.jp>). The CDT-DB project has opened a new door to elucidating cerebellar development from a transcriptomic standpoint and indicates that cerebellar circuit development is programmed by thousands of different genes that exhibit differential spatiotemporal expression patterns in the developing brain. In addition, we have been able to identify many brain development genes by CDT-DB mining. Deficits in the expression or function of brain development genes are likely to affect the development of neuronal circuits and may cause impaired brain function and behavior. We found that Ca²⁺-dependent activator protein for secretion 2 (CAPS2), one of the genes we identified, is involved in release of brain-derived neurotrophic factor (BDNF) and that CAPS2 knockout mice not only show reduced BDNF release activity but also exhibit autistic-like behavioral phenotypes such as decreased social interaction. Moreover, we identified increased expression of a rare splicing variant, which is never transported to axons, as well as single-nucleotide polymorphisms in some patients with autism, a developmental disorder characterized by qualitative impairments in social interactions and communications and by restricted interests and repetitive behavior. Taken together, we hypothesize that the disturbance of CAPS2-mediated BDNF release patterns is responsible for the onset of autism.