



# CDB SEMINAR

**Speaker: Osami Kanagawa**

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**Title: “Secondary rearrangement of TcR alpha chain in vivo: shaping and reshaping T cell repertoire”**

**Date: Tuesday, December 2**

**Time: 16:00 -17:00**

**Place: 7th floor Conference Room of Building A**

## **Summary:**

The genetic structure of the TcR  $\alpha$  locus and the lack of allelic exclusion permit multiple rearrangements on a single chromosome. Using TcR  $\alpha$  chain knock-in (KI) mice, we demonstrated the frequent secondary rearrangement of TcR  $\alpha$  chain in vivo. Furthermore, we provided evidence that  $\alpha$  locus rearrangement occurs initially using downstream V $\alpha$  and upstream J $\alpha$  and continue to use upstream V $\alpha$  and downstream J $\alpha$  for secondary rearrangement. These results indicate that single T cell can express TcRs with different specificity during thymic selection. These results together with our current analysis of the mouse line that lacks capacity to undergo TcR  $\alpha$  chain secondary rearrangement clearly demonstrate that multiple TcR  $\alpha$  chain rearrangements play a crucial role in the formation of functional T cell repertoire.

Analysis of TcR  $\alpha$  chain KI mouse lines revealed that the interaction between certain antigen/superantigen and mature peripheral T cells induces RAG dependent TcR  $\alpha$  locus secondary rearrangement. In non-immunized TcR KI mice, this change occurs in an age-dependent fashion and significantly affects peripheral T cell repertoire. Thus, secondary rearrangement of the TcR  $\alpha$  chain in mature T cells may play an important role in the age dependent-immune dysfunction.

**Host: Shin-ichi Nishikawa** <Stem Cell Biology, CDB>

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