## **NEKKEN Special Seminar**



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  Institute of Molecular and Cellular Biosciences, Department of Medical Genome Sciences,
  Institute of Molecular and Cellular Biosciences, Department of Medical Genome Sciences,
  The University of Tokyo, Japan

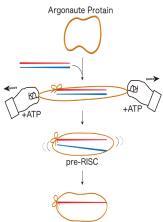
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## **Chaperoning RISC assembly**







Date: November 21 Mon, 2011 Time:17:00-18:30 Place: NEKKEN 1F Main meeting room
Daisuke Hayasaka, Institute of Tropical Medicine (NEKKEN), Tel:095-819-7828 E-mail:hayasaka@naqasaki-u.ac.jp

Mature RISC

## [Abstract]

MicroRNAs and small interfering RNAs induce posturanscriptional gene silencing of their target mRNAs via the RNA-induced silencing complex (RISC). These small RNAs are bond oduble-stranded. Loading of small RNA duplexes into Argonaute proteins, protein. Disease (RISC) and of the small RNAs are bond to the small RNA duplex are separated within Ago protein. Disease(edle), this unwinding process does not require ATP.

Why XTP hydrolysis is equived for RISC loading—seemingly a more binding of a small RNA duples and Ago protein—but not or unwinding a process that diepuse, 20 last go prin—has been undear We recently show that the 200 billion contains the containing a special product of RISC loading of small RNA duplexes, but not for unwinding, target cleavage, or release of the cleavage products. We envision that the chapterone machinery consumes ATP and mediates a conformational opening of top proteins so that they can receive bulky and rigid small RNA duplexes.

Will discuss such actions of the chaperone machinery consumes ATP and actions to the chaperone machinery consumes ATP and receive the such as the consumer and the co