

Cellular aging and abnormal regulation of telomeres in skin fibroblasts from a patient with an accelerated aging disorder.

Kenshi Komatsu

Radiation Biology Center, Kyoto University

Nijmegen Breakage Syndrome (NBS) is an autosomal recessive disorder characterized by chromosome instability, defect in repair of double strand break and high incidence of lymphoid cancers. Here, we show that NBS1 is recruited to telomere ends by alternative mechanism, in which NBS1 interacts with TRF2, possibly through telomeric DNA, and forms discrete foci in ATM-independent manner. This is demonstrated by the evidences that NBS1 foci are formed in ATM-defective cells, and that both NBS cells lacking FHA/BRCT domains and S278/343A clone, mutated in ATM-phosphorylated sites, can interact with TRF2. Moreover, C-terminus of NBS1, MRE11 nuclease-binding region, was essential for the interaction with TRF2. It is consistent that NBS1 lacking the C-terminus was failed to elongate the short telomere length in NBS cells, although transfection both with full length of the NBS1 cDNA and of the S278/343A cDNA restored the telomere length. Furthermore, the short G-tail length of NBS cell line has not changed significantly following transfection with NBS1 cDNA lacking of the C-terminus region, while the full length of cDNA complements the G-tail length. These findings suggest that NBS1 is recruited to telomere by MRE11/RAD50/NBS1 complex formation in ATR- and ATM-independent manner and is implicated in telomere length maintenance, possibly through G-tail regulation.