Analysis of DNA damage in the radiation resistant microbe Deinococcus radiodurans R1 exposed to space in Tanpopo mission

*矢田部 純 1 、河口 優子 1 、木下 伊織 1 、藤原 大佑 1 、青木 元秀 1 、谷口 紀恵 1 、鳴海 一成 3 、岩井 成憲 2 、山元 淳平 2 、橋本 博文 4 、横堀 伸一 1 、山岸 明彦 1

*Jun Yatabe¹, Yuko Kawaguchi¹, Iori Kinoshita¹, Daisuke Fujiwara¹, Motohide Aoki¹, Kie Taniguchi¹, Issay Narumi³, Shigenori Iwai², Junpei Yamamoto², Hirofumi Hashimoto⁴, Shin-ichi Yokobori¹, Akihiko Yamagishi¹

- 1. 東京薬科大学、2. 大阪大学、3. 東洋大学、4. 宇宙航空研究開発機構
- 1. Tokyo University of Pharmacy and Life Science, 2. Osaka University, 3. Toyo University, 4. ISAS/JAXA

Tanpopo mission is a Japanese astrobiology experiment addressing basic questions on the origin of terrestrial life and panspermia hypothesis (Yamagishi et al., 2009; Kawaguchi et al., 2016). We have started the space experiments at the Exposure Facility of the Japan Experiment Module on the International Space Station (ISS). Capture experiment investigates existence of terrestrial microbes in space. Exposure experiment investigates the microbial survival and DNA damage caused in space. We analyze degree and types of DNA damage in *Deinococcus radiodurans* using following methods: 1) comparison of survival fractions of mutant strains deficient in each of DNA repair systems, 2) analysis of DNA double-strand breaks using pulsed-field gel electrophoresis, 3) estimation of DNA damage using quantitative-PCR (q-PCR), 4) detection of mutation in *rpoB* gene and 5) analysis of DNA base damage using LC-MS/MS. In this work, we quantified DNA damage (double-strand breaks, single-strand breaks, hydrolysis of base, modified base, and so on) in part of the *rpoB* gene using q-PCR.

Methods

Dried deinococcal cell-aggregates with different thickness were exposed to space (space samples) for about one year (space samples). The cells were also stored in the ground laboratory (ground references) and in ISS cabin (ISS references). After exposure or storage, genomic DNA was extracted from each sample and an 887-bp region in the rpoB gene was amplified by q-PCR. Intact DNA (%) was determined from the quotient N/N₀, where N = copy number of rpoB gene amplified from DNA of exposed or stored cells and N₀ = copy number of rpoB gene amplified from freshly prepared DNA.

Results and Discussion

Intact DNA (%) of the cell-aggregates with 100 μ m-thickness exposed to space was less than 1% and all cells were dead. Pyrimidine dimer was major DNA damage caused by UV. On the other hand, DNA damage in those with 1000 μ m-thickness was similar between the ground references and the space samples (Fig. 1). The result indicates that UV affected only the surface of the cell-aggregates. Intact DNA (%) in the ground references and the space samples (UV> 170 nm) with 500 μ m-thickness were about 54%, and that in space samples (UV> 120 nm) with 500 μ m-thickness was 46%. Although a significant difference is not recognized between the two samples, UV with shorter wavelength tended to induce more damage in DNA. Intact DNA (%) showed a good correlation with surviving fraction. We will also report the types and degrees of DNA damage using other methods.

Yamagishi, A., et al., (2007) *Biol. Sci. Space* 21: 67–75. , Kawaguchi, Y., et al., (2016) *Astrobiology* 16: 363–376.

キーワード: パンスペルミア仮説、宇宙曝露実験、凝集体、DNA損傷、たんぽぽ計画、定量PCR Keywords: Panspermia hypothesis, Space exposure experiments, Cell aggregate, DNA damage, Tanpopo mission, Quantitative-PCR

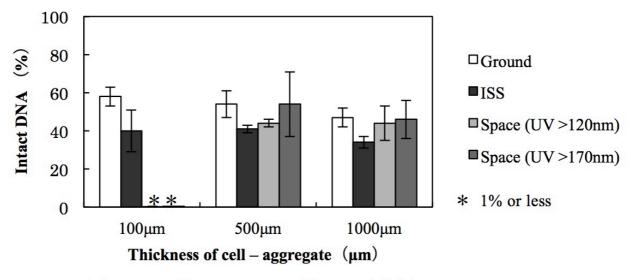


Figure 1 Percentage of Intact DNA