

JSBBA West 2nd Student Forum

要旨集正誤表 (List of Corrigenda)

O-03

誤) 【Introduction】

To address the notion on how their gut microbiota associates with their metabolic mellitus, we analyzed fecal microbiota of healthy (n=114), obese (n=49), T2D (n=47), and obese + T2D (n=40) subjects, in comparison with 317 healthy subjects from Japan, China, Indonesia, Korea, and Thailand.

【Methods and Results】

To investigate the bacterial composition of Mongolian and five Asian countries, we collected stool samples from 567 participants aged from 24 to 90 years old. The V3-V4 region of the 16S rRNA gene was amplified from stool samples and subjected to the amplicon sequencing.

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【M3thods and Results】

To investigate the bacterial composition of Mongolian and five Asian countries, we collected stool samples from 603 participants aged from 24 to 90 years old. The V3-V4 region of the 16S rRNA gene was amplified from stool samples and subjected to the amplicon sequencing.

P-09

誤)

Functional analysis of the vacuolar protein sorting gene SPBC1709.03 unique to fission yeast.

正)

Identification of a novel gene required for vacuolar protein transport in fission yeast.

P-11

誤)

Analysis of substrate specificity of α -L-arabinofuranosidase candidate enzymes in *Aspergillus nidulans*

正)

Analysis of substrate specificity of putative α -L-arabinofuranosidases in *Aspergillus nidulans*.