



## Correction to: *Glucerbacter canisensis* gen. nov., sp. nov., isolated from dog feces and its effect on the hydrolysis of plant glucosylceramide in the intestine of dogs

Misho Kawata<sup>1</sup> · Ami Tsukamoto<sup>1</sup> · Ryohei Isozaki<sup>1</sup> · Shohei Nobukawa<sup>1</sup> · Natsuki Kawahara<sup>1</sup> · Shoko Akutsu<sup>1</sup> · Masato Suzuki<sup>1</sup> · Narito Asanuma<sup>1</sup>

Published online: 7 September 2021  
© Springer-Verlag GmbH Germany, part of Springer Nature 2021

**Correction to: Archives of Microbiology (2018) 200:505–515**  
<https://doi.org/10.1007/s00203-017-1463-1>

There are few corrections in the original publication. The detail is given below.

Original P.505 L.1 (In title)

*Glucerbacter canisensis*

Correction

*Glucerberacter canis*

Original P.505 L.15 (In Abstract)

*Glucerbacter canisensis*

Correction

*Glucerberacter canis*

Original P.505 L.20 (In Keywords)

*Glucerbacter canisensis*

Correction

*Glucerberacter canis*

Original P.505 Left paragraph L.33

*Glucerbacter canisensis*

Correction

*Glucerberacter canis*

Original P.506 Left paragraph L.27

*Glucerbacter canisensis*

Correction

*Glucerberacter canis*

Original P.512 Left paragraph L.4

*Glucerbacter canisensis*

Correction

*Glucerberacter canis*

Original P.513 Right paragraph L.1

### Description of *Glucerbacter* gen. nov.

*Glucerbacter* (glu.cera.bac'ter. N.L. gen. n. glucera, Glu-Cer; N.L. masc. n. bacter a rod; N.L. masc. n. *Glucerbacter*, a GluCer-utilizing rod).

Correction

The original article can be found online at <https://doi.org/10.1007/s00203-017-1463-1>.

✉ Narito Asanuma  
asanuma@meiji.ac.jp

<sup>1</sup> Department of Life Science, Meiji University, Higashimita, Tama-ku, Kawasaki 214-8571, Japan

**Description of *Glucerbacter* gen. nov.**

*Glucerbacter* (Glu.ce.ri.bac.ter'. N.L. neut. n. glucerum GluCer; N.L. masc. n. bacter a rod; N.L. masc. n. *Glucerbacter*, a GluCer-utilizing rod).

Original P.1513 Right paragraph L.11

The genus is a member of the family *Lachnospiraceae*.

Correction

The genus is a member of the family *Lachnospiraceae*. The type species is "*Glucerbacter canis*".

Original P.513 Right paragraph L.12

**Description of *Glucerbacter canisensis* sp. nov.**

*Glucerbacter canisensis* (canis.ensis. N. masc. canis, dog; L. masc. ensis, from; N.L. masc. n. *canisensis*, from a dog)

Correction

**Description of *Glucerbacter canis* sp. nov.**

*Glucerbacter canis* (ca'nis. L. gen. n. canis of a dog)

Original P.514 Left paragraph L.2

Whole-cell fatty acids are C<sub>12:0</sub>, C<sub>14:0</sub>, C<sub>14:0</sub> DMA, unknown fatty acid with equivalent length 14.762/C<sub>15:2</sub>/C<sub>15:1</sub>, C<sub>16:0</sub> aldehyde, C<sub>16:1</sub> w7c, C<sub>16:0</sub>, C<sub>16:1</sub> w7c DMA, C<sub>16:0</sub> DMA, C<sub>17:1</sub> w8c/C<sub>17:2</sub>, C<sub>18:1</sub> w9c, C<sub>18:1</sub> w7c/unknown fatty acid with equivalent length 17.834, C<sub>18:0</sub>, C<sub>18:1</sub> w9c DMA, C<sub>18:1</sub> w7c DMA, and C<sub>18:0</sub> DMA (representing more than 1.0%).

Correction

Whole-cell fatty acids are C<sub>12:0</sub>, C<sub>14:0</sub>, C<sub>14:0</sub> DMA, unknown fatty acid with equivalent length 14.762/C<sub>15:2</sub>/C<sub>15:1</sub>, C<sub>16:0</sub> aldehyde, C<sub>16:1</sub> ω7c, C<sub>16:0</sub>, C<sub>16:1</sub> ω7c DMA, C<sub>16:0</sub> DMA, C<sub>17:1</sub> ω8c/C<sub>17:2</sub>, C<sub>18:1</sub> ω9c, C<sub>18:1</sub> ω7c/unknown fatty acid with equivalent length 17.834, C<sub>18:0</sub>, C<sub>18:1</sub> ω9c DMA, C<sub>18:1</sub> ω7c DMA, and C<sub>18:0</sub> DMA (representing more than 1.0%).

Original P.514 Left paragraph L.9

JCM31739<sup>T</sup>

Correction

JCM 31739<sup>T</sup>

Original Table 2

Line 4, Column 2 (Line; Spore formation, Column; Characteristic 1)

+

Correction

Line 4, Column 2 (Line; Spore formation, Column; Characteristic 1)

–

**Publisher's Note** Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.