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OPEN Author Correction: A novel Asfarvirus-like virus identified as a potential cause of mass mortality of abalone

Tomomasa Matsuyama, Tomokazu Takano, Issei Nishiki, Atushi Fujiwara, Ikunari Kiryu, Mari linada, Takamitsu Sakai, Sachiko Terashima, Yuta Matsuura, Kiyoshi Isowa & Chihaya Nakayasu

Correction to: Scientific Reports https://doi.org/10.1038/s41598-020-61492-3, published online 12 March 2020

This Article contains typographical errors.

In the Results under subheading 'Gap closure and genome analysis',

"A draft genome sequence consisting of 1,551,181 bp with a GC content of 31.7% was obtained by the assembly of the NGS sequence and Sanger sequencing of the 9 gap regions."

should read:

"A draft genome sequence consisting of 155,181 bp with a GC content of 31.7% was obtained by the assembly of the NGS sequence and Sanger sequencing of the 9 gap regions."

In the Discussion,

"The gap-closed sequence consisted of 1,551,181 bp with a GC content of 31.7% and 159 predicted ORFs."

should read:

"The gap-closed sequence consisted of 155,181 bp with a GC content of 31.7% and 159 predicted ORFs."

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