



유사도 확인 프로그램 turn it in Self-Check 가이드

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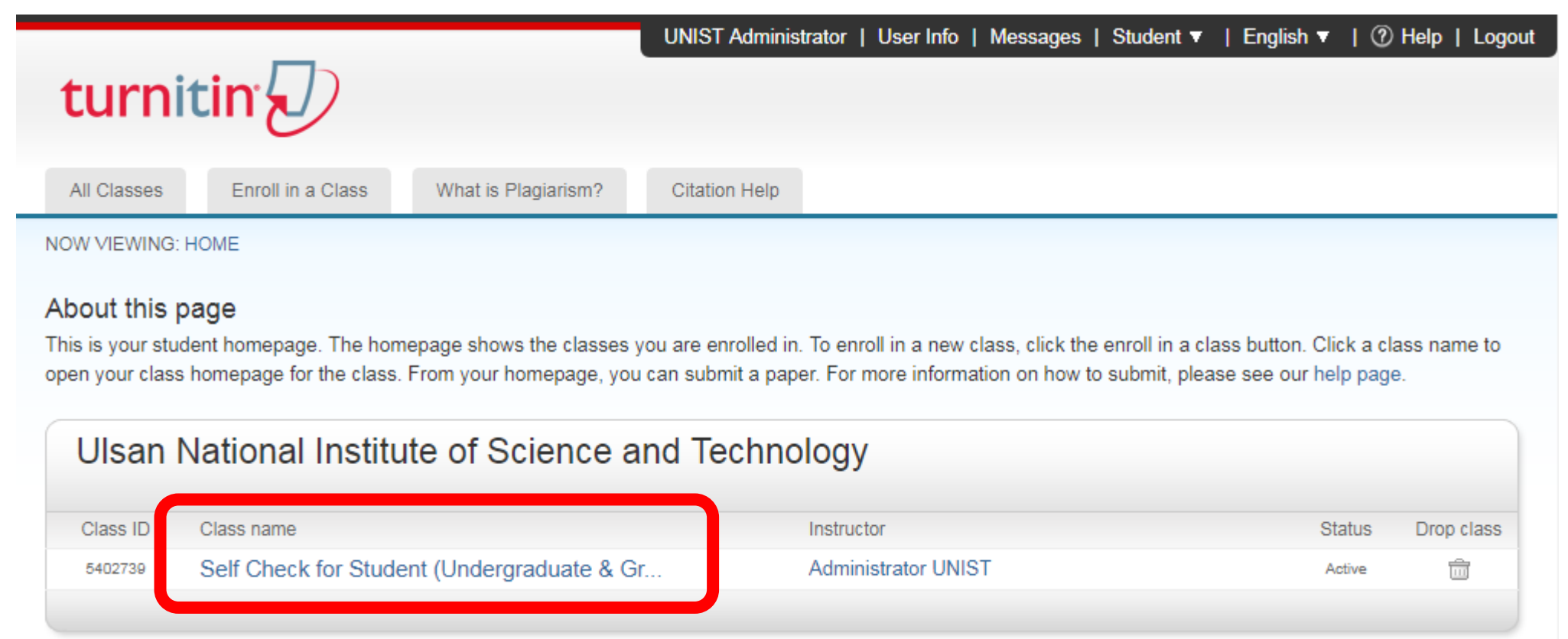
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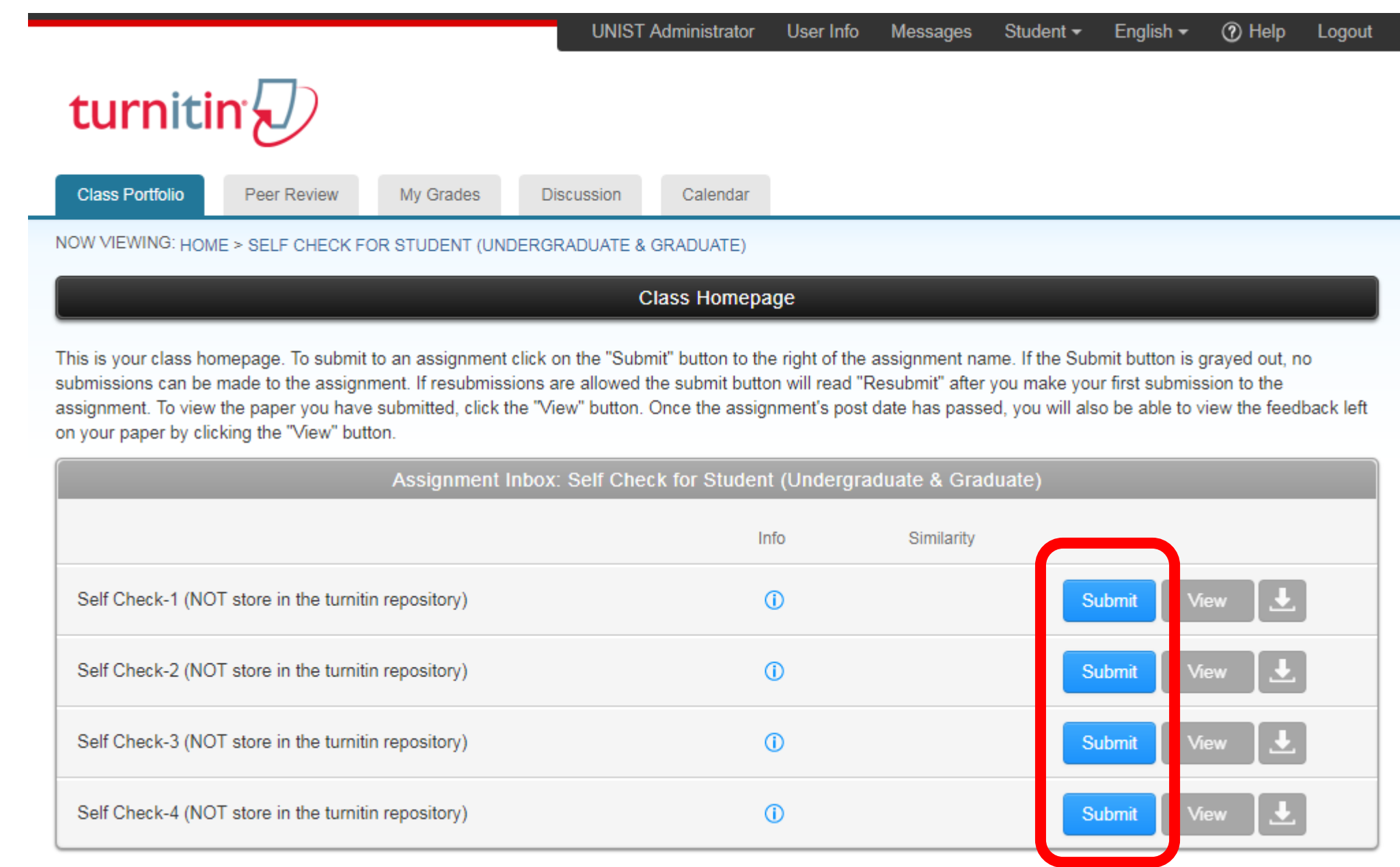
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1 Youngjune Bhak, Yeonsu Jeon, Sungwon Jeon, Oksung Chung et al. 20% >
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2 phys.org 7% >
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PRIMARY SOURCES

1 Youngjune Bhak, Yeonsu Jeon, Sungwon Jeon, Oksung Chung et al. "Myotis rufoniger genome sequence and analyses: M. rufoniger's genomic feature and the decreasing effective population size of Myotis bats", PLOS ONE, 2017 20%
Publication

<유사도 검사 결과 예시>

자동 영문법 검사

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UNIST Reveals the Whole Genome Sequences of Rare Red Bat

Their findings appeared in the July issue of the world's largest scholarly journal, PLoS ONE.

A recent study, affiliated with the Korean Genomics Industrialization and Commercialization Center (KOGIC) at UNIST has presented the first whole genome sequence and analyses of the Myotis rufoniger, one of the most well-known and iconic protected wild animals in South Korea, known as the Sp. den bat.

This breakthrough comes from a research, conducted by Professor Jong Bhak of Life Science at UNIST and Professor Doug-Young Ryu of Veterinary Medicine at Seoul National University in collaboration with the Korean Cultural Heritage Administration.

Recent studies have indicated that bats live longer than any other mammals of their sizes on earth. Myotis rufoniger is a species of vesper bat in the family Vespertilionidae. It is a rare bat species that face imminent threat of disappearance on the face of Earth. Being designated as a Korean natural monument, only 450 these bats survive in the wild in South Korea, presently. The research team expects that this study will provide a genetic foundation for the restoration and conservation of the critically endangered M. rufoniger.

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Bats are typically brown or black in color, but they also occur in a variety of color schemes. In the study, the research team found specific genetic variations that are likely responsible for the M. rufoniger's rusty orange fur color, which distinguish it from the other bats. Moreover, they also found an elemental analysis in the tissues from the M. rufoniger individual analyzed also showed a very high concentration of (As) in its intestinal tissue. This suggests an evolutionary correlation that M. rufoniger can survive in a cave, contained a high level of As.

A genome contains all of the genetic information of a given organism, including its evolutionary origins. The demographic history analysis in the present study found that the population size of the M. rufoniger was dramatically decreased during the latter part of the last glacial period. It is also shown there was a consistent decline of Myotis bat family's

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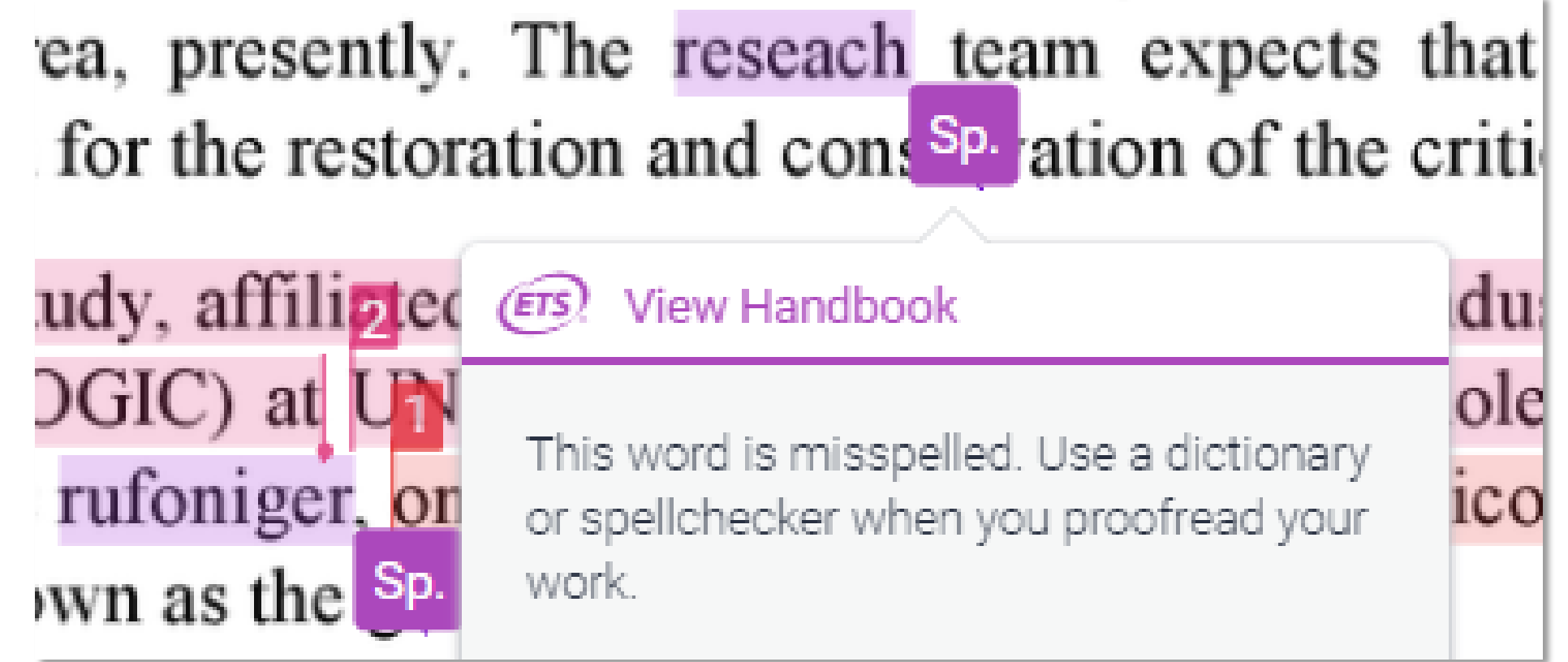
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- ※ 최대 64,000글자까지 검사 가능
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