



유사도 확인 프로그램 turn it in Self-Check 가이드 (학생용)

(2024.03.)

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The first of these, Hong Kong Martial Arts Living Archive (HKMALA), was instigated in 2012 and is an ongoing research collaboration between the International Guoshu Association, City University of Hong Kong, and the Laboratory for Experimental Museology (eM+) at EPFL, and has resulted in seven international exhibitions, including *Kung Fu Motion* at EPFL's ArtLab (2018)¹ and the Immigration Museum Melbourne in 2017, and *300 Years of Hakka Kung Fu* (2016)² at the Heritage Museum and CityU Gallery³, Hong Kong, China. The archiving project responds to the decline of Southern Chinese Kung Fu in mainland China, where a significant portion of traditional martial arts have already vanished. Hong Kong remains a vibrant center for elite practitioners and is home to some of the most prominent martial artists in the world; however rapid urban development, population growth, cultural transformation and the aging of the masters are endangering these practices.

HKMALA brings together historical materials with creative visualizations derived from advanced documentation processes, including motion capture, motion-over-time analytics, 3D reconstruction, and panoramic video (Figures 1 and 2). These archival materials are re-interpreted and re-performed through the mediums of augmented virtual reality and interactive media art, such as *Kung Fu Visualization*³ (2016). As a panoptic virtual reality environment, the Re-ACTOR system shows the intricate dynamics of the kung fu master's reenacted performances via serial 3D motion-captures from six different points of view, with an interactive control panel that allows visitors to select six different visualization styles that elucidate the underlying dynamics of the master's movements (Figures 3 and 4).

The HKMALA 'living archive' also uses new immersive and interactive display paradigms to perpetuate the performance of past masters for future generations. The *Kung Fu Weapons Archive*⁴ (2016) is a linear navigator that provides a sliding panorama of Hakka Kung Fu weapons and training tools, as well as interactively located video demonstrations of their use by Kung Fu masters. Whenever the viewer slides the screen over one particular object, it triggers a short video clip showing the Kung Fu master's handling of that respective weapon or training tool (Figure 5). With these new approaches HKMALA creates practical strategies for encoding, retrieving, and reenacting

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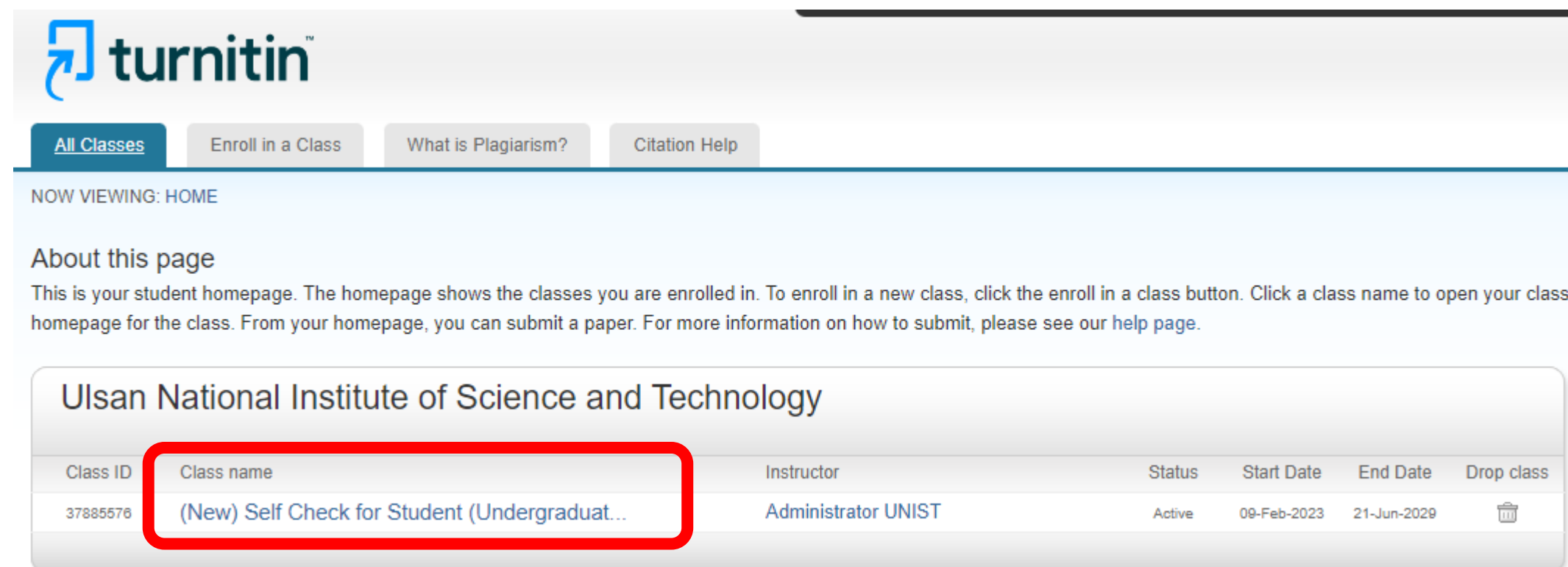
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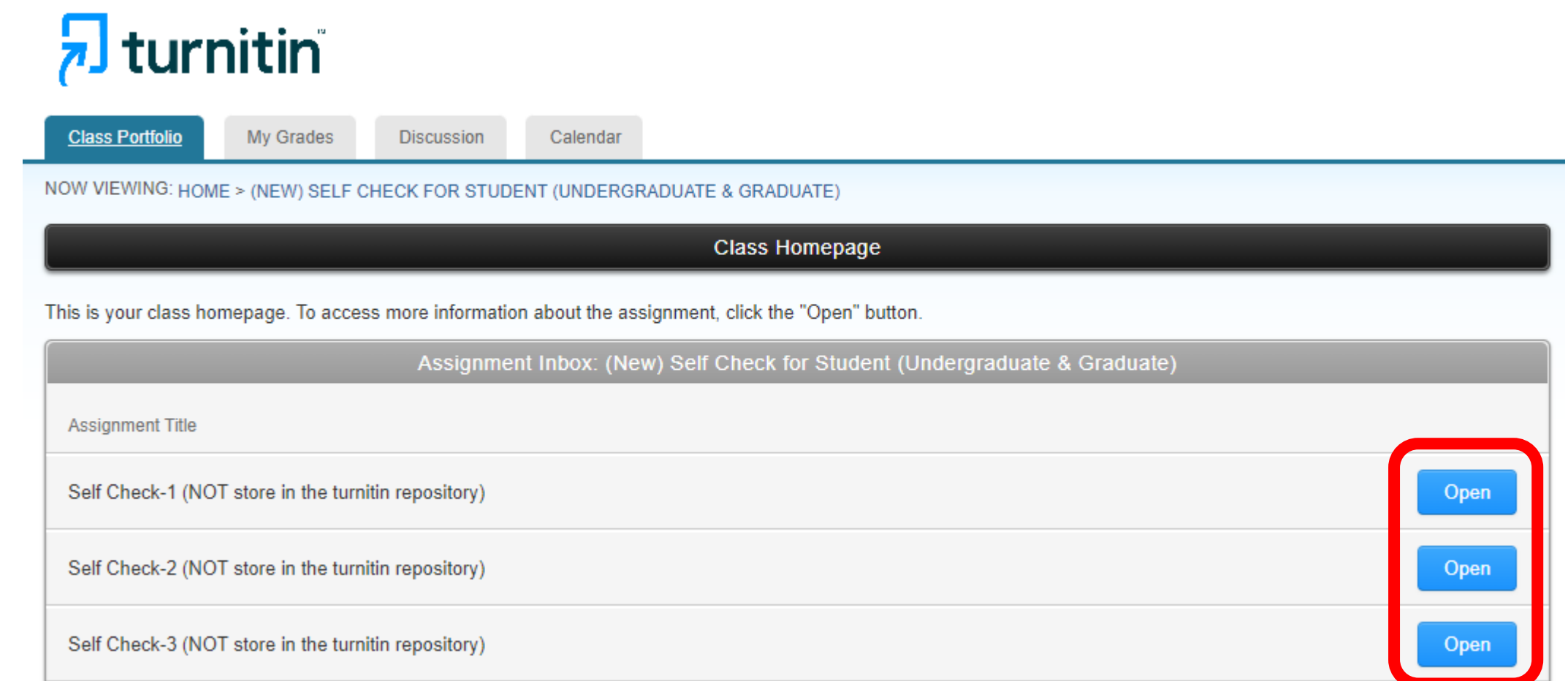
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UNIST Reveals the Whole Genome Sequences of Rare Red Bat

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 from 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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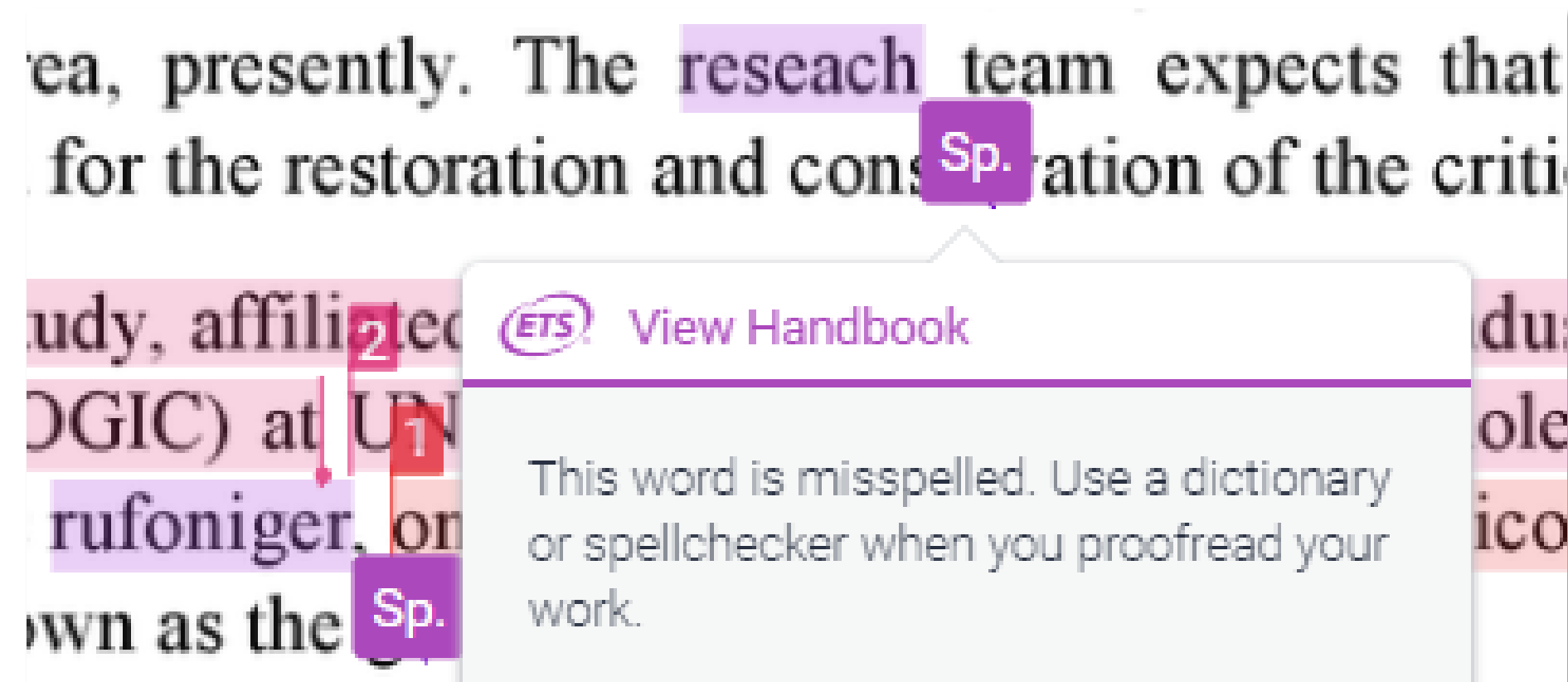
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