



Similarity Checker 'turn it in' Guide for self-checking (Students)

(Mar. 2024)

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UNIST Library

Turn it in ?

- www.turnitin.com
- Turn it in is a **similarity checking program** to **check originality** of students' works and **prevent plagiarism**.
- It **compares** papers against Internet pages, subscription repository of periodicals, journals, publications and repository of previously submitted papers.
- Related information and guides
: UNIST Library → Research Supports
→ [Plagiarism Checking \(Turn it in\)](#)

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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28%		
1개(총 1개 중) 일치		
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※ Turn it in Student quick start guide (Eng)
: <https://help.turnitin.com/feedback-studio/turnitin-website/student/quickstart.htm>

Create an Account: **Self-Checking**



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- ① Turn it in (www.turnitin.com) → Log In → New user?
- ② Create a User Profile: [Student](#)
- ③ Enter a given **Class ID** and your information

Create a New Student Profile

Class ID Information

All students must be enrolled in an active class. To enroll in a class, please enter the class ID number and class enrollment key that you were given by your instructor.

Please note that the key and pincode are case-sensitive. If you do not have this information, or the information you are entering appears to be incorrect, please contact your instructor.

Class ID

Class enrollment key

✂ Class ID & enrollment key are available on the library website [‘Plagiarism Checking \(Turn it in\)’](#) menu. (Log in is required)

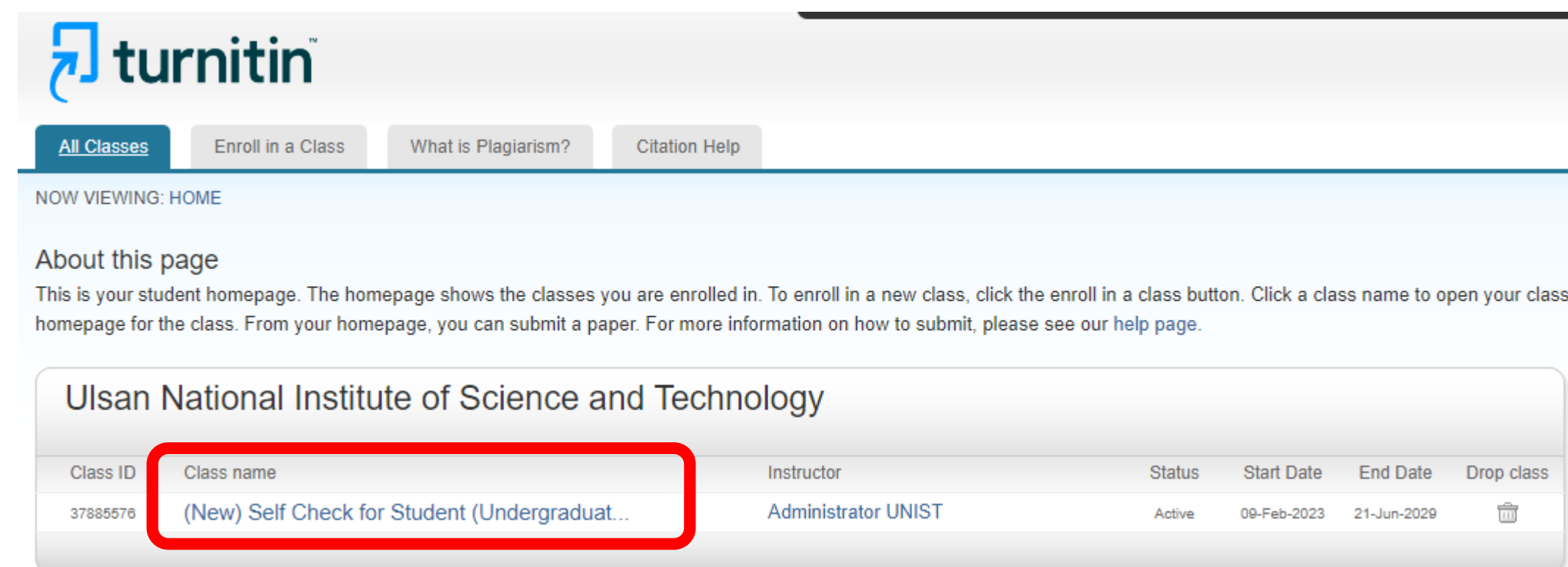
✂ e-mail Address (= account): Enter your UNIST e-mail

✂ Only UNIST members can use this service. Invalid accounts such as graduates and resignees will be deleted regularly.

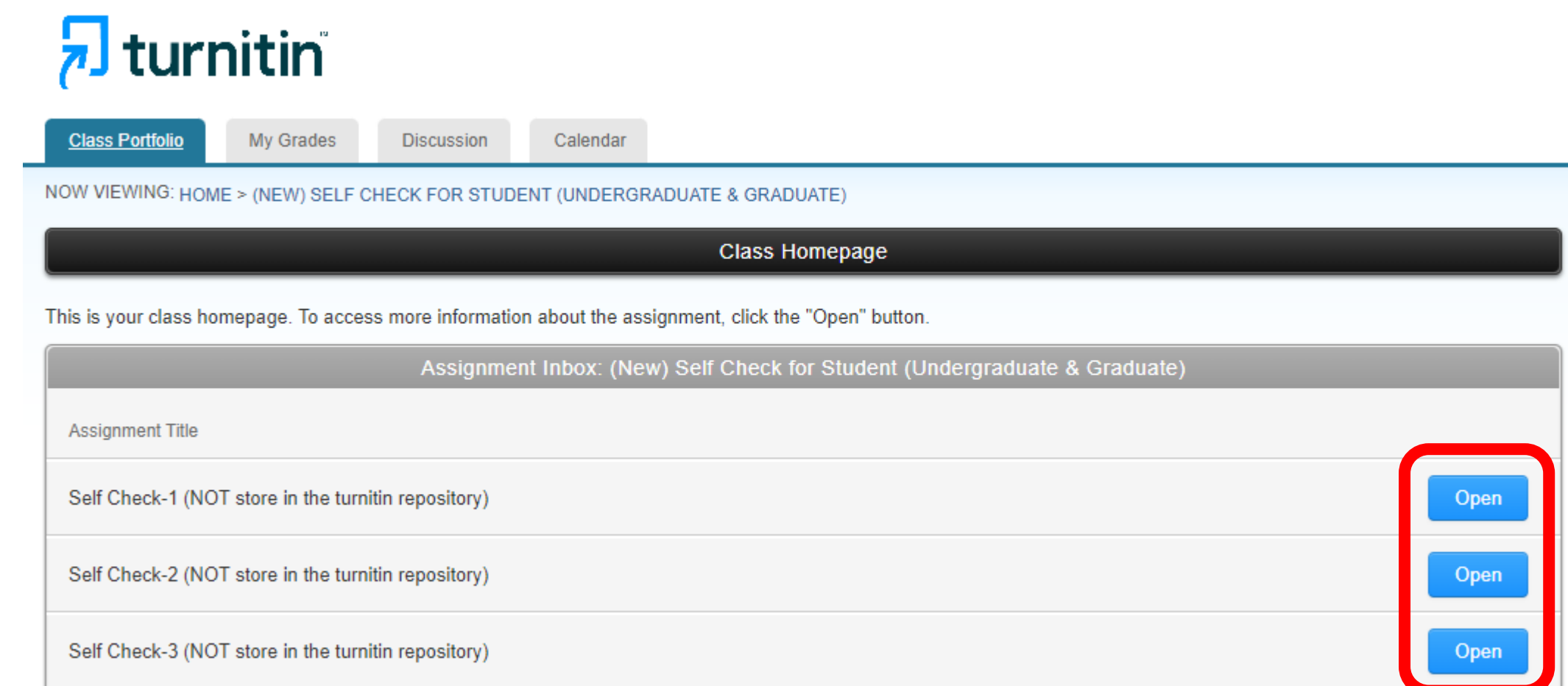
✂ Account for instructor role: Contact the library (ext. 1405)

Submitting a Paper

① Select a class for self-check



② Click the **Open** button to submit a file



※ 10 Self-Check menus are available to submit files. Click on any 'open' button to check the revised or new paper's originality index.

※ All files for self-check will not be stored in the Turn it in student paper repository. It means your paper will not be checked against other students' submissions.

Submitting a Paper

③ Submit your paper: Upload Submission

- Types of submission: 1) Upload Submission
2) Text Input 3) Cloud Submission

- Submission File: Select a file → Upload and Review

※ File types allowed: Text, MS Word, MS PowerPoint, PostScript, PDF, RTF, HTML, WordPerfect, Hangul (HWP), OpenOffice, Excel

※ The file size may not exceed 100 MB. Files of larger size may be reduced in size by removal of non-text content. Files that are password protected, encrypted, hidden, system files, or read only files cannot be uploaded or submitted to Turnitin.

The screenshot shows the Turnitin interface. At the top, there's a navigation bar with 'Class Portfolio', 'My Grades', 'Discussion', and 'Calendar'. Below it, a breadcrumb trail reads: 'NOW VIEWING: HOME > (NEW) SELF CHECK FOR STUDENT (UNDERGRADUATE & GRADUATE) > SELF CHECK-8 (NOT STORE IN THE TURNITIN REPOSITORY)'. The main heading is 'About this page', followed by a description: 'This is your assignment dashboard. You can upload submissions for your assignment from here. When a submission has been processed you will be able to download a digital receipt, view any grades and similarity reports that have been made available by your instructor.' Below this is a link: '> Self Check-8 (NOT store in the turnitin repository) ?'. A red box highlights the 'Upload Submission' button. Below the button is a 'Submit File' modal. The modal has a progress bar with three stages: 'Upload', 'Review', and 'Complete'. Below the progress bar, there are three options: 'Upload Submission' (highlighted with a red box), 'Text Input', and 'Cloud Submission'. To the right of these options is the text 'Types of submission'. Below the options is a light blue box with the text 'Drag and drop or select a file from your device.' Below that is a 'Submission Title' field with the text 'Untitled'. Below the title field is a 'Submission File' field with a red box around it and the text '파일 선택' (File Select). At the bottom right of the modal is a red box around the 'Upload and Review' button.

Submitting a Paper

- ④ Review the submission
→ **Submit to Turnitin**

Submit File

Upload Review Complete

Title

File Size
13.45 KiB

Word Count
505

Deoxyribonucleic acid (DNA) stands as one of the most pivotal molecules in the realm of life sciences. Its discovery and subsequent understanding have revolutionized the fields of genetics, molecular biology, and biotechnology, unraveling the intricate mechanisms underlying the inheritance and expression of genetic information. DNA serves as the molecular blueprint of life, carrying the genetic instructions necessary for the growth, development, and reproduction of all living organisms.

The exploration of DNA structure and function has been a remarkable journey marked by key milestones and groundbreaking discoveries. From the early work of pioneers such as Friedrich Miescher and Rosalind Franklin to the transformative research of James Watson, Francis Crick, and Maurice Wilkins, the elucidation of the double helix structure and the identification of the base-pairing mechanism have provided the foundation for our understanding of DNA.

The chemical composition of DNA is intricately woven with its structural integrity. Comprised of nucleotides, DNA consists of a sugar molecule called deoxyribose, a phosphate group, and one of four nitrogenous bases—adenine (A), cytosine (C), guanine (G), and thymine (T). The base-pairing rules, A pairs with T, and C pairs with G, forming the double helix.

Page 1 of 2

Preview Submission Cancel Submission **Submit to Turnitin**

- ⑤ View a similarity report

> Self Check-8 (NOT store in the turnitin repository) ?

Paper Title	Uploaded	Grade	Similarity	
Similarity checking	07 Mar 2024 11:25	--	31%	<div> <div>Resubmit paper</div> </div>

- **Result: Percentage of similarity**

✂ A first similarity reports is usually available in 5~10 minutes, it depends on the length of the loaded paper. If you can not get a result report, please contact the library.

- ✂ **Resubmit paper**

- You can upload a paper up to three more times consecutively.
- If you resubmit it for the fourth time or more, it will take 24 hours to generate the results. If you wish to view the results immediately after using the same button 3 more times, you can click on another 'open' button on the previous screen to submit your paper. (Note: Each individual can submit up to a maximum of 50 times using one 'open' button.)

Viewing Originality Reports

feedback studio UNIST Administrator Originality check_test

Open layers

Match Overview

27%

27

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% >

2 phys.org Internet Source 7% >

Similarity results

Grammar checking

Download results

Download

Current View

Digital Receipt

Originally Submitted File

Page: 1 of 2 Word Count: 731

- **Similarity index (%)**
: The percentage of the text in your paper that matched sources in turn it in database.
- **Result paper: Download → Current View**
✕ The result pages are on the last section of the downloaded PDF file.

Originality check_test

ORIGINALITY REPORT

27%	7%	22%	0%
SIMILARITY INDEX	INTERNET SOURCES	PUBLICATIONS	STUDENT PAPERS

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	

Example
: Similarity report

Grammar Check

The screenshot shows the UNIST feedback studio interface. The main document area displays a text snippet about the whole genome sequences of a rare red bat. The text contains several grammar and spelling errors highlighted in red. A green callout bubble labeled "Grammar checking" points to the text. On the right side, the "e-rater® Results" panel is open, showing a list of errors and their counts. The "Style" section has 0 errors, the "Usage" section has 7 errors, and the "Spelling" section has 13 errors. A yellow callout bubble labeled "Open layers" points to the "e-rater® Results" panel. The bottom status bar shows "Page: 1 of 2" and "Word Count: 731".

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.

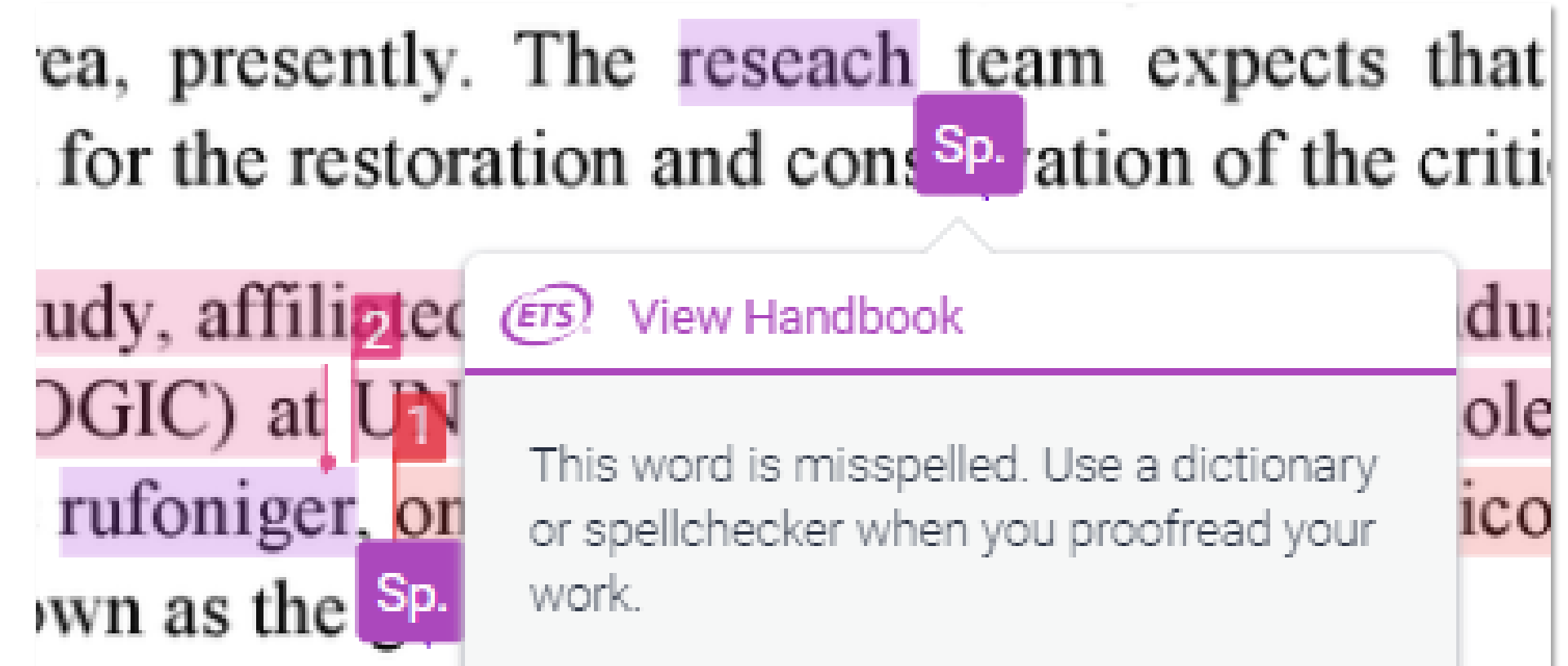
A genome contains all of the genetic information of a given organism, including its evolutionary origins. The demographic history analysis in the present study found that the population size of the *M. rufoniger* was dramatically decreased during the latter part of the last glacial period. It is also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

e-rater® Results

Category	Count
Missing " "	0
Missing "?"	0
Missing Apos.	0
Proper Nouns	0
Dup.	0
Compound	0
Hyph.	0
Fused	0
Style	0
Tone	0
Coord. Conjunction	0
P/V	0
Long	0
Short	0
Usage	7
Negation	0
Nonstandard	0
Confused	0
Article Error	7
Faulty Comparison	0
Wrong Article	0
Prep.	0
Wrong Form	0
Spelling	13
Sp.	13

Page: 1 of 2 Word Count: 731

- **e-rater Grammar Check** (Developed by ETS)
: It automatically checks submissions for grammar, usage, mechanics, style and spelling errors. Results are available on the 'e-rater' tab on the originality report.



- ✂ Maximum 64,000 characters are available to check automatically.
- ✂ e-rater settings: Advanced level, US & UK English Dictionaries



Question?

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