



# 유사도 확인 프로그램 turn it in Self-Check 가이드

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# Turn it in 이란?

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- 저작물의 **독창성**을 확인하고 **표절을 예방**하기 위한 수단으로,
- 각종 학술자료 및 웹 자료와 비교하여 **저작물의 유사도를 확인**할 수 있는 프로그램
- 유사도 검사 대상 자료  
: 학술지 논문, 도서, 백과사전 등 참고자료, 웹 자료, turn it in에 제출된 학생 과제 등
- 관련 정보 및 가이드  
: UNIST Library  
→ 연구 & 학습 지원 (Research Supports)  
→ [표절예방 \(Plagiarism Prevention Tool\)](#)



Turn it in Quick Start Training Video (English)

<https://youtu.be/AC3GB-FOMvY>

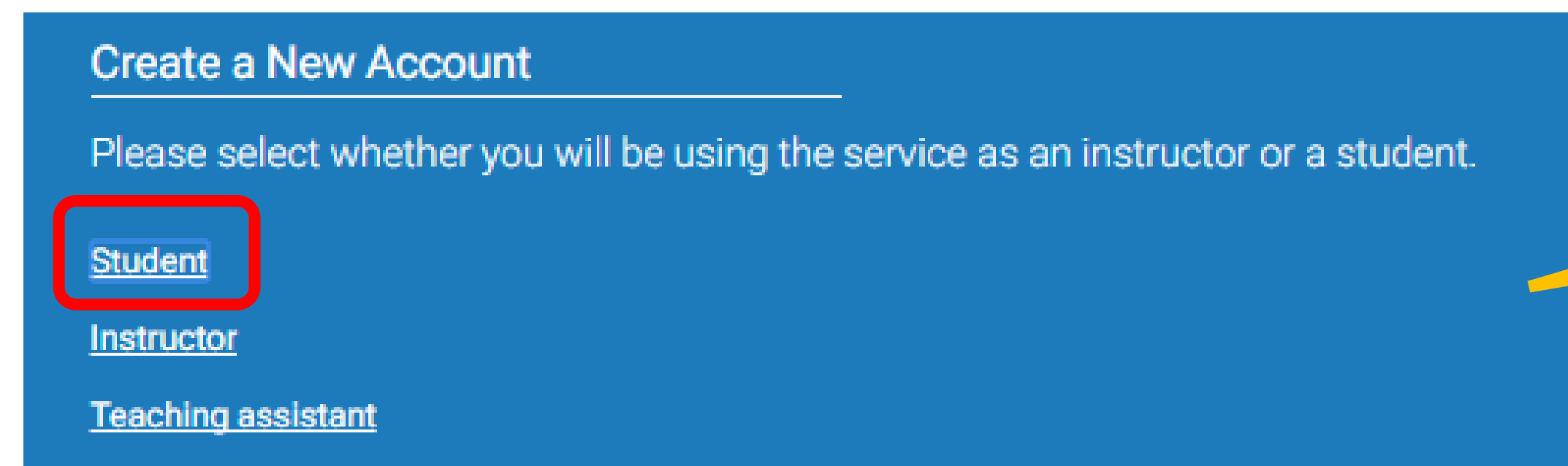
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- ① Turn it in 접속 ([www.turnitin.com](http://www.turnitin.com)) → [Create Account](#)
- ② Create a New Account → [Student](#) 선택
- ③ 계정 생성을 위한 **Class ID** 및 이용자 정보 입력

※ **Class ID 및 Enrollment key:** 도서관 웹사이트 > 연구&학습지원 > **‘표절 예방’** 메뉴에서 확인 (웹사이트 로그인 시 확인 가능)

※ **e-mail Address:** UNIST e-mail 계정 입력

※ 강사용 계정: 도서관 문의 (내선 1405, kyl7539@unist.ac.kr)



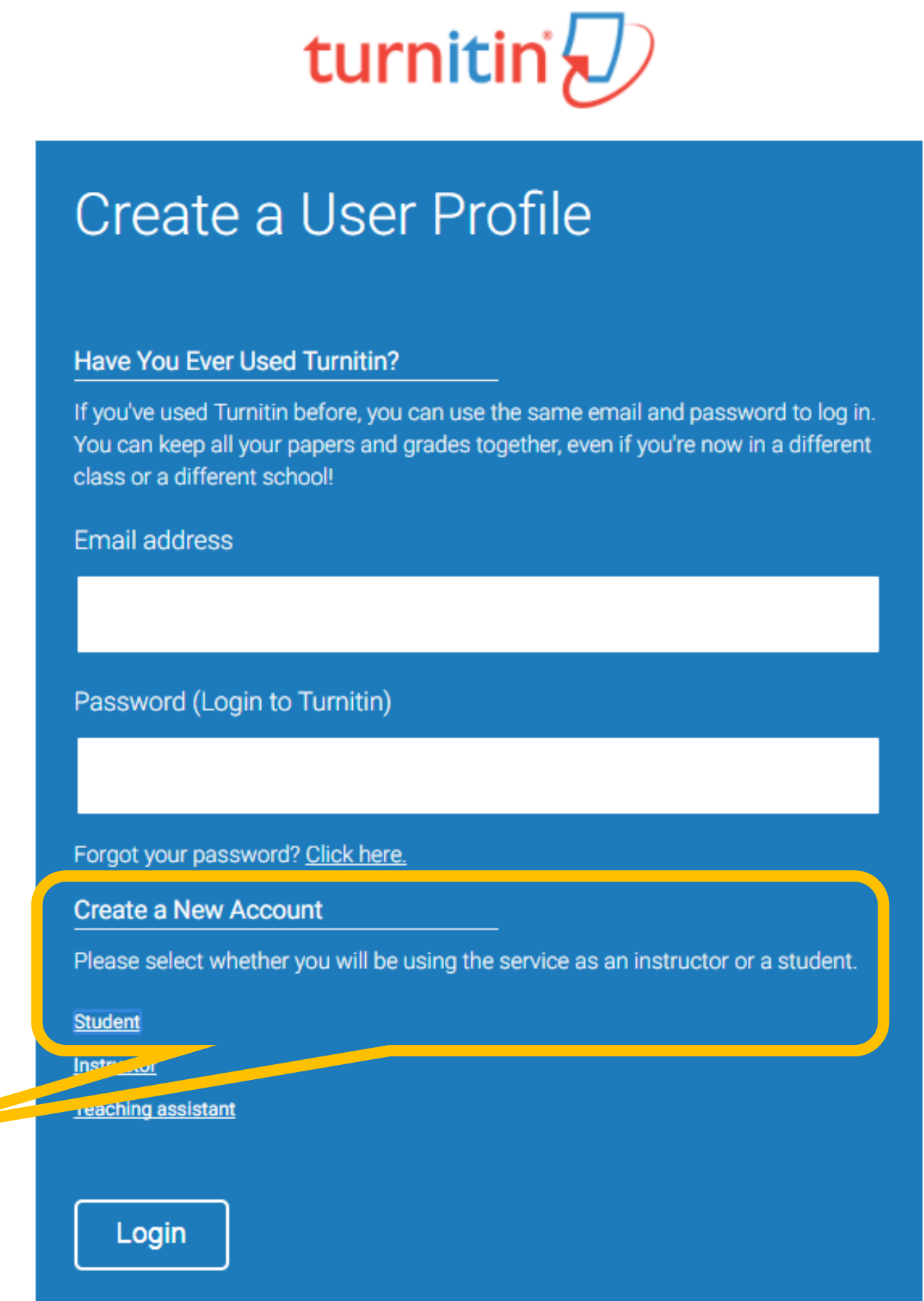
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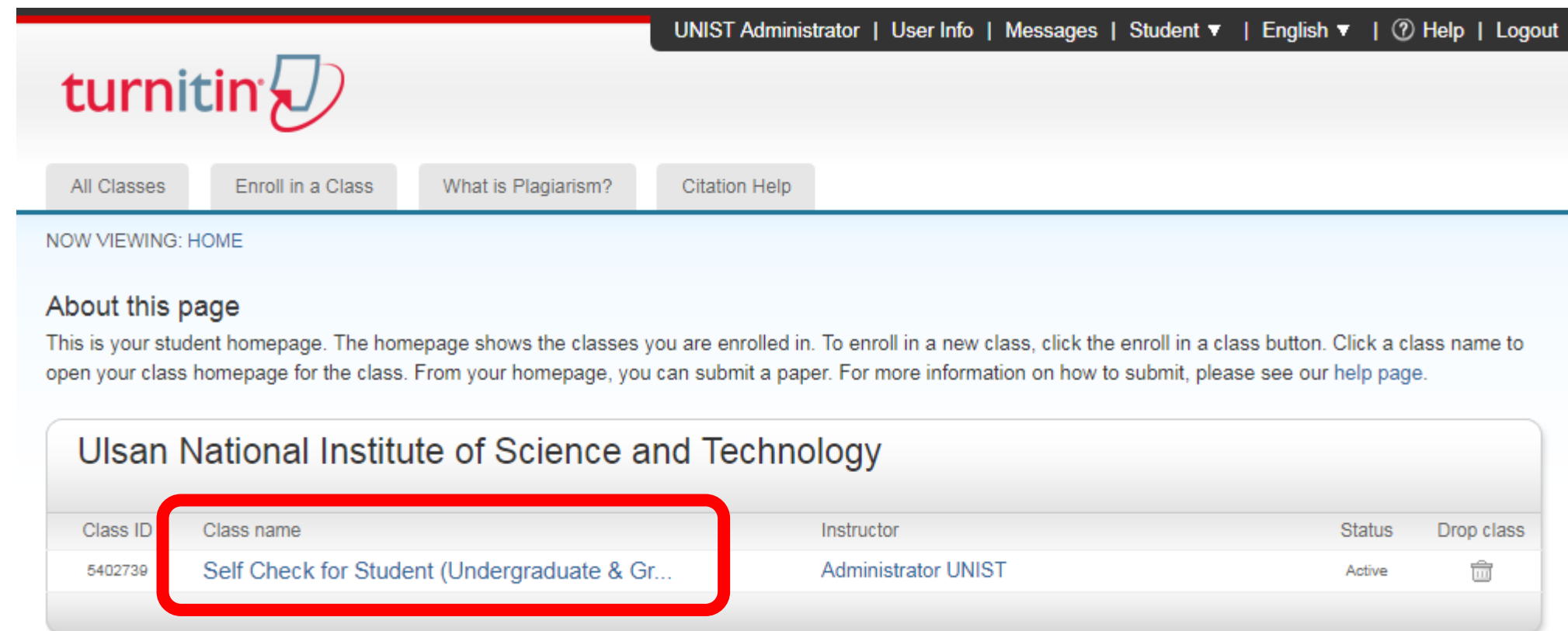
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# 유사도 검사를 위한 파일 제출

## ① Self-Check용 Class 선택



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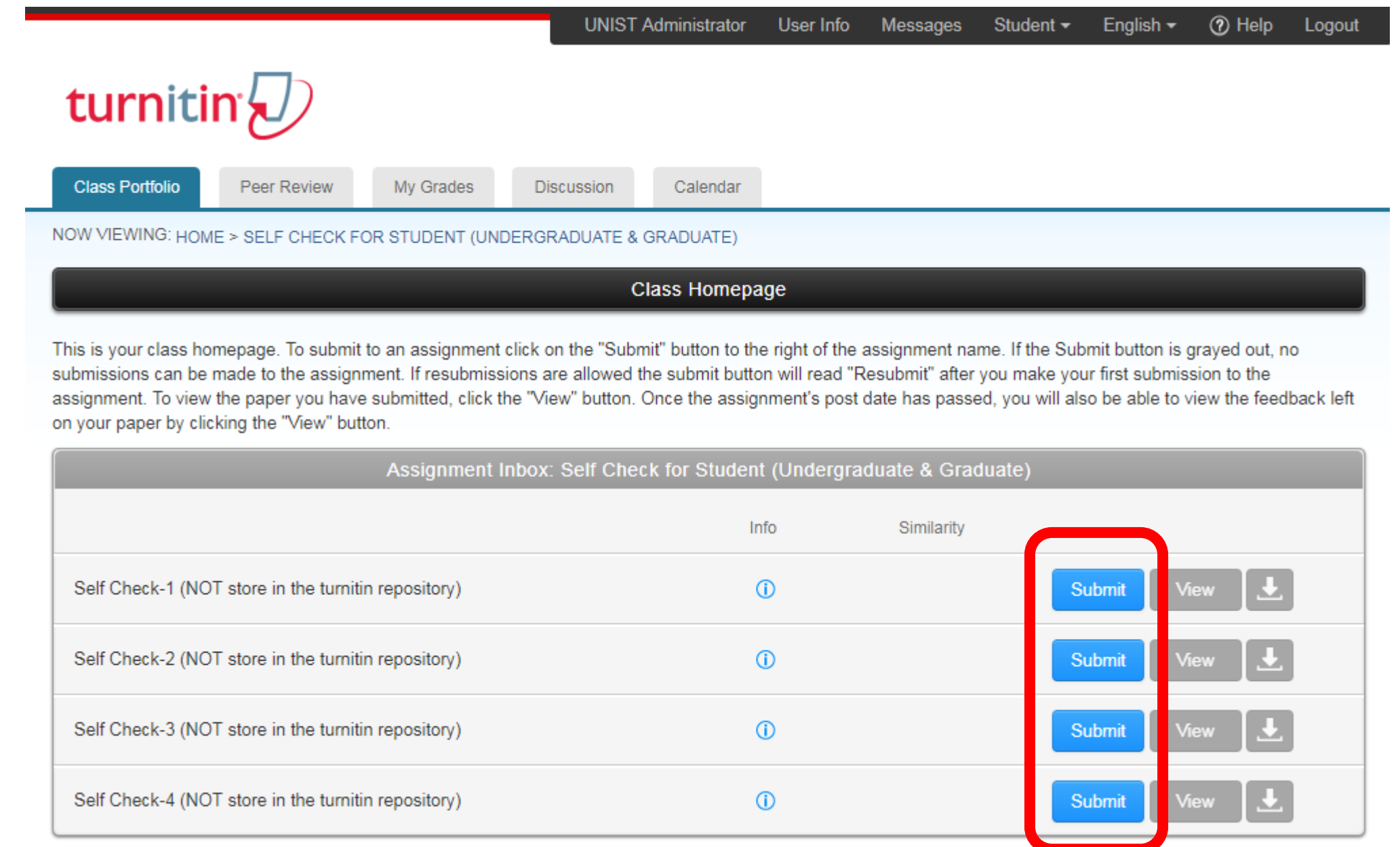
NOW VIEWING: HOME

About this page  
This is your student homepage. The homepage shows the classes you are enrolled in. To enroll in a new class, click the enroll in a class button. Click a class name to open your class homepage for the class. From your homepage, you can submit a paper. For more information on how to submit, please see our help page.

Ulsan National Institute of Science and Technology

Class ID	Class name	Instructor	Status	Drop class
5402730	Self Check for Student (Undergraduate & Gr...	Administrator UNIST	Active	

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Assignment Inbox: Self Check for Student (Undergraduate & Graduate)		
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# 유사도 검사를 위한 파일 제출

## ③ 검사 대상 파일 제출

- Submission title: 유사도 검사명 (자유 기입)
- 검사 대상 조건  
: 40MB 이하, 20단어 이상 수록, 최대 400페이지
- 등록 가능 파일 유형  
: **Microsoft Word**, Excel, PowerPoint, WordPerfect, PostScript, **PDF**, HTML, RTF, OpenOffice (ODT), 한글 (HWP), Google Docs, and plain text

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Submission title:  
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File name:  
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File size:  
14.81K

Page count:  
2

Word count:  
731

Character count:  
3948

«Page 1»

UNIST Reveals the Whole Genome Sequences of H5N1 Bird Flu

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

A recent study, affiliated with the Korean Genomics Information and Communication Center (KGIACC) at UNIST, has presented the first whole genome sequence and analysis of the H5N1 influenza virus, one of the most well-known and widely spread viral diseases in South Korea, known as the avian flu.

The study, which was conducted by Professor Jong-Wook Kim of the School of Life Science at UNIST and Professor Dong-Wook Kim of the School of Medicine at Seoul National University, is published in the journal PLoS ONE.

Recent studies have indicated that the H5N1 virus has a higher than any other members of their class in South Korea, and is a source of concern for the health of humans. It is a new type of virus that has emerged from a combination of the H5N1 virus, which was first discovered in 1997, and the H5N1 virus, which was first discovered in 2003. The study also revealed that the H5N1 virus is a new type of virus, which is a combination of the H5N1 virus and the H5N1 virus.

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Assignment Inbox: Self Check for Student (Undergraduate & Graduate)

	Info	Similarity	
Self Check-1 (NOT store in the turnitin repository)	🔍	27%	<div>ResubmitView📄</div>

- **Similarity: 유사도 검사 결과**
- ※ 첫 제출 시 약 5~10분 후 결과 확인 가능, 저작물에 따라 다름
- ※ 첫 제출 후 24시간 이내에는 최대 3회까지 재 제출건의 유사도 검사 결과를 즉시 확인 가능, 그 이후 재 제출 시 결과는 24시간 후 확인 가능
- ※ 장시간 경과 후에도 검사 결과가 미 않을 경우 문헌정보팀 문의 (파일 오류, 삭제 후 재 제출 필요)



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# 유사도 검사 결과 확인

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Match Overview

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% >

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검사 결과 출력

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Page: 1 of 2 Word Count: 731

## 유사도 검사 결과(%)

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전체 단어를 100단어로 가정 시 일치 정도

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ORIGINALITY REPORT

27% 7% 22% 0%

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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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**e-rater® Results**

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

**영문법 검사 결과**

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 of 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

Page: 1 of 2 Word Count: 731

- **e-rater Grammar Check**  
: ETS 제공 자동 영문법 검사 시스템,  
문법, 구조 규칙, 문체, 맞춤법 등 검사  
turn it in 유사도 검사 결과에서 확인 가능

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**ETS View Handbook**

This word is misspelled. Use a dictionary or spellchecker when you proofread your work.

- ※ '영어'로만 작성된 파일만 검사 가능, 한/영 혼용 시 검사 불가
- ※ 최대 64,000글자까지 검사 가능
- ※ 영문법 검사 환경: Advanced level (최고 수준),  
US & UK English Dictionaries





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