

Guideline

Released: Tue, May 31, 2022

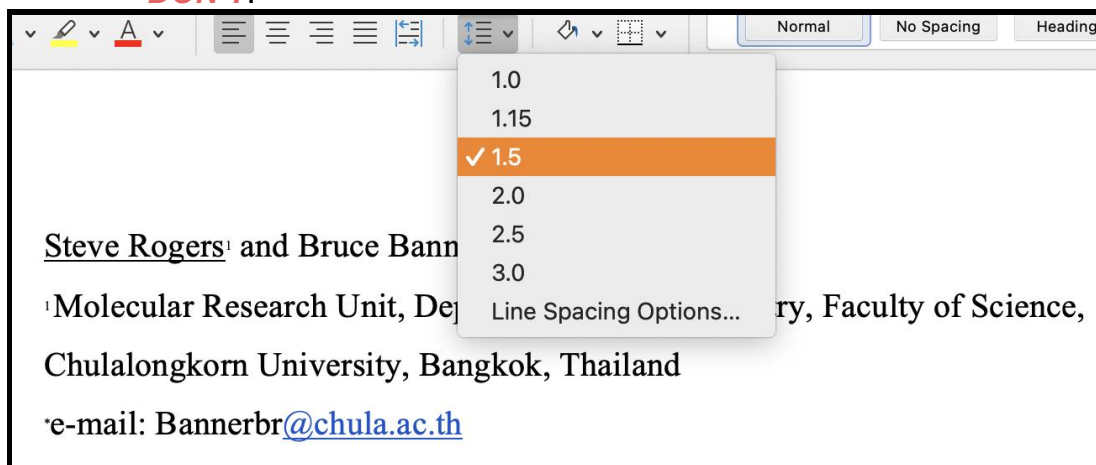
In order to make the whole abstract book, we are recommended that the author will use an example template to prepare the abstract. So, this is to clarify the **DOs** and **DON'Ts** for the abstract.

1. Make the title **bold** and **uppercase** for every word, if possible.
In case of nomenclatures, names of genes and other exceptions, you must follow their formats.
 - **DON'T:**
 - Identification of target proteins of the xyz proteins associated with cold stress in *Parkia speciosa* by using y2h screening
 - Identification of Target Proteins of the Xyz Proteins Associated with Cold Stress in Parkia Speciosa by Using y2h Screening
 - **DO:**
 - IDENTIFICATION OF TARGET PROTEINS OF THE XYZ PROTEINS ASSOCIATED WITH COLD STRESS IN *Parkia speciosa* BY USING Y2H SCREENING
Parkia speciosa is not in uppercase as it follows the format of nomenclature, but the rest of the title is in upper case.
2. Underline only the name and lastname of the presenter
 - **DON'T:** Steve Rogers¹ and Tony Stark^{2,*}
 - **DO:** Steve Rogers¹ and Tony Stark^{2,*}
3. Put **and** in the name section
 - a. In case of 2 authors, put **and** without a comma between the authors
 - **DON'T:** Steve Rogers¹, Tony Stark^{2,*}
 - **DON'T:** Steve Rogers¹, and Tony Stark^{2,*}
 - **DO:** Steve Rogers¹ and Tony Stark^{2,*}
 - b. In case of more than 2 authors, put a comma between the authors, and put **and** after a comma before the name of the last author
 - **DON'T:** Steve Rogers¹, Tony Stark², Bruce Banner^{3,*}
 - **DON'T:** Steve Rogers¹, and Tony Stark² and Bruce Banner^{3,*}
 - **DO:** Steve Rogers¹, Tony Stark², and Bruce Banner^{3,*}
4. Put a comma (if any) after a number of the author, not before.
 - **DON'T:** Steve Rogers¹, Tony Stark², and Bruce Banner^{3,*}
 - **DO:** Steve Rogers¹, Tony Stark², and Bruce Banner^{3,*}
5. A symbol **asterisk** of the corresponding author in the name section is not superscript, just type * as usual after a superscript comma with no space.
 - **DON'T:** Steve Rogers¹ and Tony Stark^{2,*}
 - **DON'T:** Steve Rogers¹ and Tony Stark^{2,*} (there's space between , and *)

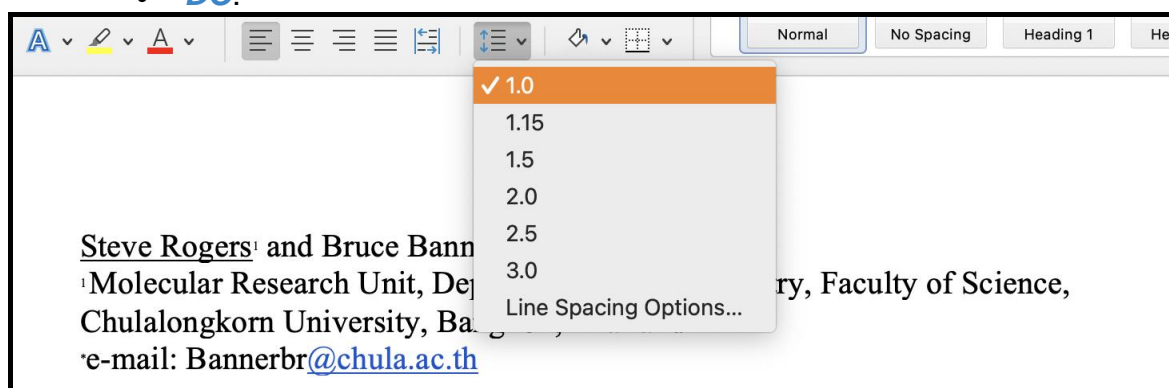
- **DO:** Steve Rogers¹ and Tony Stark^{2,*}
- **DO:** Steve Rogers¹, Tony Stark², and Bruce Banner^{3,*}

6. Make the author names and affiliation **single (1) spaced**, not double nor 1.5.

- **DON'T:**

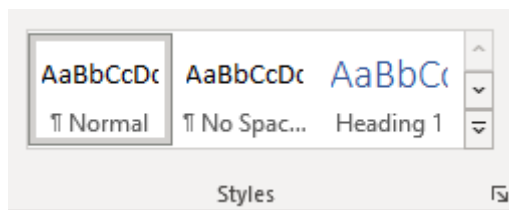


- **DO:**

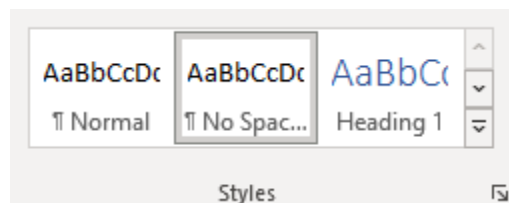


7. Make the No Spacing styles, not Normal styles

- **DON'T:**



- **DO:**



8. The name and affiliation are not separated by space between the lines.

- **DON'T:**

Steve Rogers¹ and Bruce Banner^{2,*}

¹Molecular Research Unit, Department of Biochemistry, Faculty of Science,
Chulalongkorn University, Bangkok, Thailand

*e-mail: Bannerbr@chula.ac.th

- **DO:**

Steve Rogers¹ and Bruce Banner^{2,*}

¹Molecular Research Unit, Department of Biochemistry, Faculty of Science,
Chulalongkorn University, Bangkok, Thailand

*e-mail: Bannerbr@chula.ac.th

9. Put a colon after the word **Abstract** with no space in between

Example:

- **DON'T:** Abstract
- **DON'T:** Abstract
- **DON'T:** Abstract :
- **DO:** Abstract:

10. Make **indentation** for the abstract with **no space** between the lines from the word **Abstract:**

- **DON'T:**

Abstract:

Ubiquitin (UBQ)-mediated proteolysis is one of the highly-conserved main regulatory metabolisms among eukaryotes. The XX enzyme, the most diverse group of enzymes in this process, is a ligase binding to UBQ-tagged target protein via a substrate-adapter protein YY

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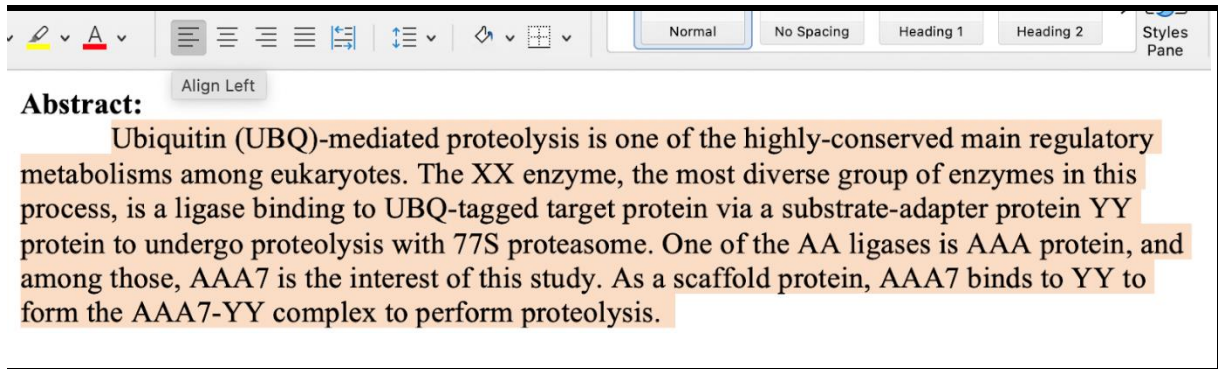
- **DO:**

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11. Justify the abstract, not Align Left nor Align Right

- **DON'T:**

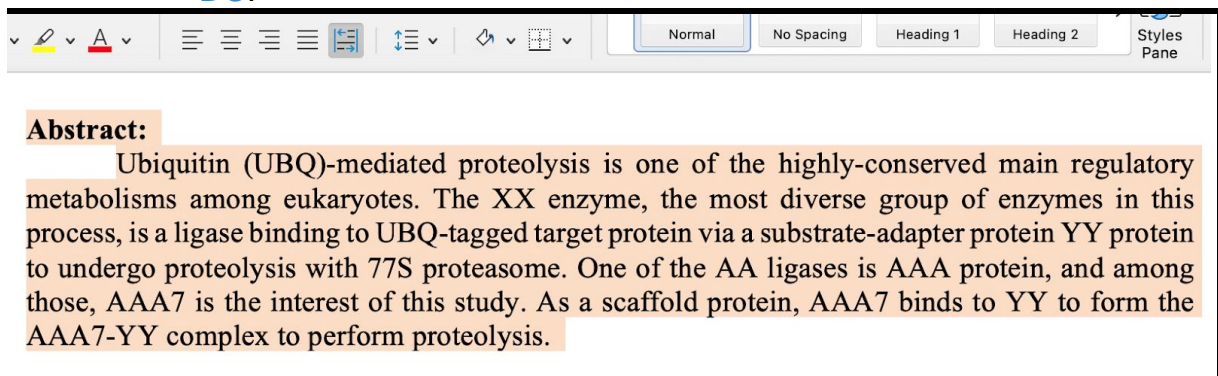


The screenshot shows a word processor interface with a toolbar at the top. The 'Align Left' button is highlighted. Below the toolbar, the word 'Abstract:' is followed by a paragraph of text. The text is left-aligned, and the 'Align Left' button is visible in the toolbar.

Abstract:

Ubiquitin (UBQ)-mediated proteolysis is one of the highly-conserved main regulatory metabolisms among eukaryotes. The XX enzyme, the most diverse group of enzymes in this process, is a ligase binding to UBQ-tagged target protein via a substrate-adapter protein YY protein to undergo proteolysis with 77S proteasome. One of the AA ligases is AAA protein, and among those, AAA7 is the interest of this study. As a scaffold protein, AAA7 binds to YY to form the AAA7-YY complex to perform proteolysis.

- **DO:**



The screenshot shows a word processor interface with a toolbar at the top. The 'Justify' button is highlighted. Below the toolbar, the word 'Abstract:' is followed by a paragraph of text. The text is justified, and the 'Justify' button is visible in the toolbar.

Abstract:

Ubiquitin (UBQ)-mediated proteolysis is one of the highly-conserved main regulatory metabolisms among eukaryotes. The XX enzyme, the most diverse group of enzymes in this process, is a ligase binding to UBQ-tagged target protein via a substrate-adapter protein YY protein to undergo proteolysis with 77S proteasome. One of the AA ligases is AAA protein, and among those, AAA7 is the interest of this study. As a scaffold protein, AAA7 binds to YY to form the AAA7-YY complex to perform proteolysis.