

〔大学院総合医薬学研究科総合医薬学専攻〕 博士後期課程先端薬科学プログラム・博士課程臨床薬学プログラム  
[Graduate School of Medicine and Pharmaceutical Sciences (Doctoral Course)]

小論文・適性検査 (Short Essay and Aptitude Test)

受験番号 (Examinee's No.)

氏 名 (Name)

(裏面にわたる場合は、この線より下に解答すること。)

(If your answer is longer than the space provided, you can write on the back of this page, but please write below this line.)

- (1) 薬科学分野へのデータサイエンスの貢献について具体的に論じなさい。

Discuss in detail the contribution of data science to the field of pharmaceutical sciences.

- (2) 最近読んだ研究論文の内容を要約し、その研究の新規性と重要性を述べなさい。

Summarize the contents of a research article that you have read recently and describe the novelty and importance of the study.

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Discuss in detail the contribution of data science to the field of pharmaceutical sciences.

出題の意図： 博士後期課程では研究を多角的に考察して推進する力を養う。いろいろな分野で重要性を増すデータサイエンスについて、薬科学分野への活用に関する考察を問う。

- (2) 最近読んだ研究論文の内容を要約し、その研究の新規性と重要性を述べなさい。  
Summarize the contents of a research article that you have read recently and describe the novelty and importance of the study.

出題の意図： その研究の意義を理解した上で、学術論文の内容を的確に説明できる能力を評価する。

小論文・適性検査 (Short Essay and Aptitude Test)受験番号 (Examinee's No.)分野名 (Educational area) Medicinal Resource Science 氏名 (Name)

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Q Describe the research you intend to pursue in your doctoral program, addressing the following five points. Please include hand-drawn diagrams, sketches, or flowcharts to make your explanation clearer. You may use both sides of the paper if needed.

1. Objective
2. Background
3. Goal
4. Methods
5. Significance

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【出題の意図 (Intention of the question)】

自らが取り組もうとする研究課題についてどれだけ深く考えているのかを評価する。

『考える力』は今まさに求められる力である。

【解答例 (Sample Answer)】

1. Research Objective

This study aims to elucidate the genetic regulatory mechanisms involved in the biosynthesis of nicotine and other alkaloids in tobacco plants. Secondary metabolites produced by plants have long played a crucial role in human society, serving as medicines, stimulants, dyes, and more. Among these, nicotine in tobacco exhibits a strong insect-repellent effect and is regulated through a jasmonic acid-mediated signaling pathway within the plant. However, the detailed mechanisms of this complex regulation remain poorly understood. This research focuses on the role of the transcription factor NtMYC and the NIC genes, employing molecular genetics techniques to uncover their functions.

2. Research Background

While the biosynthesis and regulatory pathways of flavonoids, such as anthocyanins, have been extensively studied and largely clarified at the molecular level, much remains unknown about the regulation of other secondary metabolites like alkaloids. Alkaloids, which evolved during the later stages of plant speciation, are believed to have unique regulatory mechanisms distinct from those of flavonoids. Nicotine, a key alkaloid found in Solanaceae plants, serves as a prominent example. Tobacco, which has been cultivated and genetically manipulated for commercial purposes over many years, is particularly well-suited for genetic and genomic research due to its ease of genetic transformation and the availability of comprehensive genome data.

3. Research Goals

The first objective of this study is to clarify the role of the transcription factor NtMYC, which binds to the E-box sequence in the promoter region of nicotine biosynthetic genes. By using RNA interference (RNAi) and CRES-T techniques, we will generate transformed tobacco plants with suppressed NtMYC function and assess the impact on gene expression and alkaloid accumulation. Since multiple NtMYC genes have been identified, it is necessary to determine which members are involved in the regulation of the biosynthetic pathway. Additionally, we will investigate the relationship between NtMYC and the NIC regulatory genes to facilitate the molecular cloning of the NIC genes.

4. Research Methods

To pinpoint the location of the NIC genes, we will perform genetic mapping. Using an F2 segregating population obtained through hybridization, we will analyze the linkage between known SSR markers and the low-nicotine phenotype to estimate the chromosomal position of the NIC1 gene. Subsequently, we will develop high-density molecular markers based on genomic information to narrow down the NIC1 gene locus to within 100 kilobases. Once candidate genes are identified, we will sequence them in mutant strains and conduct RNAi-based expression suppression experiments to assess their impact on nicotine accumulation.

5. Significance of the Research

This study leverages molecular genetic approaches to elucidate the regulatory factors that control the accumulation of secondary metabolites in plants, providing a valuable complement to traditional biochemical research. The findings are expected to offer insights applicable not only to tobacco but also to other plant species that produce alkaloids. Furthermore, this research may contribute to the development of improved methods for the efficient and safe production of valuable natural products, benefiting both academic and industrial fields.

(Figures omitted)