

BioscriptAI FAQ

Last Updated: 7/1/2025

1. Why aren't my graphs being generated?

- You must have Python and/or R installed on your computer to generate graphs on BioscriptAI. If you're running code client-side, please ensure:
 - Python and/or R is correctly installed and added to your system's PATH.
 - Required libraries (e.g., matplotlib, ggplot2) are installed.
- If you're using the web version and still not seeing previews, try refreshing the page or contacting support.

2. Is BioscriptAI compatible with mobile devices?

While BioscriptAI can be opened on mobile devices, we recommend using our platform on Windows/MacOS for the best experience.

3. What programming languages does BioscriptAI support?

BioscriptAI currently supports Python and R, specifically optimized for bioinformatics workflows. More languages may be added in the future based on user demand.

4. Can I save my chat history?

Yes! If you're logged in via Google, your chat history is saved and accessible at any time. Free accounts may have limits on history length or retention.

5. Do I need to know how to code?

Not necessarily! BioscriptAI is built to help both beginners and advanced users. You can start with natural language instructions, and the assistant will help you write or debug R/Python code.

6. Is BioscriptAI safe to use with sensitive data?

We do not store or share your private code or data inputs beyond what's necessary to provide the Service. For enterprise or research-grade data security, contact us about private deployment options.

7. I found a bug or need help. How do I get support?

Please contact us via our [Feedback Form](#)