

# FAQ

## General

**What data is available in CellBase?**

**How can I search for data?**

**What's a "client"? What's the difference between "clients", the RESTful API, and the command line? Which one should I use?**

**Do I need to specify which database to search? Do I need my own database?**

**I want a list of variants associated with my gene of interest. How can I do that?**

**How can I visualise the data?**

**Can I use any visual tool?**

Yes, you can use Genome Maps to browse variants.

**None of these questions helped me. Help!**

## PyCellbase

**How can I use PyCellbase? How can I install?**

**Does pycellbase work on Python 2?**

PyCellBase requires **Python 3.x**, although most of the code is fully compatible with Python 2.7.

**I'm getting an error message. What do I do?**

### Table of Contents:

- [General](#)
  - [What data is available in CellBase?](#)
  - [How can I search for data?](#)
  - [What's a "client"? What's the difference between "clients", the RESTful API, and the command line? Which one should I use?](#)
  - [Do I need to specify which database to search? Do I need my own database?](#)
  - [I want a list of variants associated with my gene of interest. How can I do that?](#)
  - [How can I visualise the data?](#)
  - [Can I use any visual tool?](#)
  - [None of these questions helped me. Help!](#)
- [PyCellbase](#)
  - [How can I use PyCellbase? How can I install?](#)
  - [Does pycellbase work on Python 2?](#)
  - [I'm getting an error message. What do I do?](#)