

First of all, I have to say that to build this API I used the Django Python Framework and, specifically, I used the Django REST Framework, which is a popular framework for building APIs. The database I used is the default database engine of Django: SQLite.

To test the HTTP requests, I used POSTMAN.

It means that in order to run this project, you must have installed Django and Django REST Framework:

- `pip install django`
- `pip install djangorestframework`

And then, run the server:

- `python manage.py runserver`
  - Type: *python manage.py runserver 80* if you want to run on port 80.

Once you browse <http://127.0.0.1:8000>, you will see the Django Admin Panel. The credentials are:

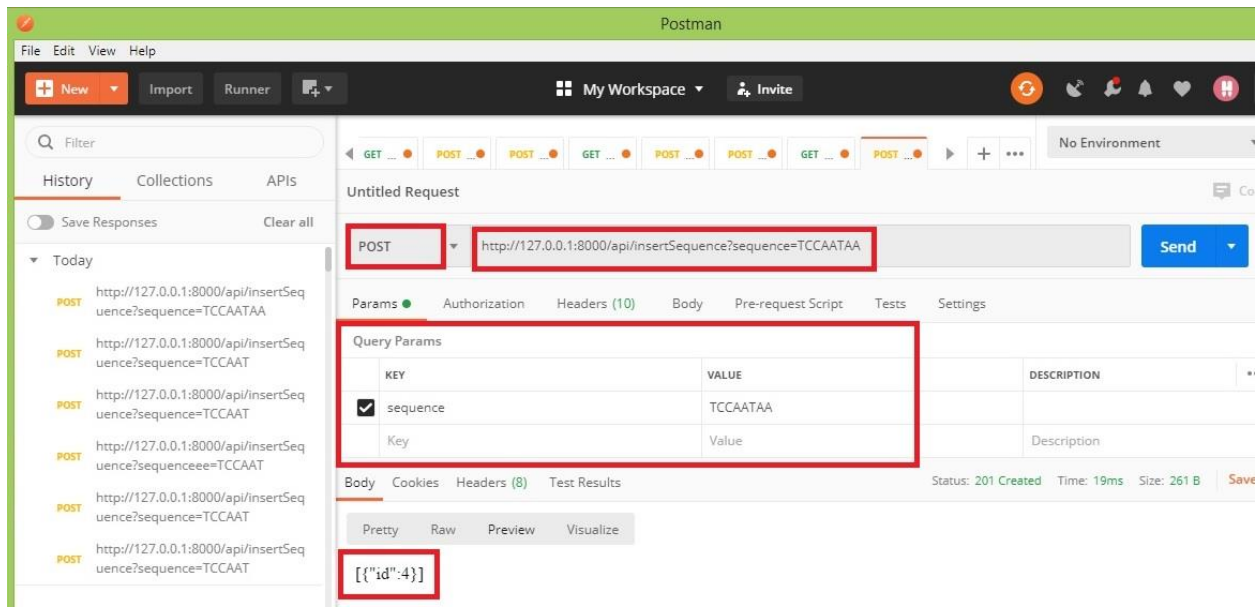
- User: **antonio**
- Password: **dna12345**

Once you log in, you will see site administration panel, and in <http://127.0.0.1:8000/api/dnasequence/> you will find all the sequences inserted in the database, as well as the possibility to manipulate them (add, delete, modify). Anyway, for adding, the idea is to use the *insert* method through an HTTP POST request in POSTMAN.

- **Method *insert***

This method can be tested generating an HTTP POST request via POSTMAN to: <http://127.0.0.1:8000/api/insertSequence> with the query param *sequence*, as follows:

<http://127.0.0.1:8000/api/insertSequence?sequence=TCCAATAAA>

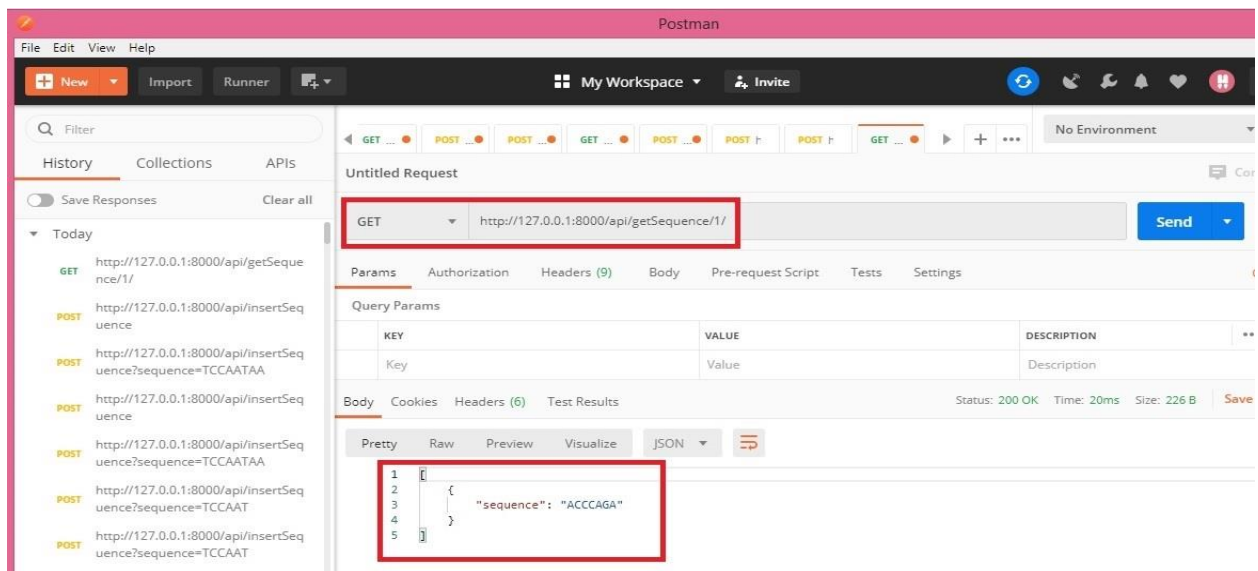


- **Method *get***

This method can be tested generating an HTTP GET request via POSTMAN or through the browser:

<http://127.0.0.1:8000/api/getSequence/<uid>/> where `<uid>` is the Unique Identifier of the sequence, which is the field `id` of the database. The `id` attribute is handled automatically by the Django's ORM and increases by 1 for each insertion, starting by the id 1. Then, for example we have:

<http://127.0.0.1:8000/api/getSequence/1/>

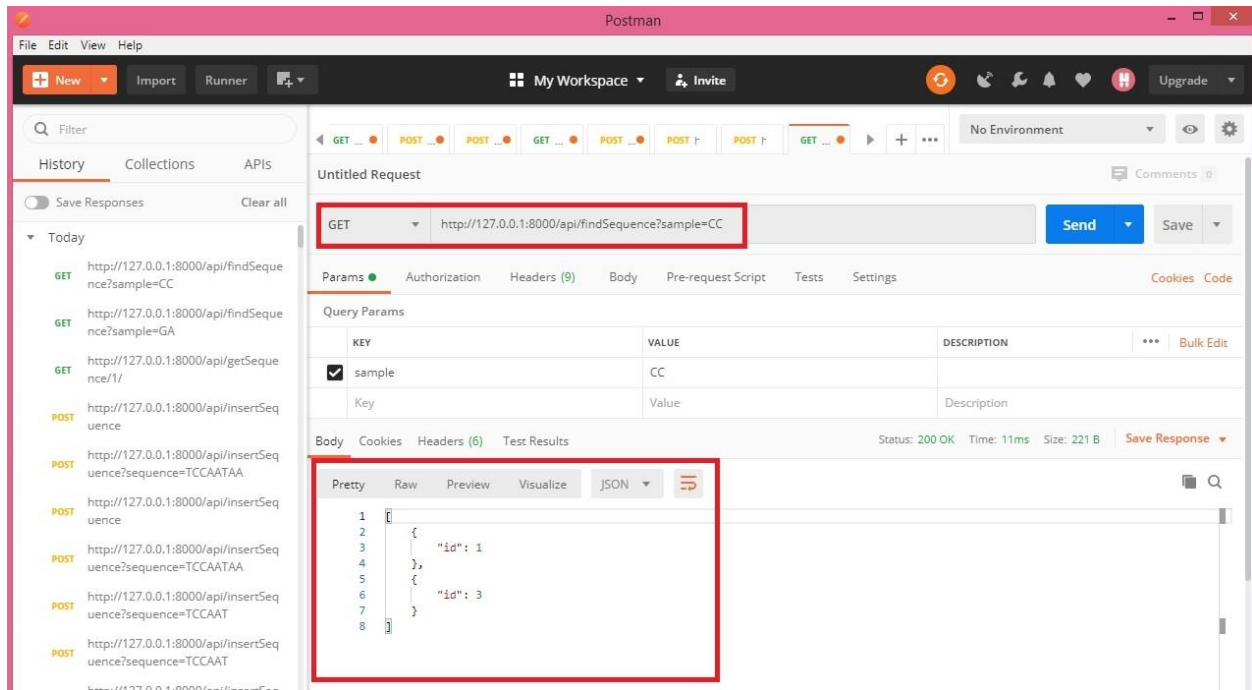


- **Method find**

This method can be tested generating an HTTP GET request via POSTMAN or through the browser:

<http://127.0.0.1:8000/api/findSequence?sample=<sample>> where <sample> is the sample input.  
For example:

<http://127.0.0.1:8000/api/findSequence?sample=CC>



- **Method overlap**

This method can be tested generating an HTTP GET request via POSTMAN or through the browser:

<http://127.0.0.1:8000/api/overlapSequence?sample=<sample>&uid=<uid>> where <sample> is the sample input, and <uid> is the Unique Identifier of the sequence. Then, for example we have:

<http://127.0.0.1:8000/api/overlapSequence?sample=GAGA&uid=1>

Postman

File Edit View Help

New Import Runner My Workspace Invite

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Today

- GET http://127.0.0.1:8000/api/overlapSequence?sample=GAGA&uid=1
- GET http://127.0.0.1:8000/api/findSequence?sample=CC
- GET http://127.0.0.1:8000/api/findSequence?sample=GA
- GET http://127.0.0.1:8000/api/getSequence/1/
- POST http://127.0.0.1:8000/api/insertSequence
- POST http://127.0.0.1:8000/api/insertSequence?sequence=TCCAATAA
- POST http://127.0.0.1:8000/api/insertSequence
- POST http://127.0.0.1:8000/api/insertSequence?sequence=TCCAATAA

Untitled Request

GET http://127.0.0.1:8000/api/overlapSequence?sample=GAGA&uid=1

Send Save

Params Authorization Headers (9) Body Pre-request Script Tests Settings Cookies Code

Query Params

KEY	VALUE	DESCRIPTION
<input checked="" type="checkbox"/> sample	GAGA	
<input checked="" type="checkbox"/> uid	1	
Key	Value	Description

Body Cookies Headers (6) Test Results Status: 200 OK Time: 12ms Size: 219 B Save Response

Pretty Raw Preview Visualize JSON

```
1
2  {
3    "overlap": true
```