MuG - 3D model RESTful API Documentation

Release 0.1

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Requirements and Installation

1.1 Requirements

1.1.1 Software

- Python 2.7.10+
- pyenv
- pyenv virtualenv
- pip

1.1.2 Python Modules

- h5py
- NumPy
- Flask
- Flask-Restful
- json
- pytest
- Waitress
- Sphinx
- sphinx-autobuild

1.2 Installation

1.2.1 Basics

Directly from GitHub:

```
1 git clone https://github.com/Multiscale-Genomics/mg-rest-3d.git
2 cd mg-rest-3d/
3 pip install -e .
4 pip install -r requirements.txt
```

Using pip:

```
pip install git+https://github.com/Multiscale-Genomics/mg-rest-3d.git
```

1.2.2 Setting up a server

```
git clone https://github.com/Multiscale-Genomics/mg-rest-3d.git
2
  cd mg-rest-3d
3
  pyenv virtualenv 2.7.12 mg-rest-3d
4
  pyenv activate mg-rest-service
5
  pip install git+https://github.com/Multiscale-Genomics/mg-dm-api.git
6
  pip install -e .
7
  pip install -r requirements.txt
8
  pyenv deactivate
9
```

Starting the service:

```
nohup ${PATH_2_PYENV}/versions/2.7.12/envs/mg-rest-3d/bin/waitress-serve --listen=127.

↔0.0.1:5002 rest.app:app &
```

1.3 Testing

Test scripts are located in the *test*/ directory. Run *pytest* to from the root repository directory to ensure that the API is working correctly.

The scripts require a valid hdf5 file generated using the scripts from mg-storage-hdf5 and a matching datasets.json file located in the *rest/* directory

1.4 Documentation

To build the documentation:

```
pip install Sphinx
pip install sphinx-autobuild
cd docs
make html
```

mg-rest-3d server

2.1 Methods

2.1.1 Get Endpoints

class rest.app.GetEndPoints

Class to handle the http requests for returning information about the end points

static get()

GET list all end points

List of all of the end points for the current service.

Example

curl -X GET http://localhost:5001/mug/api/3dcoord

2.1.2 Get Resolutions

class rest.app.GetResolutions

Class to handle the http requests for returning information about the resolutions that models have been generated for

get (*args, **kwargs)

GET List available resolutions from dataset

Parameters

- user_id (*str*) User ID
- file_id (str) Identifier of the file to retrieve data from

Returns file – JSON file listing the available resolutions within the dataset

Return type json

Examples

2.1.3 Get Chromosomes

class rest.app.GetChromosomes

Class to handle the http requests for returning information about the chromosomes that the models have been generated across

```
get (*args, **kwargs)
```

GET List available chromosomes from dataset

Parameters

- user_id (str) User ID
- file_id (str) Identifier of the file to retrieve data from
- res (int) Resolution

Returns file – JSON file listing the available chromosomes within a dataset at a given resolution

Return type json

Examples

```
curl -X GET http://localhost:5001/mug/api/3dcoord/chromosomes?user_id=test&

→file_id=test_file
```

2.1.4 Get Regions

class rest.app.GetRegions

Class to handle the http requests for returning information about the regions that are available in a given region and level of resolution

get (*args, **kwargs)

GET List available models from dataset

Parameters

- user_id (str) User ID
- file_id (str) Identifier of the file to retrieve data from
- res (int) Resolution
- **chrom** (*str*) Chromosome identifier (1, 2, 3, chr1, chr2, chr3, I, II, III, etc) for the chromosome of interest
- **start** (*int*) Start position for a selected region
- end (int) End position for a selected region

Returns file – JSON file listing the available models within a dataset at a given resolution and chromosomal region

Return type json

Examples

2.1.5 Get Models

class rest.app.GetModels

Class to handle the http requests for returning information about the models that are available within a given region.

get (**args*, ***kwargs*) GET List available models from dataset

Parameters

- user_id (str) User ID
- file_id (str) Identifier of the file to retrieve data from
- **res** (*int*) Resolution
- region (str) Region ID

Returns file – JSON file listing the available models within a dataset at a given resolution and chromosomal region

Return type json

Examples

2.1.6 Get Model

```
class rest.app.GetModel
```

Class to handle the http requests for returning the models from a given region. The list of models is a comma separated list that can return multiple models from the same region

```
get (*args, **kwargs)
```

GET List available model from dataset

Parameters

- user_id (str) User ID
- file_id (str) Identifier of the file to retrieve data from
- res (int) Resolution

- region (str) Region ID
- model (*str*) model ID

Returns file – JSON file listing the available models within a dataset at a given resolution and chromosomal region

Return type json

Examples

```
u curl -X GET http://localhost:5001/mug/api/3dcoord/model?user_id=test&file_

→id=test_file&region=1&model1
```

2.1.7 Ping

class rest.app.Ping

Class to handle the http requests to ping a service

static get() GET Status

List the current status of the service along with the relevant information about the version.

Example

curl -X GET http://localhost:5001/mug/api/3dcoord/ping

Chapter $\mathbf{3}$

Architectural Decision Record (ADR)

This file is a record of the choices that have been made about the choice of software, packages, pipelines and data structures that have been made in this repository. This document should serve the help future developers (including the original authors) understand what certain choices were made.

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