

# Article

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## 1 Initializations

Script 1.0.1 (python)

```
1 import requests, sys
2 import json
3 import numpy as np
4 import matplotlib.pyplot as plt
5 %matplotlib inline
6
7 server = "https://grch37.rest.ensembl.org"
8 headers={ "Content-Type" : "application/json", "Accept" : "application/json"}
```

## 2 Lookup

Script 2.0.1 (python)

```
1 ext = "/lookup/id"
2 # Get lookup data
3 r = requests.post(server+ext, headers=headers, data='{ "expand" : 1, "ids" :
  ↳ ["ENSG00000157764"] }')
4
5 if not r.ok:
6     r.raise_for_status()
7     sys.exit()
8
9 found = json.loads(r.text)
```

## 3 Get sequences

Script 3.0.1 (python)

```
1 # Create list of ID
2 IDs = []
3 for transcripts in found.values():
4     for transcript in transcripts["Transcript"]:
5         IDs.append(transcript["id"])
6
7 dquotes_ids=json.dumps({"ids" : IDs})
8 print(dquotes_ids)
9
10 # Get sequence data
11 ext = "/sequence/id"
12
13 # Make request and check status
14 r = requests.post(server+ext, headers = headers, data = dquotes_ids)
15 if not r.ok:
```

```
16     r.raise_for_status()
17     sys.exit()
18
19 json_data = json.loads(r.text)
```

### Output

```
{"ids": ["ENST00000496384", "ENST00000288602", "ENST00000479537", "ENST00000497784",
↪ "ENST00000469930"]}
```

## 4 Get oligos

### Script 4.0.1 (python)

```
1 count_oligos = {}
2 for record in json_data:
3     oligo = record["seq"][102:105]
4     if oligo in count_oligos:
5         count_oligos[oligo] += 1
6     else:
7         count_oligos[oligo] = 1
8 print(count_oligos)
```

### Output

```
{'AGT': 1, 'GGG': 1, 'AAA': 1, 'GGC': 1, 'GCC': 1}
```