Train Model

Run Model

Train Model **Run Model**

Use random data

Load data...

Select file

Supported filetypes: .csv, .txt, and .npy
Go to url for github docs for example file formats.

sample filepath here

Train model with this data

Train Model   Run Model

Use random data

Load data...

Training model progress

% training complete

Train Model

Run Model

Load inputs from file.

Enter manually...

Model Inputs

DNA sequence inputs can only be a max length of 300 b.p.

DNA sequence 1

```
ACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTG
CTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGA
TGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTG
```

DNA sequence 2

```
ACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTG
CTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTGA
TGACTGACTGACTGACTGACTGACTGACTGACTGACTGACTG
```

Run model with these inputs

Select file

Supported filetypes: .csv, .txt, and .npy
Go to url for github docs for example file formats.
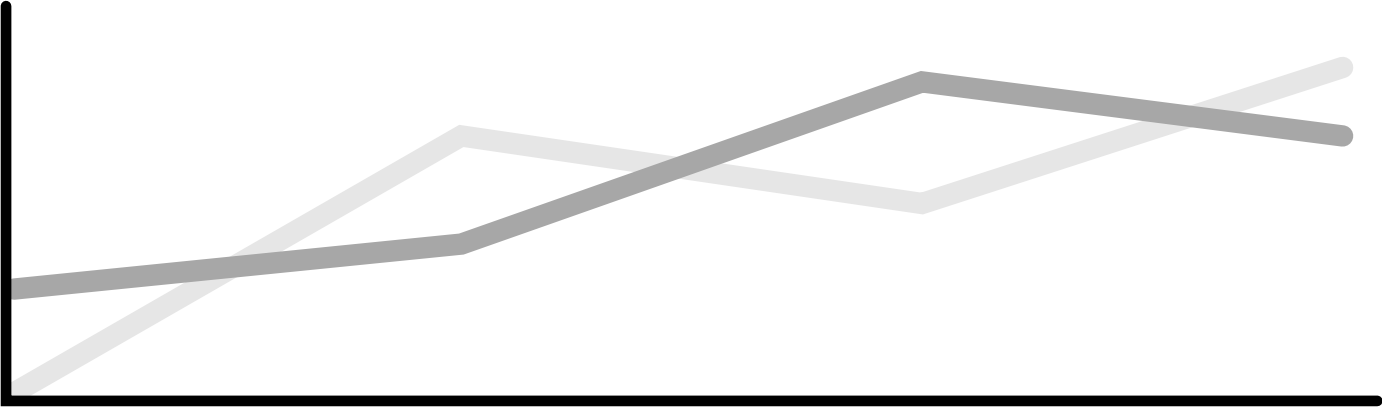
[sample filepath here]
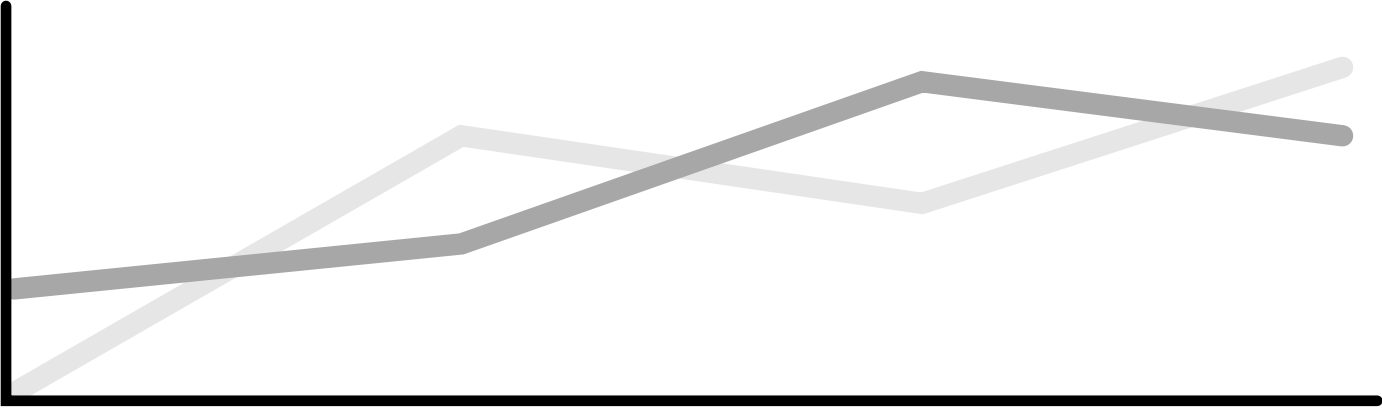
Run model with these inputs

Output graph

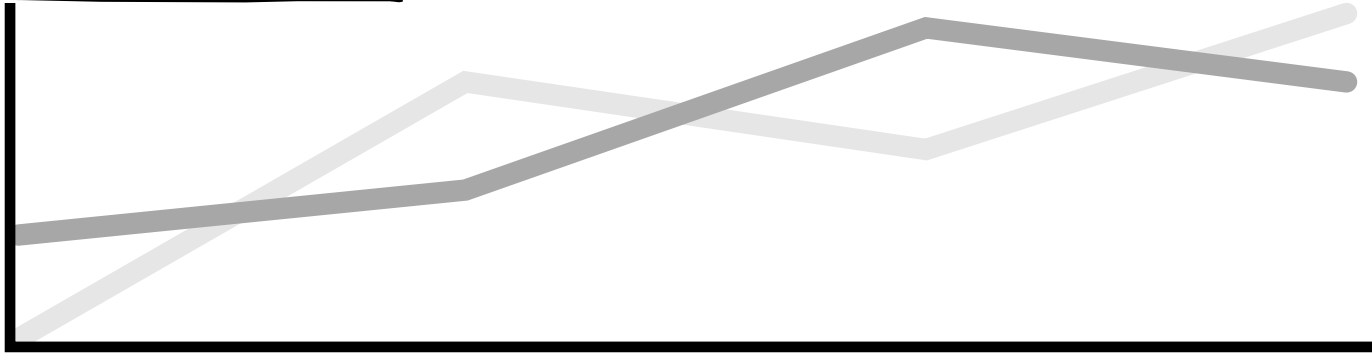Export As...

DNA sequence 1

DNA sequence 2

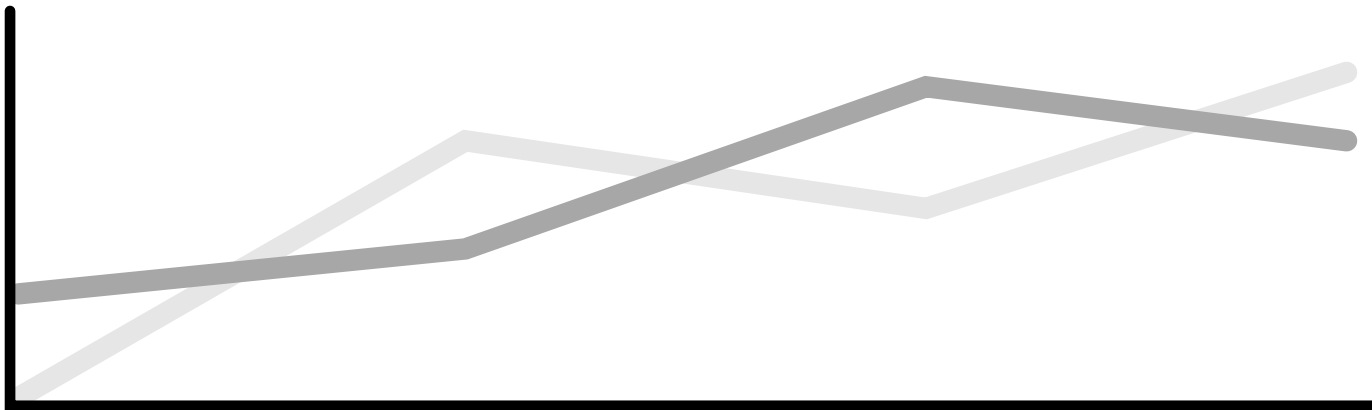# Output graph

Export As...

.png

.pdf

.svg

DNA sequence 2

Output graph

Export As...
- .png
- .pdf
- .svg

Confirmation

Output saved to current directory.

OK

DNA sequence 2