

Previously clustered datasets

SART1 x

SART2 x

SART3 x

BrainEx

Ver 1.0.3

 Query from a previously clustered dataset

 Generate clusters from a new dataset

BrainEx I File Browser



Navigation icons: back, forward, computer icon, address bar: > Computer, refresh icon, search icon

Organize System Properties Uninstall or change a program Map network drive Open Control Panel

- ★ Favorites
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Name	Date Modified	File Type
Loreem ipsum dolor	1/13/2012 9:00	File Folder
Loreem ipsum dolor	1/13/2012 9:00	Text File
Loreem ipsum dolor	1/13/2012 9:00	Text File
Loreem ipsum dolor	1/13/2012 9:00	Text File
Loreem ipsum dolor	1/13/2012 9:00	Text File
Loreem ipsum dolor	1/13/2012 9:00	Text File
Loreem ipsum dolor	1/13/2012 9:00	Text File

Upload a CSV file:

dataset4.csv

Upload

Cancel



BrainEx I File Browser



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Upload a CSV file:

dataset4.csv

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Cancel



BrainEx I Clustering Options



Similarity Threshold: 0.1 0.2 Custom: ____

Type of Distance:

Euclidean	▼
Euclidean	
City Block (manhattan)	
Minkowski	
Chebyshev	

Sequence Length of Interest: between

200	▲▼
-----	----

 and

1000	▲▼
------	----

Back

Start Processing

BrainEx I Loading Clusters



Cluster Progress...51%



Sequences Clustered: 102/200

Cancel

Next

Notes:

This page is intended to automatically load and display the finished loading clusters page. However, you need to click next because of Balsamiq's (prototype website that we used for creating this mockup) limitations on transitions.

Cluster loading is complete!

Sequences Clustered: 200/200

Go to:

Explore all data

Explore data by clusters

Start Querying

Cluster loading is complete!

Sequences Clustered: 200/200

Next

Load Dataset



Dataset Explorer



Cluster Explorer



Query Finder



BrainEx Menu

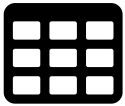
Query Finder



Dataset Explorer



Cluster Explorer



Load Another
Dataset

Current Selection



Filter

Channel

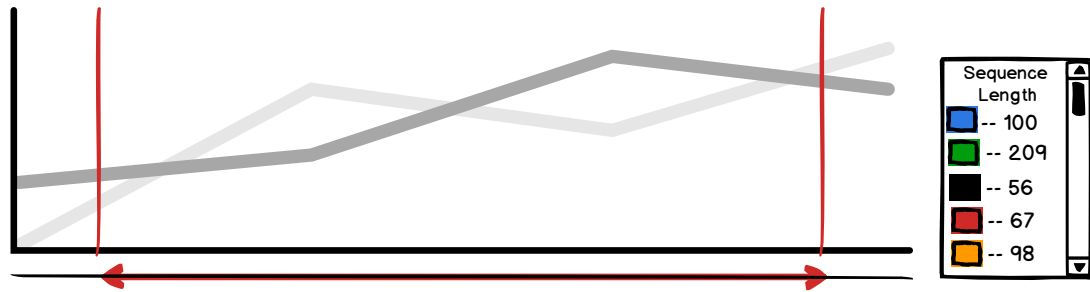
- Channel1
- Channel2
- Channel3
- Channel4
- Channel5

SubjectID

Labels

Statistics

Data Visualizer



Data Viewer

SubjectID	Event Name	Channel Name	Start Time	End Time
101 HART	target correct	Channel-1 HBO	300	9000
93 SART	target incorrect	Channel-4 VBO	789	45756
34 TART	target no response	Channel 18 JVO	210	45900

Save file

Query Selected Sequence

Notes:

Sort the data here in descending order of (end time - start time)

Design Justification:

We wanted to make the screen straightforward, simple to look at and easy to navigate. Keeping these goals in mind, we put all the user parameter selections on one side (the left) and visualizations on the other (the right).

After performing necessary functions, the user can save the file and then proceed to the next step which is querying the sequence.

At the top, there is a navigation bar that allows the user to go to the main page as well as each of the explorer/query pages.

We chose a slider instead of brushing on chart region so it does not interfere with selecting an actual sequence from the chart.

BrainEx Homepage Dataset Explorer **Cluster Explorer** Query Finder

Clusters

Number: 5

Number: 60

Number: 45

Number: 16

Filter

Number of Clusters to Display:
 1 20

Range of sequence length in each cluster:
 0 100

Statistics

Data Visualizer

Data Viewer

Sequence Length (in seconds)	Number of Sequences
100	5
209	60
56	45
67	16
98	78
509	109
290	32
876	12
487	9
1067	10

➔ Next

Notes:
 Data Visualizer will display the user specified amount of cluster representatives
 Data Table will have length (time) of sequences and number of sequences
 There will never be a cluster with the same length so can just use length as unique cluster ID -> this is false
 There can be multiple clusters with the same length as long as they are not similar in shape
 All sequences in a cluster will have the same length
 Selecting cluster and representative will highlight it on the data visualizer graph

Filters:
 Number of clusters (top 15 or 20)
 Range of length (slider)

Statistics:
 Show the cluster with the most/least sequences
 Show the top 3/least 3 with the greatest/least length of sequences

Question:
 How do we decide how many representatives to display to the user (maybe ask to input a number from 1-10) and on what basis (show representatives of the top 10 clusters with most data points)?
 If we have 30,000 clusters, how do we filter them down to approx 20? Should we ask for user input or another screen?
 How to identify each cluster? Should we assign an id to each?

Design Justification:
 To keep the UI simple, this cluster explorer screen as well as the rest of the explorer/query finder screens have the same layout with user inputs on the left and visualizations on the right
 The view data shapes area displays the data shape of the representative of each cluster which would make it simpler for the user to view and easier to pick which area to investigate more
 The number of sequences in each cluster is displayed as the label for the view data shapes area to make it easier for users to view that information. The filters are meant to allow the user to narrow down the number of clusters to view at a time



Clusters

Number: 5

Number: 60

Number: 45

Number: 16

Filter

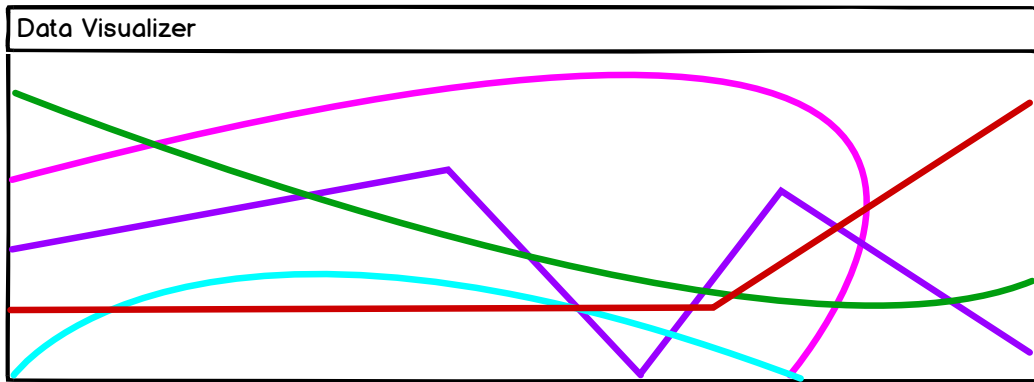
Number of Sequences from this Cluster to Display:

1 20

Range of sequence length:

0 500

Statistics



Legend

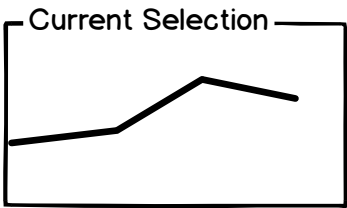
- -- 100
- -- 209
- -- 56
- -- 67
- -- 98

Data Viewer

SubjectID	Event Name	Channel Name	Start Time	End Time	Sequence Length (in seconds)
101 HART	target correct	Channel-1 HBO	300	9000	100
93 SART	target incorrect	Channel-4 VBO	789	45756	67
34 TART	target no response	Channel 18 JVO	210	45900	45
101 HART	target correct	Channel-1 HBO	300	9000	90
93 SART	target incorrect	Channel-4 VBO	789	45756	78

Notes:
 Data Visualizer will display the sequences in the cluster selected
 Data Table will have data information for each sequence in the cluster selected

Design Justification:
 In order to make the cluster explorer more understandable and easy to use, we have a graph that displays the current user selection at the top of the screen.
 The user can filter some options and select the number of sequences to view in the cluster as well as the range of the sequence length. This part is to allow the user to narrow down the number of sequences to view at a time



Query sequence ^

- Upload a sequence file
- Use selection from dataset explorer
- Use selection from cluster explorer

Enter parameters

Show query result

Statistics

Notes:
Show query result is disabled on this page because parameters have not been entered.
Gray out selection if selection not made in cluster explorer/dataset explorer.

Design Justification:
The input required for query is setup as an accordion because we wanted to show the sequential order of input that is required for this step.

Current Selection



Query sequence ^

Enter parameters

Number of best matches: 5

 Exclude data from queried sequence

Overlap allowed: 10%

Show query result

Statistics

Notes:

Design Justification: A default overlap percentage of 10 is pre-set, this value has been confirmed with Leo.



Query sequence ^

Enter parameters

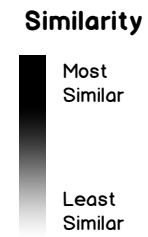
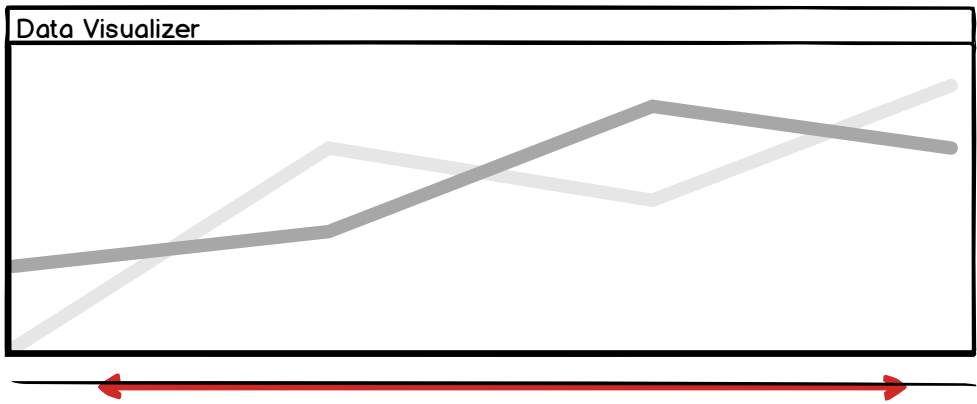
Number of best matches: ▲ ▼

Exclude data from queried sequence

Overlap allowed:

Show query result

Statistics



Data Viewer

SubjectID	Best Matches	Overlap	Similarity	Event Name	Channel Name	Start Time	End Time
101 HART	5	20%	65%	target correct	Channel-1 HBO	300	9000
93 SART	6	40%	75%	target incorrect	Channel-4 VBO	789	45756
34 TART	4	50%	85%	target no response	Channel 18 JVO	210	45900
101 HART	2	36%	95%	target correct	Channel-1 HBO	300	9000
93 SART	3	48%	55%	target incorrect	Channel-4 VBO	789	45756

Save file

Notes:
 Data Table: Best matches, overlap, similarity (95% similar), subject ID, channel, start, end
 Visualization: data points in each explorer are displayed in graph (same as graph in cluster explorer copy)

Design Justification:
 A slider is used here instead of the brushing on chart region for consistency with the previous screens, also making it easier for the user to select one specific sequence.