

INTERDICTIONary

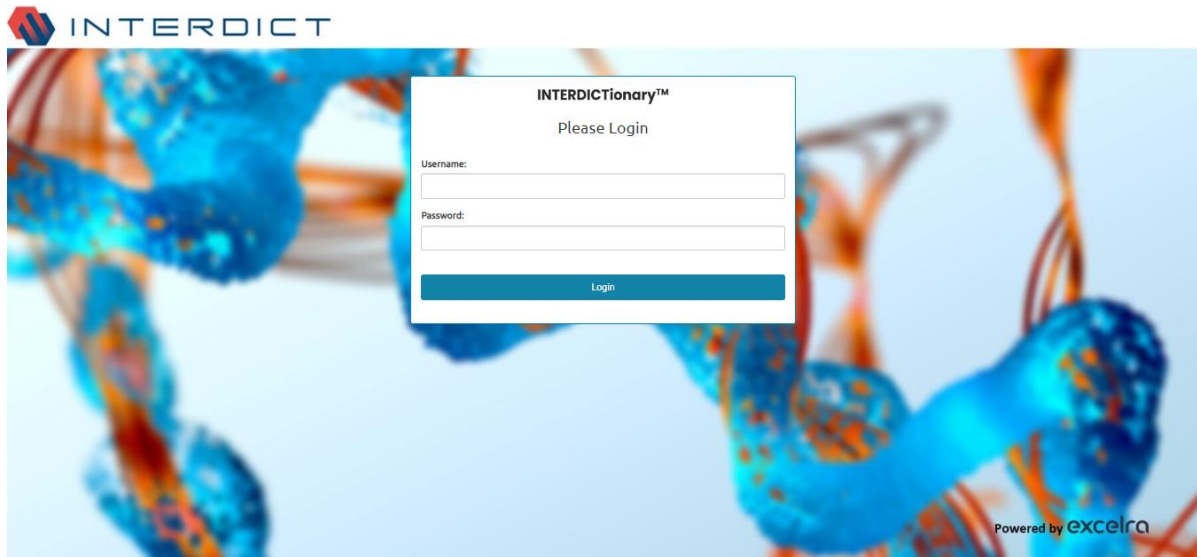
User Manual

Version 1.0

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Accessing the INTERDICTIONary Application



Home page after logged in

INTERDICTIONary™

6 7 8 9

Help User Exit

1 Choose Regex or Normal
Normal
Regex

2 Enter the sequences
Multiple sequences should be comma separated with no spaces.
(E.g. AAAA,AAAT)

3 Filter by Sequence(s)

4 Find overlaps

5 UniProt

10 Results Summary Selected Data Dashboard Admin

11

12 Search:

Column visibility Copy Print Download current page Download all data

Showing 1 to 7 of 10 entries

| EntryName | Entry | ProteinName | GeneNames | Organism | Length | Sequence | Position_List |
|--------------|--------|--|------------------|----------------------|--------|----------|---|
| HXA13_HUMAN | P31271 | Homeobox protein Hox-A13 (Homeobox protein Hox-1J) | HOXA13 HOX1J | Homo sapiens (Human) | 388 | AAAAA | 38,39,40,41,42,43,44,45,46,47,62,73,74,75,76,77,78,79,80,116,117,118,119,120,121,122,123, |
| SOX21_HUMAN | Q9V651 | Transcription factor SOX-21 (SOX-A) | SOX21 SOX25 SOXA | Homo sapiens (Human) | 276 | AAAAA | 137,138,139,140,141,142,154,155,156,157,158,159,160,161,162,208,209,210,211,212,213,21 |
| PHOX2B_HUMAN | Q99453 | Paired mesoderm homeobox protein 2B (Neuroblastoma Phox, NBPhox) (PHOX2B homeodomain | PHOX2B PHOX2B | Homo sapiens (Human) | 314 | AAAAA | 159,160,161,162,163,241,242,243,244,245,246,247,248,249,250,251,252,253,254,255,256 |

Query

a. Normal

User can enter the n number of sequences with comma separated

- E.g. AAAA, AANA, AACA

b. Regex

In order to search manually for multiple sequences using normal search, providing user to filter data using Regex so that user can filter multiple sequences data to analyse.

E.g. **`^PP[G,A,V].`****\$**

Input box

After selected the query option, user can enter the sequence in the provided input box.

Filter by sequence

User can filter the data table using this action button.

Find overlap

User can click on this action button to find the overlapped sequences based on Entry.

UniProt

User can click on this UniProt logo, it redirects to UniProt website

<https://www.uniprot.org/id-mapping>

In this application there are 4-tab panels

Results Summary

Shows the filtered data which is generated using query

Query to filter data

Filtered data

Choose Regex or Normal
Normal
Regex

Enter the sequences
AANA,AAAA,AACA

Multiple sequences should be comma separated with no spaces.
(E.g. AAA,AAA)

Filter by Sequence(s)

Find overlaps

Results Summary Selected Data Dashboard Admin

Refine your search
Add Filter Condition

Column visibility Copy Print Download current page Download all data

Search:

Showing 1 to 7 of 910 entries

| EntryName | Entry | ProteinName | GeneNames | Organism | Length | Sequence | Position_List | Count |
|-------------|----------|--|--|----------------------|--------|----------|---------------|-------|
| RIPP1_HUMAN | Q0D2K3-2 | Protein ripp1 | RIPPL1 | Homo sapiens (Human) | 104 | AACA | 4 | 1 |
| RIPP2_HUMAN | Q5TAB7 | Protein ripp2 | RIPPL2 C6orf159 | Homo sapiens (Human) | 128 | AACA | 15 | 1 |
| RIPP1_HUMAN | Q0D2K3 | Protein ripp1 | RIPPL1 | Homo sapiens (Human) | 151 | AACA | 4 | 1 |
| CL16A_HUMAN | Q2KH73-3 | Protein CLEC16A (C-type lectin domain family 16 member A) | CLEC16A KIAA0350 | Homo sapiens (Human) | 140 | AACA | 129 | 1 |
| CK02T_HUMAN | Q96N53 | Putative uncharacterized protein encoded by LINC00167 (PRDM10-DT divergent transcript) | PRDM10-DT C11orf127 LINC00167 NCRNA00167 | Homo sapiens (Human) | 147 | AACA | 69 | 1 |

Selected Data

- Shows the selected rows which are selected from **Results Summary**, for this user must click on the **Extract Data**

Selected two rows

INTERDICTIONary™

Choose Regex or Normal
Normal
Regex

Enter the sequences
AANA,AAAA,AACA

Multiple sequences should be comma separated with no spaces.
(E.g. AAA,AAA)

Filter by Sequence(s)

Find overlaps

Results Summary Selected Data Dashboard Admin

Refine your search
Add Filter Condition

Column visibility Copy Print Download current page Download all data

Search:

Showing 1 to 7 of 910 entries

| EntryName | Entry | ProteinName | GeneNames | Organism | Length | Sequence | Position_List | Count |
|-------------|----------|--|--|----------------------|--------|----------|---------------|-------|
| RIPP1_HUMAN | Q0D2K3 | Protein ripp1 | RIPPL1 | Homo sapiens (Human) | 104 | AACA | 4 | 1 |
| RIPP2_HUMAN | Q5TAB7 | Protein ripp2 | RIPPL2 C6orf159 | Homo sapiens (Human) | 128 | AACA | 15 | 1 |
| RIPP1_HUMAN | Q0D2K3 | Protein ripp1 | RIPPL1 | Homo sapiens (Human) | 151 | AACA | 4 | 1 |
| CL16A_HUMAN | Q2KH73-3 | Protein CLEC16A (C-type lectin domain family 16 member A) | CLEC16A KIAA0350 | Homo sapiens (Human) | 140 | AACA | 129 | 1 |
| CK02T_HUMAN | Q96N53 | Putative uncharacterized protein encoded by LINC00167 (PRDM10-DT divergent transcript) | PRDM10-DT C11orf127 LINC00167 NCRNA00167 | Homo sapiens (Human) | 147 | AACA | 69 | 1 |

- Go to **Selected Data** tab-panel and click on **Extract Data** button then user can see the data like below

INTERDICTIONary™

Enter Column Name

Create Brw Done

Results Summary Selected Data Dashboard Admin

Refine your search
Add Filter Condition

Column visibility Copy Print Download current page Download all data

Search:

Showing 1 to 2 of 2 entries

| EntryName | Entry | ProteinName | GeneNames | Organism | Length | Sequence | Position_List | Count |
|-------------|----------|---|------------------|----------------------|--------|----------|---------------|-------|
| RIPP2_HUMAN | Q5TAB7 | Protein ripp2 | RIPPL2 C6orf159 | Homo sapiens (Human) | 128 | AACA | 15 | 1 |
| CL16A_HUMAN | Q2KH73-3 | Protein CLEC16A (C-type lectin domain family 16 member A) | CLEC16A KIAA0350 | Homo sapiens (Human) | 140 | AACA | 129 | 1 |

Show 2 entries

Previous 1 Next

Extract Data

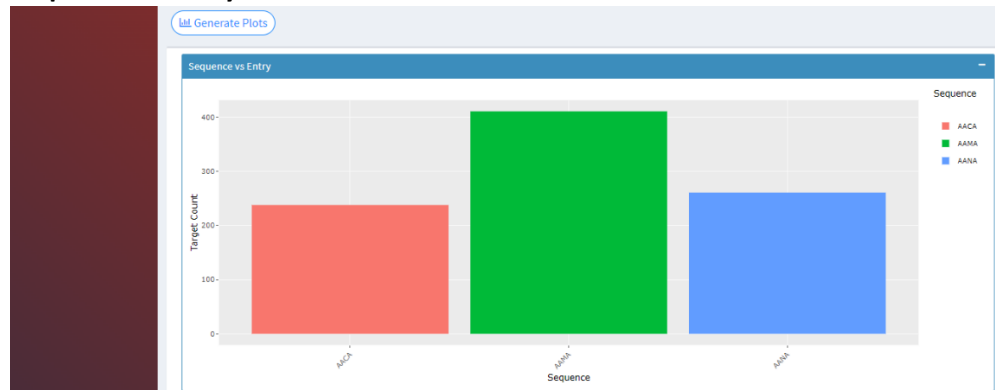
Dashboard

- Dashboard will generate the plots and summary counts based on filtered data
 - Summary counts will be displayed automatically

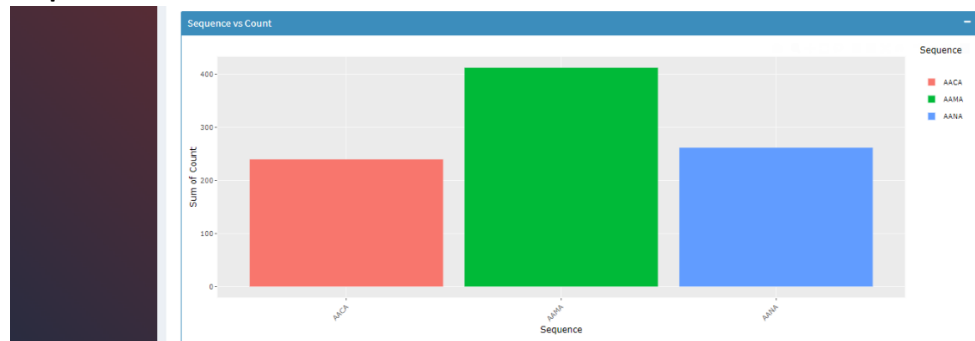
- ii. To generate the plots user must click on the Generate Plots button
- Dashboard will be changed dynamically after changed the query in Results Summary



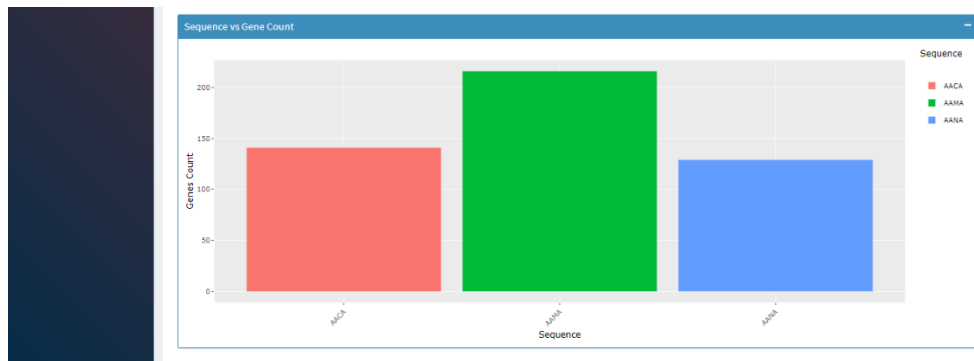
Sequence vs Entry



Sequence vs Count

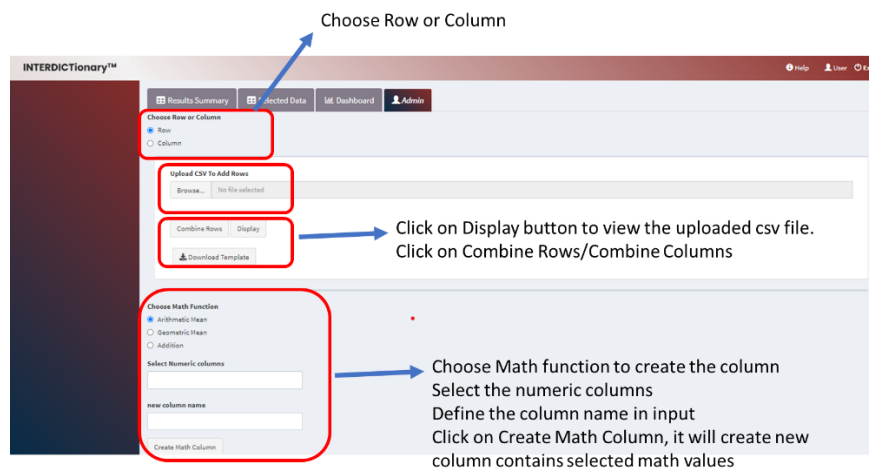


Sequence vs Gene Count



Admin

- Admin tab-panel will be accessible only by admin
- Admin panel will contain multiple options to manipulate the data which is presented in **Selected Data** tab-panel



i. Add Row

1. Download the template and add data values to that template and upload it click on the Combine Rows button
2. Results will be displayed in the **Selected Data** tab-panel

ii. Add Column

1. Download the template and add data values to that template and upload it click on the Combine Column button
2. Results will be displayed in the **Selected Data** tab-panel

iii. Add Match Column

1. Choose Math Function
 - a. Arithmetic Mean
 - i. Will calculate AM of two or more columns and creates the new column
 - b. Geometric Mean
 - i. Will calculate GM of two or more columns and creates the new column
 - c. Addition

- i. Will calculate Addition of two or more columns and creates the new column
- 2. New column name should be mentioned in the input box by the user.
- 3. Manipulated data will be displayed in the **Selected Data** tab-panel