**Example Table of SF2 Helicases**

| **Species** | **Group** | **Protein** | **Localization** | **Biological Process** |
| --- | --- | --- | --- | --- |
| S.Cerevisiae | eIF4A | [TIF1p / YKR059W TIF2p / YJL138C](https://rnahelicasedb.github.io/euk_dead/D_Eukaryota_F_DEAD-box_G_eIF4A_sG_eIF4A.html) | Cytoplasm  Plasma membrane | Translational initiation |
| C.Elegans | eIF4A | [eIF4A F57B9.6](https://rnahelicasedb.github.io/euk_dead/D_Eukaryota_F_DEAD-box_G_eIF4A_sG_eIF4A.html) |  | Translational initiation |
| D.Melanogaster | eIF4A | [CG9075](https://rnahelicasedb.github.io/euk_dead/D_Eukaryota_F_DEAD-box_G_eIF4A_sG_eIF4A.html) | Cytoplasm  P-granule | Translational initiation |
| M.Musculus | eIF4A | [Eif4a-1 Ddx2a](https://rnahelicasedb.github.io/euk_dead/D_Eukaryota_F_DEAD-box_G_eIF4A_sG_eIF4A.html) |  | Translational initiation |
| H.Sapiens | eIF4A | [EIF4A-1 DDX2A](https://rnahelicasedb.github.io/euk_dead/D_Eukaryota_F_DEAD-box_G_eIF4A_sG_eIF4A.html) | Cytoplasm  Extracellular exosome | Translational initiation  Host-virus interaction |
| M.Musculus | eIF4A | [Eif4a-2 Ddx2b](https://rnahelicasedb.github.io/euk_dead/D_Eukaryota_F_DEAD-box_G_eIF4A_sG_eIF4A.html) |  |  |
| H.Sapiens | eIF4A | [EIF4A-2 DDX2B](https://rnahelicasedb.github.io/euk_dead/D_Eukaryota_F_DEAD-box_G_eIF4A_sG_eIF4A.html) | Cytoplasm | Translational initiation  Host-virus interaction |

Each links in the **Protein** column will take you to Uniprot page, which contain all the information that we need.

The information is then gathered and organized into the HTML file (which you can open with any text editor).

You could manually copy and paste the information, OR you could make script to download the information as txt file (using uniprot’s public API) and parse the text file to our needs.

The HTML page is essentially a HTML table with rows and columns. Play around by deleting or replicating a line within the HTML table to see how the table is generated.

Note that for this protein group, because there are two isoforms (similar proteins) of EIF4A in both human and mouse (h.sapiens and m.musculus), there are additional rows for the human and mouse columns.

By carefully studying the HTML file, you can easily see how an additional row spanning is added in html table. From this, you will be able to learn how to remove or add new rows

Also, by looking at the HTML file, you will be able to learn how hyperlinks are added to the text in html file (<a> tag and href attribute).