



Open for Innovation

# KNIME

## KNIME Pros Learnathon

*Building Reliable and Reusable Components*

Group 2: Life Sciences

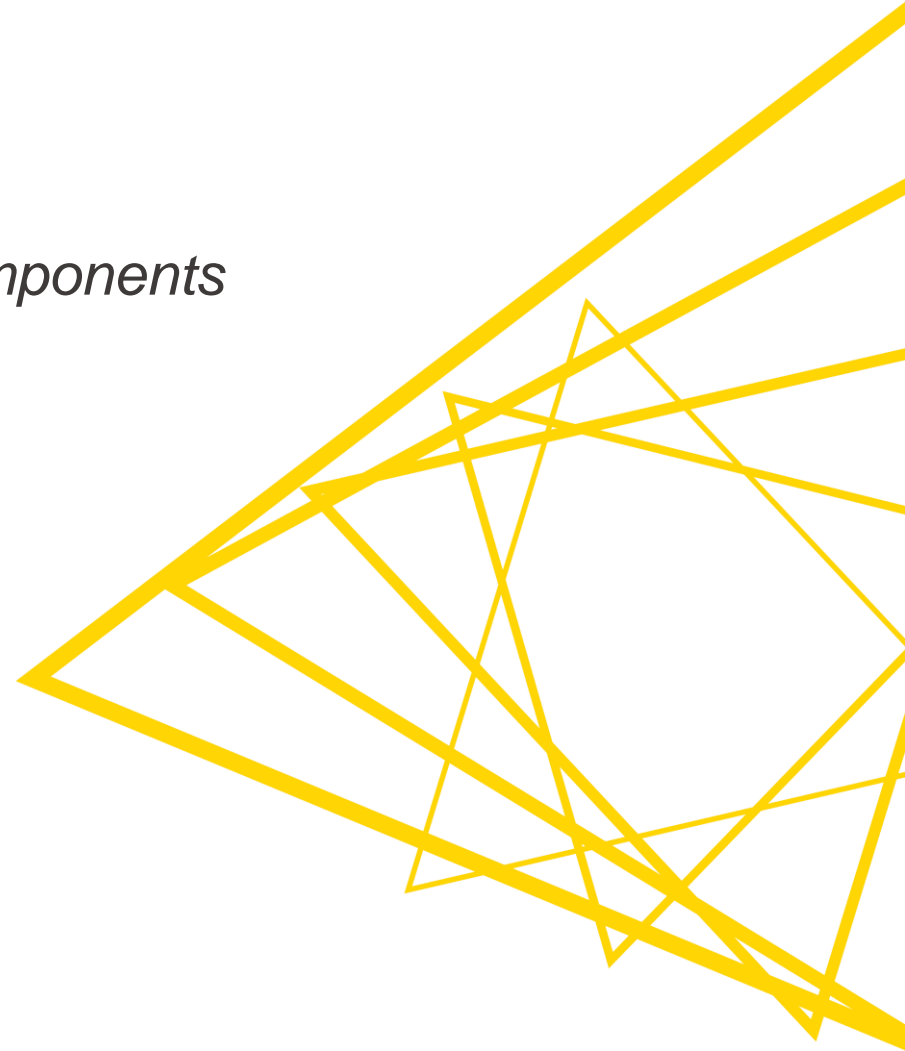
KNIME Team



@KNIME

#Learnathon

[tinyurl.com/KNIME-Pros-Stuff](https://tinyurl.com/KNIME-Pros-Stuff)



# See you soon in a Breakout Room!

## Main Zoom Session

### Group 1 Financial Analysis



**Maarit**

KNIME Team Member



**Lada**

KNIME Team Member

### Group 2 Life Sciences



**temesgen-dadi**

KNIME Team Member



**Francosinus**

KNIME Team Member

### Group 3 Automation



**paolotamag**

KNIME Team Member



**Mpattadkal**

KNIME Team Member

# Three Parallel Tracks via Zoom Breakout Rooms!

Main Zoom Session

Welcome to:  
**Group 2.**  
**Life Sciences**

Group 2  
Life Sciences



temesgen-dadi  
KNIME Team Member



Francosinus  
KNIME Team Member

▼ Group\_2-Life\_Sciences  
▲ Group\_2-Building\_a\_Component  
▲ Group\_2-Building\_a\_Component-Solution


- Download exercises from:  
**[tinyurl.com/KNIME-Pros-Stuff](https://tinyurl.com/KNIME-Pros-Stuff)**

OR


- Search **hub.knime.com** :  
“KNIME Pros Learnathon Group 2”

# Life Sciences Verified Components


**Molecular Properties Filter**




**PDB Data Extractor**




**ChEMBL Bioactivities**




**Download FASTQ files from ENA**




**ChEMBL Structure Search**




**PDB Chemical Structure Search**



**Pathway Enrichment Analysis**







## Life Sciences




Available on [hub.knime.com](https://hub.knime.com)

Home > 00\_Components > Life Sciences

←

-  **ChEMBL Bioactivities**
-  **ChEMBL Details**
-  **ChEMBL Lookup**
-  **ChEMBL Structure Search**

## Life Sciences

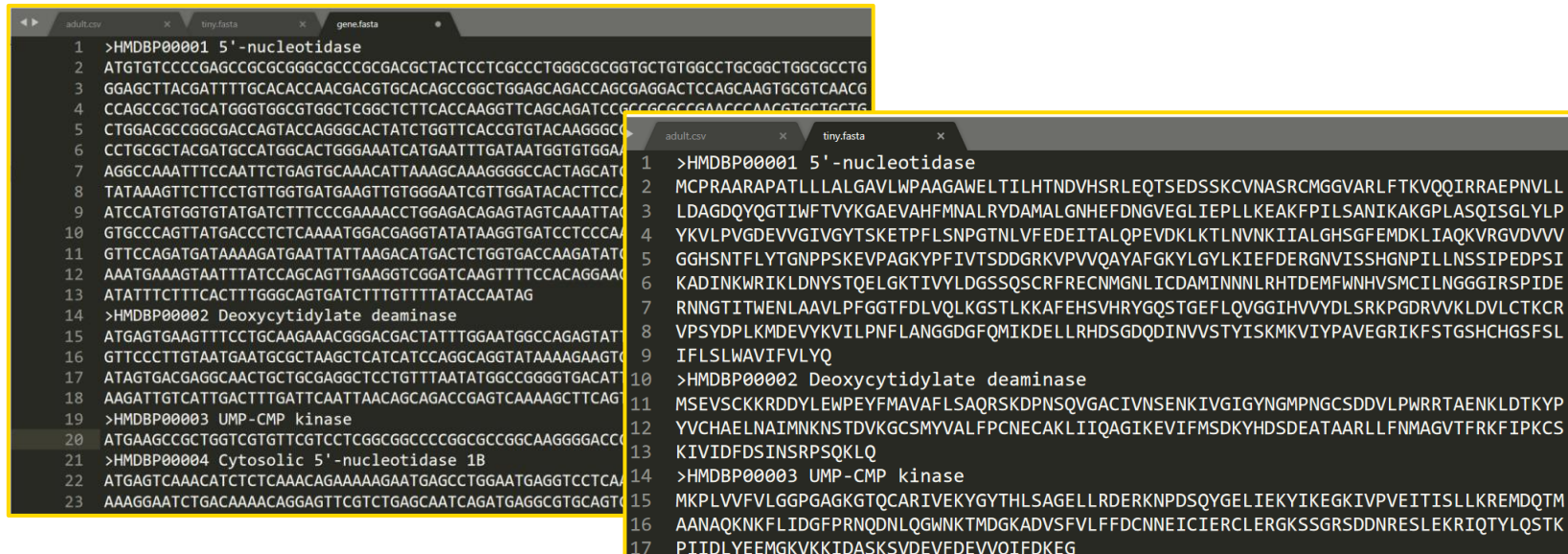


Gather and analyse Life Science data with shared Components. For example, extract data from the [European Nucleotide Archive](#), [ChEMBL](#) or [PDB](#) or perform a [Pathway Enrichment Analysis](#) by simply dragging and dropping the Component of your choice.

[View on KNIME Hub](#)

# FASTA File Format

- common file format in bioinformatics and biochemistry
- text-based format for representing either nucleotide sequences or amino acid (protein) sequences
- allows for sequence names and comments to precede the sequences

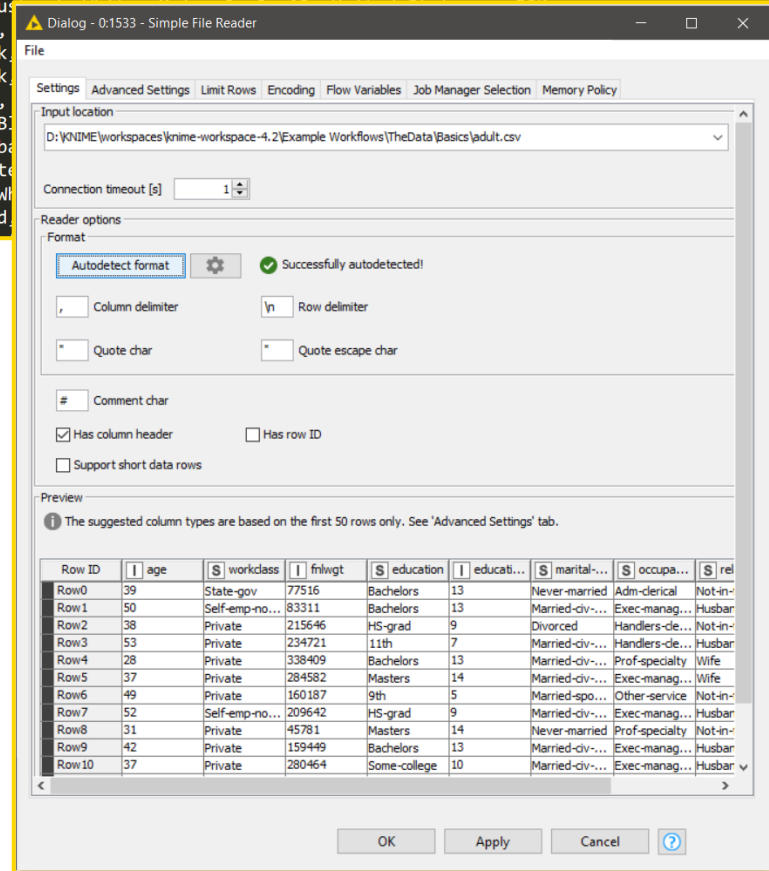


```
1 >HMDBP00001 5'-nucleotidase
2 ATGTGTCCTCCGAGCCGCGCGGGCGCCCGGACGCTACTCCTCGCCCTGGGCGCGGTGCTGTGGCTGCGGCTGGCGCCTG
3 GGAGCTTACGATTTTGCACACCAACGACGTGCACAGCCGGCTGGAGCAGACCAGCGAGGACTCCAGCAAGTGCCTCAACG
4 CCAGCCGCTGCATGGGTGGCGTGGCTCGGCTCTTCACCAAGGTTCAAGCAGATCCGCGCGGCGGCAAGGCGTGGCTGCTG
5 CTGGAGCGCCGCGGACCACTACCAAGGGCACTATCTGGTTCAACCCTGTACAAGGGG
6 CCTGCGCTACGATGCCATGGCACTGGGAAATCATGAATTTGATAATGGTGTGGAA
7 AGGCCAAATTTCCAATTCTGAGTGCAAAACATTAAAGCAAAGGGGCCACTAGCATC
8 TATAAAGTTCTTCTGTTGGTGATGAAGTTGTGGGAATCGTTGGATACACTTCCA
9 ATCCATGTGGTGATGATCTTTCCCGAAAACCTGGAGACAGAGTAGTCAAATTA
10 GTGCCAGTTATGACCTCTCAAAATGGACGAGGTATATAAGGTGATCTCCCAA
11 GTTCCAGATGATAAAAGATGAATTATTAAGACATGACTCTGGTGACCAAGATATC
12 AAATGAAAGTAATTTATCCAGCAGTTGAAGTCGGATCAAGTTTTCACAGGAAG
13 ATATTTCTTTCACTTTGGGCGAGTGATCTTTGTTTATACCAATAG
14 >HMDBP00002 Deoxycytidylate deaminase
15 ATGAGTGAAGTTTCTGCAAGAAACGGGACGACTATTTGGAATGGCCAGAGTAT
16 GTTCCCTTGAATGAATGCGCTAAGCTCATCATCCAGGCAGGTATAAAAGAAGT
17 ATAGTGACGAGGCAACTGCTGCGGAGGCTCCTGTTTAAATAGGCGGGGTGACAT
18 AAGATTGTCAATTGACTTTGATTCAATTAAACAGCAGACCGAGTCAAAGCTTCAG
19 >HMDBP00003 UMP-CMP kinase
20 ATGAAGCCGCTGGTGGTCTGCTGCTCGGCGGCGCCGCGGCAAGGGGACCG
21 >HMDBP00004 Cytosolic 5'-nucleotidase 1B
22 ATGAGTCAAAACATCTCTCAAAACAGAAAAGAAATGAGCCTGGAATGAGGCTCTCA
23 AAAGGAATCTGACAAAACAGGAGTCTGCTGAGCAATCAGATGAGGCGTGCACT
```

```
1 >HMDBP00001 5'-nucleotidase
2 MCPRAARAPATLLALGAVLWPAAGAWELTILHTNDVHSRLQTSSESSKCVNASRCMGGVARLFTKVQQIRRAEPNVLL
3 LDAGDQYQGTIWFVYKGAEVAHFNMALRYDAMALGNHEFDNGVEGLIEPLLKEAKFPILSANIKAKGPLASQISGLYLP
4 YKVLPGDEVVGVGYTSKETPFLSNPGTNLVEDEITALQPEVDKLTNLNVNKIIALGHSGFEMDKLIAQKVRGVDVVV
5 GGHSNTFLYTGNNPSKEVPAGKYPFIVTSDDGRKVPVQYAFGKYLGYLKEIFDERGNVSSHGNPILLNSSIPEDPSI
6 KADINKWRIKLDNYSTQELGKTIYVLDGSSQSCRFRFCNMGNLICDAMINNNLRHTDEMFWNHVSMCILNGGGIRSPIDE
7 RNNGTITWENLAALVPGGTFDLVQLKGSTLKKAFEHSVHRYGQSTGEFLQVGGIHVVYDLSRKPGRDVRVKLDVLCTKCR
8 VPSYDPLKMDEVYKVIILPNFLANGGDGFMKIDELLRHDSGDQDINVVSTYISKMKVIYPAVEGRIFKFTSGSHCHGSFSL
9 IFLSLWAVIFVLYQ
10 >HMDBP00002 Deoxycytidylate deaminase
11 MSEVSCCKRDDYLEWPEYFMAVAFLSAQRSKDPNSQVGACIVNSENKIVGIGYNGMPNGCSDDVLPWRRTAENKLDTKYP
12 YVCHAEALNAIMNKNSDVKGCMSYVALFPCNECAKLIIQAGIKEVIFMSDKYHDSDEATAARLLFNMAAGVTFRKFIPKCS
13 KIVIDFDSINSRPSQKLQ
14 >HMDBP00003 UMP-CMP kinase
15 MKPLVVFVLGGPGAGKGTQCARIVEKYGYTHLSAGELLRDERKNPDSQYGELIEKYIKEGKIVPEITISLLKREMDQTM
16 AANAQKNKFLIDGFPRNQDNLQGWNKTMGDKADVSVFLFFDCNNEICIERCLERGKSSGRSDDNRESLEKRIQTYLQSTK
17 PIIDLYEEMGKVKKIDASKSVDEVFDEVQIFDKEG
```

# KNIME Nodes (File Readers)

```
adult.csv      X      tiny.fasta      X
1 age,workclass,fnlwgt,education,education-num,marital-status,occupation,relationship,race,sex,capital-gain,capital-loss,hours-per-week,native-country,income
2 39, State-gov, 77516, Bachelors, 13, Never-married, Adm-clerical, Not-in-family, White, Male, 2174, 0, 40, United-States, <=50K
3 50, Self-emp-not-inc, 83311, Bachelors, 13, Married-civ-spouse, Exec-managerial, Husband, White, Male, 1562, 0, 45, United-States, <=50K
4 38, Private, 215646, HS-grad, 9, Divorced, Handlers-cleaners, Not-in-family, White, Male, 1601, 0, 40, United-States, <=50K
5 53, Private, 234721, 11th, 7, Married-civ-spouse, Handlers-cleaners, Husband, Black, Male, 1501, 0, 40, United-States, <=50K
6 28, Private, 338409, Bachelors, 13, Married-civ-spouse, Prof-specialty, Wife, Black, Female, 1501, 0, 40, United-States, <=50K
7 37, Private, 284582, Masters, 14, Married-civ-spouse, Exec-managerial, Wife, White, Female, 1501, 0, 40, United-States, <=50K
8 49, Private, 160187, 9th, 5, Married-spouse-absent, Other-service, Not-in-family, Black, Male, 1501, 0, 40, United-States, <=50K
9 52, Self-emp-not-inc, 209642, HS-grad, 9, Married-civ-spouse, Exec-managerial, Husband, White, Male, 1501, 0, 40, United-States, <=50K
10 31, Private, 45781, Masters, 14, Never-married, Prof-specialty, Not-in-family, White, Male, 1501, 0, 40, United-States, <=50K
11 42, Private, 159449, Bachelors, 13, Married-civ-spouse, Exec-managerial, Husband, White, Male, 1501, 0, 40, United-States, <=50K
12 37, Private, 280464, Some-college, 10, Married-civ-spouse, Exec-managerial, Husband, White, Male, 1501, 0, 40, United-States, <=50K
```



File Reader



CSV Reader



CSV Reader (Labs)



Simple File Reader



- Well suited to read tabular data

# FASTA File as Tabular Data

```

1 >HMDBP00001 5'-nucleotidase
2 MCPRAARAPATLLALGAVLWPAAGAWELTILHTNDVHSRLEQTSSESSKCVNASRCMGGVARLFTKVQQIRRAEPNVLL
3 LDAGDQYQGTIWFVYKGAEVAHFMMALRYDAMALGNHEFDNGVEGLTEDLLKEAKERTLSAMTKAKGDLASOTSGIYLD
4 YKVLPGDEVVGIVGYTSKETPFLSNPGTNLVFEDEITALQPEV
5 GGHSNTFLYTGNNPSKEVPAGKYPFIVTSDGRKVPVVQAYAFG
6 KADINKWRIKLDNYSTQELGKTIYVLDGSSQSCRRECNGMNLIT
7 RNNGTITWENLAALVPFGGTFDLVQLKGSTLKKAFEHSVHRYGQ
8 VPSYDPLKMDEVYKVLIPNPLANGGDFQMIKDELLRHDSGDQD
9 IFLSLWAVIFVLYQ
10 >HMDBP00002 Deoxycytidylate deaminase
11 MSEVSCKKRDDYLEWPEYFMAVAFLSAQRSKDPNSQVGACIVNS
12 YVCHAEINAIMNKNSTDVKGCSMYVALFPCNECAKLIIQAGIKE
13 KIVIDFDSINSRPSQKLQ
14 >HMDBP00003 UMP-CMP kinase
15 MKPLVVFVLGGPGAGKGTQCARIVEKYGYTHLSAGELLRDERKN
16 AANAQKNKFLIDGFPNRQDNLQGWKNTMDGKADVSVLFFDCNN
17 PIIDL YEEMGKVKKIDASKSVDEVFDEVVQIFDKEG

```

File Table - 4:1523:264 - Simple File Reader

File Edit Hilite Navigation View

Table "default" - Rows: 22 Spec - Columns: 34 Properties Flow Variables

Row ID	S Column0	S Column1	S Column2	S Column3	S Column4
Row0	HMDBP00001 5'-nucleotidase	MCPRAARAPATLLALGAV...	LDAGDQYQGTIWF...	YKVLPGDEVVGIVGYTSK...	GGHSNTFLYTGNNPS
Row1	HMDBP00002 Deoxycytidylat...	MSEVSCKKRDDYLEWPEYF...	YVCHAEINAIMNKN...	KIVIDFDSINSRPSQKLQ	?
Row2	HMDBP00003 UMP-CMP kinase	MKPLVVFVLGGPGAGKGTQ...	AANAQKNKFLIDGFP...	PIIDL YEEMGKVKKIDASKS...	?
Row3	HMDBP00004 Cytosolic 5'-nu...	MSQTSLKQKKNEPGMRSSK...	RIYIHMCLLWEQGG...	PSLHDSPPPLSGQPSLPQ...	TEWKSSSQRRGIYP
Row4	HMDBP00005 Cytosolic 5'-nu...	MEPGQPREPQEPREPQPG...	RYQLEHENEFPSPGP...	CYLKAYHTNLVLSADAQK...	ENKPLAQGPLKGFLE
Row5	HMDBP00006 5'(3')-deoxyrib...	MARSVRVLVMDGVLADFE...	LDVAVREMDLDPDTQV...	ILFTCCHNRHLVLPPTRRR...	?
Row6	HMDBP00007 Deoxycytidine ...	MATPKRSCPSFSASSEGT...	VLQMMYEKPERWSF...	WMNNQFGQSLELDGIIYL...	DFKDKEYSLVEKVE
Row7	HMDBP00008 5'(3')-deoxyrib...	MIRLGGWCARRLCSAAVP...	YGRRLRPLSEKAISI...	LTRDKTVVSADLLIDDRPD...	?
Row8	HMDBP00009 Hydroxymethyl...	MAAMRKALPRRLVGLASL...	WVPQMGDHEVLKG...	NISVRGYVSCALGCPYEGK...	ALANTLMALQMGVS
Row9	HMDBP00010 ATP-ditrate sy...	MSAKAISEQTGKELLYKFC...	VKSWLKPRLQGEATV...	NPEIDIKHLLVHAPEDKKEI...	PFGREAYPEEAYIAD
Row10	HMDBP00011 Histone acetyl...	MAENVVEPGPPSAKRPKLS...	LSELLRSGSSPNLNM...	PAMGMNTGMNAGMNP...	PLKMGMMNPNPYG
Row11	HMDBP00012 Pyruvate dehy...	MAAVSGLVRRPLREVSGLL...	RIIDTPISEMGFAGIA...	CFAAWYGHCPGLKVVPSP...	SHSRPVGHCLAAAI
Row12	HMDBP00013 Acetyl-CoA ac...	MNAGSDPVVIVSAARTIIGS...	YSVPAWSCQMIGCS...	GITAENVAKKQVQVSREDQ...	DGTGTVTPANASGI
Row13	HMDBP00014 CREB-binding p...	MAENLLDGPNNPKRAKLS...	SINPGIGNVASSPV...	LATSPATSQTGPICMN...	ETLTQVSPQMTGHA
Row14	HMDBP00015 Diamine acetyl...	MASVRIREAKEGDCGDLR...	YSTWKGRITYLEDIYV...	GEATRKLAKG	?
Row15	HMDBP00016 Malonyl-CoA d...	MRFGGPLTARRLLPLRLP...	TAQRAELLGRLARGF...	LLEAQALKVEGPDVREMN...	SHCSTPGEPLVVLHV
Row16	HMDBP00017 5-aminolevulin...	MVTAAMLLQCCPVLAGPT...	KSKIVQKAAPEVQED...	TYRVFKTVNRWADAYPFA...	QELAEHLQKDSALLF
Row17	HMDBP00018 Acetyl-CoA car...	MVLLCLSLCFLSCLTFSWL...	GPKDAGRNRSLPPS...	LMTNFILGSFDDYSSDEDS...	DFTVASPAEFVTRFG

SequenceID

← Sequence →

- Re-think Row/Column delimiters

- “>” Row delimitr
- “\n” Column delimiter

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tinyurl.com/KNIME-Pros-Stuff

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**KNIME**



# Today's Challenge: FASTA Reader Component

- Download Exercises from **link** and import .knar file to your KNIME Analytics Platform LOCAL Workspace!

OR

- Download from KNIME Hub!

[hub.knime.com/knime/spaces/Education/latest/Learnathons](https://hub.knime.com/knime/spaces/Education/latest/Learnathons)  
or Search for “**KNIME Pros Learnathon - Group 2**”

