Stepwise guideline for installation of K-PAM standalone version in Ubuntu

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1. Download the compressed file from the link given on the page: <u>http://iith.ac.in/K-PAM/sk_pam.php</u>

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K-PAM: <i>Klebsiella</i> specie	s serotype predictor	and surface an	tigens modeler	
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rotype predictor Virulence predictor K-PAM Standalor	ne version K-antigen structures O Documentation	-antigen structures Multi	mer generation ▼ Schematic ge	enerator
	K-PAM stand-alone vers	ion		
A standalone version (application program interface (API)) of files correspond to Mac and Linux operating system. This si generates the serotype prediction summary in the CSV file for	K-PAM is also available for analyzing the Landalone version has the K-typing, O-t nat.	large datasets. It has the opt yping and hypervirulent stra	on to upload multiple files. The ex in identification features. The K-	xecutable ·PAM API
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For mac OS: KPAM_for_ubuntu.tar.gz For max OS: KPAM_for_ubuntu.tar.gz				
TR Lab. Departme	© Indian Institute of Technology Hyderab nt of Biotechnology, Indian Institute of T	ad, India echnology Hyderabad, India		
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	Opening KPAM_for_ubuntu.tar.gz	8		
	You have chosen to open: KPAM_for_ubuntu.tar.gz which is: Gzip archive (2.3 MB) from: http://iith.ac.in			
	What should Firefox do with this file?	_		
A standalone version (application program interface (API)) (files correspond to Mac and Linux operating system. This generates the serotype prediction summary in the CSV file fo	Open with Archive Manager (default) Osave File Do this automatically for files like this from now of the second	v ts. It has the op ypervirulent str	tion to upload multiple files. The ain identification features. The F	executable K-PAM API
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 For mac OS: <u>KPAM_for_mac.tar.gz</u> For linux OS: <u>KPAM_for_ubuntu.tar.gz</u> 				

2. Move the downloaded *tar* file from the default download directory ("Downloads" folder) to the desired location.



3. Extract the content in the same location by double click on the *tar* file or if you are doing it from terminal give any of the following commands.

\$ tar -xvfz KPAM_for_ubuntu.tar.gz \$ tar -xvf KPAM_for_ubuntu.tar.gz

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4. Go inside the extracted directory "KPAM_for_ubuntu". A instruction manual for KPAM installation is also given in a text file ("README") or you can follow the steps written in this document.



5. Open the terminal for installation of K-PAM in local machine. Then, go to the "KPAM_for_ubuntu" directory. You can also directly go to terminal from the folder "KPAM_for_ubuntu" by clicking right button on mouse any black space in the directory and select "Open in Terminal".



- 6. To run the installation, the account must be a *sudo* user (not applicable if the *sudo* user account is not setup and the system has only one account) and confirm that the current directory is inside the directory "KPAM_for_ubuntu".
- 7. Give the command to start the installation:

\$ bash setup1.sh



- 8. Give the *sudo* user password or the password used for log in to the system (if the *sudo* user account is not setup and the system has only one account).
- During installation, the user permission would be asked sometimes to continue the installation, like "*Press [ENTER] to continue*" or "*select <OK> to continue*". Click [ENTER] or select <OK> and click [ENTER] to continue the installation process.

ŵ	File Edit View Search Terminal Help		
	Reporting Issues		
□ ⊹	Issues can be reported in the master issue tracker at: https://github.com/deadsnakes/issues/issues		
99	Supported Ubuntu and Python Versions		
ø			
, Î	- Ubuntu 16.04 (xenial) Python 2.3 - Python 2.6, Python 3.1 - Python3.4, Python 3.6 - Python3.9 - Ubuntu 18.04 (bionic) Python2.3 - Python 2.6, Python 3.1 - Python 3.5, Python3.7 - Python3.9 - Note: Python2.7 (both), Python 3.5 (xenial), Python 3.6 (bionic) are not provided by deadsnakes as upstream ubuntu provides those pack ages.		
	The packages may also work on other versions of Ubuntu or Debian, but that is not tested or supported.		
	Packages =======		
	The packages provided here are loosely based on the debian upstream packages with some modifications to make them more usable as non-def ault pythons and on ubuntu. As such, the packages follow debian's patterns and often do not include a full python distribution with jus t `apt install python#.#`. Here is a list of packages that may be useful along with the default install:		
Terr	 `python#.#-dev`: includes development headers for building C extensions inalpython#.#-venv`: provides the standard library `venv` module `python#.#-distutlis`: provides the standard library `distutlis` module `python#.#-lib2to3`: provides the `2to3-#.#` utility as well as the standard library `lib2to3` module `python#.#-gdbm`: provides the standard library `gnu.dbm` module `python#.#-tki: provides the standard library `kinter` module 		
	Third-Party Python Modules		
	Python modules in the official Ubuntu repositories are packaged to work with the Python interpreters from the official repositories. Acc ordingly, they generally won't work with the Python interpreters from this PPA. As an exception, pure-Python modules for Python 3 will w ork, but any compiled extension modules won't.		
	To install ord-party Python modules, you should use the common Python packaging tools. For an introduction into the Python packaging ec osystem and its tools, refer to the Python Packaging User Guide: https://pack.ging.python.org/installing/		
	Sources		
	The package sources are available at: https://github.com/deadsnakes/ Mace_dmcv.mittps./jiceuchpad.net/~deadsnakes/+archive/ubuntu/ppa		
	Press [ENTER] to continue of Ctrl-c to cancel adding it.		



10. When first part of the installation completes, a message stating following will appear in terminal.

"Download the latest vesrsion of ncbi blast+ (ncbi-blast-2.10.0+x64-linux.tar.gz, as of December 2019) file from ftp://ftp.ncbi.nlm.nih.gov/blast/executables/blast+/LATEST/ using a web browser for installing blast+ in the local machine. Then copy the downloaded file to current location and execute setup2.sh"

[Note: if

ftp://ftp.ncbi.nlm.nih.gov/blast/executables/blast+/LATEST/ is giving error then use following address:

http://ftp.ncbi.nlm.nih.gov/blast/executables/blast+/LATEST/

Documentation X Documentation X Sindex of /blast/et	recutabl X +		
$ \longleftrightarrow \Rightarrow $	les/blast+/I ATEST/		〒 (200%) ••• ▽ ☆
Index of /blast/e	xecutal	oles	s/blast+/LATEST
Name	Last modified	Size	
Parent Directory		-	
ChangeLog	2019-12-03 21:52	85	
ncbi-blast-2.10.0+-4.src.rpm	2019-12-03 21:50	19M	
<pre>ncbi-blast-2.10.0+-4.src.rpm.md5</pre>	2019-12-03 21:53	63	
<u>ncbi-blast-2.10.0+-4.x86_64.rpm</u>	2019-12-03 21:50	175M	
<u>ncbi-blast-2.10.0+-4.x86_64.rpm.md5</u>	2019-12-03 21:53	66	
<u>ncbi-blast-2.10.0+-src.tar.gz</u>	2019-12-03 21:53	24M	
<u>ncbi-blast-2.10.0+-src.tar.gz.md5</u>	2019-12-03 21:53	64	
<u>ncbi-blast-2.10.0+-src.zip</u>	2019-12-03 21:53	29M	
<u>ncbi-blast-2.10.0+-src.zip.md5</u>	2019-12-03 21:53	61	
<u>ncbi-blast-2.10.0+-win64.exe</u>	2019-12-03 21:49	87M	
<u>ncbi-blast-2.10.0+-win64.exe.md5</u>	2019-12-03 21:53	63	
<u>ncbi-blast-2.10.0+-x64-linux.tar.gz</u>	2019-12-03 21:52	222M	
<pre>ncbi-blast-2.10.0+-x64-linux.tar.gz.md5</pre>	2019-12-03 21:53	70	
<pre>ncbi-blast-2.10.0+-x64-macosx.tar.gz</pre>	2019-12-03 21:53	141M	
<pre>ncbi-blast-2.10.0+-x64-macosx.tar.gz.md5</pre>	2019-12-03 21:53	71	
<u>ncbi-blast-2.10.0+-x64-win64.tar.gz</u>	2019-12-03 21:50	86M	
<u>ncbi-blast-2.10.0+-x64-win64.tar.gz.md5</u>	2019-12-03 21:53	70	
<u>ncb1-blast-2.10.0+.dmg</u>	2019-12-03 21:52	143M	
<u>ncbi-blast-2.10.0+.dmg.md5</u>	2019-12-03 21:53	57	



Index of /blast/executables/blast+/LATI

Name	Last modified Size
Parent Directory	
<u>ChangeLog</u>	2019-12 03 21.52 85
<pre>ncbi-blast-2.10.0+-4.src.rpm</pre>	2019 - Opening http://diasc-2.10.04-x84-linux.car.gz
<pre>ncbi-blast-2.10.0+-4.src.rpm.md5</pre>	2019 - You have chosen to open:
ncbi-blast-2.10.0+-4.x86 64.rpm	2019 - 🔤 ncbi-blast-2.10.0+-x64-linux.tar.gz
ncbi-blast-2.10.0+-4.x86_64.rpm.md5	2019 - Krom: https://ftp.ncbi.plm.pib.gov
ncbi-blast-2.10.0+-src.tar.gz	2019 - https://ftp.pchi.olm.phi.org/blast/everytables/blast/
ncbi-blast-2.10.0+-src.tar.gz.md5	2019 - What should Firefox do with this file?
ncbi-blast-2.10.0+-src.zip	2019 - Open with Archive Manager (default)
ncbi-blast-2.10.0+-src.zip.md5	2019 - Save File
ncbi-blast-2.10.0+-win64.exe	2019 - Do this automatically for files like this from now on.
ncbi-blast-2.10.0+-win64.exe.md5	2019-
ncbi-blast-2.10.0+-x64-linux.tar.gz	2019- Cancel OK
ncbi-blast-2.10.0+-x64-linux.tar.gz.md5	2019-12-03 21.33 70
ncbi-blast-2.10.0+-x64-macosx.tar.gz	2019-12-03 21:53 141M
ncbi-blast-2.10.0+-x64-macosx.tar.gz.md5	2019-12-03 21:53 71
ncbi-blast-2.10.0+-x64-win64.tar.gz	2019-12-03 21:50 86M
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 Transfer the downloaded *tar* file of blast+ from the default download directory ("Downloads" folder) to the K-PAM folder ("KPAM_for_ubuntu" folder).



12. Ensure that only one blast+ tar file is there in the folder. Run the following command in the terminal to complete the final part of the installation.

\$ bash setup2.sh



13. Once the installation of K-PAM completes, the following message will appear in the terminal.



14. Now, just type "kpam" in the terminal from any location, the GUI window will pop-up. Submit your query sequences to predict serotype or to identify virulence of the strains corresponding to submitted query sequences.

