Python VS R: Comparison Chart for Data Processing

	Python	R
Usability and Flexibility	• more natural for people with a software engineering background • easy to code and debug • indentation matters • any piece of functionality is always written the same way with Python • flexible for creating something that has never been done before. • requires users to install packages for data analysis, and these packages have greatly improved in recent years	• easier to learn if you have no coding experience • the same piece of functionality can be written in several ways with R • easy to use complex functions in R, since all kinds of statistical tests and models are readily available and easily used • handle basic data analysis without needing to install packages.
Advantages	• general-purpose programming languages are useful beyond just data analysis • great for mathematical computation and learning how algorithms work	• best tool for making beautiful graphs and visualizations • has many functionalities for data analysis
Dis-advantages	• doesn't have as may libraries • visualizations are more convoluted, and results are not as eye-pleasing or informative	• finding the right packages to use may be time consuming • there are many dependencies between R libraries • not as popular for deep learning and NLP
Popular Libraries and Packages	• Pandas to easily manipulate data • SciPy and NumPy for scientific computing • Scikit-learn for machine learning • Matplotlib and seaborn to make graphics • statsmodels to explore data, estimate statistical models, and perform statistical tests and unit test	• dplyr,tidyr and data.table to easily manipulate data • stringr to manipulate strings • zoo to work with regular and irregular time series • ggplot2 to visualize data • caret for machine learning

import	
import library_name	library("package_name")
tips: (python)must put package name when calling functions (r)package_name::function is necessary only when the namespace collisions.	

read dataset	
df = pd.read_csv(file.csv/url)	data(df) df <- read.csv("file.csv")
tips: In r, some packages require us to load the data separately, while for others we can directly use the data	

get dataset information	
df.describe()	summary(df), str(df)

get row/column information	
df.columns, pd.index	colnames(df), rownames(df)
df.shape[1], df.shape[0] or len(df.columns), len(df.index)	ncol(df), nrow(df), dim(df),

create a dataframe	
df = pd.DataFrame(*[["val1", "val2"],["val3", "val4"]]*, columns = [*col1,col2*])	df <- data.frame (*col1*=c("*val1*", "*val2*"), *col2*=c("*val3*", "*val4*"))

change index names	
df.set_index(*"index column, e.g Columbia UNI"*, inplace = True)	rownames(df) <- df$*"Columbia UNI"*

change data type	
df['*column*']=df.column.astype('*float*')	df$col1 <- as.*numeric*(df$col1)

slicing/subset	
df.loc([*col1,col2*], [*row1,row2*]); df.iloc(*m:n, j:k)*	df[c(*row1,row2*), c(*col1,col2*)], df[*m:n,j:k*]
tips: col1, row1,etc. are variable names. m, n, j, k are indices.	

merge and concat	
merge: pd.merge(*df1*, *df2*, on='*key*') concat: pd.concat([*df1*, *df2*,..], join='*outer*')	merge: merge(*df1*, *df2*, by = '*key*') concat: cbind(*df1*, *df2*), rbind(*df1*, *df2*)

group by	
df.groupby(by=["*col*"]).*sum()*	df %>% group_by(*col*) %>% summarize(*Count = n()*)

filter/query	
df[*boolean conditions*] df.query(*boolean conditions*)	df[*boolean conditions*]; filter(df,*boolean conditions*)

sort	
df.sort_values(by=['*col1*'], ascending=False)	df[order(df$*col1*),]

handle NAs	
df.dropna()	na.omit(df)

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References: https://www.datacamp.com/community/tutorials/r-or-python-for-data-analysis ; https://shiring.github.io/r_vs_python/2017/01/22/R_vs_Py_post