	(36848, 12) 1. Find rows with missing null / na - no value NaN - Not a Number - the value isnull() is a pandas function, so	is missing. This value w	_			
[29]: t[29]:	location iso_code date total_vaccinations people_vaccinated people_fully_vaccinated daily_vaccinations_raw daily_vaccinations	0 0 0 15328 16203 19253 18696 277				
n [4]: ut[4]:	total_vaccinations_per_hun people_vaccinated_per_hund people_fully_vaccinated_pe daily_vaccinations_per_mil month dtype: int64 call it through pandas: pd.isnull(vacc_df).sum() location	lred 16203 er_hundred 19253 lion 277 0				
. [5].	iso_code date total_vaccinations people_vaccinated people_fully_vaccinated daily_vaccinations_raw daily_vaccinations total_vaccinations_per_hun people_vaccinated_per_hund people_fully_vaccinated_pe daily_vaccinations_per_mil dtype: int64 View specific columns:	r_hundred 13585 lion 241				
n [5]: ut[5]: n [6]: ut[6]:	<pre>vacc_df[['daily_vaccination daily_vaccinations 2466 total_vaccinations 1495 dtype: int64 vacc_df[['daily_vaccination daily_vaccinations 241 dtype: int64 Using numpy: isnan is a numpy</pre>	ons']].isnull().su		().sum()		
n [7]: ut[7]: n [6]:	np.isnan(vacc_df[['daily_daily_daily_vaccinations 241 dtype: int64 2. Remove missing value Look at Zimbabwe for example. Zimbabwe = vacc_df.loc[value daily_vaccinations 241 dtype: int64	ues using dropn Zimbabwe contains mi	a() ssing values:			
n [9]: ut[9]: [10]: t[10]:	<pre>zimbabwe[['total_vaccinat total_vaccinations</pre>					
[11]: t[11]:	We can see the difference when of zimbabwe.count() location iso_code date total_vaccinations people_vaccinated people_fully_vaccinated daily_vaccinations_raw daily_vaccinations	110 110 110 107 107 78 105	of values per row:			
[12]: t[12]:	daily_vaccinations total_vaccinations_per_hun people_vaccinated_per_hund people_fully_vaccinated_pe daily_vaccinations_per_mil dtype: int64 Remove all rows that contain one zimbabwe.dropna()	e or more missing valu		_fully_vaccinated daily_va	nccinations_raw dail	y_vaccinations
	24823 Zimbabwe ZWE 2021 03-2 24824 Zimbabwe ZWE 2021 03-2 24825 Zimbabwe ZWE 2021 03-2 24826 Zimbabwe ZWE 2021 03-2 24827 Zimbabwe ZWE 2021 03-2	2 43574.0 - 45197.0 - 51893.0 - 58987.0	43294.0 44135.0 49404.0 54892.0 61093.0	280.0 1062.0 2489.0 4095.0 6569.0	845.0 1623.0 6696.0 7094.0 8675.0	845.0 807.0 1755.0 2712.0 3711.0
		3 1048504.0 - 1056238.0 - 1061951.0	 684164.0 685564.0 686636.0 687321.0	364340.0 370674.0 375315.0 380786.0	 8290.0 7734.0 5713.0 6156.0	 13588.0 11349.0 8498.0 8019.0
[13]:	24900 Zimbabwe ZWE 2021 06-07 78 rows × 12 columns Note: dropna(), like most other so you should assign it back if you zimbabwe.head()	7 functions in the panda	688696.0 as API returns a new Dat	385275.0 aFrame (a copy of the or	5864.0 riginal with changes	7699.0) as the result,
t[13]:	Iocation iso_code date 24791 Zimbabwe ZWE 2021 02-13 24792 Zimbabwe ZWE 2021 02-13 24793 Zimbabwe ZWE 2021 02-23 24794 Zimbabwe ZWE 2021 02-23 24794 Zimbabwe ZWE 2021 02-23	- 0.0 9 NaN 0 NaN	eople_vaccinated people 0.0 NaN NaN NaN	NaN NaN NaN NaN NaN	NaN NaN NaN NaN NaN	y_vaccinations NaN 328.0 328.0 328.0
n [7]: ut[7]:	2021	a () e total_vaccinations po		NaN _fully_vaccinated daily_va	NaN nccinations_raw dail	
	36719 Zimbabwe ZWE 2021	2 43574.0 2 45197.0 3 45197.0 5 58987.0	43294.0 44135.0 49404.0 54892.0 61093.0	280.0 1062.0 2489.0 4095.0 6569.0	845.0 1623.0 6696.0 7094.0 8675.0	845.0 807.0 1755.0 2712.0 3711.0
	36843 Zimbabwe ZWE 2021 07-2 36844 Zimbabwe ZWE 2021 07-2 36845 Zimbabwe ZWE 2021 07-2 36846 Zimbabwe ZWE 2021 07-2	4 2116664.0 - 2127402.0 - 2178709.0	 1438890.0 1447342.0 1491493.0 1522150.0	 677774.0 680060.0 687216.0 694685.0	 44604.0 10738.0 51307.0 38126.0	 49319.0 48838.0 50153.0 45643.0
[15]: t[15]:	36847 Zimbabwe ZWE 2021 07-2. 127 rows × 12 columns Remove all values for a specific columbabwe.dropna (subset = location iso_code date)	column - using subset	ns'])	713131.0 fully_vaccinated daily_va	58581.0	46563.0 y_vaccinations
	24791 Zimbabwe ZWE 2021 02-1 24795 Zimbabwe ZWE 2021 02-2 24796 Zimbabwe ZWE 2021 02-2 24797 Zimbabwe ZWE 2021 02-2 24798 Zimbabwe ZWE 2021 02-2 24798 Zimbabwe ZWE 2021 02-2	8 0.0 2 1314.0 2 4041.0 3 7872.0	0.0 1314.0 4041.0 7872.0 11007.0	NaN NaN NaN NaN	NaN NaN 2727.0 3831.0	NaN 328.0 808.0 1312.0
	02-2	5 11007.0 	684164.0 685564.0 686636.0 687321.0	364340.0 370674.0 375315.0 380786.0	3135.0 8290.0 7734.0 5713.0	13588.0 11349.0 8498.0
[16]:	24900 Zimbabwe ZWE 2021 06-01 107 rows × 12 columns For more columns: zimbabwe.dropna(subset =	6 - 1073971.0 7 1073971.0	688696.0	385275.0	5864.0).head()	7699.0
t[16]:	Iocation iso_code date 24795 Zimbabwe ZWE 2021 02-2 24796 Zimbabwe ZWE 2021 02-2 24797 Zimbabwe ZWE 2021 02-2 24798 Zimbabwe ZWE 2021 02-2 24798 Zimbabwe ZWE 2021 02-2	1314.0 2 1314.0 3 4041.0 4 7872.0	1314.0 4041.0 7872.0	NaN NaN NaN NaN NaN	NaN 2727.0 3831.0 3135.0	328.0 808.0 1312.0
	.notnull() - displa.dropna() - Remov	ns so far: y rows that contain missay rows that don't contain we rows with missing va	•		1572.0	1750.0
	.dropna(how='.dropna(thresleast 3 non-null v	all') - drops rows on the k) - k how many values) 1) - drop columns ins	nly if all of its columns h non-null values you wa	ntain missing values in th nave NaNs nt to keep (k=3 means tl		
[17]:	 Use .fillna() to fill missing da Whatever value you choose Mean, median, mode This is called <i>imputation</i> Replace all NaNs with 0s vacc_df.fillna(0, inplace vacc_df 					
t[17]:	location iso_code da 0 Afghanistan AFG 202 02- 1 Afghanistan AFG 202 02- 2 Afghanistan AFG 202 02-	21- 222 0.0 21- 23 NaN 21- 24 NaN	0.0 NaN NaN	le_fully_vaccinated daily_v NaN NaN NaN	NaN NaN NaN	NaN 1367.0 1367.0
	4 Afghanistan AFG 02- 4 Afghanistan AFG 202- 24896 Zimbabwe ZWE 202- 24897 Zimbabwe ZWE 202- 06- 203-	NaN 21- 226 NaN 21- 03 1048504.0 21- 04 1056238.0	NaN NaN 684164.0 685564.0	NaN NaN 364340.0 370674.0	NaN NaN 8290.0 7734.0	1367.0 1367.0 13588.0 11349.0
	24898 Zimbabwe ZWE 202 06- 206- 206- 206- 206- 206- 206- 20	1061951.0 21- 206 1068107.0 21- 207 1073971.0	686636.0 687321.0 688696.0 hange the vacc_df data	375315.0 380786.0 385275.0 frame.	5713.0 6156.0 5864.0	8498.0 8019.0 7699.0
	To change it you need: vacc_df.fillna(0 , in or to assign: vacc_df = vacc_df.fil But we won't do that! This total_vaccinations w	llna(0) is where some busine	ss understanding com	es in: it's not a good idea	ı to fill a column like	2
[18]: t[18]:	See what happens: vacc_df.fillna(0).head(10 location iso_code date to the second date of the second date o		le_vaccinated people_ful 0.0 0.0	lly_vaccinated daily_vaccin 0.0 0.0	nations_raw daily_va 0.0 0.0	0.0 1367.0
	 2 Afghanistan 3 Afghanistan 4 Afghanistan 5 Afghanistan AFG 2021- 02-25 AFG 2021- 02-26 AFG 2021- 02-27 2021- 02-27 	0.0 0.0 0.0 0.0	0.0 0.0 0.0 0.0	0.0 0.0 0.0 0.0	0.0 0.0 0.0	1367.0 1367.0 1367.0
	6 Afghanistan AFG 2021- 02-28 7 Afghanistan AFG 2021- 03-01 8 Afghanistan AFG 2021- 03-02 9 Afghanistan AFG 2021- 03-03	8200.0 0.0 0.0 0.0	0.0 0.0 0.0 0.0	0.0 0.0 0.0 0.0	0.0 0.0 0.0 0.0	1367.0 1580.0 1794.0 2008.0
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	58 Afghanistan 2021-04-21 59 Afghanistan 2021-04-22 60 Afghanistan 2021-04-23 61 Afghanistan 2021-04-24 For the total_vaccinations was vacc_df[['total_vaccinations was vacc_df['total_vaccinations was vacc_df['newTotal'] = vacc_df[NaN 240000.0 NaN NaN ve'll use ffill which what you think it does ons']].fillna (methons'][52:62] night be NaN good enought! We need and is is more advanced and in newTotal - so we can be compared to the constant of the	ed to aggregate by cour id we will return to it shan see the difference in	ntry!! ortly) total_vaccinations		
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[24]: t[24]: t[25]: t[27]: t[27]: t[30]: t[30]: t[31]: t[31]: t[31]:	### Arghanistan 2021-04-21 ### Arghanistan 2021-04-22 ### Arghanistan 2021-04-24 ### For the total_vaccinations will receive and argument and argument ar	NaN	code 'ffill') [52:62] ad to aggregate by cour desire the difference in the strong of	sting dataframe that occurs before it that occurs after it sale (method = 'linear' check way to avoid it is to cre check might be a good thin	ambda x: x.fill: pywarning: /user_guide/ind. eate a copy of the g, if you didn't plan	exing.html# e dataframe: to change it,
t[23]: t[23]: t[24]: t[24]: t[26]: t[27]: t[30]: t[31]: t[31]: t[32]: t[32]: t[33]:	### Aghanistan	NaN 240000.0 NaN NaN NaN We'll use ffill which what you think it does Ons']].fillna (methons')/52:62/ Anight be NaN good enought! We need is is more advanced and newTotal - so we ca af.groupby ('loca ,'12]] NaN 120000.0 NaN 120000.0 NaN 120000.0 NaN 120000.0 NaN 120000.0 NaN 240000.0 NaN 240000.0 NaN 240000.0 NaN 120000.0 NaN 12000.0 NaN 12000.0 NaN 12000.0 NaN 12000.0 NaN 12000.0	code 'ffill') [52:62] and to aggregate by count of the will return to it shall be seen the difference in tion') [['total_vaccolor 192000.000000 200000.000000 200000.000000 224000.000000 224000.000000 224000.000000 224000.000000 232000.00000 232000.000000 232000.00000 232	sting dataframe that occurs before it that occurs after it will cause a warning to a data (method = 'linear') which is to come the man of each of the mean of each come and	ambda x: x.fill: pyWarning: //user_guide/ind eate a copy of the g, if you didn't plan ach country each	exing.html# e dataframe: to change it,
t[23]: t[24]: t[24]: t[25]: t[27]: t[29]: t[30]: t[31]: t[31]: t[32]: t[33]: t[34]:	### Applanistan 2021-04-21 ### Applanistan 2021-04-22 ### Applanistan 2021-04-24 ### For the total_vaccinations will follow the following and in the foll	NaN	code 'ffill') [52:62] and to aggregate by count of the will return to it shall be seen the difference in tion') [['total_vaccolor 192000.000000 200000.000000 200000.000000 224000.000000 224000.000000 224000.000000 224000.000000 232000.00000 232000.000000 232000.00000 232	sting dataframe that occurs before it that occurs after it will cause a warning to a data (method = 'linear') which is to come the man of each of the mean of each come and	ambda x: x.fill: pyWarning: //user_guide/ind eate a copy of the g, if you didn't plan ach country each	exing.html# e dataframe: to change it,
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