import pandas as pd
covidData = pd.read\_csv("Datasets/covid.csv")

Out[2]: Country/Region Continent Population TotalCases NewCases TotalDeaths NewDeaths TotalRecovered NewRecovered ActiveCases Serious, Critical Tot Cases/1M pop Deaths/1M pop TotalTests Tests/1M pop WHO Region 18296.0 USA North America 3.311981e+08 5032179 162804.0 2576668.0 2292707.0 15194.0 492.0 63139605.0 190640.0 NaN NaN NaN Americas Brazil South America 2.127107e+08 2917562 98644.0 2047660.0 771258.0 8318.0 13716.0 464.0 13206188.0 62085.0 1 NaN NaN NaN Americas 2 India Asia 1.381345e+09 2025409 41638.0 NaN 1377384.0 606387.0 8944.0 1466.0 30.0 22149351.0 16035.0 South-EastAsia NaN NaN 14606.0 676357.0 180931.0 2300.0 5974.0 100.0 29716907.0 203623.0 Russia Europe 1.459409e+08 871894 NaN NaN NaN Europe South Africa Africa 5.938157e+07 538184 9604.0 NaN 387316.0 141264.0 539.0 9063.0 162.0 3149807.0 53044.0 Africa NaN NaN

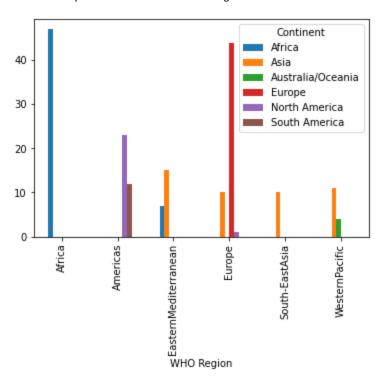
## Expectations

In [2]: covidData.head()

I expect these 2 to be very similar, since theyre almost the same. This dataset didnt have that many non numerical columns, so these were the best i could compare.

In [3]: covidData.groupby(['Continent','WHO Region']).size().unstack('Continent', fill\_value=0).plot(kind='bar')

Out[3]: <AxesSubplot:xlabel='WHO Region'>



In [4]: covidData.groupby(['Continent','WHO Region']).size().unstack('Continent', fill\_value=0)

Out [4]: Continent Africa Asia Australia/Oceania Europe North America South America
WHO Region

	WHO Region						
	Africa	47	0	0	0	0	0
	Americas	0	0	0	0	23	12
	EasternMediterranean	7	15	0	0	0	0
	Europe	0	10	0	44	1	0
	South-EastAsia	0	10	0	0	0	0
	WesternPacific	0	11	4	0	0	0

In [5]: from scipy.stats import chi2\_contingency

In [6]: chi2\_contingency(covidData.groupby(['Continent','WHO Region']).size().unstack('Continent', fill\_value=0))

Out[6]: (479.55382395382395, 1.3152901775915486e-85, , 1.02173913, 11.23913043, 6.13043478, array([[13.79347826, 11.75 3.06521739], [10.27173913, 8.75 , 0.76086957, 8.36956522, 4.56521739, 2.2826087 ], [ 6.45652174, 5.5 , 0.47826087, 5.26086957, 2.86956522, 1.43478261], [16.14130435, 13.75 , 1.19565217, 13.15217391, 7.17391304, 3.58695652], [ 2.93478261, 2.5 , 0.2173913 , 2.39130435, 1.30434783, 0.65217391], [ 4.40217391, 3.75 , 0.32608696, 3.58695652, 1.95652174, 0.97826087]]))

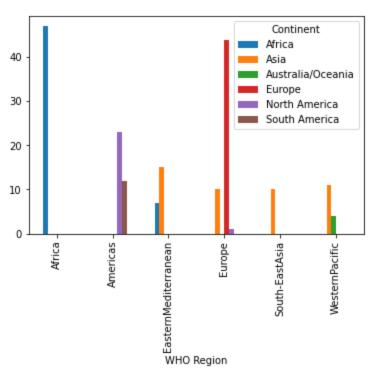
There seems to be a 132% chance the columns arent the same. Since this number is higher than 100% I pressume this means

Out[8]: Continent Africa Asia Australia/Oceania Europe North America South America

	WHO Region						
	Africa	47	0	0	0	0	0
	Americas	0	0	0	0	23	12
	EasternMediterranean	7	15	0	0	0	0
	Europe	0	10	0	44	1	0
	South-EastAsia	0	10	0	0	0	0
	WesternPacific	0	11	4	0	0	0

In [9]: contingencyTableCovid.plot(kind='bar')

Out[9]: <AxesSubplot:xlabel='WHO Region'>



I understand how this works. In the example notebooks there was an extra example which showed multiple rolls of a dice. In my dataset this isnt something that has to be done, but this shows you howmany times each continent would have gotten a certain amount.