(b)

Hypothesis: California(CA) and Texas(TX) have the most significant difference in their charges in DRG Charges 190. And CA's average charges is greater than TX's average charges.

H0: Average charges in DRG Charges 190 in CA and TX are the same.

H1: CA's average charges in DRG Charges 190 is greater than TX's average charges.

In [17]:

```
import scipy. stats as st

df1_hypo = df_6states[['DRG Charges 190', 'Provider State']]
  df1_CA = df1_hypo[df1_hypo['Provider State']=='CA']['DRG Charges 190']
  df1_TX = df1_hypo[df1_hypo['Provider State']=='TX']['DRG Charges 190']
  df1_CA = df1_CA. dropna()
  df1_TX = df1_TX. dropna()

t, p = st. ttest_ind(df1_CA, df1_TX)
  print("t statistic: " + str(t))
  print("p-value: " + str(p))

# Proform one-sided test and use significant value 0.05
  if p < 0.05/2:
        print("Reject HO.")
  else:
        print("Accept HO.")</pre>
```

```
t statistic: 9.804105998394789 p-value: 1.3330879399800308e-20 Reject HO.
```

Proform one-sided test and use significant value 0.05.

Since p-value < 0.05/2, null hypothesis H0 is rejected. Therefore, we tentatively conclude H1 to be the case, which support the claim.

(c)

Hypothesis: Pennsylvania(PA) and Georgia(GA) have the significant difference in their charges in DRG Charges 190.

H0: Average charges in DRG Charges 190 in PA and GA are the same.

H1: Average charges in DRG Charges 190 in PA and GA are the different.

In [25]:

```
df2 hypo = df 6states[['Provider State', 'DRG Charges 190', 'DRG Charges 392', 'DRG Charges 871']]
df2 PA = df2 hypo[df2 hypo['Provider State']=='PA']
df2 PA = df2 PA. dropna()
df2_PA = pd. concat([df2_PA['DRG Charges 190'], df2 PA['DRG Charges 392'], df2 PA['DRG Charges 87
1']], ignore index=True)
df2 GA = df2 hypo[df2 hypo['Provider State']=='GA']
df2 GA = df2 GA. dropna()
df2 GA = pd. concat([df2 GA['DRG Charges 190'], df2 GA['DRG Charges 392'], df2 GA['DRG Charges 87
1']], ignore index=True)
length = min(len(df2_PA), len(df2_GA))
df2 GA = df2 GA. sample (n = length, random state=3)
df2_PA = df2_PA.sample(n = length, random_state=3)
t_rel, p_rel = st.ttest_rel(df2_GA, df2_PA)
print("Two sample paired Student's t-test.")
print("t statistic(paired): " + str(t_rel))
print("p-value(paired): " + str(p_rel))
if p_rel < 0.05:</pre>
   print("Reject HO.")
else:
    print("Accept HO.")
```

```
Two sample paired Student's t-test.
t statistic(paired): -3.2213719626126287
p-value(paired): 0.0014578800424925055
Reject HO.
```

Proform two sample paired Student's t-test and use significant value 0.05.

Since p-value < 0.05, null hypothesis H0 is rejected. Therefore, we tentatively conclude H1 to be the case, which support the claim.

In [24]:

```
t_ind, p_ind = st.ttest_ind(df2_GA, df2_PA)
print("\nTwo sample unpaired t-test(two sided).")
print("t statistic(unpaired): " + str(t_ind))
print("p-value(unpaired): " + str(p_ind))
if p_ind < 0.05:
    print("Reject HO.")
else:
    print("Accept HO.")</pre>
```

```
Two sample unpaired t-test(two sided). t statistic(unpaired): -3.133704313100077 p-value(unpaired): 0.0018350152260808122 Reject HO.
```

Proform two sample unpaired t-test(two sided) and use significant value 0.05.

Since p-value < 0.05, null hypothesis H0 is rejected. Therefore, we tentatively conclude H1 to be the case, which support the claim.

As is shown above, paired t-test gets p-value slightly less then unpaired t-test, which means it's more likely to reject H0.