In [2]:

```
from sklearn.datasets import load_boston
from sklearn.model_selection import train_test_split
from sklearn.cluster import KMeans
from sklearn import preprocessing
from yellowbrick.cluster import KElbowVisualizer
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
%matplotlib inline

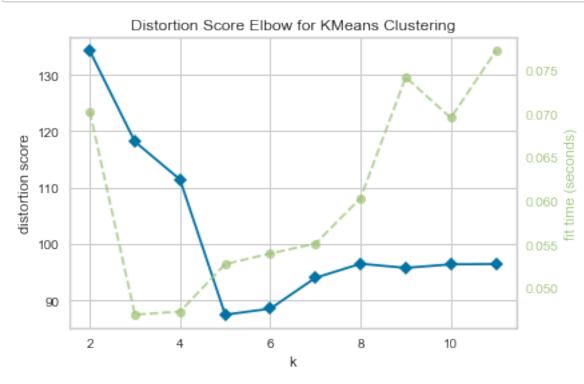
pd.set_option('display.max_columns', 500)
boston_data = load_boston()
df = pd.DataFrame(boston_data.data,columns=boston_data.feature_names)
df['target'] = pd.Series(boston_data.target)
```

In [3]:

#plot the prediction to a cross validation set evaluate the difference in pred iction and reality add the average to the model.

In [20]:

```
L = df[['LSTAT','target']]
# L = preprocessing.scale(L)
mets = ['silhouette', 'calinski_harabaz', 'distortion']
km = KMeans()
visualize = KElbowVisualizer(km, k = (2,12),metric=mets[2])
visualize.fit(L)
visualize.poof()
```



```
In [21]:
```

```
def centroid sort(centroids):
    new centroids = []
    c0s = np.sort(centroids[:,0])
    for j in c0s:
        for i in range(len(centroids)):
            if j == centroids[i][0]:
                new centroids.append([j, centroids[i][1]])
            else:
                continue
   new_centroids = np.array(new_centroids)
    return new centroids
def generate points(poly,start,stop,number of points=100):
    dummys = np.linspace(start,stop,number of points)
    for i in range(len(dummys)):
        y_points.append(poly(dummys[i]))
    return dummys, y_points
def get ave coefficients(dimension of eq, scribbling):
    ave coefficients = [0 for k in range(dimension of eq)]
    for j in range(dimension of eq):
        for i in range(len(scribbling)):
            ave coefficients[j] += scribbling[f'run {i}'][j]
        ave coefficients[j] = ave coefficients[j]/len(scribbling)
    return ave coefficients
```

In [6]:

```
# # print(centroids)

# colors = ['azure', 'darkolivegreen', 'darkblue', 'm', 'y', 'green', 'r', 'pa
payawhip' ]

# labels = km.labels__

# plt.scatter(L[:,0], L[:,1])

# plt.scatter(centroids[:, 0], centroids[:, 1], marker = 'x',c='m', s=75, line
widths = 5, zorder=10)

# # thats the line of best fit
# x, y = generate_points(p, -1, 2.5, 100)

# plt.plot(x, y, 'go-')
```

In [22]:

```
model_testing = {}
```

```
# feeding through KMeans on sample, generating the centroids again and again w
ith the best fit line
# Average over the result and return
# Assumptions data is in format already prepared for KMeans ie.: preprocess, o
perations, aggregations and so on.
def rigorous test(Var col, no of clusters, leading order=3, iterations=23, pre
proc = False):
    X = df[[Var col]]
    y = df[['target']]
    ave_eq = []
    ave_centroids = []
    if preproc:
        X = df[[Var col]]
        X = preprocessing.scale(X)
    for i in range(iterations):
        X train, X test, y train, y test = train test split(X, y, test size =
0.35)
        data = np.column stack((X train, y train))
        km = KMeans(n clusters = no of clusters)
        km.fit(data)
        centroids = km.cluster centers
        centroids = centroid_sort(centroids) #sorting centroids in order of x
coord
        # fiting current centroids
        cx = centroids[:,0]
        cy = centroids[:,1]
        ploy = np.polyfit(cx, cy, leading order)
        ave eq.append(ploy)
        ave centroids.append(centroids)
    ave coefficients = np.mean(ave eq, axis = 0)
    average line = np.poly1d(ave coefficients)
    ave_centroids = np.mean(ave_centroids, axis = 0)
    return average line, ave centroids
```

In [24]:

```
line, centroids = rigorous_test('LSTAT', no_of_clusters=5, leading_order=2, pr
eproc=False )
print('the line is: \n ', line)
print('\n \n' , 'Centroids are:\n',centroids)

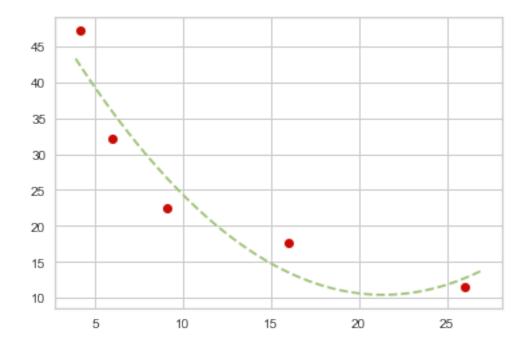
plt.plot(centroids[:,0], centroids[:,1], 'ro')
xz, yz = generate_points(line ,3.9, 27, number_of_points=100)
plt.plot(xz, yz, 'g--')
```

```
the line is: 2
0.1076 x - 4.603 x + 59.6
```

```
Centroids are:
[[ 4.16940125 47.18531813]
[ 5.98285427 32.09676768]
[ 9.06745139 22.53030121]
[16.01402642 17.5566861 ]
[26.00057608 11.58485563]]
```

Out[24]:

[<matplotlib.lines.Line2D at 0x12c689cf8>]



In [25]:

```
line.c
```

Out[25]:

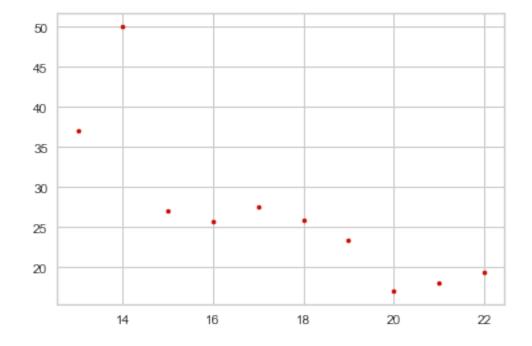
```
array([ 0.10764865, -4.60304943, 59.60211847])
```

In [26]:

```
# I forgot to mention that PTRATIO was calculated with averaging each rounded
instance of student ratio
# per housing value to deliver a distribution that is not a "radiator" distrib
ution
# it follows the pattern of regressing to the mean like all other categories.
roundy = round(df['PTRATIO'])
uniques = list(set(roundy))
yz = df['target']
dfesse = pd.concat([roundy, yz], axis=1)
uniques
cases = \{\}
for u in uniques:
    asdf = dfesse[dfesse['PTRATIO']==u]
    altering = asdf['target'].mean()
    cases[u] = altering
pt x = cases.keys()
pt_y = cases.values()
plt.plot(pt_x,pt_y, 'r.')
```

Out[26]:

[<matplotlib.lines.Line2D at 0x12c6f4080>]



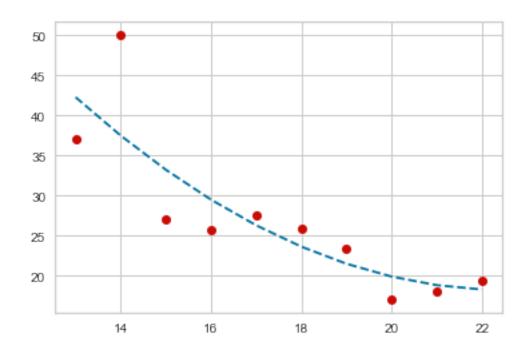
In [11]:

```
#PLotted PTRATIO
pt_x = list(pt_x)
pt_y = list(pt_y)
pt_fit = np.polyfit(pt_x, pt_y, 2)
pt_eq = np.polyld(pt_fit)
print(pt_eq.c)
pt_fit_y = pt_eq(pt_x)
plt.plot(pt_x,pt_y, 'ro')
plt.plot(pt_x,pt_fit_y, 'b--')
```

[0.26609764 -11.97890043 153.01549492]

Out[11]:

[<matplotlib.lines.Line2D at 0x12be450b8>]



In []:			

```
In [ ]:
```

```
In [ ]:
```