Dominant colors in images

CLUSTERING METHODS WITH SCIPY

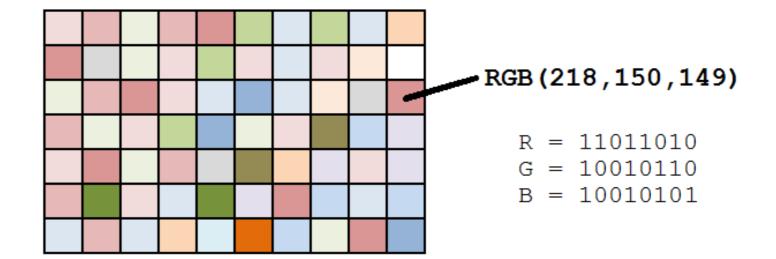


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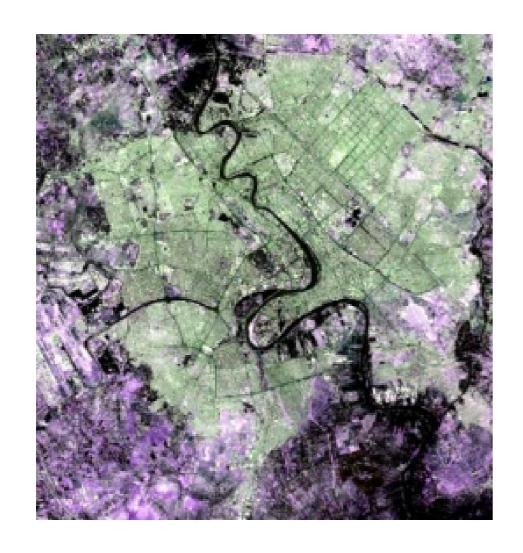
Dominant colors in images

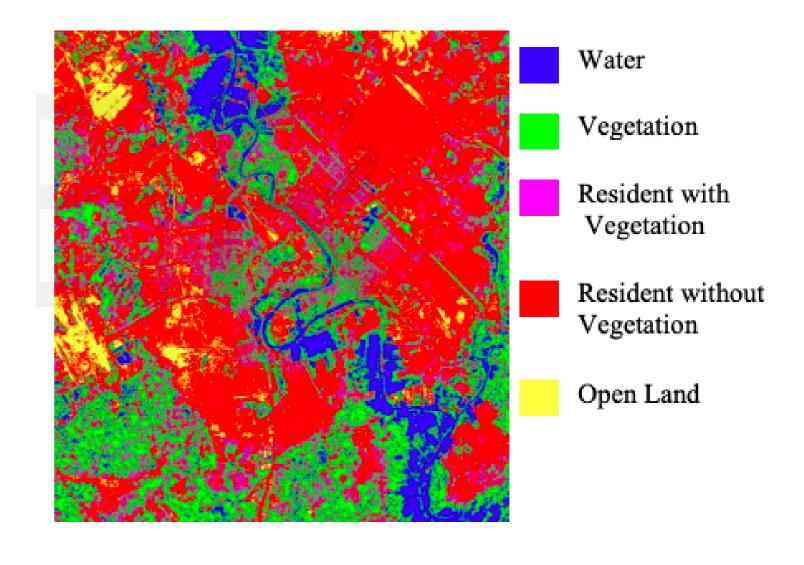
- All images consist of pixels
- Each pixel has three values: Red, Green and Blue
- Pixel color: combination of these RGB values
- Perform k-means on standardized RGB values to find cluster centers
- Uses: Identifying features in satellite images



Source

Feature identification in satellite images





Source

Tools to find dominant colors

- Convert image to pixels: matplotlib.image.imread
- Display colors of cluster centers: matplotlib.pyplot.imshow



Convert image to RGB matrix

```
import matplotlib.image as img
image = img.imread('sea.jpg')
image.shape
```

```
(475, 764, 3)
```

```
r = []
g = []
b = []

for row in image:
    for pixel in row:
        # A pixel contains RGB values
        temp_r, temp_g, temp_b = pixel
        r.append(temp_r)
        g.append(temp_g)
        b.append(temp_b)
```

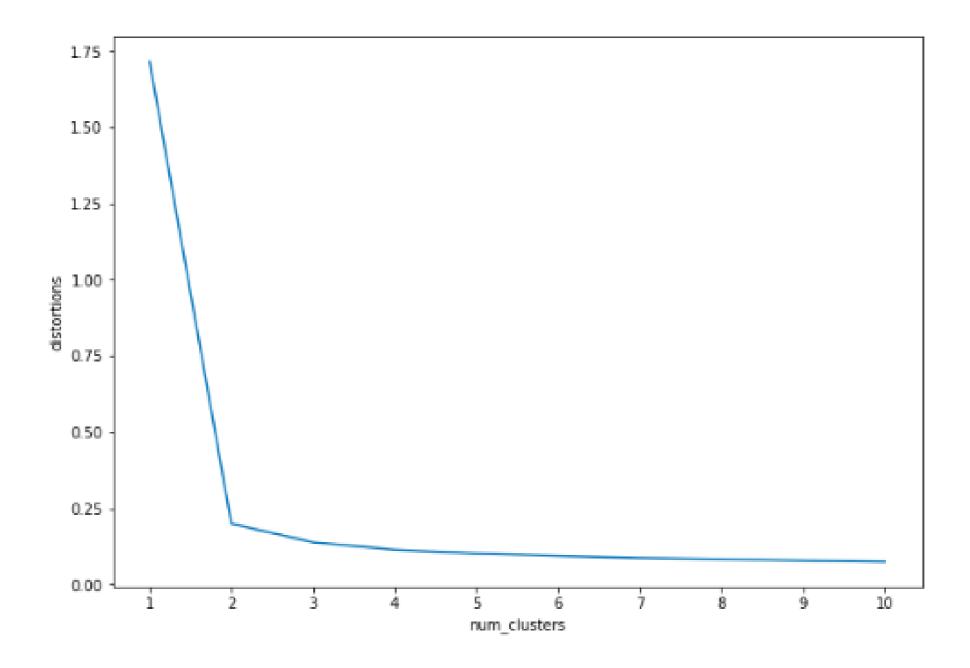
Data frame with RGB values

| red | blue | green |
|-----|------|-------|
| 252 | 255 | 252 |
| 75 | 103 | 81 |
| ••• | ••• | ••• |

Create an elbow plot

```
distortions = []
num_clusters = range(1, 11)
# Create a list of distortions from the kmeans method
for i in num clusters:
    cluster_centers, _ = kmeans(pixels[['scaled_red', 'scaled_blue',
                                        'scaled_green']], i)
    distortions.append(distortion)
# Create a data frame with two lists - number of clusters and distortions
elbow_plot = pd.DataFrame({'num_clusters': num_clusters,
                           'distortions': distortions})
# Creat a line plot of num_clusters and distortions
sns.lineplot(x='num_clusters', y='distortions', data = elbow_plot)
plt.xticks(num_clusters)
plt.show()
```

Elbow plot



Find dominant colors

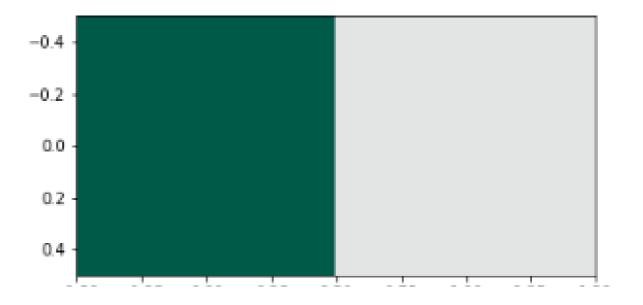
```
colors = []
# Find Standard Deviations
r_std, g_std, b_std = pixels[['red', 'blue', 'green']].std()
# Scale actual RGB values in range of 0-1
for cluster center in cluster centers:
    scaled_r, scaled_g, scaled_b = cluster_center
   colors.append((
        scaled_r * r_std/255,
        scaled_q * q_std/255,
        scaled_b * b_std/255
```

Display dominant colors

```
#Dimensions: 2 x 3 (N X 3 matrix)
print(colors)
```

```
[(0.08192923122023911, 0.34205845943857993, 0.2824002984155429),
(0.893281510956742, 0.899818770315129, 0.8979114272960784)]
```

```
#Dimensions: 1 \times 2 \times 3 (1 X N x 3 matrix) plt.imshow([colors]) plt.show()
```





Next up: exercises

CLUSTERING METHODS WITH SCIPY



Document clustering

CLUSTERING METHODS WITH SCIPY



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Document clustering: concepts

- 1. Clean data before processing
- 2. Determine the importance of the terms in a document (in TF-IDF matrix)
- 3. Cluster the TF-IDF matrix
- 4. Find top terms, documents in each cluster

Clean and tokenize data

Convert text into smaller parts called tokens, clean data for processing

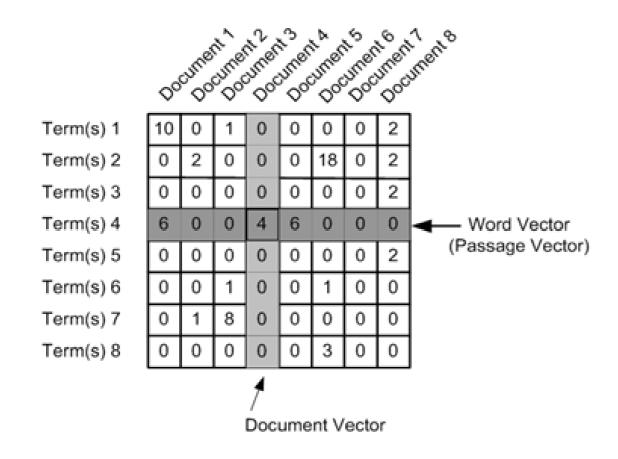
```
from nltk.tokenize import word_tokenize
import re
def remove_noise(text, stop_words = []):
    tokens = word_tokenize(text)
    cleaned_tokens = []
    for token in tokens:
        token = re.sub('[^A-Za-z0-9]+', '', token)
        if len(token) > 1 and token.lower() not in stop_words:
           # Get lowercase
           cleaned_tokens.append(token.lower())
    return cleaned tokens
remove_noise("It is lovely weather we are having.
             I hope the weather continues.")
```

```
['lovely', 'weather', 'hope', 'weather', 'continues']
```

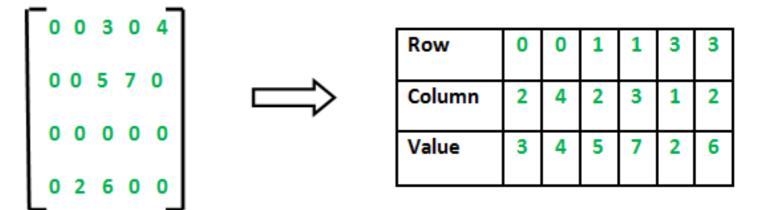


Document term matrix and sparse matrices

- Document term matrix formed
- Most elements in matrix are zeros



Sparse matrix is created



Source

Source



TF-IDF (Term Frequency - Inverse Document Frequency)

• A weighted measure: evaluate how important a word is to a document in a collection

Clustering with sparse matrix

- kmeans() in SciPy does not support sparse matrices
- Use .todense() to convert to a matrix

```
cluster_centers, distortion = kmeans(tfidf_matrix.todense(), num_clusters)
```

Top terms per cluster

- Cluster centers: lists with a size equal to the number of terms
- Each value in the cluster center is its importance
- Create a dictionary and print top terms

```
terms = tfidf_vectorizer.get_feature_names()

for i in range(num_clusters):
    center_terms = dict(zip(terms, list(cluster_centers[i])))
    sorted_terms = sorted(center_terms, key=center_terms.get, reverse=True)
    print(sorted_terms[:3])
```

```
['room', 'hotel', 'staff']
['bad', 'location', 'breakfast']
```

More considerations

- Work with hyperlinks, emoticons etc.
- Normalize words (run, ran, running -> run)
- .todense() may not work with large datasets

Next up: exercises!

CLUSTERING METHODS WITH SCIPY



Clustering with multiple features

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Basic checks

| cluster_labels | scaled_heading_accuracy | scaled_volleys | scaled_finishing |
|----------------|-------------------------|----------------|------------------|
| 0 | 3.21 | 2.83 | 2.76 |
| 1 | 0.71 | 0.64 | 0.58 |

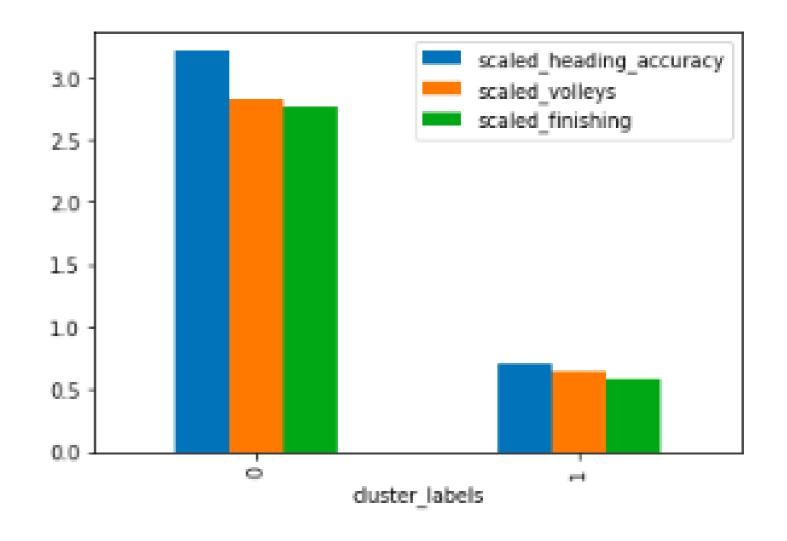
```
# Cluster sizes
print(fifa.groupby('cluster_labels')['ID'].count())
```

| cluster_labels | count |
|----------------|-------|
| 0 | 886 |

Visualizations

- Visualize cluster centers
- Visualize other variables for each cluster

```
# Plot cluster centers
fifa.groupby('cluster_labels') \
   [scaled_features].mean()
   .plot(kind='bar')
plt.show()
```



Top items in clusters

```
# Get the name column of top 5 players in each cluster
for cluster in fifa['cluster_labels'].unique():
    print(cluster, fifa[fifa['cluster_labels'] == cluster]['name'].values[:5])
```

| Cluster Label | Top Players | |
|---------------|--|--|
| 0 | ['Cristiano Ronaldo' 'L. Messi' 'Neymar' 'L. Suárez' 'R. Lewandowski'] | |
| 1 | ['M. Neuer' 'De Gea' 'G. Buffon' 'T. Courtois' 'H. Lloris'] | |

Feature reduction

- Factor analysis
- Multidimensional scaling

Final exercises!

CLUSTERING METHODS WITH SCIPY



Farewell!

CLUSTERING METHODS WITH SCIPY



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What comes next?

- Clustering is one of the exploratory steps
- More courses on DataCamp
- Practice, practice, practice!

Until next time

CLUSTERING METHODS WITH SCIPY

