

```
In [1]: from utils.data import Dataset
from utils.conllu import read_conllu_dataset
import numpy as np

In [2]: dataset = Dataset()

train_sentences = read_conllu_dataset("data/ro_rrt-ud-train.conllu")
test_sentences = read_conllu_dataset("data/ro_rrt-ud-test.conllu")

# Fit on training data
X_train, y_train = dataset.fit(train_sentences, mode="word_id", categorical_y=False)

# Encode test data (fixed shape)
X_test, y_test = dataset.encode(test_sentences)

X_train.shape, y_train.shape, X_test.shape, y_test.shape

Out[2]: ((8043, 163), (8043, 163), (729, 163), (729, 163))

In [3]: X_train.dtype, y_train.dtype

Out[3]: (dtype('float64'), dtype('int64'))

In [4]: def np2unohomogeneouslist(X,y):
    l_X, l_y = [], []
    for i in range(len(X)):
        zeros = np.argwhere(X[i]==0)
        seq_len = zeros[0,0] if len(zeros)>0 else len(X[i])
        l_X.append(X[i][0:seq_len])
        l_y.append(y[i][0:seq_len])
    return l_X, l_y

X, y = np2unohomogeneouslist(X_train.astype('int'), y_train)
```

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In [5]: from hmmlearn import hmm
from sklearn.metrics import confusion_matrix, classification_report
import seaborn as sns
import matplotlib.pyplot as plt
import numpy as np

# Get the number of unique tags and words
n_tags = len(dataset.label2id)
n_words = len(dataset.word2id)

# Convert sequences to format suitable for hmmlearn
def prepare_sequences(X, y):
    # Convert lists to numpy arrays first
    lengths = [len(seq) for seq in X]
    # Ensure the sequences are integer type
    X_flat = np.concatenate([np.array(seq, dtype=np.int32).reshape(-1, 1) for seq in X])
    y_flat = np.concatenate([np.array(seq, dtype=np.int32) for seq in y])
    return X_flat, y_flat, lengths

# Initialize HMM
model = hmm.CategoricalHMM(n_components=n_tags, n_iter=1000, init_params='')

# Initialize emission probabilities from labeled data
emission_counts = np.zeros((n_tags, n_words))
for x_seq, y_seq in zip(X, y):
    for word, tag in zip(x_seq, y_seq):
        emission_counts[int(tag), int(word)] += 1

# Add-1 smoothing and normalize emissions
model.emissionprob_ = (emission_counts + 1) / (emission_counts.sum(axis=1, keepdims=True) + n_words)

# Initialize transition probabilities from labeled data
transition_counts = np.zeros((n_tags, n_tags))
for y_seq in y:
    for i in range(len(y_seq) - 1):
        transition_counts[int(y_seq[i]), int(y_seq[i+1])] += 1

# Add-1 smoothing and normalize transitions
model.transmat_ = (transition_counts + 1) / (transition_counts.sum(axis=1, keepdims=True) + n_tags)

# Initialize start probabilities from labeled data
start_counts = np.zeros(n_tags)
for y_seq in y:
    start_counts[int(y_seq[0])] += 1
# Add-1 smoothing and normalize start probabilities
model.startprob_ = (start_counts + 1) / (start_counts.sum() + n_tags)

# Prepare test data
X_test_list, y_test_list = np2unohomogeneouslist(X_test, y_test)
X_test_flat, y_test_flat, test_lengths = prepare_sequences(X_test_list, y_test_list)

# Predict on test data using Viterbi algorithm
predicted_tags = model.decode(X_test_flat, test_lengths)[1]

# Get unique labels that actually appear in the data
unique_labels = np.unique(np.concatenate([y_test_flat, predicted_tags]))
label_names = [list(dataset.label2id.keys())[i] for i in unique_labels]

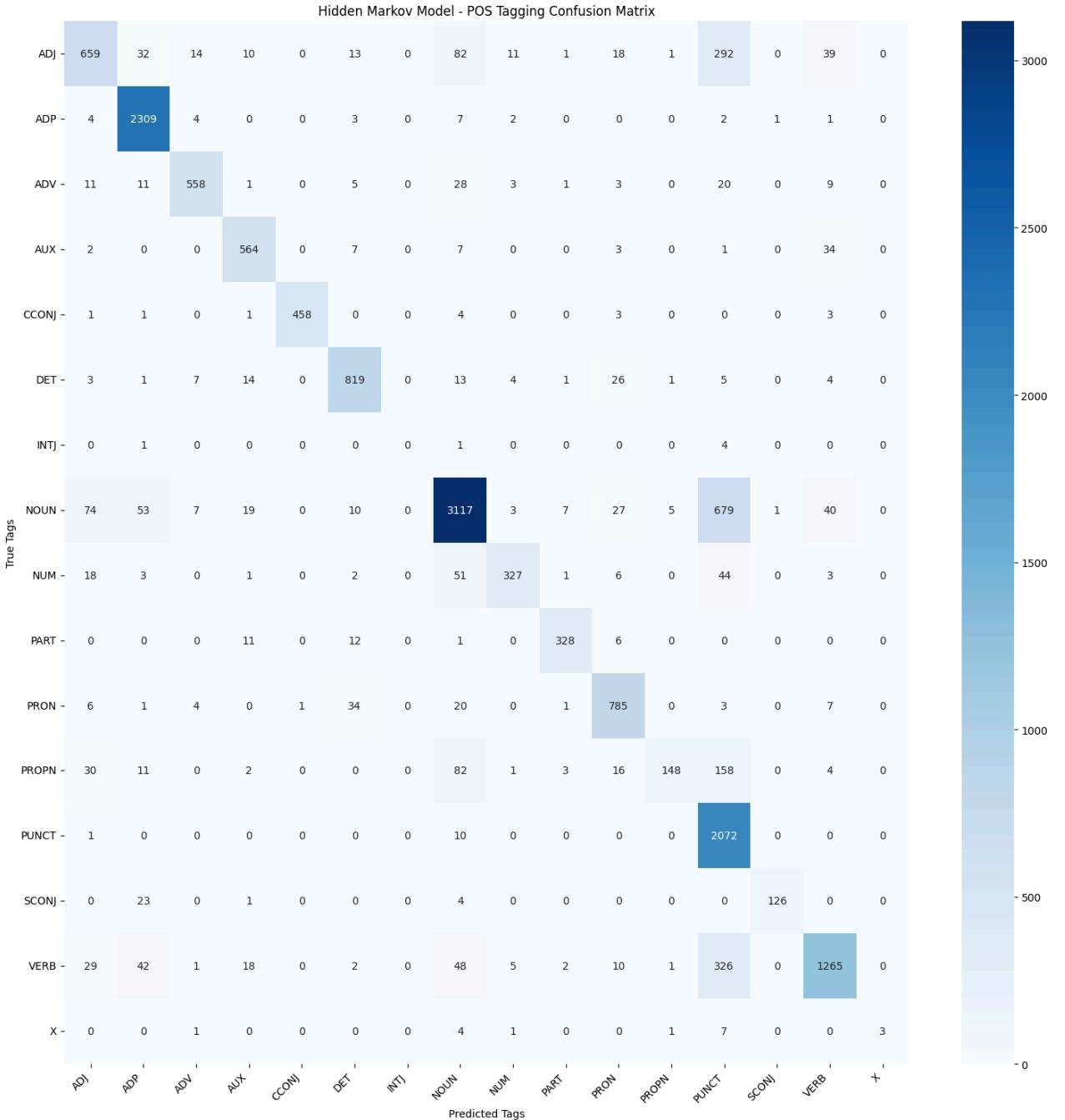
# Calculate and print metrics
print("Classification Report:")
print(classification_report(y_test_flat, predicted_tags,
                            labels=unique_labels,
                            target_names=label_names))

# Create confusion matrix visualization
cm = confusion_matrix(y_test_flat, predicted_tags,
                      labels=unique_labels)
plt.figure(figsize=(15, 15))
sns.heatmap(cm, annot=True, fmt='d', cmap='Blues',
            xticklabels=label_names,
            yticklabels=label_names)
plt.title('Hidden Markov Model - POS Tagging Confusion Matrix')
plt.xlabel('Predicted Tags')
plt.ylabel('True Tags')
plt.xticks(rotation=45, ha='right')
plt.yticks(rotation=0)
plt.tight_layout()
plt.show()

# Print overall accuracy
accuracy = np.mean(y_test_flat == predicted_tags)
print(f"\nTest Accuracy: {accuracy:.4f}")
```

Classification Report:		precision	recall	f1-score	support
ADJ	0.79	0.56	0.66	1172	
ADP	0.93	0.99	0.96	2333	
ADV	0.94	0.86	0.90	650	
AUX	0.88	0.91	0.90	618	
CCONJ	1.00	0.97	0.98	471	
DET	0.90	0.91	0.91	898	
INTJ	0.00	0.00	0.00	6	
NOUN	0.90	0.77	0.83	4042	
NUM	0.92	0.72	0.80	456	
PART	0.95	0.92	0.93	358	
PRON	0.87	0.91	0.89	862	
PROPN	0.94	0.33	0.48	455	
PUNCT	0.57	0.99	0.73	2083	
SCONJ	0.98	0.82	0.89	154	
VERB	0.90	0.72	0.80	1749	
X	1.00	0.18	0.30	17	
accuracy			0.83	16324	
macro avg	0.84	0.72	0.75	16324	
weighted avg	0.86	0.83	0.83	16324	

```
d:\anu1m\sem2\NLP\soft\PosTagging\.venv\lib\site-packages\sklearn\metrics\_classification.py:1565: UndefinedMetricWarning: Precision is ill-defined and being set to 0.0 in labels with no predicted samples. Use 'zero_division' parameter to control this behavior.
    _warn_prf(average, modifier, f"{{metric.capitalize()}} is", len(result))
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    _warn_prf(average, modifier, f"{{metric.capitalize()}} is", len(result))
```



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Test Accuracy: 0.8293
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In [6]: import os
import joblib

# Create models directory if it doesn't exist
os.makedirs('models', exist_ok=True)

# Save the trained HMM model
model_path = os.path.join('models', 'pos_hmm_model.joblib')
joblib.dump(model, model_path)

# Also save the label mapping for future use
mapping_path = os.path.join('models', 'pos_hmm_mappings.joblib')
label_mapping = {
    'label2id': dataset.label2id,
    'id2label': {v: k for k, v in dataset.label2id.items()},
    'word2id': dataset.word2id,
    'id2word': {v: k for k, v in dataset.word2id.items()}
}
joblib.dump(label_mapping, mapping_path)

print(f"Model saved to: {model_path}")
print(f"Mappings saved to: {mapping_path}")

Model saved to: models\pos_hmm_model.joblib
Mappings saved to: models\pos_hmm_mappings.joblib
```

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In [7]: #model = joblib.load('pos_hmm_model.joblib')
#mappings = joblib.load('pos_hmm_mappings.joblib')
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