## eval simcan

## September 2, 2024

```
[1]: import numpy as np
     import os
     import re
     import joblib
     from pathlib import Path
     import matplotlib.pyplot as plt
     import matplotlib.cm as cm
     import matplotlib.colors as plt_colors
     import tensorflow as tf
     import tensorflow.keras as keras
     from tensorboard.plugins import projector
     import sys
     from PIL import Image
     import smfret.fit as fit
     import smfret.finetuning as finetuning
     from smfret.tf_layers import Attention
     from smfret.tf_layers import Conv
     from smfret.tf_layers import Summary
     from smfret.tf layers import PrependTaskToken
     from smfret.tf_layers import Embedding
     from smfret.tf_layers import PositionEmbedding
     from smfret.trace_simulator import Simulator
     from smfret.trace_simulator import ParameterGenerator
     from smfret.trace_simulator import SimulatedTraceSet
     from smfret.multi_task_learning import FRETStateTraceSet
     from smfret.multi_task_learning import MultistepPhotobleachingTraceSet
     from smfret.multi_task_learning import NoiseLevelTraceSet
     from smfret.dataset import MatlabTraceSet
     from smfret.dataset import FRETTrace
     from smfret.dataset import FRETTraceSet
     from smfret.multi_task_learning import SavedTraceSet
```

```
from sklearn.manifold import TSNE
from sklearn import metrics
from sklearn.linear_model import LogisticRegression
from sklearn import mixture

from enum import Enum
import json
import umap
import scipy

from sciplotlib import style as spstyle
from tqdm.auto import tqdm
```

/Users/leyou/miniforge3/lib/python3.9/site-packages/umap/distances.py:1063: NumbaDeprecationWarning: The 'nopython' keyword argument was not supplied to the 'numba.jit' decorator. The implicit default value for this argument is currently False, but it will be changed to True in Numba 0.59.0. See https://numba.readthedocs.io/en/stable/reference/deprecation.html#deprecation-of-object-mode-fall-back-behaviour-when-using-jit for details.

@numba.jit()

/Users/leyou/miniforge3/lib/python3.9/site-packages/umap/distances.py:1071: NumbaDeprecationWarning: The 'nopython' keyword argument was not supplied to the 'numba.jit' decorator. The implicit default value for this argument is currently False, but it will be changed to True in Numba 0.59.0. See https://numba.readthedocs.io/en/stable/reference/deprecation.html#deprecation-of-object-mode-fall-back-behaviour-when-using-jit for details.

@numba.jit()

/Users/leyou/miniforge3/lib/python3.9/site-packages/umap/distances.py:1086: NumbaDeprecationWarning: The 'nopython' keyword argument was not supplied to the 'numba.jit' decorator. The implicit default value for this argument is currently False, but it will be changed to True in Numba 0.59.0. See https://numba.readthedocs.io/en/stable/reference/deprecation.html#deprecation-of-object-mode-fall-back-behaviour-when-using-jit for details.

@numba.jit()

/Users/leyou/miniforge3/lib/python3.9/site-packages/umap/umap\_.py:660:
NumbaDeprecationWarning: The 'nopython' keyword argument was not supplied to
the 'numba.jit' decorator. The implicit default value for this argument is
currently False, but it will be changed to True in Numba 0.59.0. See
https://numba.readthedocs.io/en/stable/reference/deprecation.html#deprecationof-object-mode-fall-back-behaviour-when-using-jit for details.

```
@numba.jit()
```

```
[2]: NN_K = 50 # how many nearest neighbors are considered for the score
    def calculate_distance(embedding):
         """Calculates the distance matrix from the embeddings."""
        distance = (
            tf.expand_dims(tf.einsum('ik,ik->i', embedding, embedding), axis=-1)
            + tf.expand_dims(tf.einsum('ik,ik->i', embedding, embedding), axis=0)
            - 2.0 * tf.einsum('ik,jk->ij', embedding, embedding))
        return distance
    def get_perplexity(embedding, label, branching_factor,_
      """Calculates the perplexity of a trace's neighbors."""
        distance = calculate_distance(embedding)
        top_k_results = tf.math.top_k(
            -distance, k=branching_factor + 1, sorted=True, name=None
        top_k_idx = top_k_results.indices.numpy()
        n = embedding.shape[0]
        perplexity = []
        for i in range(n):
            this_label = label[i]
            neighbour_label = label[top_k_idx[i, :].flatten()]
            unique_labels = set(neighbour_label)
            shannon_entropy = 0
            for 1 in unique_labels:
                p = np.mean(neighbour_label == 1)
                shannon_entropy -= p * np.log(p)
            perplexity.append(shannon_entropy)
        return np.array(perplexity)
[3]: keras.mixed precision.set global policy('mixed bfloat16')
     encoder = keras.models.load_model('../../saved_models/best_model/
      ⇔encoder-20240111-045226.h5', compile=False)
[4]: def read simcan tracesets(file, label=None, start frame=0, max frames=8000):
         """Converts .MAT data into model-compatible trace sets."""
        key = 'data'
        conv_width = 100
        dialation_factor = 1
        simcan_data = scipy.io.loadmat(file)[key][0]
        trace_sets = []
        labels = []
        for data in simcan_data:
            label = data[0][0][0][0]
            donor = data[0][0][2][:, 1]
```

```
acceptor = data[0][0][2][:, 2]
trace_set = FRETTraceSet()
trace_set.donor = np.expand_dims(donor, axis=0)
trace_set.acceptor = np.expand_dims(acceptor, axis=0)
trace_set.time = np.arange(len(donor)) + 1
trace_set.size = 1
trace_set.label = tf.zeros_like(trace_set.donor)
if len(donor) >= 100:
    trace_sets.append(trace_set)
    labels.append(label)
return trace_sets, labels
```

```
[5]: trace_sets, labels = read_simcan_tracesets('../../saved_dataset/downstream/

SiMCAn/converted_all_data.mat')
```

```
[6]: print(f"There are {len(trace_sets)} traces in the dataset.")
```

There are 6805 traces in the dataset.

```
[7]: # Export to the embedding projector
     def trim(x, precision=4):
         """Trims a float to a given precision."""
         if int(x) == x:
             return x
         else:
             return float(np.format_float_positional(x, precision=4, unique=False, __

¬fractional=False, trim='k'))
     def generate_projector_data(log_dir, embedding, trace_set, plot_label,_u
      →predicted_label, k_means_label, sprite=False):
         Path(log_dir).mkdir(parents=True, exist_ok=True)
         with open(os.path.join(log_dir, 'metadata.tsv'), "w") as f:
           f.write('name\tground truth\tpredicted__
      ⇔label\tk-means-clustering\tdonor\tacceptor\n')
           for index in range(embedding.shape[0]):
             donor = np.array([trim(x) for x in trace_set.traces[index].donor])
             acceptor = np.array([trim(x) for x in trace_set.traces[index].acceptor])
             f.write(f"trace__
      →{index}\t{plot_label[index]}\t{predicted_label[index]}\t{k_means_label[index]}\t{json.
      →dumps(donor.tolist())}\t{json.dumps(acceptor.tolist())}\n")
         with open(log_dir + 'tensor.bytes', 'wb') as f:
             f.write(np.array(tf.cast(embedding, tf.float32)).tobytes())
         np.savetxt(log_dir + 'tensor.tsv', embedding, delimiter='\t')
```

```
def generate_tensor_and_metadata(trace_sets, start_frame, output_dir, encoder, __
 oreducer, max_size=4000, max_frame=2000, balance_label=True,
 →logistic_regression=True, class_weight=None):
    """Generates the files for embedding projector."""
    embeddings = []
    labels = []
    colors = []
    color_count = 1
    image_files = []
    traces = []
    count = 0
    for trace_set in trace_sets:
        trim_size = len(trace_set.time) // 100 * 100
        trace_set.trim(trim_size)
        trace_set.trim(max_frame, start_frame=start_frame)
        trace_set.broadcast_data_to_traces()
        print(trace set.size)
        if balance_label:
          size = 2 * np.sum(np.max(trace_set.label, axis=-1))
        else:
          size = trace_set.size
        if count + size > max_size:
            break
        else:
            count += size
        with tf.device('/CPU:0'):
            embedding = reducer.transform(encoder.predict(trace_set.
 →to_tensor()))
            embeddings.append(embedding)
            labels.append(np.max(trace_set.label, axis=-1))
            traces.extend(trace_set.traces)
    labels = np.concatenate(labels, axis=0)
    n_included = np.sum(labels == 1)
    n_excluded = np.sum(labels == 0)
    indices = np.arange(len(labels))
    if balance_label:
      ratio = int(n_excluded / n_included)
      if ratio > 1:
        indices = np.concatenate([indices[labels == 1], indices[labels ==0][::
 →ratio]])
        labels = labels[indices]
```

```
trace_set = FRETTraceSet()
  trace_set.traces = [traces[i] for i in indices]
  trace_set.size = len(trace_set.traces)
  embedding = np.concatenate(embeddings, axis=0)[indices, :]
  if np.max(labels) == 1:
    labels = np.where(labels == 1, 'Included in Analysis', 'Excluded from_u
Analysis')
  kmeans_labels = cluster.KMeans(n_clusters=2).fit_predict(embedding)
    clf = LogisticRegression(random_state=0, max_iter=5000,_u
class_weight=class_weight).fit(embedding, labels)
    predicted_label = clf.predict(embedding)
    # kmeans_labels = hdbscan.HDBSCAN(min_cluster_size=15,_u
    gen_min_span_tree=True).fit_predict(clusterable_embedding)
    generate_projector_data(output_dir, embedding, trace_set, labels,_u
    predicted_label, kmeans_labels, sprite=False)
```

```
[8]: def generate embedding and label(trace_sets, start_frame, encoder, u
      max_frame=2000, max_traces=4000, balance_labels=True):
         """Generates the embeddings and labels."""
         embeddings = []
         labels = []
         colors = []
         color_count = 1
         image_files = []
         traces = []
         count = 0
         for trace_set in trace_sets:
             trim_size = len(trace_set.time) // 100 * 100
             trace_set.trim(trim_size)
             trace_set.trim(max_frame, start_frame=start_frame)
             trace_set.broadcast_data_to_traces()
             if count + trace_set.size > max_traces:
                 break
             else:
                 count += trace_set.size
             with tf.device('/CPU:0'):
                 label = np.max(trace_set.label, axis=-1)
                 if balance_labels:
                     n_included = np.sum(label == 1)
                     n_excluded = np.sum(label == 0)
                     if n_included == 0 or n_excluded == 0:
                         continue
                     indices = np.arange(len(label))
                     ratio = int(n_excluded / n_included)
```

```
# ratio = 0
                      if ratio > 1:
                          indices = np.concatenate([indices[label == 1],__
       →indices[label ==0][::ratio]])
                          label = label[indices]
                          trace set = trace set.get subset(indices)
                  embedding = encoder.predict(trace set.to tensor(), batch size=8,,,
       ⇒verbose=0)
                  embeddings.append(embedding)
                  labels.append(label)
                  traces.extend(trace set.traces)
          return embeddings, labels
 [9]: embeddings_spot, labels_spot = generate_embedding_and_label(trace_sets, 0,__
       Gencoder, max_frame=2000, max_traces=20000, balance_labels=False)
[10]: final_embeddings = []
      final_labels = []
      human_labels = []
      for e, l in zip(embeddings_spot, labels_spot):
          final_embeddings.append(e)
          final_labels.append(1)
      embedding_spot = np.concatenate(final_embeddings, axis=0)
      human_label_spot = np.concatenate(final_labels)
      label_spot = np.array(labels)
[13]: new_label_spot = np.array([l.replace('Late', '').replace('Middle', '').
       Greplace('Early', '').replace('Mid', '').replace('early', '') for l in □
       →label_spot])
[14]: sorted(list(set(new_label_spot.tolist())))
[14]: ['ATP3SScompiled',
       'NoATPcompiled',
       'Prp16DN3SS',
       'Prp16DNWT',
       'Prp2dep3SScompiled',
       'Prp2depWTcompiled',
       'U6depCompiled',
       'WTATPcompiled']
[15]: ordered_label_spot = [
          'NoATPcompiled',
          'U6depCompiled',
          'Prp2depWTcompiled',
          'Prp2dep3SScompiled',
```

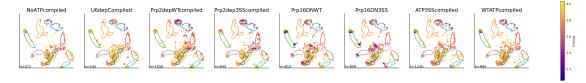
```
'Prp16DNWT',
          'Prp16DN3SS',
          'ATP3SScompiled',
          'WTATPcompiled',
      ]
[16]: perplexity_spot = get_perplexity(embedding_spot, new_label_spot,_u
       ⇔branching_factor=50)
[17]: reducer = joblib.load('../../saved_models/UMAP/atlas_pp.joblib')
      saved_atlas_density_models = joblib.load('../../saved_models/UMAP/
      →atlas_pp_density_models.joblib')
      n = len(set(label_spot.tolist()))
      cutoff = np.quantile(perplexity_spot, 1.0)
      counter = 0
      z h = reducer.transform(embedding spot)
      ax_min = np.min(z_h)
      ax max = np.max(z h)
      with plt.style.context(spstyle.get_style('nature-reviews')):
          fig, axes = plt.subplots(ncols=len(ordered_label_spot), nrows=1, figsize=(5_

state = * len(ordered_label_spot), 5));

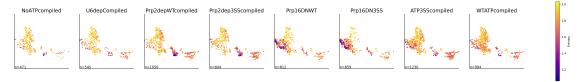
      for 1 in ordered label spot:
          idx = (new_label_spot == 1) & (perplexity_spot <= cutoff)</pre>
          z = z h[idx, ...]
          with plt.style.context(spstyle.get_style('nature-reviews')):
              ax = axes[counter]
              counter += 1
              s = ax.scatter(z[:, 0], z[:, 1], c=perplexity_spot[idx], s=30, alpha=0.
       -6, linewidths=0, cmap='plasma', vmin=np.quantile(perplexity_spot, 0.01),
       →vmax=np.quantile(perplexity_spot, 0.99))
              ax.set_xlim([ax_min, ax_max])
              ax.set_ylim([ax_min, ax_max])
              ax.set_xticks([])
              ax.set yticks([])
              ax.set_aspect('equal')
              ax.set title(f'{1}')
              ax.text(x=ax.get_xlim()[0]+0.1, y=ax.get_ylim()[0]+0.1, s=f'n=\{z.
       ⇒shape[0]}', fontdict={'size': 12})
          contour_counter = 0
          for l in saved_atlas_density_models:
              color = plt.cm.tab20.colors + plt.cm.tab20b.colors[::2]
              x = np.linspace(np.min(z_h[:, 0]) - 1, np.max(z_h[:, 0]) + 1, num=200)
              y = np.linspace(np.min(z_h[:, 1]) - 1, np.max(z_h[:, 1]) + 1, num=200)
              X, Y = np.meshgrid(x, y)
```

```
clf = saved_atlas_density_models[1]
        XX = np.array([X.ravel(), Y.ravel()]).T
        Z = clf.score_samples(XX)
        Z = np.exp(Z.reshape(X.shape))
        cmap = plt_colors.LinearSegmentedColormap.from_list("",__
 Golor[contour_counter], color[contour_counter], color[contour_counter]])
        if 'n' not in 1:
            CS = ax.contour(
                X, Y, Z, cmap=cmap, levels=[1e-2]
        contour_counter += 1
color_bar = fig.colorbar(s, ax=axes, label="Entropy")
color_bar.set_alpha(1)
color_bar.draw_all()
if cutoff >= np.quantile(perplexity_spot, 0.9):
   plt.savefig('figures/simcan_perplexity_umap_atlas.svg')
else:
   plt.savefig('figures/simcan_perplexity_umap_atlas_threshold.svg')
plt.show()
```

OMP: Info #276: omp\_set\_nested routine deprecated, please use omp\_set\_max\_active\_levels instead.



```
z = z_h[idx, ...]
    with plt.style.context(spstyle.get_style('nature-reviews')):
        ax = axes[counter]
        counter += 1
        if cutoff >= np.quantile(perplexity_spot, 0.9):
            s = ax.scatter(z[:, 0], z[:, 1], c=perplexity_spot[idx], s=8,__
 ⇒alpha=1, linewidths=0, cmap='plasma', vmin=np.quantile(perplexity_spot, 0.
 ⇔01), vmax=np.quantile(perplexity_spot, 0.99))
        else:
            s = ax.scatter(z[:, 0], z[:, 1], c=perplexity_spot[idx], s=30,_{\bot}
 alpha=0.5, linewidths=0, cmap='plasma', vmin=np.quantile(perplexity_spot, 0.
 →01), vmax=np.quantile(perplexity_spot, 0.99))
        \# s = ax.scatter(z[:, 0], z[:, 1], c=perplexity\_spot[idx], s=30, alpha=.
 →5, linewidths=0, cmap='plasma', vmin=np.quantile(perplexity_spot, 0.01),
 →vmax=np.quantile(perplexity_spot, 0.99))
        ax.set xlim([ax min, ax max])
        ax.set_ylim([ax_min, ax_max])
        ax.set_xticks([])
        ax.set_yticks([])
        ax.set_aspect('equal')
        ax.set_title(f'{1}')
        ax.text(x=ax.get_xlim()[0] + 0.1, y=ax.get_ylim()[0] + 0.1, s=f'n={z.}
 ⇔shape[0]}', fontdict={'size': 12})
color_bar = fig.colorbar(s, ax=axes, label="Entropy")
color_bar.set_alpha(1)
color_bar.draw_all()
if cutoff >= np.quantile(perplexity_spot, 0.9):
    plt.savefig('figures/simcan_perplexity_umap.svg')
else:
    plt.savefig('figures/simcan_perplexity_umap_threshold.svg')
plt.show()
```



[]: