

Aligned SOMs Implementation Details

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Github:

<https://github.com/Znerual/AlignedSOM>

Implementation Details

Aligned SOMs aims at training multiple layers of n SOMs with differently weighted subsets of attributes. The implementation of the SOM training is closely modelled after the description in the paper Aligned self-organizing maps by Pampalk, Elias [1]. The Aligned SOM implementation uses the well known MiniSom package and trains multiple layers of the MiniSom [2]. A Layer has in extension to the normal MiniSom implementation the possibility to set initial codebook weights. Further, the update method is adapted to model the distance between layers. We implemented an online-training algorithm which iteratively trains all layers.

In the following sections we will briefly describe the major concepts of the Aligned SOM and how we implemented it. We will describe the implementation and effect of the parameters when there is no difference to the normal SOM as implemented in the MiniSom library. All parameters and "public" methods of the algorithm have docstrings describing the parameter and method in more detail if some aspects are still unclear from the description.

Layer Weighting

Two aspects of features in a dataset are differently weighted in different layers of the Aligned SOMs. The first layer uses a weighting ratio between aspect A and aspect B features of 1:0. The middle or center layer, weights both aspects equally. The last layer uses a weighting ratio of 0:1.

We create the weights by layer in the **AlignedSom** class using the method `_create_weights_by_layer`. The **AlignedSom** accepts a parameter **aspect_selection** which has to be a boolean List indicating if the feature belongs to aspect A (True or 1) or if it belongs to aspect B (False or 0). The weights of features assigned to aspect A are a linear interpolation of 0 to 1 and the weights of features assigned to aspect B are a linear interpolation from 1 to 0.

Layer/Codebook initialization

We create N SOM layers initializing them identically using the same common codebook but weighting them by the respective layer weight vector.

The initialization of the layers in the **AlignedSom** class is done in the method **_create_layers**. The number of layers created can be defined by the parameter **num_layers**. We either create the common codebook randomly or train the center SOM (trained with unweighted data) and use it as basis for all layer initializations. This can be changed by the parameter **codebook_initialization_type** ("random" or "pretrained"). The weighting of the layers is done by the **weights_by_layer** as explained in the previous section. One Layer is represented by the **Layer** class which extends the MiniSom algorithm by overwriting the **update** method.

Training

We train multiple layers of SOMs iteratively with an online-training algorithm. The algorithm is implemented as follows:

1. select a random layer and a random observation from the dataset
2. select the winning unit in the selected layer based on the weighted feature vector
3. train all layers updating the weights based on the same winning unit
 - the selected layer is updated as in the normal SOM training
 - all other layers update the weights similarly but the update is further influenced by the distance to the selected layer
 - all layers use the weighted feature vector based on their respective layer weights
4. iterate steps 1-3 N times

Instead of directly calling the **train** method of one **Layer** the **AlignedSom** iteratively updates the codebook of different layers by calling the **update** method. We archive the updates of the codebook in each **Layer** by changing the **update** method of the MiniSom which now also accepts a parameter **layer_dist** which represents the distance between **Layers** in the stack.

Layer distances

The distance of the layers is defined as follows.

- the distance to the layer to itself is 1.0 which equals the normal SOM update rule
- the distance to the neighboring layer is a fraction (**layer_distance_ratio** (default 1/10)) of the distance between neighboring units in one layer
- the distance is defined by a gaussian function with $\sigma=1.0$

We initially create the distances between layers in the **AlignedSom** in the method **_create_layer_distances**. We do not reduce the distance between layers during the training time as the neighborhood in one layer because we noticed that neighboring layers would have very different codebook weights.

Visualization

The visualizations are based on the ones in [PySOMVis](#). The only change inside the visualizations is that we resize the resulting matrix to get a smoother visualization. In the **visualize.py** module the wrapper function **plot_aligned_som** can be used to plot multiple layers of the SOM next to each other. We always show the first and last layer and desired number of SOMs based on the **num_plots** parameter. The type of visualization can be selected by the **visualization_function** parameter. The implemented options are Hit Histogram (**HitHist**), U-Matrix (**UMatrix**) and Smoothed-Data-Histogram (**SDH**) which are all implemented in the same module.

References

[1] Pampalk, Elias. "Aligned self-organizing maps." Proceedings of the Workshop on Self-Organizing Maps. 2003.

URL: https://www.researchgate.net/publication/2887633_Aligned_Self-Organizing_Maps

[2] Vettigli, Giuseppe. "MiniSom: minimalistic and NumPy-based implementation of the Self Organizing Map." (2018).

URL: <https://github.com/JustGlowing/minisom>

```
In [ ]: import os, sys
import numpy as np
from typing import Tuple, List
from minisom import MiniSom
from random import randrange, seed
from tqdm import tqdm
import matplotlib.pyplot as plt
import seaborn as sns

module_path = os.path.abspath(os.path.join('../'))
if module_path not in sys.path:
    sys.path.append(module_path)

from src.data import load_dataset
from src.config import config
from src.visualize import SDH, HitHist, UMatrix
```

Implementation of one Layer

```
In [ ]: class Layer(MiniSom):
    """One layer of the Aligned SOM extending the MiniSom library"""
    def __init__(self,
                 dimension: Tuple[int, int],
                 input_len: int,
                 initial_codebook: np.ndarray,
                 sigma: float = 1.0,
                 learning_rate: float = 0.5,
                 neighborhood_function: str = 'gaussian',
                 activation_distance: str = 'euclidean',
                 random_seed: int = None) -> None:
        """Constructs one layer of the Aligned SOM
```

```

    Args:
        dimension (Tuple[int, int]): x and y dimensions of the result
        input_len (int): Dimension of the training data
        initial_codebook (np.ndarray): Weight vectors of the initial
        sigma (float, optional): Initial spread of the neighborhood f
        learning_rate (float, optional): initial Learning rate. Defau
        neighborhood_function (str, optional):
            Type of function to use for computing the neighborhood.
            Possible values: 'gaussian', 'mexican_hat', 'bubble', 'tr
            Defaults to 'gaussian'.
        activation_distance (str, optional):
            Type of function used for computing the distances between
            Possible values: 'euclidean', 'cosine', 'manhattan', 'che
            Defaults to 'euclidean'.
        random_seed (int, optional):
            Random seed used for all operations which use randomness.
            Defaults to None
    """

    super().__init__(
        x=dimension[0],
        y=dimension[1],
        input_len=input_len,
        sigma=sigma,
        learning_rate=learning_rate,
        neighborhood_function=neighborhood_function,
        topology='rectangular',
        activation_distance=activation_distance,
        random_seed=random_seed)

    # after initialization of the weights by MiniSom override them wi
    self._weights = initial_codebook

    # changed update to include the distance to the layer in the neighbor
    def update(self,
        input_vector: np.array,
        winner_position: Tuple[int, int],
        layer_dist: float,
        time_point: int,
        max_iteration: int) -> None:
        """Update the SOM codebook similar to normal SOM update including
        (extended update function of the MiniSom library)

    Args:
        input_vector (np.array): 1d input vector used for training
        winner_position (Tuple[int, int]): Tuple indicating the posit
        layer_dist (float):
            fraction representing the distance between layers
            max 1.0 which is normal SOM update for same layer
        time_point (int):
            current number of iteration of the training algorithm use
            determining the decay of learning rate and neighborhood s
        max_iteration (int):
            number of iterations used for training the map used for
            determining the decay of learning rate and neighborhood s
    """
    eta = self._decay_function(self._learning_rate, time_point, max_i
    # sigma and learning rate decrease with the same rule
    sig = self._decay_function(self._sigma, time_point, max_iteration

```

```
# improves the performances
g = self.neighborhood(winner_position, sig) * eta * layer_dist
# w_new = eta * neighborhood_function * (x-w)
self._weights += np.einsum('ij, ijk->ijk', g, input_vector-self._
```

Implementation of the Aligned SOM

```
In [ ]: class AlignedSom():
    """Aligned SOM implementation

    Details of the algorithm can be found in:
    Pampalk, Elias. "Aligned self-organizing maps." Proceedings of the Wo
    URL: https://www.researchgate.net/publication/2887633\_Aligned\_Self-Or

    """
    def __init__(self,
                  dimension: Tuple[int, int],
                  data: np.ndarray,
                  aspect_selection: List[bool],
                  num_layers: int = 100,
                  layer_distance_ratio: float = 0.1,
                  sigma: float = 1.0,
                  learning_rate: float = 0.5,
                  neighborhood_function: str = 'gaussian',
                  activation_distance: str = 'euclidean',
                  codebook_inizialization_type: str = 'random',
                  random_seed: int = None):
        """Construction of Aligned SOM

        Args:
            dimension (Tuple[int, int]): x and y dimensions of the result
            data (np.ndarray): 2d input data used for training the SOM
            aspect_selection (List[bool]):
                Selection if feature belongs to aspect A or aspect B
                True -> aspect A, False -> aspect B
                Needs to have the same dimension as the number of columns
            num_layers (int, optional):
                Number of layers trained.
                Defaults to 100.
            layer_distance_ratio (float, optional):
                The ratio used for computing the distance between layers.
                The distance between two neighboring layers is
                Defaults to 0.1.
            sigma (float, optional): Initial spread of the neighborhood f
            learning_rate (float, optional): initial Learning rate. Defau
            neighborhood_function (str, optional): _description_. Default
            neighborhood_function (str, optional):
                Type of function to use for computing the neighborhood.
                Possible values: 'gaussian', 'mexican_hat', 'bubble', 'tr
                Defaults to 'gaussian'.
            activation_distance (str, optional):
                Type of function used for computing the distances between
                Possible values: 'euclidean', 'cosine', 'manhattan', 'che
                Defaults to 'euclidean'.
            codebook_inizialization_type (str, optional):
                Type of initializing the layer codebooks.
                Possible values: 'random', 'pretrained'
                Defaults to 'random'.
```

```

        random_seed (int, optional):
            Random seed used for all operations which use randomness.
            Defaults to None
    """
    self._dimension = dimension
    self.data = data
    self._input_len = data.shape[1]
    self._aspect_selection = aspect_selection
    self._num_layers = num_layers
    self._layer_distance_ratio = layer_distance_ratio
    self._sigma = sigma
    self._learning_rate = learning_rate
    self._neighborhood_function = neighborhood_function
    self._activation_distance = activation_distance
    self._codebook_inizialization_type = codebook_inizialization_type
    self._random_seed = random_seed

    self._weights_by_layer: np.ndarray = self._create_weights_by_layer
    self.layers: List[Layer] = self._create_layers()
    self._layer_distances = self._create_layer_distances()

def train(self, num_iterations: int) -> None:
    """Online-training process of the Aligned SOM
    All layers are trained iteratively by selecting one layer and on

    Args:
        num_iterations (int): Number of Iterations the Aligned SOM is
    """
    n_observations = self.data.shape[0]
    if self._random_seed:
        seed(self._random_seed)
    for t in tqdm(range(num_iterations)):
        selected_layer = randrange(0, self._num_layers)
        selected_observation = randrange(0, n_observations)
        # print(f'selected layer: {selected_layer}')
        # print(f'selected observation: {selected_observation}')
        winner = self.layers[selected_layer].winner(
            self.data[selected_observation] * self._weights_by_layer[
        for i, layer in enumerate(self.layers):
            # print(f'current layer: {i}')
            # layer_dist = self._layer_distance(t, num_iterations, np.
            layer_dist = self._layer_distances[np.abs(selected_layer
            # print(f'distance: {layer_dist}')
            layer.update(self.data[selected_observation] * self._weig
                winner,
                layer_dist,
                t,
                num_iterations)

def get_layer_weights(self) -> List[np.ndarray]:
    """Get all layer weights (codebooks)

    Returns:
        List[np.ndarray]: Weights by layer each codebook has dimension
    """
    return [layer.get_weights() for layer in self.layers]

def set_layer_weights(self, weights_by_layer: List[np.ndarray]) -> No
    """Overrides the layers with existing codebook weights from a tra

```

```

    Args:
        weights_by_layer (List[np.ndarray]):
            A list of codebook weights where each item in the list contains a
            layer codebook with dimension (x, y, input_len)
    """
    layers = []
    for weights in weights_by_layer:
        layers.append(Layer(
            dimension=self._dimension,
            input_len=self._input_len,
            initial_codebook=np.array(weights, dtype=np.float32),
            sigma=self._sigma,
            learning_rate=self._learning_rate,
            neighborhood_function=self._neighborhood_function,
            activation_distance=self._activation_distance,
            random_seed=self._random_seed))
    self.layers = layers

# initialize the distances between layers as a fraction of the distance
# used default gaussian with sigma = 1.0 for distance between layers
# distance to layer itself (index 0) is 1.0
# shape: (num_layers)
def _create_layer_distances(self) -> np.array:
    x_mash, y_mash = np.meshgrid(np.arange(1), np.arange(self._num_layers))
    d = 2
    ax = np.exp(-np.power(x_mash-x_mash[0], 2)/d)
    ay = np.exp(-np.power(y_mash-y_mash[0], 2)/d)
    layer_distances = (ax * ay).T[0]
    layer_distances *= self._layer_distance_ratio
    layer_distances = np.insert(layer_distances, 0, 1.0)
    return layer_distances

# create a weights matrix for two aspects in a feature matrix
# the shape corresponds to shape (num_layers, input_len)
# weights are an interpolation of 0 to 1 for aspect A and 1 to 0 for aspect B
# shape: (num_layers, x, y, input_len)
def _create_weights_by_layer(self) -> np.ndarray:
    if self._aspect_selection.shape[0] != self._input_len:
        raise AttributeError('aspect_selection has to have the same dimension as input_len')
    column_weights = []
    weights_aspect_A = np.linspace(0, 1, self._num_layers)
    weights_aspect_B = np.linspace(1, 0, self._num_layers)
    for i in self._aspect_selection:
        if i:
            column_weights.append(weights_aspect_A)
        else:
            column_weights.append(weights_aspect_B)
    return np.column_stack(column_weights)

# initialize all layers of the aligned SOM
# for each layer in the Aligned SOM a Layer(MiniSom) is created
# the common weights (codebook) for all layers are created either randomly or pre-trained
# each layer is initialized with the common codebook weighted by the aspect selection
def _create_layers(self) -> List[Layer]:
    layers = []
    if self._codebook_initialization_type == 'random':
        initial_codebook = self._create_random_weights()
    elif self._codebook_initialization_type == 'pretrained':
        initial_codebook = self._create_weights_by_training_one_some()
    else:

```

```

        raise AttributeError('codebook_inizialization_type has to be
for weights in self._weights_by_layer:
    layers.append(Layer(
        dimension=self._dimension,
        input_len=self._input_len,
        initial_codebook=np.array(initial_codebook * weights, dtype
        sigma=self._sigma,
        learning_rate=self._learning_rate,
        neighborhood_function=self._neighborhood_function,
        activation_distance=self._activation_distance,
        random_seed=self._random_seed))
    return layers

# crate a random codebook with
# shape: (x, y, input_len)
def _create_random_weights(self) -> np.ndarray:
    if self._random_seed:
        np.random.seed(self._random_seed)
    return np.random.random((self._dimension[0], self._dimension[1],

# create codebook weights by training one SOM
# trained on not weighted features (same as middle layer)
# shape: (x, y, input_len)
def _create_weights_by_training_one_som(self) -> np.ndarray:
    middle_som = MiniSom(
        x=self._dimension[0],
        y=self._dimension[1],
        input_len=self._input_len,
        sigma=self._sigma,
        learning_rate=self._learning_rate,
        neighborhood_function=self._neighborhood_function,
        topology='rectangular',
        activation_distance=self._activation_distance,
        random_seed=self._random_seed)
    middle_som.train(self.data, 1000)
    return middle_som.get_weights()

```

Implementation of the Visualizations

```

In [ ]: def plot_aligned_som(asom: AlignedSom, data: np.ndarray, visualization_fu
        """Plot the aligned SOM

Args:
    asom (AlignedSom):
        trained aligned SOM to plot
    data (np.ndarray):
        input data to use for the visualization
    visualization_function (Callable, optional):
        Which visualization to use. Options are: SDH, HitHist and UMa
    num_plots (int, optional):
        How many intermediary plots to show. Defaults to 5.
    value_range (tuple, optional):
        Value range of the histogram given as tuple of min and max va
    kwargs:
        Additional arguments to pass to the visualization function

Returns:
    matplotlib figure: Figure object_

```



```

"""
assert num_plots <= asom._num_layers, "Number of plots must be less t

# calculate the histograms
visualizations = []
for layer_weights in asom.get_layer_weights():
    layer_weights = np.reshape(layer_weights, (asom._dimension[0] * a
    if visualization_function == UMatrix:
        histogram = visualization_function(
            asom._dimension[0], asom._dimension[1], layer_weights, da
    else:
        histogram = visualization_function(
            asom._dimension[0], asom._dimension[1], layer_weights, da
    visualizations.append(histogram)

# decrease figure size to increase plotting speed for larger plots
if num_plots > 32:
    figsize = (0.75 * num_plots, 0.6125)
if num_plots > 16:
    figsize = (1.5 * num_plots, 1.25)
elif num_plots > 8:
    figsize = (3 * num_plots, 2.5)
else:
    figsize = (6 * num_plots, 5)

max_value = np.max(np.array(visualizations))

# create the plot
figure, axis = plt.subplots(1, num_plots, figsize=figsize)
for i, vis_i in enumerate(np.linspace(0, asom._num_layers - 1, num_pl
    hp = sns.heatmap(visualizations[vis_i], ax=axis[i], vmin=0, vmax=
    hp.set(xticklabels=[])
    hp.set(yticklabels=[])
    axis[i].tick_params(left=False, bottom=False)
    weight_aspect_a = asom._weights_by_layer[vis_i][np.nonzero(asom._
    weight_aspect_b = asom._weights_by_layer[vis_i][np.nonzero(asom._
    if weight_aspect_a.shape[0] == 0:
        weight_aspect_a = 0
    else:
        weight_aspect_a = weight_aspect_a[0]

    if weight_aspect_b.shape[0] == 0:
        weight_aspect_b = 0
    else:
        weight_aspect_b = weight_aspect_b[0]

    hp.set(xlabel=f"A: {round(weight_aspect_a,2)}, B: {round(weight_a
figure.suptitle(visualization_function.__name__)
plt.show()
return figure

```

Example on Animals Dataset

We show here a small example on the animals dataset as in the paper "Aligned Self-Organizing Maps" to visually compare our results. The dataset comprises 16 records of different kinds of animals, described by 13 binary-valued attributes. The animals can be categorised into three classes: birds, carnivores, and herbivores.

The features are split into activity (aspect A) and appearance (aspect B) features.

activity features: hunt, run, fly, swim

appearance features: small, medium, big, 2_legs, 4_legs, hair, hooves, mane, feathers

```
In [ ]: # define params
SEED = config.SEED
N_LAYERS = 31
SOM_DIM = (3, 4)
TRAIN_STEPS = 1000

# load data
input_data, components, weights, classinfo = load_dataset('animals')
data = input_data['arr']

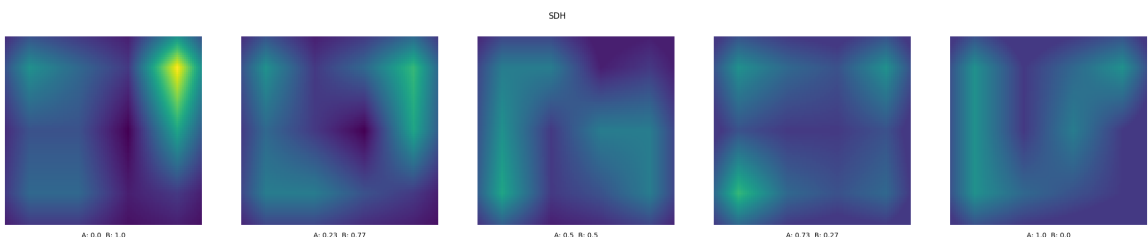
# aspect A: activity features (hunt, run, fly, swim)
# aspect B: appearance features (small, medium, big, 2_legs, 4_legs, hair, hooves, mane, feathers)
aspect_selection = np.array([0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 1, 1, 1, 1])

# create and train AlignedSom
asom = AlignedSom(
    SOM_DIM,
    data,
    aspect_selection,
    num_layers=N_LAYERS,
    random_seed=SEED)
asom.train(TRAIN_STEPS * N_LAYERS)
```

100%|██████████| 31000/31000 [00:43<00:00, 710.99it/s]

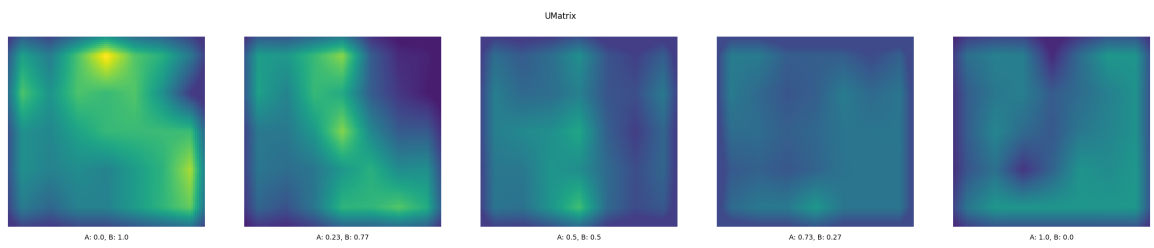
Smoothed-Data-Histogram

```
In [ ]: fig = plot_aligned_som(
    asom, data,
    visualization_function=SDH,
    num_plots=5,
    upscaling_factor=100,
    factor=2)
```



U-Matrix

```
In [ ]: fig = plot_aligned_som(
    asom, data,
    visualization_function=UMatrix,
    num_plots=5,
    upscaling_factor=100)
```



Hit Histogram

```
In [ ]: fig = plot_aligned_som(  
    asom, data,  
    visualization_function=HitHist,  
    num_plots=5,  
    upscaling_factor=100)
```

