

Spike-time Dependent Feature Clustering

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Abstract: In this paper, we present an algorithm capable of spatially encoding the relationships between elements of a feature vector. Spike-time dependent feature clustering positions a set of points within a spherical, non-Euclidean space using the timing of spiking neurons. The algorithm uses an Hebbian process to move feature points. Each point is representative of an individual element of the feature vector. Relative angular distances encode relationships within the feature vector of a particular data set. We demonstrate that trained points can inform a feature reduction process. It is capable of clustering features whose relationships extend through time (e.g., spike trains). In this paper, we describe the algorithm and demonstrate it on several real and artificial data sets. This work is the first stage of a larger effort to construct and train artificial dendritic neurons.

1 INTRODUCTION

Every data set describes a small world. The events or objects which make up these world are typically represented by an N -dimensional vector of features. When N is large the shape of these worlds becomes hard to grasp because doing so requires insight into the $N \times N$ feature relationships which are dispersed across all elements of the data set.


Dimensionality reduction (DR) aids in understanding these small worlds by reducing the number of feature relationships. In general, DR is divided into two areas: feature selection (FS) and feature extraction (FE). FS is the process of choosing a subset of the original features (Chandrashekar and Sahin, 2014). FE reduces the feature vector by casting it via a transformation into a reduced space (Khalid et al., 2014). The majority of DR methods rely on a measurement of feature importance. This measurement is used either in the selection process (FS) or is contained within the transformation of the set of features into a reduced space (FE).

Spike-time dependent feature clustering (SFC) provides a new way to measure relationships between features. SFC encodes feature relationships in the relative positions of representative points within a spherical space. Hebbian learning (Hebb, 2005), driven by individual feature values, is used to move the points toward or away from one another. Hebbian learning is a theory of neural plasticity which posits that changes

in the brain are in part caused by the temporal proximity of two or more spikes emitted by connected neurons. As a result, proximity within the spherical space is the product of (dis)similarities between features within the data set. And each feature relationship dispersed throughout the data set is compressed into a single angular distance between two points. Once a data set is encoded, traditional clustering or FS algorithms can be used to extract important features. An optimal method of identifying the resulting clusters--shape, size, or density--is beyond the scope of this paper.

2 ALGORITHM DESCRIPTION

The clustering process uses a spiking neural network (SNN) (Maass, 1997). A SNN consists of spiking neurons which emit a pulse or *spike* when an internal value reaches a threshold. A change in input does not immediately cause a change in neural output but rather an output spike is a product of input over time. Therefore, the relative timing of spikes encodes a neuron's recent input history. For this project, we use the Izhikevich spiking neuron model (Izhikevich, 2007). The Izhikevich model consists an equation for v , the membrane potential, and u , the recovery current (Eqs. (1) and (2)). A neural spike ($v \geq v_{peak}$) is modeled by a reset ($v \leftarrow c, u \leftarrow u + d$). The parameters used in the Izhikevich model are those given in (Izhikevich, 2007) for a regular spiking neuron:

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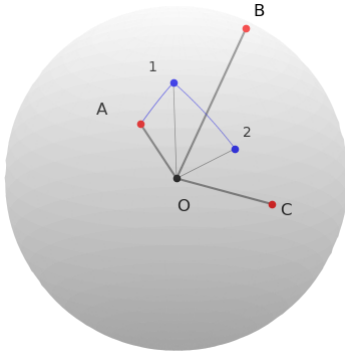


Figure 1: Movement across the spherical field. Three points (A , B , and C) correspond to three inputs. O is the field’s origin. A spike in A ’s neuron causes A to move toward B and C based on how recently their neurons emitted a spike.

$$C = 100, v_r = 60, v_l = -40, k = 0.7, v_{peak} = 35, a = 0.03, b = -2, c = -50, d = 100.$$

$$C\dot{v} = k(v - v_r)(v - v_l) - u + I \quad (1)$$

$$\dot{u} = a\{b(v - v_r) - u\} \quad (2)$$

Spike-time dependent clustering utilizes a single layer of spiking neurons. The spiking neurons transform values in the examples to a time series of spikes. Each neuron is connected to a unique point in the spherical space. For all experiments, initial point positions are uniformly randomized over a unit shell (identical radius). We refer to the spherical space (or shell) occupied by the points as the *field*. The distance between two points is then the angle of separation with respect to the origin (e.g., $\angle AOB$ in Figure 1 gives the distance between A and B).

The co-activity of input neurons causes their associated points to change position. Co-activity is defined as the temporal proximity of two (or more) spikes. *Temporal proximity* is a model hyperparameter given by W in the equations below. In Figure 1, a spike at A causes it to move toward B and C . The magnitude of change depends on how recently the neurons connected to B and C have spiked. In this example, C ’s neuron has spiked more recently than B ’s. Therefore, A ’s movement toward C is greater (or $\angle 1O2 > \angle AO1$). Lastly, point 2 would give the new location of A .

Co-activity is bounded by the *learning window*, W . The learning window determines the maximum temporal distance between two spikes that can affect positional change. The learning window for all experiments was set to 100 time steps. Furthermore, the length of the learning window partitions measurements of the field. In other words, the learning window divides the maximum angular distance between

two points, or π , giving discrete distance markers toward which points move. The goal marker of any given instance of co-activity is the temporal distance between two spikes. For example, if the learning window is set to 100 time steps, and one neuron routinely spikes 50 time steps before another, the latter’s point will move toward a location $\frac{\pi}{2}$ distance away from the former.

Point movement also depends on the sign and strength of an error signal. For all experiments, the error signal is within the range $[-1, 1]$. Negative error values cause co-activity to drive points apart (see Equation (4)). Negative error values can be used to separate points involved in unwanted co-activity during a training process. Or, as they are used in several of the following experiments, negative values can separate clusters or partition active and inactive points.

3 EXPERIMENTS

We tested the algorithm’s clustering ability on artificial and real-world data sets. The goal of these experiments was to discover whether point clusters formed by the algorithm revealed underlying patterns in the data sets. Although the types of input used across the four experiments were different, we attempted to make these differences invisible to the network. All input signals were transformed into spike trains using a Poisson process. Parameters intrinsic to the neural and point models were identical through all experiments.

Experiments 3.1 and 3.2 create patterns by using spike trains with a similar spike rate (e.g., 100 Hz). The spike trains themselves are randomly created by a Poisson process. Neurons that do not belong to a pattern are given a lower spike rate (e.g., 10 Hz) to simulate noise. Each input spike train is 1,000 time steps in length. An *iteration* consists of one spike train from each type of pattern and/or mask, and their order within an iteration is random. A *mask* is a pattern-less set of spike trains used in conjunction with a negative error signal to drive points away from each other. Spikes produced by the Poisson process decayed over time at a rate of $e^{-0.1t}$ where t is the time since the last spike. Input spikes were scaled by a weight of 500.

3.1 Two Overlapping Patterns

For this experiment, one hundred input neurons were divided into two overlapping patterns, A and B (see Figure 2). Each pattern consists of fifty-five 100 Hz and forty-five 10 Hz inputs. The inputs receiving stronger input represent the pattern and the weaker

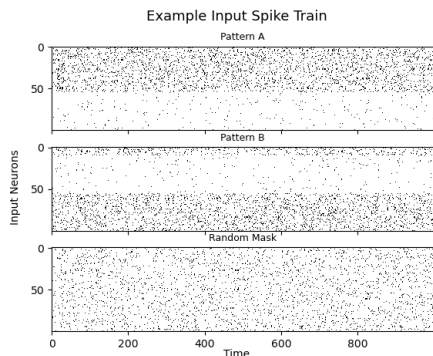


Figure 2: An example spike train for one iteration of both patterns and the random mask. Pattern A: 100 Hz inputs 0-54, 10 Hz 55-99. Pattern B: 100 Hz inputs 0-9 and 55-99, 10 Hz 10-54. Mask: 50 Hz inputs 0-99.

inputs represent noise. Ten of the stronger 100 Hz inputs are shared by both patterns. The network was shown 5,000 iterations. Included in each iteration was a random mask of identical length during which all neurons received an input of 50 Hz.

The treatment of the mask pattern differed from patterns A and B in that the accompanying error signal was negative (e.g., -1). For A and B, the error value is 1. The mask was included in this (and the following experiment) to separate clusters within the field.

Figure 3 shows the resulting point-by-point angular distances with points grouped by patterns along each axis. Points unique to a pattern cluster together (large dark blue patches). These clusters occupy disjoint areas of the dendritic field (opposing warm colors). The shared points form a separate cluster distinct from the two larger clusters of unique points. The shared cluster is approximately equidistant from the rest of patterns A and B. Two points taken from A and B are on average $\frac{\pi}{2}$ radians from one another and the shared points are $\frac{\pi}{4}$ from the unique points. In a simplified version of this experiment in which patterns A and B are composed of disjoint sets of points, clusters polarize within the field and the distance between points in opposing clusters is approximately π .

Shared points cause patterns to draw closer together. The greater the overlap, the closer the clusters. This effect can be seen to a greater extent in the experiment involving four overlapping patterns.

3.2 Four Overlapping Patterns

Next, we expanded the number of patterns and the degree and complexity with which they overlapped. Each pattern (A, B, C, and D) is comprised of twelve neurons. Four of the twelve neurons form two pairs

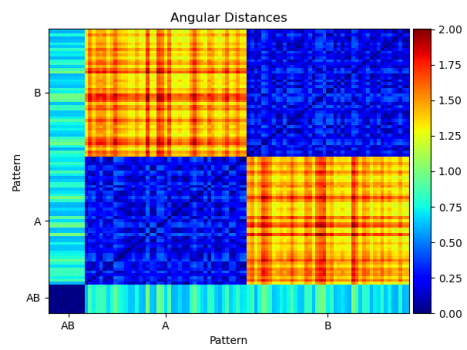


Figure 3: Resulting point-point angular distances grouped by pattern after 5,000 iterations. The overlapping points can be seen on the bottom left.

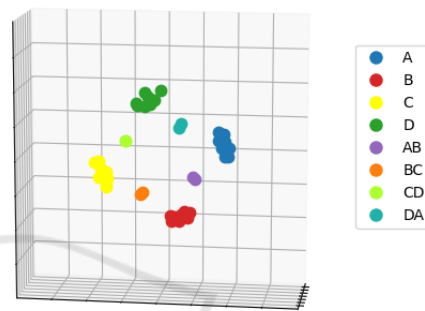


Figure 4: Point locations after 16,000 iterations. Colors indicate which pattern(s) each point belongs to.

which are shared with two other patterns (AB, BC, CD and DA). The goal of this experiment was to investigate whether resulting clusters would themselves cluster in such a way to resemble meta-patterns within the input data set. All parameters were unchanged except for the number of iterations which were increased to 16,000.

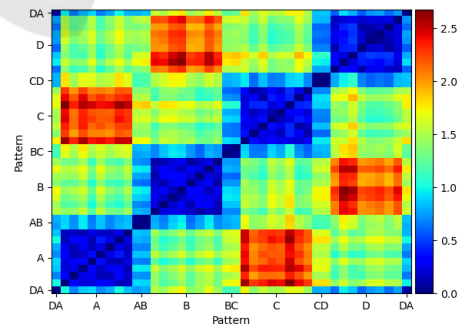


Figure 5: Resulting point-point angular distances grouped by pattern. Distance is measured in radians.

Points belonging wholly to one pattern cluster tightly together while shared points come to rest between the clusters of their dual membership. The meta-pattern formed by these clusters demonstrates a kind of adjacency via shared points. Patterns which

do not share points (i.e., A and C) oppose each other spatially. There is a clear distinction between the average angular distance separating points of adjacent versus non-adjacent patterns.

Several more versions of this experiment were run which included more patterns and more complex pattern overlap. Additionally, we examined whether SFC was capable of fuzzy membership (continuous range of rates), subpatterns (stepped rates) and additional meta-patterns (hierarchy of rates in overlapping neurons). These additional experiments reinforced the results evident in these simpler versions.

3.3 iBeacon Locations

The two previous experiments made use of artificially generated data sets. Such sets can be large enough to cover the relatively simple sample space of the entire feature vector. To understand how well spike-time dependent clustering works with sparse, real-world data sets, we attempted to reproduce the placement of iBeacons from a machine learning data set (Mohammadi et al., 2017) of RSSI readings taken from the UCI Machine Learning database (Dua and Graff, 2017).

Since SFC can train points using an unsupervised process, we were able to use the entire data set (training and testing). We removed all entries that did not contain two or more RSSI signals greater than the baseline (-200). In isolation single signal entries create no spatial association between beacons. Keeping with the other three experiments, the order of readings was randomized each iteration; therefore, associations between single signals could not be based on the historical route taken through the space by the original researcher. Figure 6 shows the grid-based placement of the iBeacons from which beacon-to-beacon distances were taken.

This modified data set was used as input to a network with thirteen neurons (one for each beacon). RSSI signals were transformed into constant rate input by the equation $I_b = \frac{S_b + 200}{200}$ for all $b \in \{0, 1, \dots, 12, 13\}$ which was then used to generate an input spike train. S_b is the RSSI signal for iBeacon b and I_b is the corresponding input. An iteration consisted of one full pass through the modified data set. Between iterations entry order within the data set was randomized.

To analyze the results we compared real-world distances to the final positions generated by the algorithm. First, both the angular distance of points and iBeacon Cartesian distances were normalized using min-max normalization. The absolute value of their difference was then taken and treated as the error

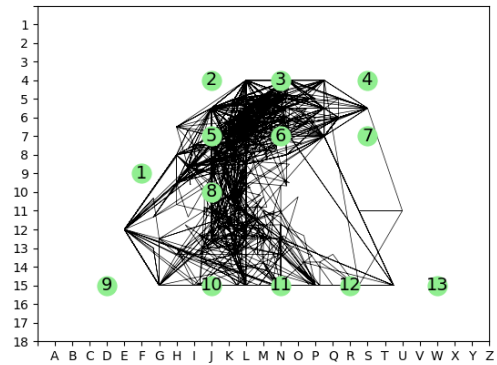


Figure 6: Reproduction of the iBeacons' physical positions (green numbered circles) with the paths produced by entries with multiple signals in both labeled and unlabeled data sets (black lines). Indoor features of the original map have been omitted. iBeacon positions have been adjusted slightly to align perfectly with the nearest row and column. Path vertices are average locations (without respect for signal strength).

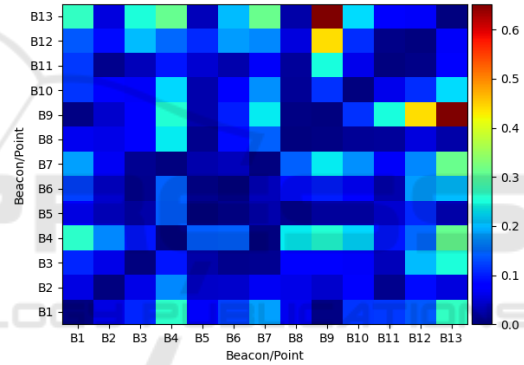


Figure 7: Absolute element-wise differences of distances between point and iBeacon positions. Point-to-point and iBeacon-to-iBeacon distances were normalized using min-max norm before the difference was taken.

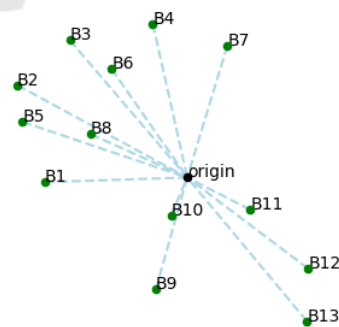


Figure 8: Point positions within the dendritic field resulting from 500 iterations. Light blue lines orient points to the origin and give a sense of depth. Labels indicate which iBeacon is associated with each point.

rate. Figure 7 depicts this error after 500 training iterations. To establish a baseline, we generated random

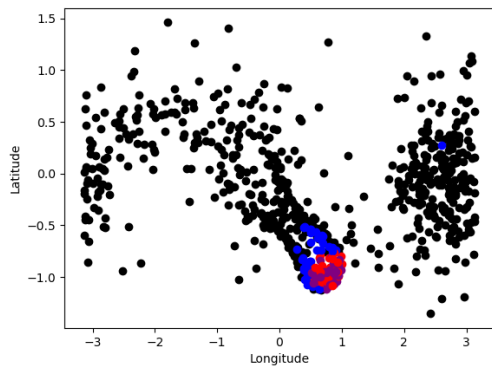


Figure 9: Digit-9 point locations. Red points indicate points used to identify digit-9 images in the test set. Blue is the same for digit-4. Purple are overlapping points.

point positions and compared them using the above procedure. The error produced by random placements (100,000 trials) had a mean of 23.7% and average median of 20.1%; whereas differences with final point position after 500 iterations had a mean of 10.5% and median of 7.1%.

The relative positions of trained points roughly correspond to actual iBeacon placement (Figure 8). Distortions are due to examples with overlapping signals that cover either a broad (> 2 beacon signals) or discontinuous (e.g., beacon 2 and 4 are heard but not 3) area. The largest error between beacons 9 and 13 appears to be caused by the frequent simultaneous hearing of beacons 11 through 13 and the lack of signal overlap between these and 6 and 7. The latter is due to a lack of direct paths between 6-7 and 11-13 owing to obstacles (see original map). The same can be seen in the points 1 and 2 with respect to 5. Within the indoor space, there is no direct path from 1 to 2 without passing 5; therefore, points 1, 5 and 2 form a straight line across the field. This suggests that the map of trained points represents obstacles as well as iBeacon proximity and is a product of both actual beacon location and available paths.

3.4 MNIST

How can the points of a trained field facilitate useful dimensionality reduction? To answer this question, we trained ten networks on the training set of images of the MNIST handwritten image database (LeCun et al., 2010) and investigated whether their points could be used to identify the categories of testing images.

The method of training was similar to that used to train the overlapping Poisson experiments. Image pixel intensities were used to generate Poisson process spike trains which were used as inputs to 784

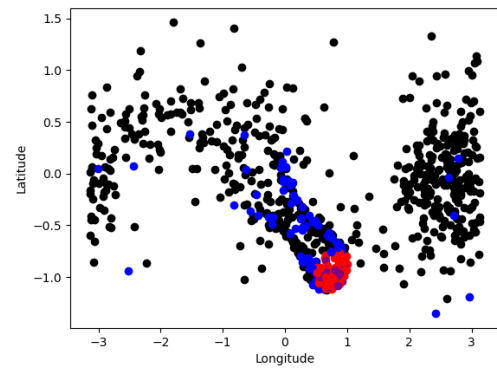


Figure 10: Digit-9 point locations. Red dots indicate points used to identify digit-9 images in the test set. Blue is the same for digit-2. Purple are overlapping points.

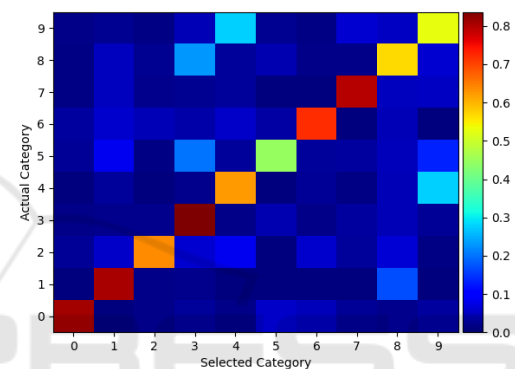


Figure 11: Accuracy matrix showing what percent of each image (by category) is identified as belonging to a particular category.

neurons. Rate was determined by $\frac{I_p}{255}$ where I_p is the intensity (0-255) of pixel p . Each network was trained on the full set of testing images of a specific category. After every ten images the networks were shown a random mask with all inputs spiking at 50 Hz and provided an error value of -1. This caused the inactive points to move away from clustered points.

Next we investigated whether tightly clustered points within the trained field could be used to recognize images from the test set. Clusters were identified by summing the angular distances of each point to all other sibling points and selecting the N points with the smallest summed angular distance. To find N , we tried several values: 20, 50, 100, 150 and 200. $N = 100$ produced the best results. It is likely that optimal results per network require a different N .

For each training image, we summed the intensities of the pixels which corresponded to the selected points of each trained field. The field with the maximum value identified the selected category. Results are shown in Figure 11.

The accuracy of each image category corresponds to a degree of overlap between two sets of selected

points. This can be seen in Figures 9 and 10. Digit-9 images were most often mistaken by this method to be digit-4 images (27.5%) and least often to be mistaken as digit-2 images (0.4%). The points belonging to the digit-9 and digit-4 clusters overlap significantly (Fig. 9). They share 61 points which correspond to the same pixels. This stands in contrast to the digit-9 and digit-2 clusters which share only 30 (Fig. 10). Interestingly, the non-overlapping points of digit-9 and digit-4 clusters show a greater degree of clustering within the field than those of digit-9 and digit-2. The greater the overlap and lower the element-wise inter-cluster distance the lower the accuracy due to misidentification.

We chose a simple cluster identification method (min of the sum of angular distances) for this experiment to emphasize that accuracy depends on field position and not additional cluster analysis. That said, more complex point selection algorithms which examine the shape and size of clusters as well as degrees of membership are likely to produce better results. As shown in the iBeacon experiment, the shape of a cluster can encode hidden information. More complex cluster analysis is left for future work.

4 METHODS

This section gives the methods governing the movement of points within the field. The process is Hebbian in nature in that it is based on the relative timing of afferent spikes.

$$t_{\delta}^i = t_c^i - t_p^i \quad (3)$$

t_{δ}^i in Eq. 3 gives the length of time between the current and previous spike of neuron i . t_c^i is the time stamp of the current input spike received by point i . t_c^i is equivalent to the current simulation time step as angular movement is enacted immediately. t_p^i is the time stamp of the previous input spike received by point i .

$$F_i = \frac{(t_{\delta}^i + W)}{W} e^{-\frac{t_{\delta}^i}{W}} E \quad (4)$$

Eq. 4 gives a scalar to positional change based on the length of time between neuron i 's two most recent spikes. W represents the effect of time. Higher values allows longer inter-spike intervals to impact point movement. W is fixed throughout the simulation at 100. Larger values slow temporal decay; smaller values increase it. t_{δ}^i is taken from Eq. 3. E is the current error value which is fixed at 1 (except for random mask trials when it is set to -1). Negative values cause co-activity to repel points.

Next, the recent activity of all sibling points is evaluated. A copy of point i 's position, \hat{P}_i , is orbited toward (or away from) all sibling points to determine the direction and magnitude of movement for point i . For all sibling points j , where $i \neq j$ we calculate the following temporal distance.

$$t_{\delta}^{ij} = t_c^i - t_c^j \quad (5)$$

t_c^j is the time of neuron j 's most recent spike and t_c^i is the same as Eq. 3. Therefore, t_{δ}^{ij} measures the interval of time between i and j 's most recent spikes.

$$D_t = \pi \frac{t_{\delta}^{ij}}{W} \quad (6)$$

D_t represents the target angular distance between point i and j . W is the same as given in Eq. 4. When values of t_{δ}^{ij} exceed W , the point is ignored.

$$D_{ij} = D_c - D_t \quad (7)$$

D_{ij} in Eq. 7 is the difference between the current angular distance (D_c) between points i and j and their target angular distance (D_t from Eq. 6). This and the value of F_i form the magnitude of angular change of point i toward point j (Eq. 8).

$$M_{ij} = F_i D_{ij} \quad (8)$$

\hat{P}_i is then orbited across the spherical plane in the direction of P_j , the spherical position of j , by the magnitude given by M_{ij} .

Once all sibling points have been evaluated for their effect on \hat{P}_i , the actual position of i , P_i , is changed. P_i is orbited toward \hat{P}_i by the angular distance between them times the positional learning rate, λ which is currently set to 0.01. This value is the result of a parameter sweep from 0.1 to 0.001.

5 DISCUSSION

Spike-time dependent clustering uses a Hebbian rule to position points within a spherical field. Each point is associated with one element of a feature vector. The relative positions of trained points encode aspects of the set of input examples. In the iBeacon experiment, we showed that from overlapping RSSI signals we can reproduce the approximate layout of the beacons themselves. And in the MNIST experiment, we can draw out a set of representative pixels which can be used to identify image categories.

There are a number of challenges in applying this approach to complex data sets. First, SFC assumes each feature has a near unique position within the

overall input space. If they do not, for instance if the centering or rotation of MNIST digits were randomized, SFC's results will be much degraded. Another process would be needed to identify bounding boxes and rotation. Alternatively, SFC could be used to construct multiple layers of fields which, similar to parts of the visual cortex, move from simple, invariant patterns (lines) into more complex ones (digits).

Another challenge is the separation of patterns or irrelevant (or inactive) features. All experiments except the one involving iBeacon signals used random masks and a negative error signal to create separation. Without a separating force, inactive features will always occupy their initial position. Clusters do not separate on their own. Cluster orientation and position across the spherical field is relative. Irrelevant features could, by their starting position, insinuate themselves into a cluster. It is our goal to find a method by which masks are not necessary or at least not for this purpose. A possible solution is the inclusion of polarity among points. A class of negative points would attract positive co-active points but repel each other.

Next, the order in which points move toward other points matters. In Figure 1, a movement first toward *C* and then *B* would produce a different new *A*. To combat this, we randomize update order between each iteration. The impact of update order is unknown. However, without randomization, points have a tendency to orbit the field while maintaining relative distances.

Co-activity depends on the length of the learning window. If it is too short, co-activity over longer intervals will be missed. If the window is too long, there is a tendency for all points to form a single cluster. The latter also depends on the shape of the Hebbian function. One possibility is to use multiple discrete windows across different networks to identify long and short term relationships. This aspect is a challenge of neural plasticity algorithms and credit assignment in general.

A significant problem in data analysis concerns the identification of the number of classes within a data set. In the MNIST experiment, we presupposed the number of classes by training ten different sets of points and selecting one set of features from each field. In an unlabeled context, this is not possible. One possibility for future research is to use SFC as a wrapper where results from each iteration are used to partition the data set for subsequent iterations.

In conclusion, we note that SFC is part of a larger project to create artificial neurons with fully trainable dendrites. Neuroscience evidence suggests that dendritic neurons are computationally attractive (Mel,

1999). Such dendritic neurons would have expansive dendritic trees in which synaptic (i.e., point) position, tree shape, branch compartmentalization and synaptic weights are all trainable parameters. Spike-time dependent feature clustering describes the first stage of this work: synaptic position on a spherical plane. SFC has a second part not detailed in this paper. The second half uses co-activity of input and output neurons to adjust the radius of a point (or synapse). In other words, this paper describes clustering due to input-input co-activity, while the other half describes clustering due to input-output co-activity. The second half was not included because the goal of this work was to show that input co-activity can be used to encode small world information in the position of synapses in a dendritic field.

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