# Contextual Saliency for Nonrigid Landmark Registration and Recognition of Natural Patterns

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Abstract:

In this paper we develop a method for injecting within-pattern information into the matching of point patterns through utilising the shape context descriptor in a novel manner. In the domain of visual animal biometrics, landmark distributions on animal coats are commonly used as characteristic features in the pursuit of individual identification and are often derived by imaging surface entities such as bifurcations in scales, fur colouring, or skin ridge minutiae. However, many natural distributions of landmarks are quasi-regular, a property with which state-of-the-art registration algorithms have difficulty. The method presented here addresses the issue by guiding matching along the most distinctive points within a set based on a measure we term *contextual saliency*. Experiments on synthetic data are reported which show the contextual saliency measure to be tolerant of many point-set transformations and predictive of correct correspondence. A general point-matching algorithm is then developed which combines contextual saliency information with naturalistic structural constraints in the form of the thin-plate spline. When incorporated as part of a recognition system, the presented algorithm is shown to outperform two widely used point-matching algorithms on a real-world manta ray data set.

#### 1 INTRODUCTION

Registering sets of image locations between various visual captures or models is an important stage in many computer vision applications including optical character recognition (Belongie, Malik, & Puzicha, 2002) and medical image registration (Rueckert & Schnabel, 2011).

Solving this registration problem essentially requires the derivation of a correspondence mapping between two point sets which are related by some initially unknown geometric transform. This is difficult due to the combinatorial explosion of possible between-image point correspondences and usually very large transformation parameter spaces, especially in cases of non-rigid transformation, position noise and partial occlusion.

Here we consider the point set registration problem in the context of animal biometrics (Kühl & Burghardt, 2013), specifically with regard to the identification of individual animals utilizing their characteristic coat markings as identifiers. Prominent examples of unique coat patterns include spot or blob configurations on cheetahs or manta

rays, and the alignment of stripes on zebras.

Animal biometrics is widely used to enable non-invasive ecological monitoring and conservation-relevant population modelling (e.g. Gamble et al., 2008). Such systems often combine landmark distribution information with local appearance and textural cues to form descriptors of animal identity. However, here we analyse natural point-set registration in isolation of these other factors that may also contribute to identity recovery.

The difficulty in natural point-set registration is two-fold: firstly, transformations between point-sets are generally non-rigid and sometimes incomplete due to the wide range of possible animal poses and occlusion: and secondly, the stochastic morphogenesis (Turing, 1952) of coat patterning can often lead to a dense, widely homogenous layout of landmarks from which points are extracted. We address these difficulties through introducing a transform-persistent and noise-robust intra-pattern ordering of points, which is used to guide the registration process alongside traditional interpattern distance measures. We formally define this ordering measure using the shape context descriptor

(SC, Belongie et al, 2002) and name it *contextual* saliency,  $\Psi$ .

First, experiments on synthetic point-sets are reported which investigate the robustness of point ordering on  $\Psi$  to noise and perspective transformation and whether this ordering is associated with correct correspondence. A general non-rigid point-matching algorithm,  $\Psi$ -Match, is then described which incorporates an ordering of points on  $\Psi$  in iteratively building a correspondence set between point-sets. Finally, results from an identification experiment on a real-world manta-ray data set are reported, comparing the performance of a recognition setup using the  $\Psi$ -Match algorithm to the same setup when employing two widely-used registration algorithms.

### 2 RELATED WORK

In the general case of gauging similarity between two sets of points in  $\mathbb{R}^2$ ,  $\mathbf{X} = \{x_1, x_2, ..., x_n\}$  and  $\mathbf{Y} = \{y_1, y_2, ..., y_m\}$ , a common approach is to apply a normalising transform to one or both sets, e.g.  $\mathbf{X} \to f(\mathbf{X})$ , followed by the application of a similarity measure in that space,  $d(f(\mathbf{X}), \mathbf{Y})$ . This value then quantifies the variation between the sets which was not captured by the transformation. Deriving a suitable normalising transformation between the sets can be seen as a counterpart to the correspondence problem: to indicate for points in one set the corresponding points in the other. Pointmatching has therefore been approached from both perspectives.

Methods which solve for the transformation, for example the Hough transform (Ballard, 1981), have proven successful in estimating rigid deformations, however the concept is unsuitable for highly non-rigid domains due to the associated dimensional explosion of the parameter space. Other methods concerned with deriving a correspondence set aim to construct higher-order structures from the point-sets (e.g. curves or surfaces; Metaxas et al., 1997), drastically reducing the parameter space at the expense of curve-fitting complexity when data are noisy due to reliance on accurate feature extraction and curve smoothness.

Shape contexts (SC, Belongie et al., 2002) are a different approach to the construction of higher order features; here each point in a set is represented by its spatial relationship to all other points, and correspondences are computed by comparing these descriptors between sets. SC utilises a circular histogramming approach where bins are uniform in

log-polar space, making the descriptor more sensitive to local structure, capturing the intuition that more proximate point pairs are likely to undergo more similar transformations when compared to more distal pairs.

Other approaches have adopted an iterated estimation framework. The iterated closest point algorithm (ICP; Zhang, 1994) operates on roughly aligned sets and assigns correspondence based on a simple nearest-neighbour criterion which is then used to update the transformation between the sets. The thin-plate spline robust point matching algorithm (TPS-RPM; Chui & Rangarajan, 2003) combines correspondence and transformation into a single objective function, and uses an EM-like optimisation routine to iteratively solve for both The routine leverages softassign (Rangarajan et al., 1997), a loosening of the binary correspondence criterion such that correspondences can take on probabilistic values; and course-to-fine deterministic annealing, where constraints on transformation non-rigidity and correspondence binarity are gradually reduced as the algorithm progresses. Other methods in this vein have approached point-matching through relaxationlabelling as a graph-matching problem constrained by neighbourhood relations (Zheng & Doermann, 2006) or as probability density estimation where point positions define centroids in Gaussian mixture models (Myronenko & Song, 2010).

# 3 THE CONTEXTUAL SALIENCY HYPOTHESIS

There are limitations with the above approaches in the domain of landmark-based animal biometrics due to the stochastic morphogenesis that underpins animal patterning (Turing, 1952). This process often leads to a dense, quasi-regular arrangement of landmarks across the animal surface. The lack of discernible shape in these patterns makes correspondence mapping between instances difficult, which becomes especially clear in the case of partial occlusion, where the aim is superimposing a subset of a point pattern upon the full pattern, a common occurrence in ecological imagery.

Here we demonstrate failures of TPS-RPM and SC approaches in this scenario using controlled, synthetic, quasi-regular point-sets. A first model set, Y, is depicted in Figure 1. It is a regular grid of equally spaced points with two points being slightly displaced, while the probe set is a rectangular subset,

 $X \subset Y$ , which includes those displaced points and is shown in red.

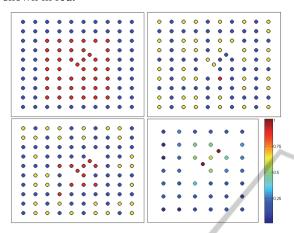


Figure 1: Top Left: Synthetic point-sets. The smaller set (red) is a subset of the larger (blue); a correct correspondence is shown as a superposition of the two sets. Results of TPS-RPM (top right) and SC (bottom left) algorithms on the synthetic data. Points in X are shown superimposed on Y according to the correspondences found: red points refer to correct and yellow points refer to incorrect assignments. Bottom Right: Contextual saliency calculated for each point, scaled to the range [0, 1], represented by colour.

Application of the TPS-RPM algorithm results in only a single correct correspondence between the sets (see Figure 1, top right). The algorithm favoured (incorrect) global characteristics and scaled **X** to cover **Y**, resulting in incorrect correspondences transformation. In this example SC performed better, yet still only aligned a few central points.

The present work aims to address this demonstrated shortcoming by developing a method of ordering points within sets across a measure we term *contextual saliency*. In the above example, the cluster of points centred near and including the displaced points is salient in comparison to the more distal points forming part of the regular grid: we therefore develop a procedure for biasing matching processes in favour of these points.

The intuitive reasoning is two-fold: points which are contextually salient are likely to remain contextually salient after distortion or deformation, and correspondences based on locally distinguishable points across patterns are more likely to be correct than correspondences drawn from the remaining pool of regular points. In Section 6 these intuitions are substantiated with experiments on controlled data.

# 4 FORMALISING CONTEXTUAL SALIENCY

We now utilise the shape context (SC) descriptor as a means for quantifying contextual saliency. For a given point  $x_i \in \mathbf{X}$ , the SC descriptor is defined as a histogram  $h_i$  of the remaining point positions, and is populated according to  $h_i(q) = \#\{x_j \neq x_i : x_j - x_i \in \text{bin}(q)\}$ , where bins are uniform in log-polar space. In the original application of SC, correspondences between point-sets  $\mathbf{X}$  and  $\mathbf{Y}$  are estimated by minimising an assignment problem of SC histogram distances based on the  $\chi^2$  statistic (Belongie et al., 2002). However, here we apply this distance measure within a set, such that it reflects similarity of a point to another point within the same set based on local structure. This similarity between two points,  $x_i \in \mathbf{X}$  and  $x_i \in \mathbf{X}$ , is given by

$$C_{ij} = rac{1}{2} \sum_{q=1}^{Q} rac{\left[h_i(q) - h_j(q)
ight]^2}{h_i(q) + h_j(q)},$$

where  $h_i(q)$  and  $h_j(q)$  represent the *Q*-bin normalised SC histograms of each point. To define the contextual saliency of a point  $x_i$  we compute the similarity measure between  $x_i$  and all other points in **X** giving a vector  $C^{\{i\}} = \{C_{i1}, C_{i2}, ..., C_{i(n-1)}\}$  of similarities. The minimum value of this vector is termed the contextual saliency of point  $x_i$ :

$$\Psi_i = \min_{x_j \in \{\mathbf{X}/x_i\}} C_{ij}.$$

A point with a large minimum similarity to other points within the set is therefore relatively dissimilar from all other points, and has high contextual saliency. Applied to the synthetic data set from Section 3, we see in Figure 1 (bottom right) that the contextual saliency measure has numerically captured this notion.

# 5 CHARACTERISTICS OF CONTEXTUAL SALIENCY

Here we investigate by controlled experiment the tolerance of the contextual saliency measure to point-set noise and perspective distortion, and then investigate whether contextual saliency is predictive of correspondence.

#### 5.1 Synthetic Data Generation

Quasi-regular point-sets were constructed by the addition of Gaussian noise to each point in a fully regular grid of points. For each initial grid point  $x_i^{init} \in \mathbf{X}^{init}$ , the regularity of the point-set X can then be controlled by the variance parameter,  $\sigma^2$ . To preclude point-set boundary information, we define a central subset of points,  $\mathbf{Q} \subset \mathbf{X}$ , such that a shape context descriptor placed at the extreme outer points of Q will not overlap edges of the complete set X. Q is the set with which inferences regarding the contextual saliency measure are made, whilst the remainder of points in X are not studied in themselves. In the following experiments, the initial point-set  $X^{init}$  is comprised of 900 points in a 30  $\times$ 30 regular grid. Set Q is a 10  $\times$  10 central subset of X. We used a shape context descriptor with 12 equally spaced orientation bins and 5 log-linear radial bins with the inner bin extending to 1/16 units and outer bin extending to 1.

#### 5.2 Distribution

The initial variance of the point-sets was varied in the range  $[1 \times 10^{-3}: 1.8 \times 10^{-2}]$  in 5 equal steps. For each setting, 200 separate point-sets were generated and the contextual saliency of the points calculated. Figure 2 presents the mean  $\Psi$ distribution across point-sets for each variance parameter.

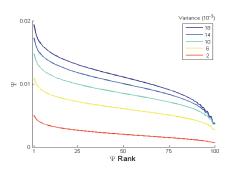


Figure 2. Mean Ψdistributions at given variances.

Critically, a pronounced peak is evident in  $\Psi$  for the most salient points at all probed variance values, indicating that a subset could be usefully isolated based on this measure.

#### 5.3 Preservation

We investigated whether contextual saliency values and point-ordering based on  $\Psi$  are tolerant to further addition of Gaussian noise and the application of perspective transformations. We calculated initial  $\Psi$ 

values and within-set rankings for point-sets with an initial variance of  $\sigma^2 = 5 \times 10^{-3}$ . Point-sets then underwent either the addition of Gaussian noise or perspective transformation, in the form of a plane rotation about the *x*-axis in 3D space, at 5 manipulation levels. For each level results were again averaged 200 point-sets.

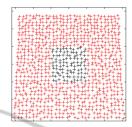


Figure 3: An example of perspective transformation; a rotation of  $10^{\circ}$  (set **Q** in black, set **X** in red).

The  $\Psi$  values of points after point-set degradation,  $\Psi_2$ , were plotted against the original ordering of points. As shown in Figure 4, for both types of set manipulation, highly salient points remain salient.

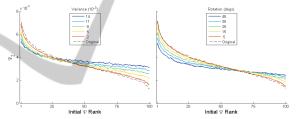


Figure 4: Ψvalues of points after (left) noise addition and (right) rotation plotted according to their initial Ψranking.

The preservation of ordering is shown in Figure 5: initially highly ranked points are more likely to remain in the same rank than other points within the pattern. We also see a marked peak at very low ranks which is explained as a form of group saliency whereby small groups of points are remarkable in their similarity and as such that points in this group remain similar after manipulation

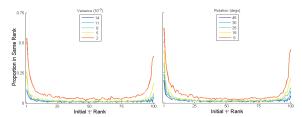


Figure 5: Average ranking by Ψof points within point-sets after (left) further noise addition and (right) rotation. The *x*-axis again refers to the point's initial ranking.

### 5.4 Predictor of Correspondence

Using the same experimental procedure, the relationship between contextual saliency ranking and correct correspondence was investigated. True correspondence between initial and transformed sets was known *a priori*, while experimental correspondences were derived as the global minimum over all pairwise SC histogram costs using the Hungarian method (Munkres, 1957).

Results are shown in Figure 6. While correspondence accuracy is generally reduced in more extreme transformations, initial contextual saliency is positively correlated with correspondence accuracy. Furthermore, this trend becomes stronger as the manipulation becomes more extreme.

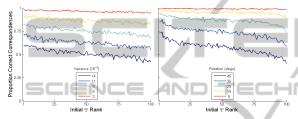


Figure 6: Correct correspondence proportion as a function of initial contextual saliency rank under (left) additional noise, and (right) perspective transformation.

# 6 A NEW ALGORITHM: Ψ-MATCH

We now introduce a novel algorithm for registering a test point-set X to a model set Y. The procedure iteratively builds a correspondence set comprised of a contextually salient subset of Y.

#### **6.1 Y-Biased Matching Approach**

Our approach is to extract subsets of the point-sets,  $\mathbf{X}_n \subset \mathbf{X}$  and  $\mathbf{Y}_n \subset \mathbf{Y}$ , such that the transformation between the two is representative of the transformation between the whole sets. Given an initial assumed correspondence set of n points  $\mathbf{X}_n^{init} \sim \mathbf{Y}_n^{init} = \{(x_1, y_1), (x_2, y_2), ..., (x_n, y_n)\}$ , the transformation mapping between them  $f_0: \mathbf{X}_n^{init} \rightarrow \mathbf{Y}_n^{init}$  is applied to the test set such that  $\mathbf{X} \leftarrow f_0(\mathbf{X})$ . An additional correspondence pair is then added to  $\mathbf{X}_n \sim \mathbf{Y}_n$ : the most contextually salient point in  $\mathbf{Y}$  is paired with that point in  $\mathbf{X}$  which elicits the lowest SC cost. The transformation is found between this expanded correspondence set, and the process repeated until some stopping criterion is reached.

For continuation, define the nested transformation function  $f_S(z) = f_{n-1}(f_{n-2} \dots f_0(z))$ , then on the *nth* iteration the subsets defining the correspondence set are given by

$$\mathbf{Y}_n = \mathbf{Y}_{n-1} \cup \left\{ argmax_{y \in \mathbf{Y}/\mathbf{Y}_{n-1}} \Psi(y) \right\}$$

where the second term represents the point  $y_n$ , the newest addition to the correspondence set, and

$$\mathbf{X}_n = \mathbf{X}_{n-1} \cup \{argmin_{x \in \mathbf{X}/\mathbf{X}_{n-1}} \mathcal{C}(y_n, f_{\mathbf{s}}(x))\},\$$

where C(a, b) represents the shape context distance between the points a and b.

In our case study, we initialise the correspondence set  $\mathbf{X}_n^{init} \sim \mathbf{Y}_n^{init}$  using manually tagged reference points common to a particular species: the edges of gills on manta ray bellies. However, a fully automatic system could be initialised using the most probable correspondences using SC, or if image data were available, a robust method such as SIFT or ASIFT (Yu & Morel, 2011). The necessary size of this initial correspondence set is dependent on the class of transformation used.

#### 6.2 Transformation Parameterisation

To ensure a naturalistic and smooth interpolation between the point-sets the transformation is parameterised by the thin-plate spline (TPS), which models a thin metal plate with certain resistance to bending. The TPS fits a mapping function, f(X), between corresponding point-sets X and Y through minimising the energy function:

$$E_{TPS}(f) = \sum_{i=1}^{N} ||y_i - f(x_i)||^2 + \lambda \iint_{\mathbb{R}^2} \left(\frac{\partial^2 f}{\partial x^2}\right)^2 + \left(\frac{\partial^2 f}{\partial x \partial y}\right)^2 + \left(\frac{\partial^2 f}{\partial y^2}\right)^2 dx dy.$$

While the first term quantifies the spatial distance between corresponding points, the second is a measure of the total curvature of the function, the importance of which can be controlled with the regularising parameter,  $\lambda$ , such that high  $\lambda$  strictly penalises non-affine warping. For a given regularisation parameter,  $\lambda$ , there exists a unique minimiser, f, of the form

$$f(x_i, d, w) = x_i \cdot d + \phi(x_i) \cdot w$$

where d is a  $(Dim + 1) \times (Dim + 1)$  affine transformation matrix, w is a  $N \times (Dim + 1)$  matrix of non-rigid warping coefficients, and  $\phi(x_i)$  is a  $1 \times N$  vector related to the TPS kernel where each entry  $\phi_j(x_i) = ||x_j - x_i||^2 \log ||x_j - x_i||$ . The TPS kernel

effectively contains information regarding the pointset's internal structure. Least-squares solutions for dand w are arrived at through minimising the QR decomposition of the energy function into separate affine and non-affine warping spaces according to the procedure described in (Wahba, 1990).

# **6.3** The Constancy Heuristic

The definite nature of the algorithm as introduced here is problematic in the case where a false correspondence is incorrectly admitted to  $\mathbf{X}_n \sim \mathbf{Y}_n$ . Therefore we admit a new correspondence pair only if that pair is chosen in two successive iterations: if in a subset of time during the dynamic fitting process we encounter a constant, this implies a meaningful correspondence since both the spline and SC matching are in agreement. There is however, the scenario where the algorithm does not produce consecutive matching correspondence pairs; to tackle this we build a history of previous candidate pairs and when a history limit is reached, the candidate pair with the most entries in the history is admitted to the permanent correspondence set (we use a limit of 5 in our case study and informal experimentation suggests that recognition performance is quite tolerant of this parameter). The number of pairs to admit to the set in total is explored experimentally in the case study

# 7 CASE STUDY: MANTA RAY RECOGNITION

Manta rays (*Manta birostris* and *Manta alfredi*) are suitable subjects for landmark-based matching systems due to a characteristic blob-pattern present on the underside which is thought to be a uniquely identifying biometric feature (Kitchen-Wheeler, 2011). Figure 7 below shows sample blob distributions of two different identity manta rays.

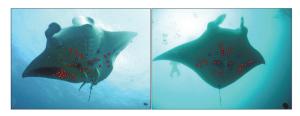


Figure 7: Examples of manually tagged manta rays. Red tags are placed on belly spots, while blue tags represent reference points placed at the end point of gills and the base of the tail; these form the initial known correspondence set.

We benchmark performance of our algorithm against TPS-RPM and SC in a recognition context.

#### 7.1 Dataset and Analysis Details

Images of 67 individual manta rays were taken in waters off the Maldives, most showing non-linear deformation and partial occlusion. Three-hundred images were used in the study; a mean of 4.5 images per individual ray, with a minimum of 2 and maximum of 7 instances per individual. Individual manta ray belly spots were tagged by experts, who also tagged reference points corresponding to the endpoints of the two most ventral gills and the base of the tail. The average number of points per pattern across all instances was  $\mu = 36.5$  with standard deviation  $\sigma = 20.4$ , and all point-sets were scaled to the range [0,1] in both the x and y-axes.

Recognition performance was compared for systems using SC, TPS-RPM, or  $\Psi$ -Match point-matching methods. For each method, the associated similarity measure was derived for each possible pair of manta instances, resulting in a 300  $\times$  300 similarity matrix; a correct recognition was declared when, for a given instance, the most similar pattern in a different image was of the same identity. We also investigated whether a same identity pattern was within the top N most similar patterns, useful information for a semi-automatic system providing the top N matches to an expert for ultimate recognition

### 7.2 Algorithm Parameters

For each comparison an initial scaling and alignment phase was conducted by recovering and applying the TPS transformation between the reference points. The a priori known correspondence between these points was enforced throughout for each algorithm (i.e. these assignments were always present in the correspondence set). Parameters for the TPS-RPM algorithm were taken from (Chui & Rangarajan, 2003). For the SC method, SC descriptors were constructed using 12 equally spaced orientation bins and 5 log-linear radial bins with inner radius of 1/16 and outer radius of 1, normalised through division by the mean pair-wise distance between points within each set. Correspondences were assigned based on the global minimum of all SC costs using the Hungarian method (Munkres, 1957). A TPS transformation was recovered between sets using these correspondences and applied to the transforming point-set before the application of a similarity measure; the TPS regularisation parameter

was set at the squared pair-wise Euclidean distance between points within the transforming set,  $\alpha^2$ , such that  $\lambda = \alpha^2$ , as in Belongie et al. (2002).

For the  $\Psi$ -Match approach, SC histograms and the TPS transformation were parameterised identically to the SC method. In this experiment we utilise the ratio of points incorporated into the correspondence set to the maximum number of points possible to incorporate (i.e. the number of points in the lowest cardinality point-set) as a means to parameterise  $\Psi$ -Match; we term this ratio  $\beta$ .

#### 7.3 Similarity

Similarity was measured following point-set transformation using the modified Hausdorff distance, a variant of the Hausdorff distance which has been shown to outperform the standard measure in the presence of position noise (Dubuisson & Jain, 1994):

$$d_{MH} = max \left\{ \frac{1}{N_{\mathbf{X}}} \sum_{\mathbf{x} \in \mathbf{X}} d(\mathbf{x}, \mathbf{Y}), \frac{1}{N_{\mathbf{Y}}} \sum_{\mathbf{y} \in \mathbf{Y}} d(\mathbf{y}, \mathbf{X}) \right\}$$

where  $d(x, \mathbf{Y})$  is the directed distance between point  $x \in \mathbf{X}$  and all points in  $\mathbf{Y}$  and is here defined as  $d(x, \mathbf{Y}) = min_{y \in \mathbf{Y}} ||x - y||$ .

### 7.4 Results

The recognition accuracy of the system using the  $\Psi$ -Match algorithm was first compared with performance without saliency information. This configuration was identical to the  $\Psi$ -Match procedure; however the choice of model point  $y_n$  to incorporate into the correspondence set was random instead of being biased by  $\Psi$ . We calculated similarity at each iteration of the matching algorithms, allowing an investigation of the tolerance to the parameter  $\beta$ .

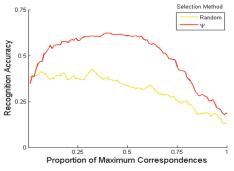


Figure 8: Recognition accuracy as a function of  $\beta$  (proportion of maximum correspondences) for the  $\Psi$ -Match algorithm and a randomly ordered variant.

Ψ-Match exhibits expected behaviour, in that recognition accuracy is rapidly improved as the correspondence set is expanded - indicating more accurate transformations for matching instances – until a fairly stable performance is attained between β = 0.1 and β = 0.6. The dip after this is likely due to the greater chance of false correspondences being introduced and degrading transformation quality. This performance pattern is not seen for the algorithm disregarding saliency information, where accuracy steadily declines throughout the process, confirming the utility of contextual saliency in pattern recognition.

Recognition performance using the SC and TPS-RPM registration algorithms was calculated in an identical manner; results are seen in Table 1 in comparison to the  $\Psi$ -Match algorithm with  $\beta = 0.25$ .

Table 1: Recognition accuracy of the biometric system across landmark-based registration methods.

Registration Method	Recognition Accuracy
SC	0.41
TPS-RPM	0.46
Ψ-Match	0.59

Ψ-Match is shown to outperform SC and TPS-RPM by wide margins. Note that although the stopping criterion was chosen here *a posteriori*, Ψ-Match performance was improved over other methods for  $\beta$  in the range [0.06, 0.71], indicating the algorithm's robustness with respect to this parameter on our data set

Performance was then assessed within the context of a semi-automatic recognition system through looking at whether a correct identity was produced within the N most similar patterns.

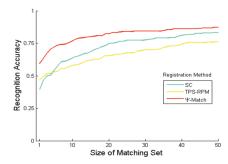


Figure 9: Recognition accuracy across validation matching set size N for systems using SC, TPS-RPM, or  $\Psi$ -Match ( $\beta=0.25$ ) registration.

 $\Psi$ -Match outperforms other methods across the range of matching set size. At a practical set size of 10, the system incorporating  $\Psi$ -Match registration

produced a correct identity instance in 75% of cases, in comparison to 61% and 54% for TPS-RPM and SC methods, respectively; confirming the usefulness of saliency-biased registration in animal biometrics.

#### 8 CONCLUSIONS

A method for ordering points in a set on a measure of distinguishability, *contextual saliency*, has been introduced in this paper. Ordering on this basis is shown to be tolerant of noise and perspective transformation, as well as be predictive of correspondence, in synthetic experiments.

This information is leveraged in an iterative non-rigid registration algorithm, Ψ-Match. A case study on a difficult real-world manta ray data set found improved performance for a recognition system using Ψ-Match registration in comparison to the same setup using either shape context (Belongie et al., 2002) or TPS-RPM (Chui & Rangarajan, 2003) registration algorithms.

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