CLOSED BOUNDARY FINDING, FEATURE SELECTION AND CLASSIFICATION APPROACH TO MULTI-IMAGE MODELING\*

by

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Remote sensing is the science and art of acquiring information about material objects from measurements made at a distance without coming into physical contact with the objects. The information may be transmitted to the observer through electromagnetic fields, in particular, through the spectral, spatial, and temporal variations of these fields [1,2]. The data is gathered from sensors mounted on an aerospace platform and provides information on existing conditions of physical objects. The sensors measure target radiance in several regions of the electromagnetic spectrum; the data is usually called multispectral data.

In order to derive information from these data sets (Multi-images), one must be able to relate these measurements to those of known objects or materials. The quantity of data is typically so large that machine oriented analysis techniques must be devised. At the same time to enhance image processing activities, an understanding of the pictorial aspects of the data to be processed is required. Also to further progress in image analysis and image processing techniques such as image coding, registration, deblurring, noise removal, etc., better image models need to be developed.

In this paper a specific multi-image analysis and processing technique is studied to develop a good image model. A boundary finding algorithm is developed involving the image characteristics such as grey level edges and neighboring second order statistical properties of these data sets. The

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technique of boundary finding narrows the method of feature selection to Karhunen-Loeve transformation and the evaluation of the boundaries has been done based on test field classification.

## Introduction

Natural scenes usually consist of objects with structures unless the picture contains objects out of focus. Structure is a departure from randomness. This structure is not usually homogeneous—different parts of a picture usually contain different kinds of structure. The study of this structure is called Image Modeling.

One approach to image modeling is to characterize what we see in the image. Generally there exists structure in the relationship between areas—an area can be defined as the largest connected set of picture elements each of which has a property P. P can represent grey level, texture, a statistic etc. Adjacent areas share boundaries so that the shape of each is determined by the shape of its neighbors. For multi-spectral images these "areas" become "volumes". For future reference these will be referred to as "blobs" rather than "areas" or "volumes". Grey level blobs can be characterized in terms of their boundaries which are related to gray level discontinuities.

Most methods for foundary finding reported in the literature are based on extensions of the classical digital gradient and laplacian operators. Several investigators have studied mathematical methods of defining edge detectors that are optimum in various senses. Heuckel [3] finds the perfect step edge that best matches the given digital picture in a certain disc shaped neighborhood of each point. Similar work has been done recently by Griffith [4] to detect edges in simple scenes using apriori information. Another approach to edge detection involves the use of both coarse and fine difference operators at each point. This approach

is suitable for detecting steps in average grey levels. Coarse operators detect the steps, while the fine ones locate them sharply [5,6]. Some approaches investigated at Purdue University were based on the gradient concept [7], clustering [8] and hypothesis testing based on first order statistics [9]. The gradient approach is inherently noisy and produces borders that are discontinuous of varying width and also produces spurious isolated points. Clustering is more stable and less noisy but very time consuming; and also closed boundaries are not guaranteed. The hypothesis testing technique guarantees closure.

The boundary finding algorithm discussed here is also based on hypothesis testing but takes into account second order statistics as well as first order statistics.

## Description of the Data Source

The multispectral source to be studied is shown in Figure 1. The process  $a(x_1,x_2,\lambda)$  is some measure of the spectral energy at wavelength  $\lambda$  for ground resolution point  $(x_1,x_2)$ . The output sample space is the set of all vectors A in  $R^N$  that are obtained from the continuous stochastic process  $a(x_1,x_2,\lambda)$  by discretizing the two spatial variables  $(x_1,x_2)$  and the spectral variable  $(\lambda)$ .

The elements of A are assumed to be jointly normal. This assumption is made for the mathematical simplification it allows. In addition it has been found experimentally that the normal distribution [10] is a reasonable approximation for multispectral data for several applications in pattern recognition.

The output vector A is then

$$\mathbf{A}_{lm} = \left[\mathbf{a}_1, \mathbf{a}_2, \ldots, \mathbf{a}_{N}\right]^{t}$$

where  $a_i = a[x_{1\ell}, x_{2m}, \lambda_i]$  and  $x_{1\ell}, x_{2m}$  and  $\lambda_i$  are the  $\ell$ th, mth, and i-th sample in  $(x_1, x_2, \lambda)$ 

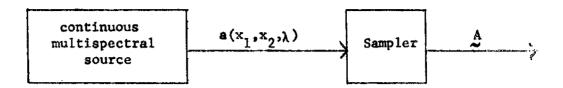


Fig. 1 Model of the Multispectral Source

The Joint density function is given by

$$g(A) = \frac{1}{N/2} \cdot \frac{1}{|\Sigma|^{\frac{1}{2}}} \exp \left[-\frac{1}{2}(A - \mu)^{t} \Sigma^{-1} (A - \mu)\right]$$

where  $\Sigma$  is the N x N covariance matrix for A and  $\mu$  is the mean vector.

In the following discussion each a will be called an element and a group of elements will be called simply a "group". The groups are squares containing equal number of elements on each side. The elements of each blob also are assumed to be Gaussian.

## Hypothesis Testing

The general problem is this: we have two normal populations. One with variate  $x_1$  which has mean  $\mu_1$  and variance  $\sigma_1^2$  and one with variate  $x_2$  which has mean  $\mu_2$  and variance  $\sigma_2^2$ . On the basis of two samples, one from each population, it is desired to test the hypotheses in both of the following:

i, 
$$H_{o1}$$
 :  $\sigma_1^2 = \sigma_2^2$ 

$$H_{a1}$$
 :  $\sigma_1^2 + \sigma_2^2$ 

ii, 
$$H_{02} = \mu_1 = \mu_2$$
,  $\sigma_1^2 = \sigma_2^2 = \sigma^2$ 

$$H_{a2} = \mu_1 + \mu_2$$
,  $\sigma_1^2 = \sigma_2^2 = \sigma^2$ 

In hypothesis i, the parameter space  $\Omega$  is four dimensional; a joint distribution of  $x_1$  and  $x_2$  is specified when values are assigned to the four quantities  $(\mu_1,\mu_2,\sigma_1^2,\sigma_2^2)$ . The subspace  $\omega$  is three dimensional because values for only three quantities  $(\mu_1,\mu_2,\sigma^2)$  need be specified in order to specify completely the joint distribution under the hypothesis that  $\sigma_1^2 = \sigma_2^2 = \sigma^2$ .

Let there be m observations  $(X_{11,\lambda_0},X_{12,\lambda_0},\dots,X_{1m,\lambda_0})$  in the sample from the first population and n observations  $(X_{21,\lambda_0},X_{22,\lambda_0},\dots,X_{2n,\lambda_0})$  from the second. Let us also assume for this analysis that  $\lambda_0$  is fixed and determines the spectral band from which the samples are being considered.

The likelihoods in parameter space  $\boldsymbol{\Omega}$  and subspace  $\boldsymbol{\omega}$  are:

$$L(\Omega) = \left(\frac{1}{2\pi\sigma_{1}^{2}}\right)^{m/2} \exp\left[-\frac{1}{2}\sigma_{1}^{2} \sum_{i=1}^{m} \left(X_{1i,\lambda_{0}} - \mu_{1}\right)^{2}\right] \left(\frac{1}{2\pi\sigma_{2}^{2}}\right)^{\frac{n}{2}} \exp\left[-\frac{1}{2\sigma_{2}^{2}} \sum_{j=1}^{n} \left(X_{2j,\lambda_{0}} - \mu_{2}\right)\right]$$

and

$$L(w) = \left(\frac{1}{2\pi\sigma^2}\right)^{\frac{m+n}{2}} \exp\left[-\frac{1}{2\sigma^2} \left\{ \sum_{i=1}^{m} \left(x_{1i,\lambda_{\beta}} \gamma_{i,1}\right)^2 + \sum_{j=1}^{n} \left(x_{2j,\lambda_{\beta}} \gamma_{i,2}\right)^2 \right\} \right]$$

Differentiating L(Ω)  $_w$ . r. t.  $_{\mu_1}$ ,  $_{\mu_2}$ ,  $_{\sigma_1}^2$ ,  $_{\sigma_2}^2$  the estimates of  $_{\mu_1}$ ,  $_{\mu_2}$ ,  $_{\sigma_2}^2$  respectively are as:

$$\hat{\mu}_{1} = \frac{1}{m} \sum_{i=1}^{m} X_{1i,\lambda_{p}} = \overline{X}_{1,\lambda_{p}}$$

$$\hat{\mu}_2 = \frac{1}{n} \sum_{j=1}^n X_{2j,\lambda_p} = \overline{X}_{2,\lambda_p}$$

$$\hat{\sigma}_{1}^{2} = \frac{1}{m} \sum_{i=1}^{m} \left( \mathbf{x}_{1i,\lambda_{\rho}} - \overline{\mathbf{x}}_{1,\lambda_{\rho}} \right)^{2}$$

$$\hat{\sigma}_{2}^{2} = \frac{1}{n} \sum_{j=1}^{n} \left( \mathbf{x}_{2j,\lambda_{\rho}} - \overline{\mathbf{x}}_{2,\lambda_{\rho}} \right)^{2}$$

Substituting these values the maximum of  $L(\Omega)$  is given by

$$L(\widehat{\Omega}) = \begin{bmatrix} \frac{m}{m} & \frac{n}{2} \\ 2\pi \sum_{i=1}^{\infty} (x_{1i,\lambda_{\rho}} - \overline{x}_{1,\lambda_{\rho}})^2 \end{bmatrix} \exp \begin{bmatrix} -\frac{m}{2} \end{bmatrix} \begin{bmatrix} \frac{n}{n} \\ 2\pi \sum_{j=1}^{\infty} (x_{2j,\lambda_{\rho}} - \overline{x}_{2,\lambda_{\rho}})^2 \end{bmatrix} \exp \begin{bmatrix} -\frac{n}{2} \end{bmatrix}$$

Similarly the maximum of L(w) can be derived. The maximum likelihood Ratio is then

$$R_{\lambda_{\rho}} = R(X_{11,\lambda_{\rho}}, X_{12,\lambda_{\rho}}, \dots, X_{1m,\lambda_{\rho}}, X_{21,\lambda_{\rho}}, \dots, X_{2n,\lambda_{\rho}})$$

$$= \frac{L(w)}{\frac{m}{2} \frac{n}{n^2}} \frac{\sum_{i=1}^{m} (X_{1i,\lambda_{\rho}} - \overline{X}_{1,\lambda_{\rho}})^2}{\sum_{j=1}^{n} (X_{2j,\lambda_{\rho}} - \overline{X}_{2,\lambda_{\rho}})^2}$$

$$= \frac{m}{\sum_{i=1}^{m} (X_{1i,\lambda_{\rho}} - \overline{X}_{1,\lambda_{\rho}})^2} \frac{m+n}{\sum_{i=1}^{n} (X_{2j,\lambda_{\rho}} - \overline{X}_{2,\lambda_{\rho}})^2}$$

Defining 
$$F_{\lambda_{\rho}} = \frac{n-1}{m-1} = \frac{\sum_{i=1}^{m} (x_{1i,\lambda_{\rho}} - \overline{x}_{1,\lambda_{\rho}})^2}{\sum_{j=1}^{n} (x_{2j,\lambda_{\rho}} - \overline{x}_{2,\lambda_{\rho}})^2}$$
 one obtains

$$R_{\lambda_0} = \frac{\frac{\underline{m+n}}{2}}{\frac{\underline{m}}{2} \frac{\underline{n}}{n^2}} \frac{\left(\frac{m-1}{n-1} F_{\lambda_0}\right)^{\frac{\underline{m}}{2}}}{\left(1 + \frac{m-1}{n-1} F_{\lambda_0}\right)^{\frac{\underline{m+n}}{2}}}$$

So if the distribution of  $F_{\lambda}$  is known, the distribution of  $R_{\lambda}$  is also known. Since  $F_{\lambda}$  has F-distribution with (m-1) and (n-1) degrees of freedom, therefore on plotting  $R_{\lambda}$  as a function of  $F_{\lambda}$  it is apparent that the critical region  $0 < R_{\lambda} < C$  corresponds to a two tailed test on F.

In the second hypothesis the parameter space  $\Omega$  becomes three dimensional with co-ordinates  $(\mu_1, \mu_2, \sigma^2)$ , while,  $\omega$  for the null hypothesis  $\mu_1 = \mu_2 = \mu$  (say) becomes two dimensional with co-ordinates  $(\mu_1, \sigma^2)$ .

The maximum likelihood ratio in this case is

$$R_{\lambda_{\rho}} = \begin{bmatrix} 1 + \frac{\frac{mn}{m+n} \left(\overline{x}_{1,\lambda_{\rho}} - \overline{x}_{2,\lambda_{\rho}}\right)^{2}}{\frac{m}{m} \left(\overline{x}_{1,\lambda_{\rho}} - \overline{x}_{1,\lambda_{\rho}}\right)^{2} + \frac{n}{j+1} \left(x_{2j,\lambda_{\rho}} - \overline{x}_{2,\lambda_{\rho}}\right)^{2}} \end{bmatrix}^{\frac{m+n}{2}}$$

Again defining a parameter

$$t_{\lambda_{\rho}} = \frac{\sqrt{\frac{mn}{m+n}} \left(\overline{x}_{1,\lambda_{\rho}} - \overline{x}_{2,\lambda_{\rho}}\right) \sqrt{m+n-2}}{\sqrt{\frac{m}{\sum_{j=1}^{m} \left(x_{1i,\lambda_{\rho}} - \overline{x}_{1,\lambda_{\rho}}\right)^{2} + \sum_{j=1}^{n} \left(x_{2j,\lambda_{\rho}} - \overline{x}_{2,\lambda_{\rho}}\right)^{2}}} \quad \text{which}$$

has a student t-distribution with m + n - 2 degrees of freedom, the likelihood ratio is

$$R_{\lambda_{\rho}} = \begin{bmatrix} \frac{1}{t^{2}} \\ \frac{1}{h^{\rho}} \\ 1 + \frac{1}{m+n-2} \end{bmatrix}$$

Its distribution is determined by the t-distribution. The test is done in terms of t rather than R . This is also a two-tailed test. A 5 percent critical region for t is  $t_{\lambda_a}^2 > (t_{0.25})^2$ .

In short the first hypothesis is based on F-test and the second on student t-test.

### Testing Algorithm

Let the group to be compared be called sample 1 and the group to be compared with it sample 2. The goal is to calculate the value of F to compare against a critical value to determine whether both of the samples have the same variance irrespective of their means and then finally to calculate the value of f to compare against another specified critical value to determine whether both of the samples belong to the same population.

The boundary is assumed to exist if the null hypothesis  $H_{01}$  fails for the critical value already specified. For this purpose, standard tables [11,12] have been used. The boundaries may be weak in some spectral bands and strong in others. To take this into account the first hypothesis is tested for all values of  $\lambda_{\beta}$ ,  $\beta=1,\ldots$ , number of spectral bands considered. After the null hypothesis  $H_{01}$  is accepted in all the spectral bands, the algorithm proceeds to test for the second hypothesis. Again the failure

of the null hypothesis  $H_{02}$  for any value of  $\lambda_{\rho}$  indicates that the two populations are not the same and hence a boundary exists between them.

A block diagram of testing algorithm is shown in Figure 2. In this block diagram,  $\lambda_{\rho}$  is not fixed anymore, while calculating the covariance matrix.

Multivariate tests can take all the spectral bands into account simultaneously but they are very time consuming. To cut down on processing time the algorithm determines the boundaries in all the spectral bands separately. Also, the experiments conducted show that for the multispectral data available at the Laboratory for Application of Remote Sensing (LARS), multispectral tests are not very advantageous.

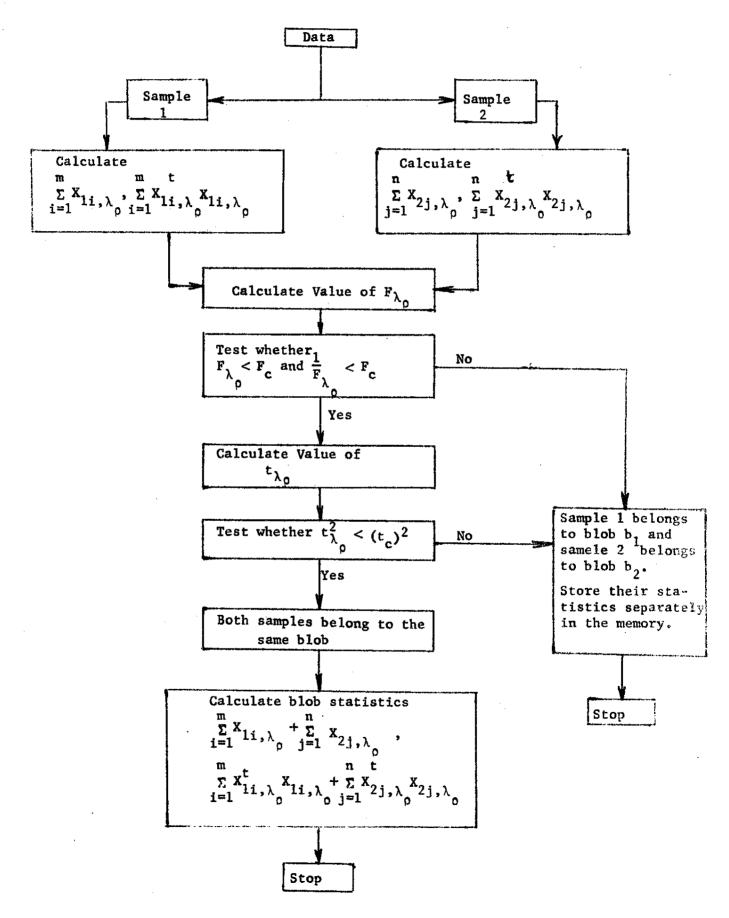
## Building of Blobs

Let us suppose g columns and g rows of digital data are from the same distribution. The g rows of the data (one row of groups) are processed at a time. Let there be W groups in a row and k'th group of the previous row is designated as group k. Let the blob to which group k is assigned be given by  $b_k$  and the blob to which group k is assigned  $b_k$ .

In the beginning of the data to be processed group  $1_c$  is arbitrarily assigned to blob  $1_c$ . Group  $2_c$  is compared with group  $1_c$  according to the Testing Algorithm. If they have the same distribution, their statistics are added and both groups  $1_c$  and  $2_c$  are assumed to be in the same blob. Next group  $3_c$  is compared with group  $2_c$  and in general the group W is compared with group W-1 at the end of the row. At every step an account is kept for the number of groups in all the blobs.

From second row of groups onward the following steps are followed: Group  $1_c$  is compared to the blob containing group  $1_c$ . If they are

the same then this group is also added to b otherwise processing continues



Pin 2 Tooking Almontate

to group  ${}^2_{\rm c}$ . Group  ${}^k_{\rm c}$  is compared to the blob which contains group  ${}^k_{\rm c}$ . If they are the same, group  ${}^k_{\rm c}$  is added to  ${}^b_{\rm k}$  and a check is made if group  ${}^k_{\rm c}$ -1 is assigned to any blob. If it is then it continues to  ${}^k_{\rm c}$ +1 otherwise group  ${}^k_{\rm c}$ -1 is tested against blob  ${}^b_{\rm k}$ . If they are different then the processing continues to  ${}^k_{\rm c}$ +1. Here again if group  ${}^k_{\rm c}$ -1 is the same it is added to blob  ${}^b_{\rm k}$  and a check is made on group  ${}^k_{\rm c}$ -2 and so on.

On the other hand if group  $k_c$  and blob  $b_k$  are different then group  $k_c$  is compared to group  $k_c^{-1}$ . If they are the same groups  $k_c$  and  $k_c^{-1}$  are added otherwise the processing continues to group  $k_c^{+1}$ . In the end of the second row group  $k_c^{-1}$  is processed in the same manner as group  $k_c^{-1}$  except that there is no other group  $k_c^{-1}$  to continue with.

After group  $W_c$  is processed a backward scanning is done on this row. The scanning starts with group  $W_c$ . Whenever the group  $J_c$  is encountered which has not been assigned to any blob, group  $J_c$  is assigned to a new blob and looks for group  $J_c$ -1. If group  $J_c$ -1 is assigned already the scanning continues further otherwise group  $J_c$ -1 is compared to blob by the limit of they are the same the group  $J_c$ -1 is added to by and a check is made whether group  $J_c$ -2 is assigned or not and so forth.

As the blob building continues to the subsequent group rows and it is determined that no group in the current row belongs to the blob, the blob is said to be closed and it is assigned to one of the pattern classes based on pattern classification techniques.

# Fidelity of Reconstruction

When the blob is determined to be closed the original data elements of this blob are replaced by the mean value averaged over all the original elements of this blob and an estimate of the percent mean square error is calculated.

If A is the original data and  $\hat{A}$  is the reconstructed data after replacing the blobs by their mean values, then the mean square error is defined as

$$MSE = E[|A - \hat{A}|]^2$$

Fidelity of reconstruction can be measured by the MSE between A and A or by % MSE  $_{(0)}$ . r. t. the variance of the data.

If  $\sigma_{\bf i}^2$  is the data variance in the i-th channel and there are o-channels in all considered for processing purposes then

% MSE = 
$$\frac{E\{||\mathbf{A} - \hat{\mathbf{A}}||^2\}}{\sum_{i=1}^{p} \sigma_i^2} * 100$$

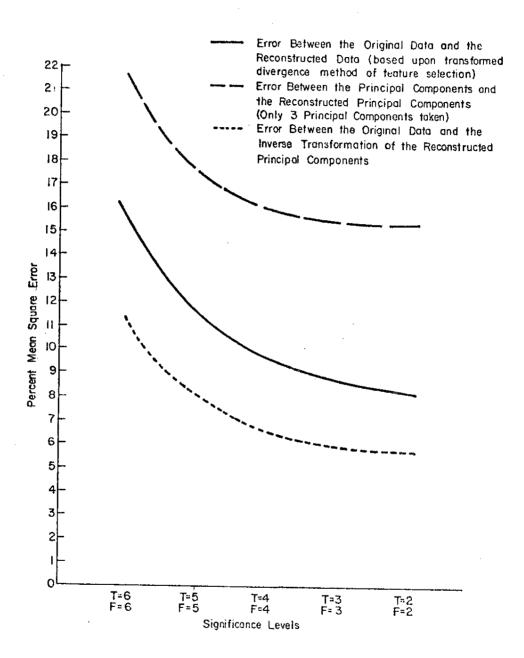
% MSE has been calculated for various significance levels of student 't' and F-tests and a plot is given in Fig. 3. Pictures using digital display system taken on the original data and the reconstructed data are also given in Fig. 4.

# Classification of Data

One immediate advantage of finding blob boundaries is in classification of the data in already known classes.

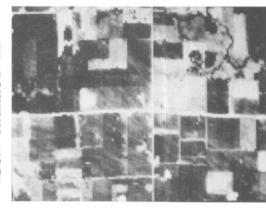
Presently the classification is usually done on the basis of a per point classification (spectral information only) and the advantage of spatial information is not taken into account. To use the spatial information it becomes absolutely necessary to define spatial boundaries which separate the regions containing the classes of interest. The boundary finding algorithm provides such a means.

Let  $\{W_i^{}\}$  be the set of pattern classes with their distributions  $\{F_i^{}\}$ ,

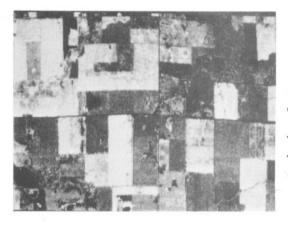




from Channel 12 Original Data



Reconstructed Data from Channel 12



Original Data from Channel 8



Reconstructed Data from Channel 8

Figure 4



Reconstructed Data from Channel 7

and let the set of points  $\{y\}$  be all the elements of the closed blob. To decide to which  $F_i$  they belong, a distance measure  $d(F_i, F_j)$  is used as a measure of separation between any two distributions  $F_i$  and  $F_j$ .

In a mathematical sense the term "distance" and "metric" are sometimes (incorrectly) used interchangeably. A metric on a set S is, of course a real valued  $\delta$  defined on S x S (x indicates cartesian product) such that for arbitrary  $F_i$ ,  $F_i$ ,  $F_k$  in S

a, 
$$\delta(F_i, F_j) \ge 0$$
  
b,  $\delta(F_i, F_j) = \delta(F_j, F_i)$   
c,i,  $\delta(F_i, F_i) = 0$   
ii, If  $\delta(F_i, F_j) = 0$  then  $F_i = F_j$   
d,  $\delta(F_i, F_j) + \delta(F_j, F_k) \ge \delta(F_i, F_k)$ 

The distance measure in this report should not be confused with metric. It obeys properties a, b, and c, only; not property d.

Let G(y) be a distribution governing  $\{y\}$ . The magnitudes of  $\{d(G, F_i)\}$  are compared; the distribution  $F_i$  which minimizes the distance  $d(G, F_i)$  is presumed to be the distribution which contains G(y).

In other words assign {y} to the ith class in case

$$d(G, F_i) = \min_{j=1, ..., k} d(G, F_j)$$

where k is the total number of classes considered.

Let  $F_i(y)$  and  $F_j(y)$  be the distributions of pattern classes  $W_i$  and  $W_j$  respectively, and let  $\rho_i(y)$  and  $\rho_j(y)$  are their probability density functions defined in  $\Omega_y$ . Then a statistical distance measure appearing in the work of Jeffreys Matusita [13,14] is defined for two distributions  $F_i(y)$  and  $F_j(y)$  as follows.

$$B = d^{2}(F_{i}, F_{j}) = \int_{\Omega} \left( \sqrt{\rho_{i}(y)} - \sqrt{\rho_{j}(y)} \right)^{2} dy$$

For convenience, B will be referred to herein as the Bhattacharyya distance.

Defining

$$\mathbf{P} = \int_{\Omega} \sqrt{\rho_{\mathbf{i}}(\mathbf{y})} \cdot \sqrt{\rho_{\mathbf{j}}(\mathbf{y})} d\mathbf{y}$$

$$d^{2}(F_{i}, F_{j}) = 2(1-P)$$

In multispectral data one is concerned with multivariate distributions and as already stated that the blobs have multivariate Gaussian distributions,  $\rho$  for two such densities  $\{\rho_i(y) = N(\mu_i, \Sigma_i), i=1,2\}$  can be reduced to the following expressions [15]

$$P = \exp[-\sigma]$$

where

$$\sigma = \frac{1}{8} \left( \mu_1 - \mu_2 \right)^t \left[ \frac{\Sigma_1 + \Sigma_2}{2} \right]^{-1} \left( \mu_1 + \mu_2 \right) + \frac{1}{2} \ln \frac{\det \left( \frac{1}{2} \left( \Sigma_1 + \Sigma_2 \right) \right)}{\sqrt{\det \left( \Sigma_1 \right) \det \left( \Sigma_2 \right)}}$$

Then for whichever class P is maximum, the vector  $\{y\}$  is decided to be a member of that class.

The multispectral data used for classification is taken from 1971 Corn Blight Watch Experiment [16]. It has eleven spectral bands in the wavelength region of 0.4 to 3 micrometers and one thermal infrared band from 9.7 to 11.3 micrometers. The strip of terrain observed is represented by 222 samples across track and 176 samples per mile along track taken at a 5000 foot altitude.

A set of training fields was obtained representing 5 classes namely corn, forage, soybeans, forest and water. On the basis of these 5 patters classes, all the 222 samples across track and 400 samples along the track

were classified using the boundary finding algorithm defining blob bound and the above mentioned distance measure.

Note that  $\sigma$  in B-distance involves the inversion of a covariance matrix and for some of the blobs the determinant of the covariance matrix is very insignificant. Such blobs are classified according to the Euclidean distance to the mean of the pattern classes.

The classified data is written on a separate data storage tape and is evaluated on the basis of given test fields.

Presently the same data at LARS is being classified on the point class. fication basis using Gaussian Maximum likelihood Ratio decision rule [1] and is getting 95.9% overall classification accuracy as compared to 98% obtained by this algorithm. A summary of the results is presented in Figure 5.

# Feature Selection

The number of spectral bands (channels) poses a problem in the sense that the computation time and computer memory requirements are increased as the number of channels increases. Also, to minimize the overall probability of misclassification the best set of features should be selected.

The results obtained in the previous section were obtained by using a subset chosen on the basis of minimum transformed divergence. The minimum transformed divergence between any two training classes was larger for this particular set of channels than all other combination of three channels. The transformed divergence is a measure of class separability in feature space [17].

Per point classification results (Based on test fields)

Channels (7,8,12)

Class	No. of Points	Percent Correct	C	Classi			
			Corn	Forage	Soybeans	Forest	Water
Corn	5950	95.6	5686	203	38	23	o
Forage	3151	97.3	78	3065	0	8	o
Soybeans	4770	96.0	59	95	4578	38	0
Forest	680	92.6	3	13	32	630	2
Water	37	97.3	0	1	0	0	36

Overall performance (13995/14588) = 95.9% correct

# Classification results obtained by Boundary Finding Algorithm Channels (7,8,12)

	No. of	Percent		Classi			
Class	Points	Correct	Corn	Forage	Soybeans	Forest	Fater
Corn	5950	99.0	5888	50	4	8	O
Forage	3151	96.1	111	3027	4	9	٥
Soybeans	4770	97.9	63	30	4669	8	O
Forest	680	99.1	2	4	0	674	0
Water	37	97.3	0	1	0	0	36

Overall performance (14294/14588) = 98% correct

Figure 5 Classification Results Using 3 Best Features
Selected according to the Minimum Transformed
Divergence Method of Feature Selection

The boundary finding algorithm is based on variance analysis therefore the important consideration in selecting such a feature set should be such that it contains the maximum variance in it.

To overcome these problems, the Karhunen-Loeve (K-L transformation technique of selecting the best feature can be implemented.

K-L or Eigenvector transformation is an NxN transformation matrix T (reconstruction transformation matrix  $T^t$ ) and is used to map A down to n<N dimensions. It is shown in [18] than n rows of T are the first n orthonormal solutions to the characteristic equation

$$\sum_{k} t = \lambda t$$

corresponding to the n largest eigenvalues  $\lambda_1>\lambda_2>\dots>\lambda_n$  of the NxN covariance matrix  $\Sigma$  of A. The transformation to n<N dimensions is given by

where  $\underline{\mathbf{T}} = \begin{bmatrix} \mathbf{t}_1, \mathbf{t}_2, \dots, \mathbf{t}_n \end{bmatrix}$ 

It is also shown in [18] that the elements of Y are uncorrelated and the variance of the n elements of Y are given by the n  $\lambda_1$ 's, that is

$$Var (Y_1) = \lambda_1$$

$$Var (Y_2) = \lambda_2$$

$$\vdots$$

$$Var (Y_n) = \lambda_n$$

In other words this transformation rotates the source output A in the N-space to a favorable orientation with respect to the coordinate systems. The favorable orientation is so-called because the average energy (variance) of the source is re-distributed over the coordinates such that a larger

percentage of the source variance is distributed over fewer coordinates.

This packing of the source variance provides a means of reducing the coordinates to determine blob boundaries and implement sample classification.

The 12 x 12 spectral covariance matrix  $\sum$  was estimated for a Corn blight Watch Experiment flight by averaging over the entire flight line. The number of data points available for processing were 2.5 x  $10^5$ . The resulting covariance matrix is shown in Figure 6. The eigenvectors and normalized eigenvectors for  $\sum$  were computed using the IBM Scientific Subroutine Package subroutine "EIGEN". Figure 7 lists the eigenvalues ranked such that

$$\lambda_1 > \lambda_2 > \cdots > \lambda_n$$

The eigenvector transformation T is constructed from the first 3 eigenvectors of  $\Sigma$ . The first Principal components contain 93.9% of the total variance contained in all the twelve features. The blob boundaries were located in the transformed data and then the blobs were classified using already mentioned pattern classification techniques. The overall classification accuracy is 98.2% as compared to 98% using minimum transformed divergence method of feature selection. The blobs data was replaced by their mean values and then an inverse transformation operation was performed on this data. An estimate of mean square error was determined between the reconstructed Principal components and original Principal components and also between the inverse transformed reconstructed Principal components and the original data. These values are also plotted in Figure 3. The digital display pictures of first three principal components and the reconstructed Principal components are given in Figure 8. The summary of the classification results is detailed in Figure 9.

***		_
CHA	NNE	T

	1	2	3	4	5	6	7	8	9	10	11	12
Ĕ	434.3											,
2	338.9	299.9										
3	320.4	265.0	266.1									
÷,	391.7	325.0	327.9	487.2								
5	447.5	375.1	366.2	491.2	552.3							
6	466.6	411.8	370.1	497.0	559.3	678.9						
7	539.7	459.8	421.3	528.9	622.3	688.4	791.7					
8	-71.6	-59.8	-27.7	21.8	-37.7	-112.9	-147.7	382.1				
9	-27.0	-22.3	-3.1	37.3	1.0	-38.6	-53.5	257.0	231.3			
1.0	295.7	250.1	249.7	370.7	385.0	406.2	454.5	159.2	203.2	694.5		
11	3 <b>37.3</b>	289.7	270.1	376.2	408.4	465.0	513.9	66.5	136.6	602.1	651.1	
12	441.0	374.2	337.1	430.8	516.3	631.4	707.8	-294,3				1062.3
				nce is								¥-, ø

Fig. 6. Spectral Covariance Matrix for Data for Corn Blight Watch Experiment Flight

Eigenvalue	Percent of Variance	Cumulative percent
4640.70 940.85 554.47 112.28 79.93 61.00 49.26 29.72 26.51 15.79 12.42 8.77	71.05 14.40 8.49 1.72 1.22 0.93 0.75 0.45 0.41 0.24 0.19 0.13	71.05 85.45 93.94 95.66 96.89 97.82 98.57 99.03 99.43 99.68 99.87

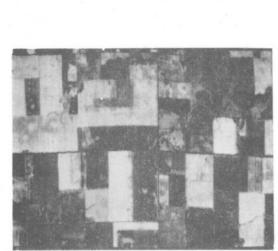
Fig. 7. 12 Spectral Eigenvalues for CBWE Flight



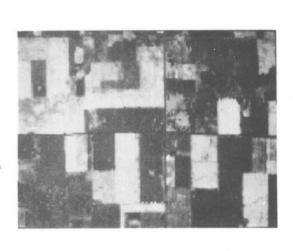
1st Principal Component



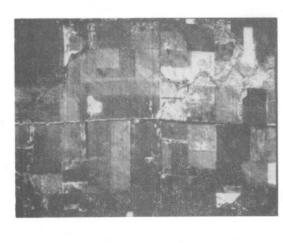
1st Reconstructed Principal Component



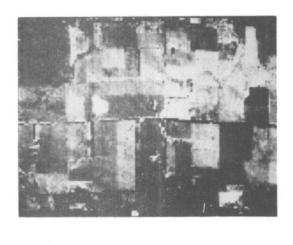
2<sup>nd</sup> Principal Component



2nd Reconstructed Principal Component



3rd Principal Component



3<sup>rd</sup> Reconstructed Principal Component

Figure 10 shows the block diagram of the different operations involto calculate the mean square error between the inverse transformed reconstructed Principal components and the original data.

The boundaries of the classified blobs were printed on a line printer of IBM 360/67 and is presented in Figure 11. An aerial photograph related to the ground truth is also presented for comparison purposes.

Class	No. Points	Percent Correct	Corn	Classif Forage	ied as Soybeans	Forest	Water
Corn	5950	99.7	5932	14	4	0	0
Forage	3151	94.6	160	2980	2	9	Ö
Soybeans	4770	98.7	21	40	4709	0	ø
Forest	680	98.2	12	0	0	668	o
Water	37	97.3	0	1	0	0	36

Overall performance (14325/14588) = 98.2% correct

Fig. 9 Classification Results Based on First Three Principal Components

## Summary

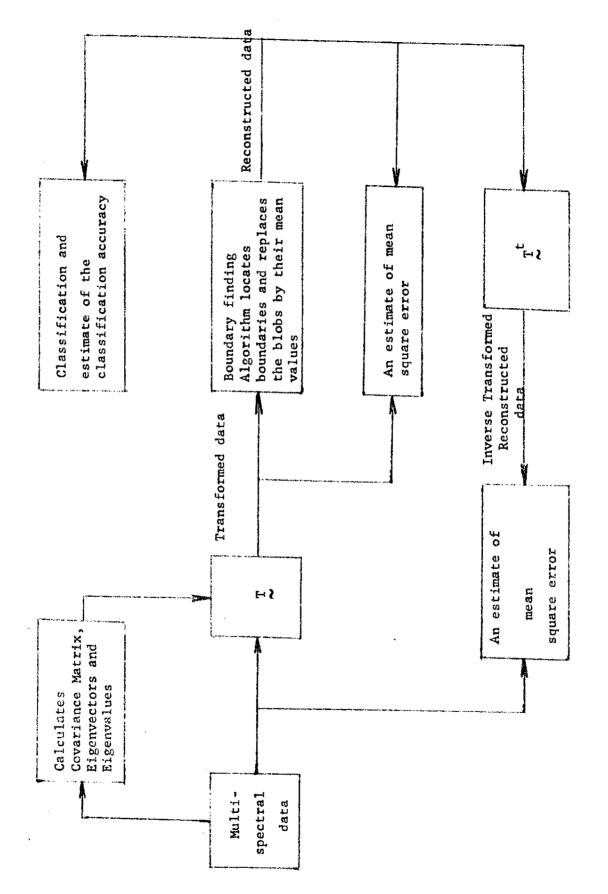
In this paper it has been shown that spatial information can be extracted from a multispectral data image by machine and effectively used in the classification process. The degree of improvement in classification accuracy depends upon the amount of spatial information contained in the data. An experiment was conducted on data from the Earth Resources Technology Satellite (ERTS).

For ERTS data there are relatively few elements per blob. Since there is less spatial information to take advantage of, the boundary finder provides only a small degree of improvement over the per point classifier as illustrated by figure 12.

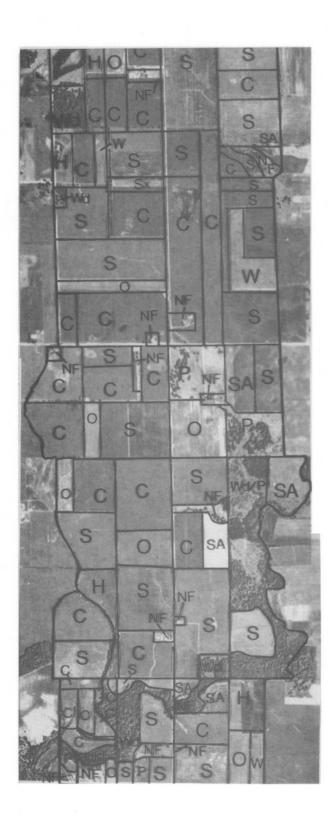
The concept of boundary finding seems to be a fairly robust one. For the aircraft data [16] any three spectral bands were chosen at random and the boundary finder in conjunction with the sample classifier was implemented to evaluate the classification results. These results are presented in Figure 13 and 14. It is evident that though this procedure involves more complex computations but it produces consistently better results than the per point classifier.

The variance analysis used in boundary finding in a way restricts our choice to the Karhunen Loeve Transformation technique of feature selection. From figures 5 and 9 we note that the Maximum likelihood Decision Rule class—ifies the data with an error rate of 4.1%. The boundary finder in conjunction with Minimum Transformed Divergence method of feature selection reduces the rate of 2% while the boundary finder implemented on the transformed data according to k-L Transformation reduces this rate even further to 1.8%.

The identification of boundaries is important for other reasons also. If the boundaries are easily defined and show a repetitive pattern, the possibility of data compression exists (19). This will help in storing the data without losing much of information by storing the field boundaries and their classification. A measure of mean square error was estimated and it is found in Figure 3 that the k-L Transformation technique of feature selection reduces the percent mean square error quite significantly over the Minimum Transformed Divergence method of feature selection.



Different Steps Involved in the Classification and the Estimate of the Fidelity Criterion Fig. 10.



RUN NUMBER\* 74053900
MAX. NO. OF CLASSES CONSIDERED FOR CLASSIFICATION\* 5
PIZEL GROUP 5126 IS 2 X 2 SIGNIFICANCE LEVEL FOR T =6 AND FOR F =6
THE CHANNELS CONSIDERED ARE\* CACCCBCCCCCCCCC AARDO BAARA AARDO 4.591 4.074

Aerial photograph and ground observations for agricultural area (Seg 210 - 1971 CBWE) On-line printer results after boundary finding and classification of the data

NO. OF POINTS CLASSIFIED W.R.T. EUCLIDEAN DISTANE ARE

Figure 11

Per point classification results (0.5-0.6, 0.6-0.7, 0.7-0.8 Micrometer bands)

Overall performance (3718/4588) = 81.0 correct

Class	No. of Points	Percent Correct	Class Corn	sified as Soybeans	Othera
Corn	3315	88.7	2941	128	246
Soybeans	593	83.8	28	497	68
Others	680	41.2	230	170	280

Classification results based on boundary finding algorithms (0.5-0.6, 0.6-0.7, 0.7-0.8 Micrometer bands)

Class	No. of Points	Percent Correct	Class Corn	sified as Soybeans	Others
Corn	3315	91.2	3023	128	164
Soybeans	593	83.6	40	496	57
Others	680	35.4	285	154	241

Figure 12 Classification Results of the ERTS Data

Overall performance (3760/4588) = 82% correct

Per point classification results (Based on test fields)

Channels (1,2,3)

	No. of Samples	Percent Correct					
Class			Corn	Forage	Soybeans	Forest	Water
Corn	5950	60.1	3577	654	762	0	957
Forage	3151	69.7	570	2195	319	1	66
Scybeans	4770	63.5	635	112	3028	648	347
Forest	680	65.0	30	1	181	442	26
Water	37	94.6	0	1	1	0	35

Classification results based on Boundary finding algorithms
Channels (1,2,3)

Overall performance (9277/14588) = 63.6% correct

Class	No. of Samples	Percent Correct	Corn	Classi Forage	fied as Soybeans	Forest	Vater
Corn	5950	68.6	4078	1740	36	4	92
Forage	3151	78.4	639	2471	41	O	D
Soybeans	4766	65.6	1556	76	3132	0	0
Forest	680	42.4	139	0	253	288	O
Water	37	97.3	0	1	0	0	36

Figure 13 Classification results using 3 original spectral bands of the aircraft data.

Overall Performance (1005/14588) = 68.6% correct

Per point classification results (Based on test fields)
Channels (5,7,9)

Class	No. Points	Percent Correct	Corn	C1 Forage	assified as Soybeans	Forest	Water
Corn	5950	79.0	4700	1170	2	71	<b>D</b>
Forage	3151	87,1	403	2744	o	4	O
Soybeans	4770	90.7	8	157	4325	280	o
Forest	680	96 5	4	6	8	656	6
Water	37	100	0	0	0	0	37

Overall performance = (12462/14588) = 85.4%

Classification results based on Boundary finding algorithm
Channels (5,7,9)

	No. Points	Percent	Classified as						
Class		Correct	Corn	Forage	Soybeans	Forest	Water		
Corn	5950	97.1	5775	167	o	8	o		
Forage	3151	80.7	607	2542	. 0	2	0		
Soybeans	4770	94.7	23	44	4516	187	0		
Forest	680	99.4	4	0	0	676	o		
Wator	37	97.3	0	1	o	0	36		

Overall performance = (13545/14588) = 92.9%

Figure 14 Classification results using 3 original spectral bands of the aircraft data

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