

DIFFUSE MATRIX

An Optimized Data Structure for the Storage and Processing of Hyperspectral Images

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Keywords: Hyperspectral data format, diffuse matrix, storage optimization, geo-rectification, AVIRIS images.

Abstract: This paper proposes a new format for storing and processing hyperspectral images captured by spectrometer AVIRIS (Airborne Visible/InfraRed Imaging Spectrometer). Obtaining such images is difficult, because the sensor that takes the images is carried in an aircraft that suffers turbulences while the camera is taking photos. So, a geo-rectification process is necessary to correct the information of different bands. The format proposed in this paper, DMF (Diffuse Matrix Format), allows a more efficient storage, because a list with the original information received in the sensor is saved for each position (X,Y) of the scanned ground. The format of the list saves space and time because no redundant information is saved using it. To show the possibilities of this new format an application that makes some thresholding and filter operations has been built. This program, firstly, creates the diffuse matrix in memory from the file that stores the image information, and then, some filter operations are executed over the diffuse matrix to check it. In this way, we prove that diffuse matrix processing is fast and simple, as well as the space used in the disk for its storage is quite less than the space used by typical formats.

1 INTRODUCTION

Typical raster hyperspectral image formats used in remote sensing (BSQ, BIL, BIP, HDF-EOS, Geo-TIFF... (HDF-EOS, 2006)) save redundant information when they store an image with some acquisition errors (which is very common).

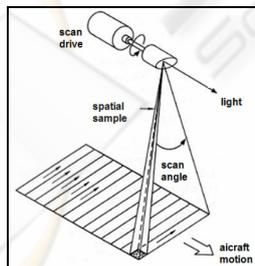


Figure 1: Diagram of hyperspectral image acquisition by AVIRIS spectrometer.

For this reason, the files that hold such images take up a lot of space in the hard disk, or at least, more space than the space necessary. Moreover, the image

processing is not very fast and requires a lot of memory, because the typical matrix that holds a hyperspectral image has to be entirely read before any processing over it.

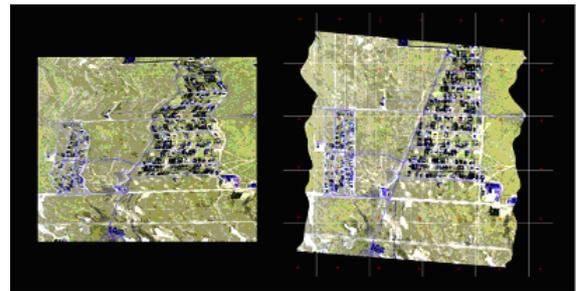


Figure 2: AVIRIS hyperspectral image before (left) and after (right) geo-rectification.

The over-information of typical hyperspectral image formats is caused because the original images, which are taken using spectrometer AVIRIS (AVIRIS, 2007) (as it can be seen in figure 1), have to be pre-processed before any usage with them (Martínez et al., 2005)(Brunn et al., 2003). This is

due to the turbulences suffered by the plane or other problems that happen while the sensor is taken the images. Figure 2 shows a hyperspectral image taken with AVIRIS before and after its geo-rectification. As we can see in that figure, before rectification, the image seems to be deformed, because some pixels of some bands are not in their appropriate place. After the geo-rectification, the imperfections in the image are corrected (misplaced pixels are interpolated using their neighbours).

If we choose an image format that applies geo-rectification we will have some remarkable disadvantages, because the process that corrects the image interpolates the missing or misplaced pixels with their neighbouring pixels, and this causes that some information is duplicated, or even erroneous, in the file that contains the image.

Therefore, the disadvantages of geo-rectification in the traditional three-dimension formats (Lx,y,l) carry us to study and develop other ways of representation, storage and analysis of AVIRIS hyperspectral images, where any interpolated information is not saved (so, any redundant or erroneous information will be not stored in the file).

Some studies to improve hyperspectral data formats have been published before. It is worth mentioning Boardman work (Boardman, 1999). He describes three types of file (IMG, GLT and GEO). These files are created in the image geo-rectification process, and they split up the image information (between data and meta-data information), but this work is different to ours, because we do not apply any geo-rectification for the image storage, but when the image is going to be processed.

The rest of the paper is organized as follows: section 2 explains the solution suggested in this work (diffuse matrix), after that we put forward the application created for this study which works with the new structure and then, in section 4, we speak about the obtained results. Finally, conclusions and future work are expounded in section 5.

2 NEW DATA STRUCTURE: DIFFUSE MATRIX

There are some criteria that make an image format to be a good format (Folk, 1998). Between this characteristics can be pointed out the following:

- The space that it occupies, both in disk and in memory.
- The easiness of the format (it has to be simple and easy to understand).

- It has to be self-describing, using a metadata file or similar.
- Allow sequential access through the image.
- Information access has to be easy to implement.
- Rigorous and perfectly clear definition.
- It has to be efficient: using the format, image processing algorithms has to run efficiently and quickly.

We try to follow all the previous requirements in our work. Besides, we have made a robust format taking into account the problems caused by the sensor characteristics (Martinez et al., 2005) (which are explained in the introduction). Diffuse matrix separates physical storage and logical processing. Thus, spectral information is compressed and saved using a file named DMF (Diffuse Matrix File). In this file no redundant information is saved, and only when the structure is loaded in memory (when the image is going to be processed, but not in its storage), the required operations to put in order the spectrum in the matrix are done.

To preserve the acquisition scheme, and to have the information necessary for each band anytime, each position in the diffuse matrix (once it is loaded in memory) is a pointer to a dynamic list, where each node contains the information shown in fig. 3.

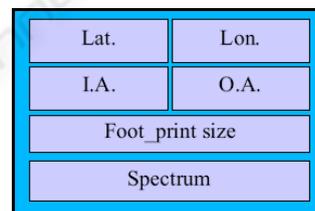


Figure 3: Data register created for each measurement taken by the sensor.

As it can be seen in figure 3, each register has 6 fields: A pair of fields of two bytes for latitude (Lat.) and longitude (Lon.) of the scanned area; another two fields for incidence angle (I.A.) and observation angle (O.A.); one field of one byte which holds the pixel spatial resolution (Foot_print size) for the captured image, and finally, a field of variable length for the spectrum (data) scanned by the sensor.

In diffuse matrix, each cell contains a pointer, which points to null if no information is hold for that coordinate in the matrix (as we said, this structure saves space, because there are not interpolation of missing information. If there are nothing, it is saved nothing); or points to a dynamic list where each element is a register like the described in fig. 3.

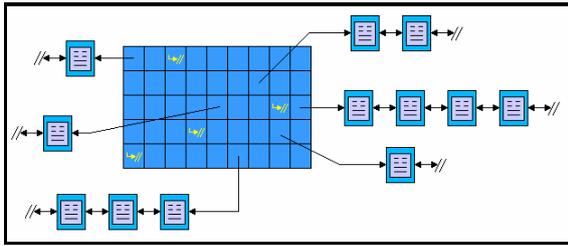


Figure 4: Diffuse matrix structure with some example nodes (empty and with information nodes).

Figure 4 shows the diffuse matrix structure. At first sight, the advantages obtained with the usage of this structure are:

- Diffuse matrix does not need any information not captured by the sensor, so, any interpolation or processing is not needed to save the information.
- Original coordinates (latitude and longitude) are saved for each measurement.
- Data distribution is independent of visualization. So, when data are shown, they have to be organized before, because the information is saved in an optimal way, without considering data spatial location. (Only in the latest stages of data processing, the interpolation of the image is necessary -to calculate missing data in the image acquisition-).

3 AN EXAMPLE OF APPLICATION FOR DIFFUSE MATRIX PROCESSING

We have developed a program just to show the possibilities and the usage of diffuse matrix. The application fulfils the following goals:

- Diffuse matrix generation from DMF file which contains it.
- Image visualization using diffuse matrix structure.
- Performance of some image processing algorithms over the diffuse matrix (we have chosen a set of thresholding and filter operations).

3.1 Diffuse Matrix Loading from DMF File

A DMF image is divided into two files: a DMF file, which contains the image, and a HDR file (header file), which contains meta-information about the

image. Fig. 5 shows the information held in a typical HDR file. HDR file contains information about the number of image samples (which will be the number of cells with information in the diffuse matrix), the number of rows and columns that the matrix will have, the number of bands per sample (bands in the image), and some values to calculate the exact position (X,Y) of each sample.

DIFFUSE MATRIX	
Samples	= 16437
Columns	= 98
Rows	= 186
Bands	= 224
GIFOVLon	= 0.06430849350967992
GIFOVLat	= 0.06411265872172466
MinLonXPoint	= 478905.8126117649
MinLatXPoint	= 478891.8524675080
MinLonYPoint	= 4169180.816114629
MinLatYPoint	= 4170156.255587298
MaxLonXPoint	= 480408.4594452424
MaxLatXPoint	= 480420.4117560154
MaxLonYPoint	= 4172073.124577305
MaxLatYPoint	= 4169282.552612827

Figure 5: HDR file with meta-information about a DMF image.

To load the diffuse matrix from a DMF file is necessary to use the HRD file (fig. 5) which is associated with that DMF file. Loading the image in the matrix basically consists in reading each sample sequentially from the DMF file (we know how many samples there are from HRD file). After reading one sample, its position is calculated and that sample is added to the right coordinate in the matrix. Interpolation will be performed to calculate the final position of each sample in the matrix if the image is loaded in memory with some association factor among the cells. The application allows 4 levels of association (the user chooses it when the image is loaded), as we will explain in the following section.

Diffuse matrix structure is hold in memory all the time while the application is working with the image. Thus, the position of each sample is only calculated once at the beginning of the process (not for every image processing operation), doing it in an efficient way.

When the process starts, the number of samples for a particular cell of the matrix is unknown. Only when the image is completely loaded, we know if in a particular cell is held one, several or no samples. So, a matrix of dynamic lists is the best structure to hold the image in memory.

The position of a particular sample in the diffuse matrix is obtained using the following equations:

$$\begin{aligned} \text{posX} &= (\text{Lat} - \text{MinLatXPoint}) * \text{GIFOVLat} \\ \text{posY} &= (\text{Lon} - \text{MinLonYPoint}) * \text{GIFOVLon} \end{aligned} \quad (1)$$

Lat (latitude) and *Lon* (longitude) values are read for each sample (fig. 3), while the other values are the same for all the samples in the image and are read from HDR file (fig. 5).

3.2 Image Visualization using Diffuse Matrix Structure

When the hyperspectral image is shown, it is considered as a set of n monochrome images (256 grey-levels each). The user chooses the band that he/she wants to see and this band is shown as a grey BMP image (in fact, the application allows saving that image as a BMP image). The user also chooses the *association factor* that the matrix cells will suffer when the image is loaded at the beginning. Association among cells is allowed because some cells in the matrix are empty (without samples); so, if the association is not done, some areas in the image will not have any information (in this case the image has some pixels without information, and the application shows them in purple colour -they are not processed-). Fig. 6 shows an image loaded with and without cell association.

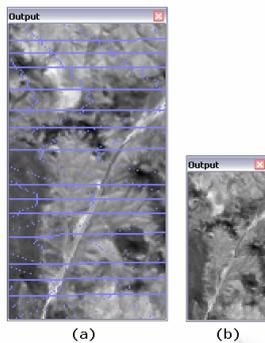


Figure 6: Image visualization with no association among cells (a) and with a 2x2 association (b).

The application allows association factors of 1x1 (no association), 2x2 (groups of 4 cells), 3x3 (groups of 9 cells) and 4x4 (groups of 16 cells). As we can see in fig. 6, the more association factor that the image has, the smaller it appears in the window when it is shown (this reduction of dimensions has some performance advantages, as we will see in the next section). For the right location of each sample in the matrix, the formulas shown in equation 1 are divided by the *association factor* chosen by the user. For example, the column occupied by a sample is calculated using the updated formula: $\text{posX} = (\text{Latitude} - \text{MinLatXPoint}) * \text{GIFOVLat} / \text{association factor}$ (the same that in equation 1 but divided by the *association factor*).

With the association, it is difficult that any cell of the matrix stays empty, because in case that the association factor is, for instance, equal to 2, each cell contains the samples of 4 neighbouring pixels.

3.3 Image Processing using Diffuse Matrix

To prove the easiness and potential of diffuse matrix, we have developed some thresholding and filter operations (González and Woods, 1992). The user uses the window shown in figure 7 to configure the different operations that can be done over the diffuse matrix.

In this window, the user can configure the band which he/she wants to work with (on the right area of the window) and the operation that he/she wants to apply over the selected band (no operation -only showing the image-, a thresholding operation or a filter operation -of smoothing, sharpening or edge detection-). In particular, we have implemented 15 filters: Average3x3, Average5x5, Average7x7, Gauss0391, Gauss0625, AverageModA, AverageModB, Laplace1, Laplace2, SobelX, SobelY, Sobel45+, Sobel45-, Prewitt1 and Prewitt2. In summary, we have implemented an important amount of operations using the structure (both thresholding operations and convolution operations) to perform a complete test of the behaviour of the diffuse matrix for these operations, which are quite important in image processing. In fig. 8, some examples of filtering are shown.

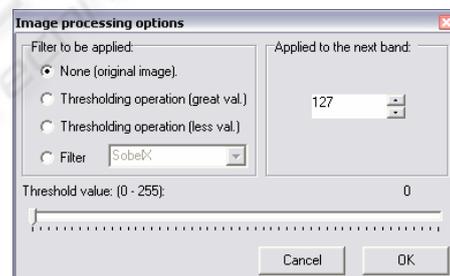


Figure 7: Image processing configuration window.

Moreover, if the user chooses association among the cells of the matrix, an important advantage is that the image which the application has to operate is "a dense image", smaller than the original image, so, the image processing is faster and more efficient. In fact, this idea is similar to the one used in Vector Architectures, with the technique called "scatter-gather" -which is included on many of the recent supercomputers-, where data are grouped to process them faster in a parallel way). In conclusion, diffuse matrix can be processed faster than traditional formats if the user chose association among the cells when the image was loaded.

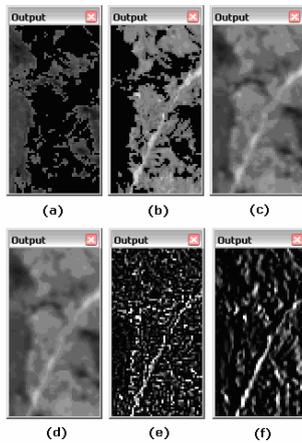


Figure 8: Some example of filters applied over the band 36 of the image.

The filters observed in figure 8 are: (a) thresholding -pixels with values less than 100 become black-, (b) thresholding -pixels with values greater than 100 become black-, (c) 3x3 average filter, (d) Gauss filter, (e) Laplace filter, (f) SobelX gradient filter.

We have used the following method to implement every filter operations over the diffuse matrix: For each cell in the matrix is calculated the centre position and the distance from this centre to every sample in the cell. Using these distances, the weight of each sample over the pixel which is going to be drawn in that position is calculated. The value of the pixel associated to a cell is obtained taking into account all the weights of all the samples in the cell after a normalization process into the range 0-255. This process is done for every cell in the selected band. Finally, the chosen operation is applied over the normalized pixel matrix and the result is shown as can be seen in fig. 8.

4 SOME RESULTS OBTAINED WITH THE STUDY

We have done some size comparisons, and we can conclude that due to the lack of redundant information included in the DMF file from georectification, the size of the image is much smaller than using traditional formats. In fact, we have done a comparison with the format proposed by Boardman in (Boardman, 1999) (which is an optimization that consist in 3 files to support an image -GEO, GLT and UTM files-). Using the same image (*IVAHOHA_BEACH_low_altitude*), the following results are obtained: our file with the

diffuse matrix (DMF) occupies 23,648,372 bytes (22.5 MB) and our HDR file takes up an almost negligible 551 bytes. On the other hand, the Boardman solution files take up: 69,898,752 bytes (66.6 MB) for the GEO file, 624,096 bytes (609 KB) for the GLT file and 2,496,384 bytes (2.4 MB) for the UTM file. If we compare the results, the compression ratio obtained is over 3:1, as we can see in fig. 9.

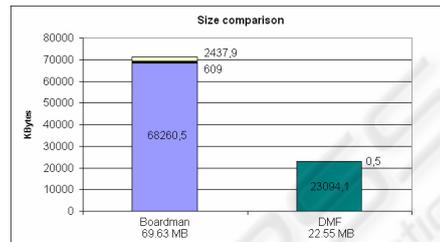


Figure 9: Size comparison between Boardman format and DMF format.

Furthermore, as we said in section 3.2, we can select an association factor when the image is loaded. If we associate cells when the matrix is loaded, the matrix will be compacted in memory (it has less dimensions) and the final size will be reduced. In fig. 10 we show the details for the same image without association factor and in figure 11 with association factor equal to 2. The diffuse matrix dimensions with an association factor of 2 are reduced to a quarter, but the cell dimensions are bigger with the association, because there are more points for each cell. Moreover, it is important to point that, as it can be seen in figures 10 and 11, with no association, the matrix has 0.9 samples per cell (there was cells with no samples), and with the association, there are 3.6 samples for each cell (at least 1 sample per cell and no more than 4 samples).

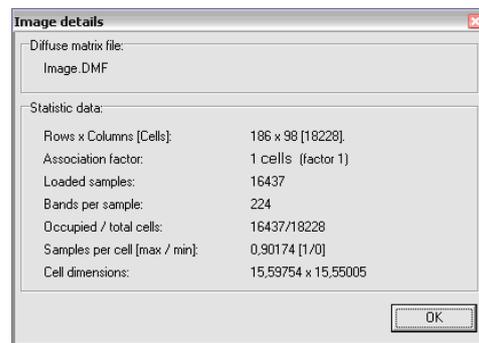


Figure 10: Image details for an image loaded without any cell association.

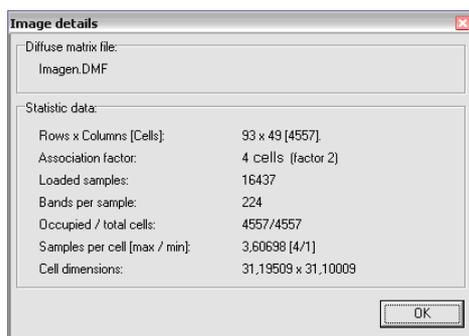


Figure 11: Image details for an image loaded with cell association (4 cells).

Anyway, with any association factor, the image is processed in the same way, because the structure of the matrix is always the same: a matrix of lists, where each node in the list contains the information of a sample acquired by the sensor, with altitude and longitude data (fig. 3), so the processing of this structure is homogeneous and efficient.

5 CONCLUSIONS AND FUTURE WORK

We have described a new format to work with AVIRIS hyperspectral images. With the DMF format, the physical storage in the disk (DMF and HDR files) and the later image processing (diffuse matrix with or without association of cells) have been clearly separated. This means that the same images take much less space in hard disk using DMF format than other typical formats. Furthermore, the image can be processed in a more efficient way, because it is loaded as a matrix of dynamic lists, taking advantage of the structure used. We have mentioned some of these advantages in the previous sections, but moreover, the diffuse matrix makes easier the parallelization of algorithms, because when the image is going to be processed, each cell is independent of the others in the matrix, and it is possible to divide the diffuse matrix and process each piece in a separate/parallel way, improving and speeding up the image processing (this aspect is very interesting because the size of hyperspectral images is quite big).

In conclusion, this work is a first step (the most important milestone) for the upcoming works in next months. Future work includes: (a) a detailed statistical comparison among different algorithms and different formats to get the concrete improvement obtained by the diffuse matrix in each

case; (b) a study in depth about the parallelization of some algorithms using the proposed structure (diffuse matrix).

ACKNOWLEDGEMENTS

This work has been developed in part thanks to the OPLINK project (TIN2005-08818-C04-03).

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