On data analysis and variable selection: the minimum entropy analysis

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Abstract

In this work, we present a minimum entropy analysis scheme for variable selection and preliminary data analysis. The variable selection can be achieved by the increasing preference of variables. We show such a preference to has a unique form, which is given by the entropy of models associated with variables. Evaluating the entropy provides a complete ranking scheme of variables. This scheme not only indicates preferred variables but also may reveal the system’s nature and properties. We illustrate the proposed scheme to analyze a set of geological data for three carbonate rock units in Texas and Oklahoma, and compare to the discriminant function analysis. The result suggests this scheme to provide a quick and robust analysis, and the use in data analysis is promising.

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1 Introduction

When one investigates an unknown system according to experimental observations made on this system, two questions are commonly addressed. How does one model the system? The model associates the experimental observations with corresponding experimental responses. Unfortunately, there is no systematical method to answer it. It is usually resolved through the methods of trials and errors, empirical regressions, and some intuitive assumptions etc. The second question is that suppose a model is found, which experimental observations shall be codified into the model. This is exactly a variable selection question. Here we will put the first question aside, and focus on resolving the second question.

A problem similar to variable selection has been tackled through different approaches in the past. It is the model selection problem. Several methods such P-value, Bayesian approach, and Kullback-Leibler distance based approach et al. are some examples. The P-values method selects model by comparing probability of model given a null model and experimental data sets to a threshold value assessed from same data sets \cite{1}. Yet since this method is restricted to two models and required ad hoc rules to assess threshold value, people has developed the Bayesian approaches to overcome these defects (\cite{1} and \cite{2}). The Bayesian method applies Bayes theorem to update our beliefs and uncertainty about models from prior distributions generated from some prior modeling rules first. A preferable model, thereafter, is chosen according to Bayes factor, ratio of posterior distributions of different models. Bayesian Information Criterion (BIC) is one of most popular Bayesian based model selection criteria (\cite{1}, Kieseppa, Forbes and Peyrard’s works in \cite{2}). Yet all of these methods require prior information generated from some ad hoc prior modeling rules that suits people’s need.

Aside from Bayesian framework, people has developed relative entropy, mutual information, or Kullback-Leibler distance based approach (\cite{3}, \cite{4}, and An Information Criterion (AIC) of Akaike \cite{5}). The rationale is to design a criterion based on aspect of entropy. The reason of employing
relative entropy for selection criterion is discussed in [6]. Recently, another criterion for model selection, CIC, is proposed by Rodriguez based on aspect of information geometry [7]. That is a generalized version of AIC and BIC. Preferred model is selected to minimize a quantity derived from Bayes’s theorem.

In the case of variable selection, Dupuis and Robert [4] proposed to model the system with different combinations of experimental observations, the variables selected from a set of variables. It generates several models associated with different combinations of variables. Thus variable selection problem becomes the problem of model selection. Evaluating the Kullback-Leibler distance between full model described by a complete set of variables for interested system and it’s approximations, submodels, described by subset of variables. When the full model is tractable, the preferred submodel is selected when it’s Kullback-Leibler distance reaches a threshold value. This threshold value is usually estimated by experiences from experimental data. Since submodels are projections of full model, there is no need the prior modeling rule to generate prior distribution for submodel. Yet one still requires prior information on full model. In addition, when there is no complete set of variables, namely, only limited experimental measurements regarding to the system can be conducted, this strategy becomes inadequately.

Our goal is to apply the method of maximum entropy (ME) to design a tool for variable selection that is free from defects in Dupuis and Robert’s approach. Following the axiomatic approach proposed in developing method of ME to a tool for assigning a probability to a system [8] and a tool for updating probability [9], Tseng has showed an entropic criterion for model selection [6]. Based on this study, we generalize it to obtain a Minimum Entropy Analysis (MEA) in Sec.2. The proposed analysis tool provides a complete ranking scheme of variables. It not only allows to select preferred variables but also to suggest an analysis scheme to reveal nature and properties of the system. To illustrate the MEA scheme, we will study a binary system in the geology in Sec.3. Some discussions are made afterward. Our demonstration shows the MEA scheme to be a promising tool in data analysis. Finally some conclusions are given.

2 The minimum entropy analysis

2.1 Basic features

Despite designing a pertinent criterion for selection, the selection also can be achieved by the increasing preference of variables. Since properties and meaning of the variables, experimental observations, are sometimes quite different, it is meaningless to compare variables directly. For example, suppose two experimental observations, mass and area, are measured for studying a system. How does one evaluate weightings of these two quantities to determine the dominant quantity in the model given to study this system? Namely, what is the preference of these variables? In Dupuis and Robert’s on variable selection problem [4], they proposed codifying the variables by a specific model such as logit model for a linear binary system. Afterward, ranking variables by evaluating the Kullback-Leibler distance between the model described by complete set of variables and it’s projections, submodels, described by subset of variables. Yet when there is no complete set of variables, namely, experimental measurements only provide limited information regarding to the system, this strategy becomes inadequately.

The approach for model selection proposed by Tseng may spells out a resolution in variable selection problem [6]. It states that suppose a family of probability models is found to interpret the system \( \{ P^m(x) \} \), where \( m \) labels the model and \( x \) denotes states of the model. The preference of models is uniquely determined, which is in the form of relative entropy of model \( P^m(x) \) and a uniform reference measure \( \mu \),

\[
S[P^m | \mu] = - \sum_x P^m(x) \ln \frac{P^m(x)}{\mu} .
\]
This scalar value measures differences between models and a uniform reference measure. Maximizing the relative entropy indicates $P_m(x)$ to equal to $\mu$. Namely, there is no information regarding to the system being codified into $P_m(x)$. Within the family, when the relative entropy is decreased, there is more information of the system being codified into $P_m(x)$ One can then rank those probability models according to decreasing $S[P_m|\mu]$ value.

2.2 Logic behind the MEA

Based on Dupuis and Robert's approach, one can determine the preference of variables by first codifying these variables into a model. This model can be any function that optimally associates the experimental observations, variables, and responses. According to Tseng's work on model selection [9], one needs to further codify this model into a probability distribution of observing the experimental responses given the variables in order to compute the preference. Thus the preference of variables is determined form the preference of those probability distributions.

Based on these aspects, the logic behind the proposed MEA scheme then involves two stages. The first stage is to determine a probability model that associates experimental observations and responses. The method of ME proposed by Jaynes [8] provides a solution. Since the method of ME requires the information that will be codified into the probability distribution to be in the form of constraint, it turns the question of probability assignment into a question of what is constraint. We will come back this point later.

Next we follow the axiomatic approach [9] to determine the form of preference of the probability distributions. The basic strategy (Skilling of [9]) is one of induction: (1) if a general rule exists, then it must apply to special cases; (2) if in a certain special case we know which is the best approximation, then this knowledge can be used to constrain the form of preference; and finally, (3) if enough special cases are known, then preference will be completely determined. The known special cases are called the “axioms” of ME. The axioms used here must reflect the conviction that one should not change one’s mind frivolously, that whatever information was originally codified into the exact probability distribution is important and should be preserved. As shown in [9], three axioms are employed: (1) local information has local effects; (2) the ranking should not depend on the coordinates of the system, and (3) consistency for independent subsystems. The functional form for the preference is therefore uniquely determined, which has the form of relative entropy, Eq.(1).

2.3 The MEA scheme

Suppose a model, function of $f(\hat{x}, \hat{\beta})$, is given to associate N experimental observations denoted by variable $\hat{x} = \{x_1, x_2, \ldots, x_N\}$ and parameters $\hat{\beta} = \{\beta_1, \beta_2, \ldots, \beta_N\}$ with experimental responses $M(\hat{x}, \hat{\beta})$. Each observations is repeated $l$ times, which give $l$ measurements $x_i = \{x_{i1}, x_{i2}, \ldots, x_{il}\}$ and corresponding responses $M(\hat{x}, \hat{\beta}) = \{M^1, M^2, \ldots, M^l\}$. The response $M$ will be either “1” for positive response or “0” for negative response. Notes that one way to determine $\hat{\beta}$ is through method of Maximum Likelihood Estimate (MLE) [10]. For example, the logit model

$$f_{\text{logit}}(\hat{x}, \hat{\beta}) = \frac{\exp \sum_{i=1}^{N} \beta_i x_i}{\exp \sum_{i=1}^{N} \beta_i x_i + 1},$$

is usually given as a regression model for a binary output system. Given these N variables, there will be $2^N - 2$ different combinations (subsets) of variables $\hat{x}_{s_i} \in \hat{x}$ to be chosen from $\hat{x}$, and form $2^N - 2$ submodels $f(\hat{x}_{s_i}, \hat{\beta}_{s_i})$. 

3
Afterward we have to determine the probability $P_s (\hat{x}_s | \beta_s)$ of observing the experimental responses $\mathbf{M}$ given the subset of variables $\hat{x}_s$. In the framework of method ME, one has to identify the relevant information to be codified into the probability distribution. In this example, it is obvious that the experimental responses are the quantity we need to know about the system, that can be written in the form of constraint

$$\langle \mathbf{M} \rangle = \sum f (\hat{x}_s, \beta_s) P_s (\hat{x}_s | \beta_s),$$

where $\langle \mathbf{M} \rangle$ is the expectation value of the responses. Thus the method of ME indicates the preferred $P_s (\hat{x}_s | \beta_s)$ to be

$$ P_s (\hat{x}_s | \beta_s) = \frac{\exp - \lambda f (\hat{x}_s, \beta_s)}{Z},$$

where partition function $Z = \sum_{\hat{x}_s \in \hat{x}} \exp - \lambda f (\hat{x}_s, \beta_s)$ and $\lambda$ is a Lagrangian multiplier, which is set to one for sake of simplicity. Alternatively, the probability distribution can be obtained by simply normalizing submodels $f (\hat{x}_s, \beta_s)$,

$$P_s (\hat{x}_s | \beta_s) = \frac{f (\hat{x}_s, \beta_s)}{Z'},$$

where normalization constant $Z' = \sum_{\hat{x}_s \in \hat{x}} f (\hat{x}_s, \beta_s)$ as well. Notes that this form of probability distribution is actually just a first order approximation of Eq. (4).

Thus the increasing ranking order of these submodel $f (\hat{x}_s, \beta_s)$ is given by the decreasing relative entropy Eq. (1) with $P^m$ being replaced by $P_s (\hat{x}_s, \beta_s)$ or $P_s (\hat{x}_s, \beta_s)$. Furthermore, one can easily rewrite Eq. (4) into

$$S[P_s | \mu] = S[P_M] + \ln \mu,$$

where $S[P_s] = - \sum_{\hat{x}_s \in \hat{x}} P_s (\hat{x}_s, \beta_s) \ln P_s (\hat{x}_s, \beta_s)$ entropy of the submodel. Because $\ln \mu$ is a constant value for a uniform reference measure, the preference of submodels is equivalent to the decreasing $S[P_s]$ value.

Evaluating the entropy of all submodels, a complete ranking scheme of different subsets of variables $\hat{x}_s$ is determined. Preferred subset of variables then can be identified that is the one that has minimum entropy value within this set of variables. Notice that the use of this scheme is not totally exhausted yet. By further analyzing the ranking scheme of different subsets $\hat{x}_s$, one may determine significance of different combinations of variables that are codified into the submodel. The nature and properties of the system may thereafter be revealed through this analysis. For example, as we know, correlation functions between two variables arbitrarily chosen within $\hat{x}$ may reveal some properties of the system. Although we did not compute the correlation functions in this scheme, the preference of different correlations is still implicitly spelled out by the ranking scheme of different subsets. One can attribute this to when the model is given to associate the variables and responses, the correlations between the variables are defined in the model. Thus determining the ranking scheme of different combinations of variables indirectly indicate the significance of different correlations.

Thus one may treat the MEA scheme as a quick data analysis tool. It provides preliminary information about the system. This use is implicitly in some other approaches mentioned previously. We will illustrate the use of the MEA scheme in detail by studying a geological problem next.
3 A special case: a binary system in Geology

3.1 The problem

Considering a geological example of sample classification (Davis [11]). We briefly address the result by means of a standard tool, the Discriminant Function Analysis (DFA), for classifying geological samples. Then we show how to extract important variables in determining the category of samples using our minimum entropy analysis. Furthermore, some information regarding the formation of sample rocks. Comparing both results from the DFA and the MEA improves our understanding and enhances our confidence in our MEA scheme.

Saltwater is trapped in sedimentary rocks at the time they are formed in the marine environment. The chemical composition of the connate water is subsequently modified by ion exchange and other reactions, by mixing with other brines, and by dilution by infiltrating surface waters. Brines recovered during drillstem tests of wells may have relict compositional characteristics that provide clues to the origin or depositional environment of their source rocks. Table 1 contains brine analyses for oil-field waters from three groups of carbonate units in Texas and Oklahoma (Davis [11]). The first column in Table 1 denotes the brine samples belonging or not belonging to some specific carbonate unit, Unit G.

3.2 The discriminant function analysis and results

The discriminant function analysis combines a rationale similar to that of analysis of data variance with computational procedures based on eigenvector calculations, e.g., the PCA (principle component analysis). Multivariate measurements made on the samples alone, such as the brine data in Table I, can be used in the DFA to find combinations of measurements that allow the various categories of samples to be distinguished. The problem of DFA is basically one of finding a set of linear weights for the variables that causes a multivariate analogue of the F-ratio to be a maximum. A succession of discriminant functions along which the samples are as distinct as possible, can be thus calculated and each represents successively the most efficient discriminator possible. For many calculation details, please refer to the book of Davis [11].

The DFA can be applied to those data in Table I to determine if they are distinctive. The first discriminant function thus calculated is an inner product of 
\[ (-0.3765, -0.0468, 0.0112, -0.0148, -0.0174, -0.0110) \cdot (\text{HCO}_3, \text{SO}_4, \text{Cl, Ca, Mg, Na})^T \]
which can clearly separates samples from Unit G and other units. Please note that the weighting factors in the first discriminant function for variables \text{HCO}_3 and \text{SO}_4, i.e. -0.3765 and -0.0468, represent the first two largest factors in magnitude among six, thus indicating those two variables play the most dominant effect in classification.

3.3 The minimum entropy analysis

According to conventional studies [4] and [10], one can pertinently associate six experimental observations with binary responses in this geological example through the logit model, Eq. 2. The variables \( \mathbf{x}_G = \{x_1, x_2, \ldots, x_6\} \) denotes observations on contents of six chemical compounds \{HCO_3, SO_4, Cl, Ca, Mg, Na\} correspondingly. As shown in Table I, nineteen measurements were made. There are five positive experimental responses, denoted by symbol “Y” and fourteen negative responses. The method of ME suggests that the preferred probability distribution of observing the experimental responses given variables \( \mathbf{x}_G \) to give

\[
P(\mathbf{x}_G; \hat{\beta}) = \frac{\exp\left(-f_{\text{logit}}(\mathbf{x}_G, \hat{\beta})\right)}{Z},
\]

(7)
Table I: Chemical analyses of brines (in ppm) recovered from drillstem tests of three carbonate rock units in Texas and Oklahoma. Adapted from Davis of [11].

| Unit G | HCO$_3$ | SO$_4$ | Cl | Ca | Mg | Na |
|--------|---------|--------|----|----|----|----|
| N      | 10.4    | 30     | 967.1 | 95.9 | 53.7 | 857.7 |
| N      | 6.2     | 29.6   | 1174.9 | 111.7 | 43.9 | 1054.7 |
| N      | 2.1     | 11.4   | 2387.1 | 348.3 | 119.3 | 1932.4 |
| N      | 8.5     | 22.5   | 2186.1 | 73.6 | 1803.4 |
| N      | 6.7     | 32.8   | 2015.5 | 75.1 | 1691.8 |
| N      | 3.8     | 18.9   | 2175.8 | 63.8 | 1793.9 |
| N      | 1.5     | 16.5   | 2367   | 95.8 | 1872.5 |
| Y      | 25.6    | 0      | 134.7  | 12.7 | 7.1 | 134.7 |
| Y      | 12      | 104.6  | 3163.8 | 95.6 | 90.1 | 3093.9 |
| Y      | 9.0     | 104    | 1342.6 | 104.9 | 160.2 | 1190.1 |
| Y      | 13.7    | 103.3  | 2151.6 | 103.7 | 70 | 2054.6 |
| Y      | 16.6    | 92.3   | 905.1  | 50.9 | 871.4 |
| Y      | 14.1    | 80.1   | 554.8  | 62.3 | 472.4 |
| N      | 1.3     | 10.4   | 6399.5 | 532.3 | 235.6 | 2642.5 |
| N      | 3.6     | 5.2    | 974.5  | 147.5 | 69 | 768.1 |
| N      | 0.8     | 9.8    | 1430.2 | 295.7 | 118.4 | 1027.1 |
| N      | 1.8     | 25.6   | 183.2  | 35.4 | 161.5 |
| N      | 8.8     | 3.4    | 289.9  | 32.8 | 225.2 |
| N      | 6.3     | 16.7   | 360.9  | 41.9 | 318.1 |

where partition function \( Z = \sum \hat{x}_G \exp \left( -f_{\logit}(\hat{x}_G, \hat{\beta}) \right) \). Or normalizing the logit model within this data set gives

\[
P'(\hat{x}_G | \hat{\beta}) = \frac{f_{\logit}(\hat{x}_G, \hat{\beta})}{Z'} = \frac{1}{Z'} \exp \sum_{i=1}^{6} \beta_i x_i + 1 \exp \sum_{i=1}^{6} \beta_i x_i ,
\] (8)

where normalization constant

\[
Z' = \sum \exp \sum_{i=1}^{6} \beta_i x_i + 1 .
\] (9)

Given these six variables, \( 2^6 - 2 = 62 \) different combinations of variables \( \hat{x}^G \in \hat{x}_G \) are obtained. Thereafter, one can generate 62 probability submodels \( P_{s_i}(\hat{x}^G) = P(\hat{x}^G | \hat{\beta}) \) or \( P'(\hat{x}^G | \hat{\beta}) \).

Evaluating the entropy of \( P_{s_i}(\hat{x}^G) \) and \( P'(\hat{x}^G | \hat{\beta}) \), with different subsets of variables \( \hat{x}^G \) gives ranking order of different submodels \( P_{s_i}(\hat{x}^G) \). The coefficient \( \beta_i \) are determined through fitting the logit model to experimental measurements by MLE method (in Appendix: software for ordinal data modeling of [10] - a MATLAB function of “Maximum likelihood estimation and model criticism”). The result is listed in Table II. Here we only list 18 out of 62 submodels. We calculate the entropy of two probability distributions, Eq. (7) and (8), in second and third column respectively. The ranking scheme is in the order of decreasing entropy value. To analyze this ranking scheme, we proceed with a two-step approach. In first step, we examine the submodel that has the minimum entropy value. In this example, there are 16 out of 62 submodels that has the minimum entropy value, 2.866 or 1.791 in the second and third column of Table II respectively. Notes that since the minimum significant figure of experimental data in Table I is three, the entropy value should also has three significant figures and forth digit is just an estimate. The preference of these 16 submodels are indistinguishable. Notes that the digits in round bracket shows a numerical result when the significant figure is not considered. It just indicates that if the significant figures are higher the resolution of entropy will
be better. Thus the preference of these 16 submodel then still can be identified. That will further aid the analysis of preference in detail.

In order to determine the most dominant variables from these 16 submodels, frequencies of six variables appeared in these 16 submodels are recorded in the present. The frequencies for observing first and second variable are 16 and 15 respectively and 8 for rest of variables. This result suggests that the ability of interpreting the experimental measurements by the logit model is strongly dominated by the first variable, HCO$_3^-$, and the second variable, SO$_4^{2-}$. The variables 3 to 6 seem to play a minor role here. This is exactly the result obtained though the DFA analysis mentioned previously. Yet the MEA scheme is more straightforward.

In second step, we analyze the ranking scheme further to identify preference of first and second variable. Since the variables 3 to 6 play a minor role here, we concentrate on first two variables. We list two more submodels that only include the first and second variable, HCO$_3^-$ and SO$_4^{2-}$, respectively in Table II. The entropy value in third column shows a dramatic changes from 2.328 of submodel \{010000\} that only has second variable, 2.060 of \{100000\} that has only first variable, to 1.791 of \{110000\} for the case of two variables being simultaneously included. The same trend is also observed in the second column although no dramatic changes is observed. The ranking scheme indicates that the first variable HCO$_3^-$ should play a more important role than the second variable SO$_4^{2-}$ in the model.

| HCO$_3^-$ | SO$_4^{2-}$ | Cl | Ca  | Mg  | Na  | $S[P_s]$  | $S[P'_s]$ |
|-----------|-------------|----|-----|-----|-----|------------|------------|
| 0         | 1           | 0  | 0   | 0   | 0   | 2.893(229075) | 2.328(713745) |
| 1         | 0           | 0  | 0   | 0   | 0   | 2.881(069331)  | 2.060(483312)  |
| 1         | 1           | 0  | 0   | 0   | 0   | 2.866(92132)   | 1.791(857378)  |
| 1         | 1           | 0  | 0   | 0   | 1   | 2.866(921309)  | 1.791(85668)   |
| 1         | 1           | 1  | 0   | 0   | 0   | 2.866(921264)  | 1.791(854715)  |
| 1         | 1           | 0  | 1   | 0   | 0   | 2.866(921168)  | 1.791(849677)  |
| 1         | 1           | 0  | 0   | 1   | 1   | 2.866(921139)  | 1.791(848879)  |
| 1         | 1           | 1  | 1   | 0   | 0   | 2.866(921109)  | 1.791(84701)   |
| 1         | 1           | 1  | 1   | 0   | 1   | 2.866(921101)  | 1.791(846531)  |
| 1         | 1           | 1  | 0   | 0   | 1   | 2.866(921112)  | 1.791(843955)  |
| 1         | 1           | 1  | 1   | 0   | 0   | 2.866(921028)  | 1.791(842415)  |
| 1         | 1           | 1  | 1   | 0   | 1   | 2.866(921084)  | 1.791(842417)  |
| 1         | 1           | 1  | 0   | 0   | 1   | 2.866(921072)  | 1.791(841769)  |
| 1         | 1           | 0   | 1   | 1   | 0   | 2.866(921005)  | 1.791(840749)  |
| 1         | 1           | 1   | 1   | 1   | 0   | 2.866(920963)  | 1.791(83836)   |
| 1         | 1           | 0   | 1   | 1   | 1   | 2.866(920964)  | 1.791(838358)  |
| 1         | 1           | 1   | 0   | 1   | 1   | 2.866(920971)  | 1.791(838291)  |
| 1         | 1           | 1   | 0   | 0   | 1   | 2.866(920965)  | 1.791(838227)  |

4 Discussions

Lithologically speaking, unit G is mainly composed of dolomite (CaMg(CO$_3$)$_2$) and anhydrite (CaSO$_4$). In ancient geological times, Unit G, which is in geology called the “Grayburg Dolomite” [11], experienced two important sedimentary processes of dolomitization, which is associated with the dissolution of calcite by acidic fluids, and evaporation. Anhydrite is one of the index products
from evaporation. Chalcraft and Ward further claimed that the principal diagenetic processes include dolomitization, anhydrite occlusion of primary porosity, and leaching [12]. The dolomitization plays a crucial role in the formation of unit G, and followed by anhydrite occlusion of primary porosity. One therefore can infer that the process of the dissolution of calcite by acidic fluids is more significant than anhydrite occlusion in the formation of unit G. Namely, the process involves with chemical compound HCO$_3^-$ is the most important one among the six compounds.

In analyzing a set of geological data to seek out the origin or depositional environment of their source rocks, the MEA suggests that HCO$_3^-$ and SO$_4^{2-}$ are two key variables in the occurrence process of Unit G. The MEA also suggests that HCO$_3^-$ plays a more important role than SO$_4^{2-}$. Therefore, we can conclude that formation of Unit G may strongly involve with chemical process associated with HCO$_3^-$. The chemical process associated with SO$_4^{2-}$ may then plays a minor factor in the formation. It is the exact result inferred previously but the MEA analysis is more straightforward. Similarly, one may conduct further analysis to extract more information, yet it is out of our scope here.

5 Conclusions

The minimum entropy analysis scheme is proposed to analyze experimental data for extracting information of the corresponding system such as which experimental observations to play a more important role etc. This is a question of variable selection, and can be resolved by determining the preference of these observations. To determine the preference, one associates the experimental responses and those observations by a probability model first. Thereafter, as shown in the context, the form of preference is uniquely determined through the axiomatic approach [9]. It is in the form of entropy of probability of observing the experimental responses given variables. The preferred variables are the one that have minimum entropy value. Furthermore, since the minimum entropy analysis present a complete ranking scheme of different combinations of experimental observations, it indirectly indicates significance of different combinations of variables in the model. This ranking scheme not only suggests the preferred variables that should be codified into the model but also may spell out a route to study the system. Besides, this design resolves two defects in Dupuis and Robert’s approach [4] mentioned previously.

We have illustrated the use of the minimum entropy analysis by analyzing a set of geological data for three carbonate rock units in Texas and Oklahoma. The MEA scheme indicates the preferred variables most relevant to the formation of unit G or Grayburg Dolomite to be HCO$_3^-$ and SO$_4^{2-}$. This result agrees with the result from another well known analysis tool, the discriminant function analysis. Furthermore, since the MEA presents a complete ranking scheme of six chemical compounds measured in the samples, it points out a principal diagenetic process obtained in [12]. Yet this conclusion is not clear in the discriminant function analysis.

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