TECHNICAL UNIVERSITY OF CRETE Mineral Resources Engineering School



Performance of multivariate clustering methods in oil families' identification

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Abstract

As science progresses, the need for analyzing multivariate data sets is growing by the minute. Multiple disciplines, either scientific or not, require the examination of large amounts of data, in a short period of time, in order to obtain useful information. During the recent few decades, multivariate statistical analysis methods have been developed, aiming to satisfy such purposes.

This dissertation deals with the implementation of multivariate data analysis methods on a given data set, derived from oil family affiliations, which originate from Williston Basin of North America. In particular, Hierarchical Clustering, k-means and Principal Component analysis have been applied on four independent models, in an attempt to extract information regarding the oil-oil correlations among the samples under study. The models used on the exploration of the compositional information were the Saturated Fraction Compositional Model, the Saturated Fraction Ratios Model, the Gasoline Range Compositional Model and the Biomarkers Compositional Model.

These standard statistical methods were found to be quite insufficient in classifying the sample set into distinct familial affiliations. For this reason, the need to examine the nature of the data set arose. Compositional data represent a category on their own as they are characterized by specific numerical properties which present significant consequences when being analyzed by standard multivariate techniques. The analysis of such type of data represents a whole new chapter in the world of statistics and the need for further examination on this matter is constantly growing.

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1. Introduction

Over the last decades an overwhelming amount of data is poured into our lives and obtaining meaningful information out if them is an imperative task for people. Multiple disciplines such as chemistry, biology, medicine etc. demand the analysis of huge amounts of data and sometimes their multivariate nature makes it difficult to analyze. For this reason, special statistical techniques have been developed in order to process information in a meaningful fashion.

In this project, multivariate clustering methods have been implemented on geochemical data concerning oil family affiliations that exist in Williston Basin, North America, in order to explore the oil-oil correlations. The methods which have been utilized consider both Supervised and Unsupervised learning phases. These include Hierarchical Clustering, k-means clustering, as well as Principal Component Analysis. The ultimate goal of this project is to test how well such multivariate analysis methods perform, as far as classification of the compositional data is concerned.

The thesis project is organized into seven chapters. In Chapter 2 a detailed description of the geological setting of Williston Basin is presented. The stratigraphy and the tectonic regime are thoroughly described and special focus is placed upon the geochemical classification of oil families which have been recognized in the area.

Chapter 3 raises the subject of Multivariate Data Analysis (MDA). It provides a brief presentation of the principles of Hierarchical Clustering, k-means clustering as well as Principal component analysis. All the main concepts that characterize each method are included.

In Chapter 4 we discuss the matter of the existing Family Affiliations of Williston Basin. In this chapter, there is an attempt to test the criteria under which the classification of the oil families was determined.

Chapter 5 deals with the application of multivariate data analysis methods on two different models; the Saturated Fraction Component Model and the Saturated Fraction Ratios Model. All MDA methods were implemented on both models and the results are discussed briefly.

In the final Chapter (6) the subject of Compositional Data, as a special type of data, is introduced. In this chapter, we analyze the properties of Compositional Data as well as the methodology with which, such kind of data should be treated.

2. Geological Setting of Williston Basin

The Williston Basin is an intracratonic, sub-circular sag basin that comprises main part of the North American craton. In particular, it forms a large depression in the western edge of the Canadian shield, occupying much of North Dakota, northwestern South Dakota, the eastern quarter of Montana, a significant part of southern Saskatchewan, and a portion of southwestern Manitoba. Among these regions major production of oil and gas occurs. Williston Basin is characterized by Phanerozoic, carbonate and clastic sedimentation of more than 16,000ft strata thickness in its central part, near Watford City, North Dakota [1, 2]. Having undergone episodic and prolonged subsidence rates, it comprises a preservational basin and it is composed by six major depositional sequences, each bounded by larger structural features [2, 3, 4] (Fig. 1). The basin is neither considered structurally complex nor tectonically active and its well -established petroleum provinces, clearly described rock succession, modest burial history and simple tectonics make this an uncomplicated area to study.



Fig. 1 Location map showing the main geological and geophysical elements of Williston Basin and environs. The region of anomalous subsidence that is Williston Basin proper (Ahern and Mrkvicka, 1984) is generally coincident with the 1 km depth contour on Carboniferous strata. The region of preserved Middle Devonian Prairie Formation salt deposited in Elk Point Basin is illustrated. The inset shows the location of Williston Basin and the extent of Elk Point Basin. Samples from petroleum pools entrapped at the subcrop of the upper Paleozoic succession in southeastern Saskatchewan and southwestern Manitoba, as well as American samples constitute the sample set for this study (following Burrus et al., 1996a).

Williston Basin is discretized into the American and the Canadian portions. The American portion of the basin is influenced by major deformational features, mainly anticlines (Fig. 2). The Canadian part of Williston Basin forms a petroleum province where oil production is quite active. Petroleum accumulations mainly occur in stratigraphic traps within the Phanerozoic succession [5]. There is, however, variety of trapping features which are structurally linked to Precambrian basement [6, 1, 7]. In southwestern Manitoba and southeastern Saskatchewan, oil exists around the Mississippian subcrop. In southwestern and west-central Saskatchewan, oil exists in stratigraphic traps within latest Devonian to Mississippian, Jurassic, and Lower Cretaceous formations.



Fig. 2 Petroleum region and crucial tectonic elements in the Williston Basin and adjacent area. Only generalized outlines of the Mississippian Madison Group Subcrop Petroleum Province and other Williston Basin petroleum provinces are indicated.

2.1 Stratigraphy of Williston Basin

The Williston Basin forms a large, roughly circular depression on the North American Craton. Its sedimentology is characterized by Paleozoic and Cenozoic – Mesozoic carbonate and clastic deposition, accordingly with a thickness of strata that exceeds 16,000 ft in the basin's core (Fig. 3).



Fig. 3 Contour map pf Williston Basin presenting the thickness of sediments. Contour interval is 1,000 ft. [8]

There are six main depositional sequences, each bounded by major unconformities [3], which can be distinguished within the Phanerozoic succession of North American portion of the basin. The formulation of unconformities resulted in numerous processes affecting its final structure, such as primary and secondary dissolution, deposition of salt and anhydrite beds, and secondary dolomitization of limestone. Clastic deposition initiated in Mesozoic and Cenozoic Eras, including mudstone, sandstone, siltstone, coal and shale. All depositional sequences are briefly described in the following paragraphs.

Sauk Sequence (Middle Cambrian – Lower Ordovician)

The Sauk sequence was deposited on the early Paleozoic miogeocline of western North America [7, 9], and is composed of Upper Precambrian sediments, interrupted by minor transgressions and regressions, which create several sub-members within the formation [10]. Saul deposition, mainly represented by Deadwood formation, includes shallow marine, coastal and alluvial plain sediments along with sandstone, mudstone and siltstone successions and finalizes due to the activity of an unconformity.

Tippecanoe Sequence (Ordovician – Silurian)

The Tippecanoe sequence marks the beginning of Ordovician clastic, carbonate and evaporitic sedimentation. From bottom to top, it consists of Winnipeg, Red River, Stony Mountain and Stonewall Formations, each unconformably overlying the other (Fig. 4). Upper Ordovician

rocks of this sequence contain important petroleum sources. Depositional processes terminate at the end of the Silurian due to major regression activity.



Fig. 4 Diagram showing geologic time scale, major stratigraphic sequences of [3], first- and second order sea level curves from [11], and ages of petroleum source and reservoir rocks in the Williston Basin. Solid black interval in source rock column are for thick accumulations; thin lines indicate association with carbonate depositional cycles. In reservoir rock column, green is for oil and red is for gas; thin lines indicate generalized reservoir rock and do not necessarily represent the full spectrum of possible reservoirs. E, Early; M, Middle; L, Late; Pal, Paleocene; Eoc, Eocen; Olig, Oligocene; Mio, Miocene; Plio, Pliocene (following Lawrence, et al., 2013).

Kaskaskia Sequence (Devonian - Mississippian)

The Kaskaskia sedimentation cycle initiated in Ordovician, continued to Jurassic and concluded due to transgressional activity. Three main transgressional events impacted on the depositional history of the sequence, during which several formations were deposited. The most significant is the Bakken Formation which represents the first major input of clastic material into the Williston Basin since the Cambrian Deadwood and Winnipeg Formations. Bakken marks a change in Kaskaskia sequence depositional patterns and sedimentation style [12, 13] and it is the most important interval for petroleum source rocks in the Williston Basin. In general, the Kaskaskia Sequence is stratigraphically characterized by subtidal, intertidal and rare supratidal depositional environments.

Absaroka Sequence (Pennsylvanian - Triassic)

The Absaroka Sequence includes the Tyler and the Minnesula formations and mainly occurs in the American portion of the Williston Basin. It is vastly affected by major unconformities, occurring near the end of Pennsylvanian, Permian and Triassic [14] and contains effective oil source rocks [15, 16].

Zuni Sequence (Jurassic – Early Tertiary [Eocene])

Two major transgressional events influenced the depositional history of the Zuni Sequence, which is characterized by shallow marine and clastic sediments. Sedimentation terminated during early Paleocene and the sands of the Dakota Group are likely the most significant targets for sequestration in the Zuni Sequence. This sequence can be locally subdivided into

two other sequences. The first includes the Jurassic, when Williston Basin changed from a large reentrant on the craton margin into an orogenic foreland [17, 18]. The lower sequence contains a time equivalent succession to the last cratonically derived miogeoclinal succession.

Tejas Sequence (Tertiary - Quaternary)

Latest Jurassic and Cretaceous successions of the Columbian and Laramide orogenic forelands [19] form the final significant depositional episode [20, 21]. Thick shales of this final sequence include significant probable source rocks, but they are all immature in the Canadian Williston Basin. The first produced hydrocarbons in North Dakota were from the youngest strata in the state, glacial drift of the Tejas Sequence. However, there is no production from glacial drift today.

2.2 Tectonic Regime of Williston Basin

In order to understand the Williston Basin's evolution, structural configuration, sedimentation, and thermal patterns, one must refer to the geological history of the Precambrian basement underlying the basin.

Two critical structures have influenced the evolution of the basin; the Trans-Hudson orogenic belt [22] and the northeast—southwest trending Proterozoic lineament and structural zones [23]. The Trans-Hudson belt sutured the Archean Superior craton to the Archean Wyoming craton (Fig. 5A, B); the resulting collision created a north—south trending strike-slip fault and shear belt. A basin center was created, caused in part by later folding of the Trans-Hudson orogenic belt and rifting [24], although Nelson et al., [25] stated that there is a lack of direct evidence of a rift.

The northeast–southwest trending Proterozoic lineament and structural zones were renamed as the Transcontinental arch, Brockton-Froid fault zone, Great Falls tectonic zone, Poplar fault, and Hinsdale fault. These Precambrian structures were reactivated during the Neoproterozoic, which resulted in the creation of new north–south and northwest–southeast trending structures.



Fig. 5 Precambrian structural configuration of the Williston Basin and surrounding area. A: Tectonic map of the northern Great Plains region [23] showing northeast-southwest strike slip faults; Williston Basin province outline is shown for scale. Ga, billion years ago. B: Map showing the configuration of Trans-Hudson orogenic belt and associated north-south trending structures of the Williston Basin (modified Nelson et al., 1993).

Numerous studies have shown that surface lineament patterns in the Northern Great Plains region, including the Williston Basin, are a result of the aforementioned reactivation of Precambrian faults during the Phanerozoic [26, 27, 28, 29]. These studies show pervasive northeast—southwest and northwest—southeast trends that are parallel to major lineaments of Proterozoic terrane. North—south trending lineaments that are parallel to the Trans-Hudson structural system are less prominent, although north—south thermal patterns are evident from present-day subsurface temperature measurements.

Based on several observations, it is believed that Precambrian tectonic events and their recurrent movement along preexisting zones of weakness played a major role in the development of most of the major fault and shear systems in the Williston Basin. Although the basin is generally reported as a depression and tectonically inactive, its final structure is thought to be mostly formed as a result of structural deformation and down-to-the-basin block faulting from Precambrian rooted structures, as well as from deformation related to the Trans-Hudson orogenic belt.

2.3 Geochemical Classification of Oil Families in Williston Basin

Classification of oil families in the Canadian portion of the Williston Basin has been attempted by a number of investigators over the past decades. Dow and Williams, in their 1974 papers, were the first researchers to apply the 'petroleum system' concept, identifying three oil systems in the Williston Basin, relying mainly on stable isotopic and gasoline range hydrocarbon composition: Tyler, Bakken, and Winnipeg [15, 16]. Each oil system is associated with a unique oil type. Type I refers to Ordovician and Silurian oils which originate from Middle Ordovician Winnipeg shale sources. Type II oils occur in Upper Devonian, Mississippian and Mesozoic reservoirs, and are probably linked to Fammenian – Tournaisian Bakken Formation Source rocks. Type III refers to Pennsylvanian oils which originate from Tyler Formation source rocks.

Most recent studies, however, have defined at least nine oil systems in the area. Zumberge [30] and Leenheer and Zumberge [31] defined five oil families based on the study of samples from the American part of the Williston Basin, while, Osadetz et al., [32] categorized oils from the Canadian part of the Basin (southeastern Saskatchewan and southwestern Manitoba) into four compositional families (Table 1). The criteria under which the classification of the latter was conducted, include pristane/phytane (Pr/Ph) ratio, n-alkane predominance, C_{23} tricyclic/ C_{30} pentacyclic terpane ratio and prominence amongst extended hopanes.

In particular, Family A oils occur in Ordovician to Middle Devonian and Upper Ordovician formations and match solvent extracts from kukersites (marine Type I rocks) of the Late Ordovician Binghorn Group [32, 33], rather than, as initially suggested, extracts from Winnipeg shales [15, 16]. Oils of this family present diagnostic saturate fraction gas chromatograms (SFGC), low C₂₃tricyclic/C₃₀ pentacyclic terpane ratios (<0.20) and a strong C₃₄ hopane prominence. They can be further subdivided into a group distinguished by low Pr and Ph, relative to faster eluting n-alkanes nC₁₇ and nC₁₈, a strong odd-even predominance among n-alkanes between C₁₅ and C₂₀, and a low relative abundance in higher carbon number n-alkane homologues [34].

Family B oils primarily occur in Bakken reservoirs [32, 35, 33], they are however, also found in early Cretaceous reservoirs. They are sourced from Type II marine organic matter in the Upper Devonian-Mississippian Bakken Formation shale members. Main characteristic of this family is that it displays the highest Pr/Ph (>1.50) and C_{23}/C_{30} (>0.80) ratios, accompanying n-alkane and hopane profiles, without any predominance and prominence respectively.

Table 1 Table showing all groups and oil families, in correlation with the according formations, present in Williston Basin (modified by Osadetz, 1994)

Zu Williams, 1974 Lee	mberge, 1983; sheer and Zumberge, 1987	Osadetz et al., 1992, 1994	Source rocks
Type III	Not studied	Not studied	Tyler Fm. (Pennsyl.)
(Pennsylvanian oils)			
not studied		Family E	Exshaw/Bakken Fm. (U. DevMiss.)
		(Bakken oils)	
Type II	Group 2	Family B	Bakken Fm. (U.DevMiss.)
(Devonian, Mississippian & Mesozoic oils)	n (Mission Canyon oils)	(Bakken oils)	
		Family C	Lodgepole Fm. (L. Miss.)
		(Miss. & Jurassic oils)	
Not studied	Group 4	Family D	Winnipegosis Fm. (M.Dev.)
	(Nisku oils)	(Winnipegosis oils)	
	Group 3		
	(Duperow oils)		
Type 1	Group 1	Family A	Winnipeg Gr. (M. Ord.)
(Ordovician-Silurian oils) (Red River oils)	(Red River oils)	and Bighorn Gr. (U.Ord.)
	Group 5 (Cambrian oil)	Not studied	unknown (?U.CamOrd)

Family C oils occur the Mississippian Madison Group and Mesozoic formations and are sourced from Type II marine rocks in the Mississipian Lodgepole formation. They present high C_{23}/C_{30} (>0.20) ratio but, compared to Bakken sources, lower Pr/Ph ratio (<1.1), a pronounced (>n C_{20}) even n-alkane predominance and a strong C_{35} prominence.

Finally, Family D oils occur in Silurian to Mississippian sediments. They originate from Middle Devonian Winnipegosis Formation marine rocks, which vary in terms of depositional background. In particular, there are two kinds of settings; the platform depositional and starved basinal. Family D oils display similar terpane compositional characteristics to kukersite derived oils (abundant Pr, Ph and generally complex SFGCs), they differ however, in that they present greater relative acyclic isoprenoid and higher carbon n-alkane abundance. Oils of D Family, are further discretized into D_1 platformal and D_2 starved basinal, based on nC_{17}/Pr and nC18/Ph ratios. They display higher nC17/Pr ratios for a given nC18/Ph ratio compared to otherwise similar oils that occur in overlying Saskatchewan and Manitoba groups' strata, and they belong to the Elk Point Group of Winnipegosis reef formulations. Group D_1 predominantly occurs in younger Devonian reservoirs, lacking however, clear source definition. Suggested possible source rocks are thin organic-rich beds in Winnipegosis platform carbonates, the Birdbear Formation, and some Upper Devonian rocks. Group D_2 occurs in pinnacle reefs of the Middle Devonian Winnipegosis Formation and the Brightholme Member comprises the source rock. Oils having similar molecular compositions to D₂ oils have been found in the Upper Cambrian Deadwood Formation, Silurian pools of the Nesson Anticline, and new discoveries in the Middle Ordovician Winnipeg Formation. They have, however, very different isotopic compositions of carbon and sulphur, suggesting that a stillundescribed petroleum system exists in Paleozoic strata [36]. Family D oils correlate to Groups 3, 4, and 5 of Leenheer and Zumberge [31].

As previously mentioned, the compositional classification of the Williston Basin petroleum, relied much on terpane, sterane, and select n- and iso- alkane characteristics. The original classification by Williams [16], however, took into consideration the gasoline range fraction

(GRH) and later studies, based on that scheme, came to agree that families A - D and families B - C were inseparable and consistent with oil Types I and II [37]. Most recent work, depends on multivariate statistical methods, such as Principal Component Analysis (PCA), combined with geological information, in an attempt to enhance the independent interpretation of GRH and SFH fractions [32, 38]. Findings show that, while Family A oils can be uniquely classified, oils from Families B, C and D present insufficient characteristics for independent classification. Especially the composition of Family C seems to be quite heterogenous, often overlapping with families B and D [39].

This is attributed to the mixing of oils derived from different sources, without however, the extent to which this process occurs, having been defined [38]. A characteristic example of that mixing is the relative effectiveness of Bakken and Lodgepole petroleum systems [40, 41]. While part of the scientific community suggests that mixing is rare in the American portion of the basin [42, 43, 44], there is another part, proposing that major mixing is possible, without an impact on the biomarker traits [45, 46]. What is to account for the inability to precisely define the extend of mixing sources, is either the neglection of current interpretive techniques or the semi-quantitative confirmation of the biomarker based classification in the GRH and SFH [34, 39].

3. Exploratory Data Analysis

Analysis of Data (DA) constitutes the science of collecting, organizing and examining raw data under the purpose of obtaining useful and usable information for decision-making by users. The analysis may describe and summarize the data, identify relationships among variables, compare and identify differences between them as well as forecast outcomes. Data analytics is distinguished from data mining, which is a particular data analysis technique, by the scope, purpose and focus of the analysis. The target of Data Mining is rather predictive than descriptive. Data miners sort through huge data sets using sophisticated software to identify undiscovered patterns and establish hidden relationships. Data analytics focuses on inference, the process of deriving a conclusion based solely on what is already known by the researcher. Statistician John Turkey defined the term "Data Analysis" in 1961 as: "Procedures for analyzing data, techniques for interpreting the results of such procedures, ways of planning the gathering of data to make its analysis easier, more precise or more accurate, and all the machinery and results of (mathematical) statistics which apply to analyzing data."

Turkey [47] distinguished in Data Analysis techniques and procedures, two major groups: Exploratory Data Analysis (EDA) and Confirmatory Data Analysis (CDA). In EDA analysts make a few assumptions under the purpose of suggesting hypotheses and according to Turkey it is a rather detective work. In contrast, CDA "quantifies the extent to which deviations from a model could be expected to occur by chance" [48]. Confirmatory Data Analysis utilizes the traditional statistical tools of inference, significance, and confidence.

As a scientific tool, DA can be further subdivided in alternate groups. Therefore, based on the quantity of variables examined, Data Analysis can be dichotomized into Univariate (UDA) and Multivariate (MDA). Univariate data analysis is conducted when one variable is used for one observation. Subsequently, it makes sense to state that Multivariate data analysis is used when more than one outcome variables are measured and it is concerned with the study of association among sets of measurements. It is referred to as any statistical technique used to analyze data that arises from more than one variable.

3.1 Multivariate Data Analysis (MDA)

This project will focus on MDA techniques that will be implemented on the given data set and the outcomes will be examined thoroughly. Multivariate Data Analysis can fall into two phases: Unsupervised learning and Supervised learning. The goal of unsupervised learning is the detection of hidden structure in unlabeled data and encompasses many techniques that seek to summarize and explain key features of the data (i.e. Clustering Analysis, PCA). Supervised learning is a task of inferring a function from labeled training data. Each example on training data is a pair consisting of an input object (typically a vector) and a desired output value (i.e. Classification Analysis). In general, supervised methods are used when the aim is the construction of a model to be used to classify future samples [49].

There are several clustering techniques established by the scientific community, all governed by some kind of taxonomy [50, 51]. A major distinction among them involves the Hierarchical and the Partitional approaches, which are based on whether the set of produced clusters is nested or unnested. A Hierarchical clustering leads to a set of nested clusters that are organized as a tree, whereas a Partitional clustering formulates a division of the set of data objects into non-overlapping subsets (clusters) such that each data object is in exactly one subset. Characteristic examples of algorithms derived from the aforementioned approaches are agglomerative or divisive, deterministic or stochastic, incremental or non-incremental, monothetic or polythetic and hard or fuzzy [51].

3.1.1 Hierarchical Clustering

Hierarchical Clustering Analysis is an unsupervised technique that examines the interpoint distances between all of the data objects and generates a tree diagram or dendrogram on which, that information is visualized. It can be considered both as a sequence of nested partitions and the similarity levels at which these change [51, 52]. Hierarchical clustering algorithms are either bottom-up (agglomerative) or top-down (divisive). At each step of the agglomerative hierarchical approach each subject is treated as a singleton cluster which is successively merged into the closest cluster [51, 49, 53]. This process is repeated until all clusters have been merged into a singleton cluster that contains all subjects. The alternate divisive approach, begins with a single cluster containing all subjects, and at each step, the cluster splits until N clusters form (each with a single subject).

The criterion under which clusters are merged or split, differentiates at each case. Since the bottom-up approach agglomerates pairs of clusters with the minimum distance, measures of similarity and dissimilarity have to be taken into account. Those measures are defined by linkage functions which have a direct impact on the whole clustering procedure. They affect the way clusters are merged together and subsequently the final cluster solution. Therefore, linkage measures will be discussed extensively in the process.

The following notation is given in order for the various linkages to be described:

- Cluster r is formed from clusters p and q.
- n_r is the number of objects in cluster r
- x_{ri} is the ith object in cluster r

Single Linkage (Nearest Neighbor) functions utilize the shortest distance between any two objects in a pair of clusters [54, 55]:

$$d(r,s) = \min\left(dist(x_{ri}, x_{sj})\right), i \in (i, \dots, n_r), j \in (1, \dots, n_s)$$

The similarity under which two clusters merge is the similarity of their most similar objects and the merge criterion is local. Single linkage is a bottom-up (agglomerative) process where the number of clusters is reduced by one at each step.

Complete Linkage (Furthest Neighbor/Maximum Method) functions utilize the furthest distance between any two objects in a pair of clusters [55]:

$$d(r,s) = \max\left(dist(x_{ri}, x_{sj})\right), i \in (i, \dots, n_r), j \in (1, \dots, n_s)$$

Accordingly, in complete linkage method, the similarity under which two clusters fuse, is the similarity of their most dissimilar objects and the merge criterion is non-local, that is, the entire structure of clustering can affect the way how clusters fuse.

Average linkage functions utilize the averaged distance between all pairs of the two clusters' members [55]:

$$d(r,s) = \frac{1}{n_r n_s} \sum_{i=1}^{n_r} \sum_{j=1}^{n_s} dist(x_{ri} x_{sj})$$

There is also, an average linkage method within groups, proposed by Sokal & Michener [55], which takes into consideration the variability present within each cluster. This method will not be further discussed.

All the three methods mentioned above (single, complete and average) use a proximity matrix as input and the inter-cluster distances used are presented in Fig. 6.



Fig. 6 Single, Complete and Average linkage graphical representations, modified after [56].

Centroid linkage (Unweighted Pair-Group Method using the centroid approach- UPGMC) utilizes the Euclidean distance between the centroids of the two clusters:

$$d(r,s) = \|\widetilde{x_r} + \widetilde{x_s}\|_2$$

where $\widetilde{x_r} = \frac{1}{n_r} \sum_{i=1}^{n_r} x_{ri}$

As single linkage method, centroid linkage also represents an agglomerative approach to hierarchical clustering. This approach uses a data matrix, in contrast to the previous ones, rather than a proximity matrix and involves merging clusters with the most similar mean vectors.

Median linkage (Weighted Pair-Group Method using the centroid approach) functions also utilize the Euclidean distance between the weighted centroids of the two clusters:

$$d(r,s) = \|\widetilde{x_r} - \widetilde{x_s}\|_2$$

where $\widetilde{x_r}$ and $\widetilde{x_s}$ are weighted centroids for the clusters r and s. If cluster r was created by combining clusters p and q, $\widetilde{x_r}$ is defined recursively as:

$$\widetilde{x_r} = \frac{1}{2} \left(\widetilde{x_p} + \widetilde{x_q} \right)$$

Apart from the Euclidean distance, other proximity measures may be used for the Centroid and the Median linkage approaches, they would, however, lack interpretation in terms of the raw data [56]. The following table (Table 2) presents a brief description of various proximity measures used in linkages.

Table 2 Several Computational methods for distance

Distance measures	Formula	
Euclidean Distance	$ a - b _2 = \sqrt{\sum_{i=1}^n (a_i - b_i)^2}$	
Squared Euclidean Distance	$ a - b _2^2 = \sum_{i=1}^n (a_i - b_i)^2$	
Manhattan/City block DIstance	$ a - b _1 = \sum_{i=1}^{n} a_i - b_i $	
Maximum Distance	$ a-b _{\infty} = max a_i - b_i $	
Mahalanobis Distance	$\sqrt{(a-b)^T S^{-1}(a-b)}$ where S is the covariance matrix	

Ward's Method aims to minimize the variance between clusters by utilizing an incremental sum of squares: that is, the increase in the total within-cluster sum of squares as a result of joining two clusters [57]. The within-cluster sum of squares is defined as the sum of the squared distances between all objects in the cluster and the centroid of the cluster. The sum of squares measure is equivalent to the following distance measure d(r,s), which is the formula linkage:

$$d(r,s) = \sqrt{\frac{2n_r n_s}{(n_r + n_s)}} \|\widetilde{x_r} + \widetilde{x_s}\|_2$$

Where:

- $\|\widetilde{x_r} + \widetilde{x_s}\|_2$ is the Euclidean distance
- $\widetilde{x_r}$ and $\widetilde{x_s}$ are the centroids of the clusters r and s
- $\widetilde{n_r}$ and $\widetilde{n_s}$ are the number of elements in clusters r and s

In some references, factor of 2 multiplying $n_r n_s$ is not utilized by Ward's method. The linkage function uses this factor so that the distance between two singleton clusters is the same as the Euclidean distance. Ward's method differs from the centroid approach in clustering, in that centroids are weighted by $n_r n_s/(n_r + n_s)$ when computing distances between centroids, where n_r and n_s are the numbers of objects in the two clusters r and s.

Finally, *Weighted Average Linkage* (WPGMA) utilizes a recursive definition for the distance between two clusters [58]. If cluster r was created by combining clusters p and q, the distance between r and another cluster s is defined as the average of the distance between p and s and the distance between q and s:

$$d(r,s) = \frac{(d(p,s) + d(q,s))}{2}$$

There are several other hierarchical approaches, related to the ones described above. There is the Sum-of-Squares Approach [59, 60] which differs from Ward's method in that it is based on the sum of squares within each cluster rather than the increase in sum of squares in the merged cluster. Another flexible method defined by values of the parameters of a general recurrence formula has also been introduced by Lance and Williams [61] but in this project, it will not be discussed any further.

3.1.2 k - means Clustering

The k-means algorithm is one of the most used clustering algorithms and it was first described by Macqueen [62]. It was designed to cluster numerical data in which each cluster has a center called the mean. k-means belongs to the partitional (non-hierarchical) clustering methods [50], which are fundamentally different from the hierarchical ones. Partitional clustering methods generate a single partition of the data in an attempt to recover natural groups in the data. While hierarchical clustering methods require only the proximity matrix among the data points, partitional techniques expect the data in the form of a pattern matrix.

k-means [62] is one of the simplest unsupervised learning algorithms, which is used to solve the well-known clustering problem. The goal of k-means method is to divide the data into k distinct groups (clusters) so that observations within a group are similar, whilst observations between groups are different. The value of k (number of clusters) may or may not be specified. In most cases, it is assumed to be fixed. As an algorithm, it is rather iterative than hierarchical, which means that at each stage of the algorithm data points are assigned to a fixed number of clusters (whereas in hierarchical clustering, the number of clusters ranges from the number of data points down to a single cluster). The method allows the reallocation of data objects from one cluster to another, which is not the case at hierarchical clustering.

There is an error function behind this reallocation of data objects. It proceeds, for a given initial k clusters, by allocating the remaining data to the nearest clusters and then repeatedly changing the membership of the clusters according to the error function until the error function does not change significantly or the membership of the clusters no longer changes. The conventional k-means algorithm [63, 64] is briefly described below.

Let D be a data set with n instances, and let C_1, C_2, \ldots, C_k be the k disjoint clusters of D. Then the error function is defined as

$$E = \sum_{i=1}^{k} \sum_{x \in Ci} d(x, \mu(Ci)),$$

where μ (Ci) is the centroid of cluster Ci. d(**x**, μ (Ci)) denotes the distance between **x** and μ (Ci), and it can be one of the many distance measures, a typical choice of which is the Euclidean distance.

Given a set of observations, k-means clustering aims to partition n observations into k clusters so that the total distance between the group's members and its corresponding centroid, representative of the group, is minimized. The component to be minimized is the withincluster sum of squares (WCSS):

$$\sum_{j=1}^{k} \sum_{i=1}^{n} \left\| x_i^j - c_j \right\|^2$$

where the term $||x_i^j - c_j||^2$ provides the distance between any data point and the cluster's centroid.

Each cluster is associated with a centroid, which is the mean of the points in the cluster. Each point is assigned to the cluster with the closest centroid. The first step of k-means is to select as initial cluster centers K, randomly selected documents, the *seeds* (initialization phase). The algorithm then moves the cluster centers around in space in order to minimize WCSS (iteration phase). This is accomplished iteratively by repeating the following steps until a stopping criterion is met: reassigning documents to the cluster with the closest centroid; and recomputing each centroid based on the current members of its cluster. Firstly, WCSS decreases in the reassignment step, since each vector is assigned to the closest centroid, so the distance it contributes to WCSS decreases. Secondly, it decreases in the re-computation step because the new centroid is the vector \vec{v} for which WCSS_k reaches its minimum. Ultimately, k-means converges for the common similarity measures to a local minimum point after a finite number of iterations (normally the first few) [65]. Convergence and some probability properties regarding the k-means algorithm are also discussed in Pollard [66, 67], and Serinko & Babu, [68]. García-Escudero and Gordaliza [69] discussed the robustness properties of the k-means algorithm.

The complexity of the whole procedure is summarized in the following expression:

O(n*K*I*d)

Where: n= number of points

K= number of clusters

I= number of iterations

d= number of attributes

Choosing the right initial number of centroids is very important as it controls the performance of the algorithm. If there are K 'real' clusters (especially when K is large), then the probability of selecting one centroid from each cluster is relatively small. Particularly, if clusters are of the same size, n, then the aforementioned probability is as follows:

$$P = \frac{number \ of \ ways \ to \ select \ one \ centroid \ from \ each \ cluster}{number \ of \ ways \ to \ select \ K \ centroids} = \frac{K! \ n^K}{(Kn)^K} = \frac{K!}{K^K}$$

There are several approaches to this problem such as multiple runs, sampling and usage of hierarchical clustering to determine the initial centroid number, selection of more than k initial centroids and re-selection among these (the most widely separated), postprocessing and/or bisecting k-means. Some methods for selecting good initial centers are proposed in Babu and Murty [70] and Bradley and Fayyad [71]. Pena et al. [72] provide a comparison of four initialization methods: a random method, Forgy's approach [56], Macqueen's approach [62], and Kaufman's approach [73]. Other initialization methods are presented in Khan and Ahmad [74].

Silhouette analysis is a method for selecting the number of clusters for k-means clustering. It can be used as a tool to study the separation distance between the resulting clusters. The silhouette plot displays a measure of how close each point in one cluster is to points in the neighboring clusters and thus provides a way to assess parameters like number of clusters visually. This measure has a range of [-1, 1].

Silhouette coefficients (as these values are referred to as) near +1 indicate that the sample is far away from the neighboring clusters. A value of 0 indicates that the sample is on or very close to the decision boundary between two neighboring clusters and negative values indicate that those samples might have been assigned to the wrong cluster. In other words, a value of +1 is ideal and -1 is the least preferred. Hence, the higher the value, the better is the cluster configuration.

The silhouette value for the ith point, Si, is defined as

$$Si = (b_i - a_i) / max(a_i, b_i)$$

where a_i is the average distance from the i^{th} point to the other points in the same cluster as i, and bi is the minimum average distance from the i^{th} point to points in a different cluster, minimized over clusters.

A disadvantage of k-means algorithm is that it is sensitive to the presence of outliers and when clusters are of different size, different density or non-globular it might be disfunctional. For this reason, pretreatment and postprocessing of data is essential when implementing k-means, especially on high-dimensional data. Also, working only on numerical data restricts some applications of the k-means algorithm.

All in all, k-means is a greedy, computationally efficient technique, being the most popular representative-based clustering algorithm.

3.1.3 Principal Component Analysis (PCA)

Principal Component Analysis (PCA) constitutes a multivariate statistical technique, probably one of the most popular in the chemometric literature, used by various scientific disciplines, in order to identify patterns and relationships within a data set [75, 76]. It is an unsupervised learning method which aims to reduce the dimensionality of a high-dimensional data set consisting of a large number of interrelated variables and at the same time to retain as much as possible of the variation present in the data set. In mathematical terms, this is accomplished by manipulating a data matrix in such a way that the variation or spread of data objects (i.e. the description of their interpoint distances) is described by as few dimensions as possible. In addition to data reduction, Principal Component Analysis forms a transformation technique

of data, also used for simplification, modelling, outlier detection (identification of their class membership), variable selection, classification, prediction and unmixing of constant sum mixtures (curve resolution) [77, 78, 79].

The information that PCA extracts from the mathematical manipulation of the data matrix, is expressed by a new orthogonal set of variables (PC axes), known as the Principal Components (PCs) [76]. These are new variables that are uncorrelated and ordered such that the first few retain most of the variation present in all of the original variables. Principal components are obtained as linear combinations of the original variables and each one of them is characterized by certain properties. For example, the first PC contains the maximum amount of possible variance in the data set, in one direction and successive PCs describe decreasing amounts of variation. Each data object has coordinates, defined by the original variables, which are relative to the new principal component axes (scores). What is more, PC axes are influenced by variables and this is because the formulation of each axis is based on combinations among the original measurement variables. Variables' contribution to PC axes depends mainly on the relative orientation between those two elements. Hence, parallel arrangement (in space) of the variable and PC axes, means that minimum variation is contained in the PC and accordingly, orthogonal arrangement of the two, means maximum variation. Finally, the maximum PC quantity to be calculated, is at the same time, the minimum quantity of data objects or variables (six habits).

The PCs are defined as follows. Let $\mathbf{v} = (v_1, v_2, \dots, v_d)'$ be a vector of d random variables, where ' is the transpose operation. The first step is to find a linear function $\mathbf{a}'_{1\mathbf{v}}$ of the elements of \mathbf{v} that maximizes the variance, where \mathbf{a}_1 is a d-dimensional vector $(a_{11}, a_{12}, \dots, a_{1d})'$, so

$$\mathbf{a'}_1 v = \sum_{i=1}^d a_{1i} u_i$$

After finding $\mathbf{a}'_1 \mathbf{v}, \mathbf{a}'_2 \mathbf{v}, \ldots, \mathbf{a}'_{j-1} \mathbf{v}$, we look for a linear function $\mathbf{a}'_j \mathbf{v}$ that is uncorrelated with $\mathbf{a}'_1 \mathbf{v}, \mathbf{a}'_2 \mathbf{v}, \ldots, \mathbf{a}'_{j-1} \mathbf{v}$ and has maximum variance. Then we will find d such linear functions after d steps. The j th derived variable $\mathbf{a}'_j \mathbf{v}$ is the j th PC. In general, most of the variation in \mathbf{v} will be accounted for by the first few PCs. To find the form of the PCs, we need to know the covariance matrix Σ of \mathbf{v} . In most realistic cases, the covariance matrix Σ is unknown, and it will be replaced by a sample covariance matrix. For $j = 1, 2, \ldots, d$, it can be shown that the jth PC is given by $z_j = \mathbf{a}'_j \mathbf{v}$, where \mathbf{a}_j is an eigenvector of Σ corresponding to the jth largest eigenvalue λ_j .

4. Family Affiliations of Williston Basin Oils

The sample set under study consists of four compositional families, A, B, C and D, each containing 44, 11, 38 and 27 oil samples, respectively (a total of 120 oil samples – see Appendix). Family A oil samples belong to Red River and Yeoman formations. Family B oil samples belong to Bakken and Lodgepole formations while samples of family D belong to Winnipegosis formations. Oil samples of family C belong to various formations, such as Midale, Tilston, Bakken, Frobisher, Ratcliffe, Lodgepole, and Madison formations. The exploration of the compositional data was conducted on the main hydrocarbons of the gasoline range, the n-alkanes in the saturated fraction of the oils, as well as the biomarker's content of this sample set.

As far as the gasoline range is concerned, it represents the number of hydrocarbons containing less than twelve carbon atoms, and are often referred to as light hydrocarbons. In highly thermally mature oils, this range constitutes almost the 100% of the oil composition and therefore geochemical characterization of such oils is carried out based on these compounds.

The saturated fraction of hydrocarbons (SFH) is comprised of either the linear, branched or cyclic hydrocarbons. SFH contains the structural group of n-alkanes (usually between C_{12} - C_{35}) as well as the pristane (Pr) and phytane (Ph) isoprenoid compounds, measured in geochemical studies along with n-alkanes, due to their geochemical significance. In the analysis, the lighter n-alkanes were excluded and only the C_{13} - C_{32} alkanes were considered.

Biomarkers are a group of compounds, found in oils and rock extracts. They have a variety of applications in petroleum exploration. Such applications are in source-rock correlation and/or in the inference of characteristics of the source rock that generated an oil, without examining the source rock itself. Specifically, biomarkers in an oil can reveal the relative amount of oil-prone vs. gas-prone organic matter in the source kerogen, the age of the source rock, the environment of deposition, the lithology of the source rock (carbonate vs. shale), and the thermal maturity of the source rock during generation. Such data may be key inputs to effective basin modelling of a prospect or block. In this study, the sterane and hopane parts of the biomarkers' range have been examined thoroughly.

Before performing Multivariate Data Analysis (MDA) on the given oil sample set, an attempt was made in order to test the criteria under which the classification of the four family affiliations of Williston Basin, was determined in previous studies. The biomarker based classification of the four families relies on various compositional criteria, including Pr/Ph ratio, tricyclic to pentacyclic C_{23}/C_{30} ratio, n-alkane predominance and prominence amongst extended hopanes and many other, extensively described in the following paragraphs. Empty spaces on the barcharts presented below, correspond either to zero component values for specific samples, or to infinite numbers, generated during the calculative ratio calculations.

The compositional character of each family is unique and this is evident from their n-alkane distributions, biomarker signature as well as their gasoline range characteristics, in general [39, 32]. Family A oils display diagnostic saturate fraction gas chromatograms (SFGC) and are fairly distinguishable from the other families by their overall n-alkane profile (centered at C_{13} - C_{17}) and CPI values (average CPI: 1.59) [39, 32]. According to Obermajer et al., (2000), they

also present a smooth extended hopane profile with a steady decrease in the concentration of C_{31} + homologues with increasing carbon numbers [39]. In addition, Family A oils, are characterized by a C_{34} hopane prominence, according to Osadetz et al., [32]. Homohopane distributions have been used to distinguish oils from different organic facies of the same source rock. Such distributions are sensitive and may be altered due to various factors such as thermal maturity and API gravity. Judging from the barchart (Fig. 7), Family A oils display a high C_{34} homohopane distribution, but, in addition, oils from Family D, present an even stronger prominence on this compound (Fig. 7). The behavior of C_{34} for Families B and C is similar to that of Family A.



Fig. 7 C_{34} barchart for the whole sample set.

Another diagnostic feature of this group is its very low concentration of acyclic isoprenoids relative to n-alkanes, presenting the lowest Pr/C_{17} and Ph/nC_{18} ratios among all families [39]. The corresponding barcharts (Fig. 14, Fig. 15), in which these ratios have been plotted, is in agreement with this fact. According to Osadetz et al., [32], the C_{23} tricyclic/ C_{30} pentacyclic terpanes ratio, especially for Families A and B, is very distinct, differentiating them from the rest family groups. From the corresponding barchart, it is indeed observed that Family A oils display very low values of C_{23}/C_{30} , whereas Family B displays the highest peaks for the same ratio (Fig. 8). What is also noticeable from the C_{23}/C_{30} barchart, is that Family D oils, similarly to Family A, present very low values for this ratio.



Fig. 8 Barchart presenting C_{23}/C_{30} ratios for the whole sample set.

Family B oils, according to Obermajer et al., [39], differ from the rest in that they present a smooth n-alkane distribution with a maximum in the C_{13} - C_{17} range, lacking any homohopane prominence, which is in agreement with Osadetz et al., [32]. According to Obermajer et al., [39], there are variations in 17a(H)-trisnorhopane (Tm) over 18a(H)-trisnorhopane (Ts), compared to the rest oils. From the respective barchart (Fig. 10) we observe that there are indeed, intense variations within this Family affiliation, the density of the specimens, however, is not adequate enough in order to confirm the clear distinction of this family from the rest. The calculative process of the code has produced the NaN notation, resulting in non-plotted samples. The Ts/Tm ratio profiles of the rest Families (A,C and D) show almost equivalent variations.

Another characteristic of Family B oils, is that they obtain values above unity for the Pr/Ph ratio [32]. This ratio is one of the most common correlation parameters, utilized as an indicator of depositional environment [80]. Variations may reflect multiple degrees of oxidation during the early stages of chlorophyll degradation. It is one of the most commonly utilized correlation parameters, indicative of the source rock's depositional environment [80]. Being sensitive to diagenetic conditions, values of Pr/Ph ratios substantially below unity are considered to indicative of petroleum origin and/or highly reducing depositional environments. Very high Pr/Ph ratios (> 3) reflect source material of terrestrial origin. Pr/Ph ratios ranging between 1-3 reflect oxidizing depositional environments [81]. According to Lijmbach [82] low Pr/Ph values (<2) reflect aquatic depositional environments including marine, fresh and brackish water (reducing conditions), intermediate values (2-4) reflect fluviomarine and coastal swamp environments, whereas very high values (up to 10) are related to peat swamp depositional environments (oxidizing conditions). From the corresponding barchart (Fig. 9), we observe that, contrary to Family C, Families A, B and D present similar, above unity values for this ratio, which is in agreement with Osadetz et al., [32]. At the same time, however, Family B oils display the highest peaks (Fig. 9).



Fig. 9 Pr/Ph ratios barchart for the whole sample set.



Fig. 10 Ts/Tm ratios barchart for the whole sample set.



Fig. 11 CPI profile for the whole sample set.



Fig. 12 Odd/Even predominance for the whole sample set.



Fig. 13 C_{35} barchart for the whole sample set.

Main characteristic of Family C is the strong C_{35} prominence [32], which is confirmed by the respective barchart (Fig. 13). The lowest C_{35} homohopane distribution is indicative of Family A oils, as shown. According to Osadetz et al., [32] and Obermajer et al., [39], these oils obtain lower Pr/Ph values in comparison to the rest, and in particular, less than unity. This fact holds true, as we observe from the corresponding barchart (Fig. 9), confirming at the same time that Family C oils display a strong and consistent predominance of Ph/Pr ratio.

Additionally, oils of this familial group also display an even/odd n-alkane predominance [32]. The composition and distribution of n-alkanes carbon numbers reflect the source of kerogenic organic matter, sedimentary environment, and maturity of the rocks. Traditional geochemists feel that the odd/even carbon number predominance of n-alkane decreases as rocks mature. The OEP (odd/even predominance) of mature source rocks is close to 1. However, the odd carbon number predominance appears in Upper Ordovician source rocks, and an even carbon number predominance is found in Cambrian - Lower Ordovician source rocks. Family C oils are characterized by an even/odd n-alkanes predominance and this is confirmed by both the CPI and OEP, respective barcharts (Fig. 11, Fig. 12).



Fig. 14 nC_{17}/Pr barchart for the whole sample set.



Fig. 15 nC_{18} /Ph barchart for the whole sample set.

Oils from Family D display a distinctive stratigraphic occurrence and have been subdivided into two separate groups D_1 and D_2 , based on nC_{17}/Pr and C_{18}/Ph ratios [32]. The corresponding barcharts (Fig. 14, Fig. 15) present the distributions of these ratios amongst the whole sample set. What is more, D_1 and D_2 oils, depending on the pools they occur, either in Madison or Birdbear, they display $Pr/Ph \le 1.0$ and Pr/Ph > 1.1, respectively. This is indeed, evident, from the corresponding barchart (Fig. 9).

Reviewing the barcharts presented before, it would be important to state that based on individual geochemical characteristics, the four families can be indeed uniquely identified at a great extent. However, it would be a challenge to investigate if a clear classification can be obtained, by applying this time, multivariate data analysis (MDA) on raw data.

In the next chapters, we implement several multivariate methods on the given data set and examine the results, that each method produces. Hierarchical clustering, k-means and Principal Component Analysis are applied on four independent models that were developed for this purpose; the Saturated Fraction Compositional Model, the Saturated Fraction Ratios Model, the Gasoline Range Compositional Model and the Biomarkers Compositional Model. All of the steps that were followed are extensively described.

5. Application of MDA methods; inputs and results

The core of this project is the investigation of the oil-oil correlations among compositional data of a sample set from Williston Basin, by using multivariate statistical analysis methods. Oil-oil correlations are based on compositional criteria and examine whether a genetic relationship exists among a group of oil samples. In particular, Hierarchical Clustering, k-means and PCA have been employed in order to explore compositional data from the gasoline range (GRH), saturated hydrocarbons (SFH) and biomarker traits of 120 oil samples from the Williston Basin Petroleum province. The samples examined in this study are from four, previously defined, compositional families (A-D) [34].

For the application of MDA methods on the sample set, a MATLAB code created in the "Hydrocarbons Chemistry and Technology Research Unit", of the School of Mineral Resources of the TUC, was utilized. All necessary adjustments and modifications were applied in order for the code to work.

From the sample set under study, four independent models were developed in order to explore different compositional information. The models used for the identification of petroleum systems were: a) Saturated Fraction Compositional Model (SFCM) b) Saturated Fraction Ratios Model (SFRM), c) Gasoline Range Compositional Model (GRCM) and d) Biomarkers Compositional Model (BCM). SFCM embodies original variables derived from the gas chromatographic analysis of the Saturated Fraction of Hydrocarbons (SFH). It takes into account peak areas of n-alkanes, nC₁₃-nC₂₄, pristane (Pr) and phytane (Ph). The SFRM contains the most commonly utilized compositional ratios and factors derived from the gas chromatographic analysis of the SFH (Pr/Ph, n-C₁₇/Pr, n-C₁₈/Ph, CPI n-C₁₄₋₂₀, CPI n-C₂₂₋₃₂). GRCM includes variables derived solely from GRH compositional data. The parameters reflect internal variations for compounds with the same number of carbon atoms to minimize possible variations due to sample handling and experimental conditions. Finally, BCM contains all variables derived from biomarkers' traits of the oil sample set.

The approach under which all statistical methods were applied, was that of trial and error in order to achieve a "clear clustering" (if possible) of the oil samples. A pretreatment scheme of the sample set was considered necessary in order to reformat the original data file and prepare data for clustering. This is because the data set consists of peak areas that are analysis-dependent. As a consequence, only by preprocessing the data, we get meaningful statistical results, since all components are put under the same scale. The idea is that if different components of data (features) have different scales, then derivatives tend to align along directions with higher variance, which leads to poorer/slower convergence. The chemometric software package that was utilized, offered various pretreatment options, all of which were originally applied on the sample set, in order to examine which one produces the best classifying solution. While only the results from one preprocessing option will be presented, all pretreatment schemes which were utilized, are briefly described below.

Command "pre_scaling_0_1": It refers to the subtraction of the minimum value and the division of each column by the range. The results of this pretreatment scheme are going to be presented in the upcoming chapters.

Command "norm_variables_0_1": It refers to the subtraction of the minimum value and the division of each variable by the range.

Command "pre_minusMean": It concerns the subtraction of the mean value from each variable.

Command "pre_PQN" (Probabilistic quotient normalization): It refers to the division of each sample with the sum of the sample's variables. The calculative process takes into consideration the median value of each column.

Command "pre_CLR" (Centered log-ratio normalization): It concerns the division of each sample with the sum of the sample's variables. It differs, however, from "pre_PQN" in that it takes into consideration the geometric mean of each column.

Command "Subtract_sample_min": It refers to the subtraction from each sample of its minimum value.

Command "pre_TSN" (total sum normalization): It concerns the division of each sample with the sum of the samples' variables.

Command "pre_max": This matlab command refers to the division of each sample with the maximum value of the samples' variables.

5.1 Saturated Fraction Compositional Model (SFCM)

5.1.1 Hierarchical Clustering on SFCM

The subtraction of the minimum value from the subset and division of each variable by the range ("pre_scaling_0_1" command) resulted in the following dendrogram (Fig. 16). Average linkage with a correlation coefficient were combined

It is evident that the oil samples from all four family affiliations overlap, presenting no clear distinction. In particular, there is a slight overlap of samples from Families B (B1014, B1993, B2121, B2179, B1879, B1874) and D (D1275, D1276, D1289, D1313, D1288, D1290, D1291) with Family A. The original clustering solution detected outlier values (samples C599, D2595 and C566), removing which from the sample set and reprocessing it under the same pretreatment, made no difference on the clustering solution.



Fig. 16 Resulting Dendrogram under the command "pre_scaling_0_1" for the Saturated fraction compositional model (SFCM).

As observed, the algorithm failed to discriminate distinct familial affiliations among the given oil sample set, under this pretreatment scheme. In order to test how Hierarchical Clustering would offer the best clustering solution, many other pretreatment schemes were also applied on the data set and as a procedure, this was also followed in the upcoming MDA methods. The following dendrogram resulted from the division of each sample with the sum of the samples' variable - Total Sum Normalization ("pre_TSN" command of the chemometric software package). City block distance and Centroid linkage were combined and the produced dendrogram displays a relatively good distinction of Family A. It fails, however, to distinguish amongst Families B, C and D, which, once again, overlap one another (Fig. 17).



Fig. 17 Dendrogram under the "pre_TSN" command for the Saturated fraction compositional model (SFCM).

An interesting feature of the dendrogram in Fig. 17, is that it displays a non-monotonic tree. This occurs when the distance from the union of two clusters, r and s, to a third cluster is less than the distance between r and s. In this case, in a dendrogram drawn with the default orientation, the path from a leaf to the root node takes some downward steps. Usually, the centroid and median methods (as in this case) can produce a cluster tree that is not monotonic and if this happens, it is better to utilize another linkage method. In our case, however (Fig. 17), the centroid linkage, which was automatically chosen by the chemometric software package, produced a dendrogram which classified sufficiently samples of Family A. All other pretreatment options failed in this task significantly.

5.1.2 k – means algorithm on SFCM

k-means clustering was then performed under the same pretreatment option ("pre_scaling_0_1") resulting in the following features (Fig. 18, Fig. 19, Table 3).

Table 3 Summary of k-means clustering under the "pre_scaling_0_1" pretreatment option for the Saturated fraction compositional model (SFCM).

K-values	Best distances sums	Average silhuette values
K=2	594.077	0,566689
K=3	41.787	0,539449
K=4	345.548	0,503212
K=5	290.114	0,467112

The silhouette plots for K=2, K=3, K=4 and K=5 clusters are shown in the following figure (Fig. 18). An insufficient choice of an initial K value would result in clusters below average silhouette scores or even wide fluctuations in the size of the silhouette plots. This is the criteria under which, each clustering solution is evaluated as sufficient or insufficient.



Fig. 18 Silhouette plots for k=2, k=3, k=4 and k=5 clusters under the "pre_scaling_0_1" pretreatment option for the Saturated fraction compositional model (SFCM).

From the silhouette plots (Fig. 18), we observe that, in general, the obtained silhouette values fall in the range of 0.1-0.9. The size of the silhouette plots does not present wide fluctuations for each case, and negative values are present in all clustering solutions. The two - cluster solution has an average silhouette value of 0.566689, being the highest amongst the others (Table 3). This is an indication that grouping into two clusters using k-means is more efficient compared to grouping into three, four or five clusters. It is not, however, sufficient enough, as we would expect, grouping into four clusters to be the best solution. In Fig. 20 we can observe, which cluster each sample is assigned to.


Fig. 19 The plot of k-means clustering for k=2 under the "pre_scaling_0_1" pretreatment option for the Saturated fraction compositional model (SFCM). The \otimes symbol represents the centroid of each cluster.

Fig. 19 represents the plot of k-means clustering, for the case of k = 2. Taking into consideration the average silhouette value, k=2 is the most efficient clustering solution. However, by observing the plot, we could say that there is no clear boundary between the two clusters and samples overlap with each other.



Fig. 20 Table displaying to which cluster each sample belongs, for each K value of the SFCM (idx2 = k:2, idx3 = k:3, etc.)

Taking into consideration that the most sufficient clustering solution is that of k=2 (idx=2) and according to Fig. 20, all samples from Family A oils are assigned to one cluster. The vast majority of Family C oil samples are assigned to a different cluster with a few exemptions (C540, C543). The discretization of these two families is relatively sufficient, but as far as Family D and B oil samples are concerned, they overlap with A and C considerably, as samples from both families are assigned to both clusters.

5.1.3 Principal Component Analysis on SFCM

Sample scores describe a position in principal component space, and each original variable has loadings that describe their contribution to each principal component. The sample score of the first two principal components and the respective loading diagrams are presented in figure (Fig. 21a, b). The percentages of variation attributed to each of the Principal Components are shown in Fig. 21c.



Fig. 21 a) Sample scores for the first to Principal Components resulting from the Saturated Fraction Compositional Model (SFCM) of selected Williston Basin petroleum oils. Colors on sample symbols indicate compositional families determined by independent analysis. Blue color represents oils of Family A, green applies for Family B oils, red for Family C oils and yellow for Family D oils. "Pre_scaling_0_1" command was used on the data set. b) Original Variable loadings for the first to Principal Components resulting from the Saturated Fraction Compositional Model (SFCM) of selected Williston Basin petroleum oils. c) Percentage of variance explained by each Principal Component.

SFCM sample scores of the first two PCs explain almost 85 per cent of the variance (Fig. 21c). There are linear gradients observed in the data by comparing the sample scores of the first two principal components of the SFCM. These gradients indicate that both distinctive family characteristics and linear compositional variations of the original variables exist within each family. Sample scores of Family A oils exhibit the most coherent grouping and are characterized by a positive gradient defined by positive PC1 scores and positive PC2 scores. Family C oils are also characterized by a positive gradient whereas Family B and D are defined by a negative gradient. The mild gradients of Families B, C and D exhibit positive PC1 scores, as Family A, but negative PC2 scores. There is a considerable overlap of Family B with Family D and a slight overlap of Family C with Family D. What is more, for a given value of PC1 Families C and D have more negative PC2 scores but this is not enough to be uniquely distinguishable.

As far as variable loadings are concerned, they are a tool used for the understanding of the role and importance of the original variables. The original variable loadings for the SFCM distinguish between a preponderance of lighter versus heavier n-alkanes (Fig. 21b). C_{13} - C_{17}

alkanes are characterized by strongly positive PC1 and PC2 loadings but C_{16} and C_{17} exhibit negative PC3 values. Probably all these variable loadings control the gradients that separates independently defined oil families. The variable with the higher weight (0.1034) among the 22 variables of the SFCM, is alkane C_{13} with strongly positive PC1, PC2 and PC3 loadings.

5.1.4 Discussion on the performance of MDA on the SFCM

To summarize, Hierarchical Clustering, k-means and Principal Component analysis were applied on the Saturated Fraction Component Model. Both in Hierarchical Clustering and PCA, Family A oils presented the most coherent group, being sufficiently separated from the rest familial affiliations. Families B and D overlapped significantly while also in both cases there appeared a slight overlap between families C and D. The method which completely failed to distinguish among the four oil families (A, B, C and D) was k – means clustering. The clustering solution produced only two clusters and according to which cluster each sample was assigned, k-means presents only 25% of success. Out of the three statistical methods, k-means was the one to produce the most insufficient results.

5.2 Saturated Fraction Ratios Model (SFRM)

5.2.1 Hierarchical Clustering on SFRM

The following dendrogram is the outcome of the "pre_scaling_0_1" command (Fig. 22). Average linkage along with Euclidean distance as a measure of proximity, were combined.

Family A is clearly distinguished from the rest. Family C considerably overlaps with Families B and D. All pretreatment schemes that were applied on the data set, behaved similarly producing almost the same results when Hierarchical Clustering was performed; all distinguished Family A quite sufficiently, but exhibited a slight overlap amongst Families B, C and D.



Fig. 22 Resulting Dendrogram under the command "pre_scaling_0_1" for the Saturated fraction ratios model (SFRM).

5.2.2 k – means algorithm on SFRM

Under the same pretreatment scheme ("pre_scaling_0_1" command), k-means algorithm was applied and below we present the results.

Table 4 Summary of k-means clustering under the "pre_scaling_0_1" pretreatment option on the Saturated Fraction Ratios Model (SFRM).

K-values	Best distances sums	Average silhuette values
K=2	184.117	0,727318
K=3	115.198	0,691804
K=4	832.004 831.205	0,715499
K=5	649.877	0,702406

The silhouette plots for K=2, K=3, K=4 and K=5 clusters are presented in the following figure (Fig. 23).



Fig. 23 Silhouette plots for k=2, k=3, k=4 and k=5 clusters under the "pre_scaling_0_1" pretreatment option for the Saturated Fraction Ratios Model (SFRM).

From the silhouette plots (Fig. 23), we observe that in all cases we obtain silhouette values above 0.6 and negative silhouette coefficients are always present. Average silhouette values are similar for all clustering solutions, with a maximum of 0,727318 for K=2 (Table 4). This is an indication that under the "pre_scaling_0_1" pretreatment scheme, grouping into two clusters using k-means is more efficient compared to grouping into three, four or five clusters. In Fig. 25 we can observe, which cluster each sample is assigned to.

In Fig. 24 the plot of k-means clustering, for the case of k = 2 is presented with different colors for sample members that belong to different clusters.



Fig. 24 The plot of k-means clustering for k=2, of the Saturated Fraction Ratios Model (SFRM). The \otimes symbol represents the centroid of each cluster.



Fig. 25 Table displaying to which cluster each sample belongs, for each K value of the SFRM (idx2 = k:2, idx3 = k:3, etc.)

Based on the average silhouette values, the most efficient clustering solution is that of k=2 (idx=2). According to Fig. 25, all samples from Family A oils are assigned to cluster one. Almost all of Family C oil samples are assigned to cluster two (only sample C1705 is assigned to cluster 1). Oil samples from family B are all assigned to cluster 2, whereas family D oil samples are assigned in both clusters.

5.2.3 Principal Component Analysis on SFRM

The original variables used in the Saturate Fraction Ratios Model (SFRM) include the compositional factors Pr/Ph, nC_{17}/Pr , nC_{18}/Ph and the carbon preference indices for both lighter (nC_{14} - nC_{20}) and heavier (nC_{22} - nC_{30}) alkanes of the saturated fraction hydrocarbons. The

sample scores of the first two principal components and the respective loading diagrams are presented in Fig. 26. The percentages of variation attributed to each of the Principal Components are shown in Fig. 26c.



Fig. 26 a) Sample scores for the first to Principal Components resulting from the Saturated Fraction Ratios Model (SFRM) of selected Williston Basin petroleum oils. Colors on sample symbols indicate compositional families determined by independent analysis. Blue color represents oils of Family A, green applies for Family B oils, red for Family C oils and yellow for Family D oils. "Pre_scaling_0_1" command was used on the data set. b) Original Variable loadings for the first to Principal Components resulting from the Saturated Fraction Ratios Model (SFRM) of selected Williston Basin petroleum oils. c) Percentage of variance explained by each Principal Component.

SFRM sample scores of the first two PCs explain 83 per cent of the variance (Fig. 26c). There are two linear gradients observed in the data by comparing the sample scores of the first two principal components of the SFRM; a dispersed positive gradient displayed by samples with positive PC1 scores and PC2 scores less than 0, and a relatively tight negative gradient consisting of both positive PC1 and PC2 scores. As in the SFCM, these gradients also indicate that distinctive family characteristics and linear compositional variations of the original variables exist within each family. When the samples are compared to the biomarker-based oil families, Family A is once again clearly distinguished by consistently positive PC1 and PC2 scores and a linear variation between them. Only sample D2626 overlaps with this group, however. Family D oils are also characterized by a general positive gradient, while Families B and D are defined by mainly a positive PC2 scores. As in the SFCM, the fields of PC1 and PC2 in Families B, C and D exhibit negative PC2 scores. As in the SFCM, the fields of PC1 and PC2 in Family C overlap those of Families B and D, effectively obscuring their separation. However, Family C samples appear to fall along a positively correlated gradient in PC1 vs PC2 space.

The original variable loadings for the SFRM indicate a lack of discriminating power of the nC_{17}/Pr and nC_{18}/Ph with respect to Families B and C, which opposes to Osadezt et al. [32], who claim that this biomarker parameter is highly effective as far as the discrimination among these affiliations is concerned.

5.2.4 Discussion on the performance of MDA on the SFRM

MDA methods on the Saturated Fraction Ratios Model seemed to perform in a similar manner as in the Saturated Fraction Compositional Model. In all three methods Family A was significantly distinguished in contrast to the rest familial affiliations. Only sample D2626, in PCA overlapped with family A samples. As far as k-means is concerned, even though it discretizes family A, as a whole, it failed in considerably in separating families B, C and D. It produced a two-cluster solution.

5.3 Gasoline Range Compositional Model (GRCM)

5.3.1 Hierarchical Clustering on GRCM

Applying the Hierarchical Clustering algorithm on GRCM, produced the following dendrogram (Fig. 27). Single linkage with Euclidean distance were combined this time.

From the figure, we notice that oil samples from all four family affiliations overlap, presenting no clear distinction. In this case, we also observe that a few samples from C and D are excluded from the clustering solution (samples B1873, B1874, B1014, C1390, and D842).



Fig. 27 Resulting Dendrogram under the command "pre_scaling_0_1" for the Gasoline range compositional model (GRCM).

These components presented zero values for all variables. To examine how the model would perform without these values, they were removed from the data set and then hierarchical clustering was implemented again. The following dendrogram is the result.



Fig. 28 Resulting Dendrogram under the command "pre_scaling_0_1" for the Gasoline range compositional model (GRCM) after removing zero values.

Implementing the algorithm produced another two outlier samples from family D (samples D1312 and D2885). Family A oil samples, however, seem to distinguish from the rest, but not sufficiently enough, as there is a slight overlap with samples from family D. As far as families B, C and D are concerned, there is a considerable overlap among them.

5.3.2 k-means algorithm on GRCM

Implementing the k-means algorithm on the Gasoline range compositional model produced the following results. Components with zero values (as mentioned before) were kept out of the analysis.

Table 5 Summary of k-means clustering under the "pre_sc	caling_0_1" pretreatment option on the Gasoline Range
Compositional Model (GRCM).	

K-values	Best distances sums	Average silhuette values
K=2	60,1718	0.4438
K=3	49,6733	0.4572
K=4	43,5093	0.4510
K=5	38,3703	0.4271

The silhouette plots for K=2, K=3, K=4 and K=5 clusters are presented in the following figure (Fig. 23).



Fig. 29 Silhouette plots for k=2, k=3, k=4 and k=5 clusters under the "pre_scaling_0_1" pretreatment option for the Gasoline Range Compositional Model (GRCM).

From the silhouette plots (Fig. 29), we observe that generally we obtain silhouette values in the range of 0.01-0.8. Negative silhouette coefficients are present in all cases. Average silhouette values are close for all clustering solutions, with a maximum of 0.4572 for K=3 (Table 5). The outcome of this analysis, infers that grouping into three clusters using k-means is more efficient compared to grouping into two, four or five clusters. In Fig. 31 we can observe, which cluster each sample is assigned to.

In Fig. 30 we observe the clustering solution of k-means for k=3. The figure shows the three clusters along with their centroids.



Fig. 30 Plot of k-means clustering for k=3, of the Gasoline Range Compositional Model (GRCM). The \otimes symbol represents the centroid of each cluster.



Fig. 31 Table displaying to which cluster each sample belongs, for each K value of the GRCM (idx2 = k:2, idx3 = k:3, etc.)

Even though the three-cluster solution seems to be the most efficient out of the analysis, from the plot we observe that the clusters present no clear boundaries from one another. The overlapping among samples is evident. Fig. 31 confirms this fact as it presents in which of the three clusters, each sample is assigned to.

5.3.3 Principal Component Analysis on GRCM

The sample scores of the first two principal components and the respective loading diagrams are presented in Fig. 32. The percentages of variation attributed to each of the Principal Components are shown in Fig. 32c.



Fig. 32 a) Sample scores for the first to Principal Components resulting from the Gasoline Range Compositional Model (GRCM) of selected Williston Basin petroleum oils. Colors on sample symbols indicate compositional families determined by independent analysis. Blue color represents oils of Family A, green applies for Family B oils, red for Family C oils and yellow for Family D oils. "Pre_scaling_0_1" command was used on the data set. b) Original Variable loadings for the first to Principal Components resulting from the Gasoline Range Compositional Model (GRCM) of selected Williston Basin petroleum oils. c) Percentage of variance explained by each Principal Component. GRCM sample scores of the first two PCs explain 82 per cent of the variance (Fig. 32c). There is generally one linear gradient observed in the data by comparing the sample scores of the first two principal components of the GRCM; a dispersed negative gradient displayed by samples with positive PC1 scores and PC2 scores both positive and negative. Family A is once again clearly distinguished by consistently positive PC1 and PC2 scores and a linear variation between them. The gradient exhibits high positive PC1 scores but for Families B, C and D exhibits also negative PC2 scores. The gradients of Families B, C and D overlap each other's scores resulting in the obscureness of their separation.

The variable loadings for the GRCM indicate that PC1 is controlled strongly by loadings attributed to the relative concentration of n-alkanes and branched and cyclic alkanes. High negative PC1 loadings are characteristic of the GRH n-alkanes, while the cyclic and branched alkanes with 6 to 8 carbon atoms are characterized by strong positive values. In our case the GRCM fails in the task of classifying the four family affiliations.

5.3.4 Discussion on the performance of MDA on the GRCM

Although in several studies (e.g. [38]) the Gasoline Range Compositional Model appears to be successful in classifying efficiently oil samples of the four family affiliations recognized in Williston Basin, in our case it substantially fails. All statistical methods that were implemented on this model, classified relatively sufficiently only family A. Families B, C and D presented a significant overlap, both one to another, but also with Family A. This is evident from the dendrogram of Fig. 28 as well as from Fig. 32a. The overlapping of oil families is incredibly apparent in the k-means plot (Fig. 30), where there is no distinct cluster.

5.4 Biomarkers Compositional Model (BCM)

The biomarkers of the given sample set were examined in multiple ways; firstly as a whole and secondly in their separate parts of steranes and hopanes. The results that each model produced were similar, as far as the classification of oil families, is concerned. For this reason, only the results from BCM will be presented in the upcoming paragraphs, as the most characteristic.

5.4.1 Hierarchical Clustering on BCM

Applying the Hierarchical Clustering algorithm on BCM, produced the following dendrogram (Fig. 27). Average linkage with Euclidean distance were combined this time.



Fig. 33 Resulting Dendrogram under the command "pre_scaling_0_1" for the Biomarkers compositional model (BCM).

Hierarchical clustering on BCM seems to separate relatively well Family C oils. Only sample B1443 (of Family B) overlaps with family C. The dendrogram illustrates an overlapping of Family D with Family A and the rest of Family B samples form a small group which is interrupted by sample D2885.

5.4.2 k-means algorithm on BCM

Implementing the k-means algorithm on the Biomarkers compositional model produced the following results.

Table 6 Summary of k-means clustering under the "pre_scaling_0_1" pretreatment option on the Gasoline Range

Compositi	onal Model (GRC	CM).	
	K-values	Best distances sums	Average silhuette values

K-values	Best distances sums	Average silhuette values
K=2	250.993	0.5503
K=3	15.438	0.6665
K=4	131.334	0.5865
K=5	113.741	0.5425

The silhouette plots for K=2, K=3, K=4 and K=5 clusters are presented in the following figure (Fig. 23).

From the silhouette plots (Fig. 34), we observe that generally the highest silhouette values we obtain almost reach the value of 0.9. Fluctuations in the width of clusters is present in all cases and so are negative silhouette coefficients. Average silhouette values fall in the range of 0.5425-0.665, with 0.665 being the maximum for K=3 (Table 6). The outcome of this analysis, infers that grouping into three clusters using k-means is more efficient compared to grouping into two, four or five clusters. In Fig. 36 we can observe, which cluster each sample is assigned to.



Fig. 34 Silhouette plots for k=2, k=3, k=4 and k=5 clusters under the "pre_scaling_0_1" pretreatment option for the Biomarkers Compositional Model (BCM).

Fig. 35 illustrates the clustering solution of k-means for k=3. The figure shows the three clusters along with their centroids.



Fig. 35 Plot of k-means clustering for k=3, of the Biomarkers Compositional Model (**B**CM). The \otimes symbol represents the centroid of each cluster.



Fig. 36 Table displaying to which cluster each sample belongs, for each K value of the BCM (idx2 = k:2, idx3 = k:3, etc.)

Fig. 35 illustrates the three-cluster solution that silhouette analysis produced as the most efficient. The clusters do not exhibit clear boundaries and overlapping is evident. Fig. 36 supports the overlapping fact as it illustrates in detail in which cluster each sample is assigned to.

5.4.3 Principal Component Analysis on BCM

The sample scores of the first two principal components and the respective loading diagrams are presented in Fig. 32. The percentages of variation attributed to each of the Principal Components are shown in Fig. 32c.

BCM sample scores of the first two PCs explain almost 90 per cent of the variance (Fig. 37c). By comparing the sample scores of the first two principal components of the BCM, we observe no clear distinction among families. All families exhibit high positive PC1 scores and all of them present both negative and positive PC2 scores. Family B (green symbols on the PC plot) exhibit solely negative PC2 scores. Family A overlaps here mainly with family D and a few samples of family D overlap with family C. Scores of family B are quite dispersed in the plot.



Fig. 37 a) Sample scores for the first to Principal Components resulting from the Biomarkers Compositional Model (BCM) of selected Williston Basin petroleum oils. Colors on sample symbols indicate compositional families determined by independent analysis. Blue color represents oils of Family A, green applies for Family B oils, red for Family C oils and yellow for Family D oils. "Pre_scaling_0_1" command was used on the data set. b) Original Variable loadings for the first to Principal Components resulting from the Biomarkers Compositional Model (BCM) of selected Williston Basin petroleum oils. c) Percentage of variance explained by each Principal Component.

5.4.4 Discussion on the performance of MDA on the BCM

The Biomarkers Compositional Model appears to be successful in classifying relatively well oil family C. All methods produced similar results as far as this classification pattern is concerned. Families A and D overlap significantly, while family B overlaps slightly with family D.

All in all, the performance of MDA methods was insufficient, failing in all models to classify the samples into four familial affiliations. Based on common compositional information, it seems that unsupervised methods fail to cluster these oils. They cannot be implemented blindly without additional information. For this reason, in the next chapter we examine the compositional character of the given data set in an alternative approach.

6. Compositional Data

As discussed in the previous chapters, MDA methods failed in the task of classifying the data set into distinct oil family affiliations. This applies to all the compositional models and is probably attributed to the nature of the data, which fall into a special category of data; the Compositional Data. The Saturates Fraction Ratios Model is excluded from this category and none of the following information concerns this model.

Compositional Data (CoDa) are a type of multivariate data, the components of which represent proportions or fractions of a whole. Such data come in a closed form, meaning that they sum to a constant value (e.g. one if measured in parts per unit or 100 if measured in percentages). However, the term Compositional Data, covers all those vectors representing parts of a whole which only carry information on the relative (and not the absolute) frequencies, with which different and positive components occur.

Typical examples of Coda are geochemical elements in geology, data corresponding to categories of sedimentary particle-size distributions, proportions of fossil species in two or more assemblages, body composition (fat, sugar, etc.) in medicine, nutrient-balance ionomics (measurement of the total elemental composition of an organism to address biological problems) in agriculture, genotype frequency in genetics, chemical compositions in chemistry, and many more other. This type of data is generally widespread in disciplines supporting modeling, classification or discrimination and is characterized by specific numerical properties that have significant consequences for any statistical analysis [85] [86] [87] [88] [89] [90] [91]. Their fundamental properties are briefly reviewed in the upcoming paragraphs.

6.1 The Constant Sum Constraint (CSC) – Impacts on the Analysis

As mentioned before, Compositional Data only convey relative information as they represent part of a whole, and their unique properties are a corollary of this fact. They concern data consisting of vectors of always positive components, often subject to a constant (unit-) sum constraint; they must sum to one because they are proportions. Their main difference to unconstrained variables is that they are never free to vary independently, which in turn imposes constraints upon their variance-covariance structure (Aitchison 1986, chapter 3). The constant sum constraint forces at least one of their covariance to obtain a negative value. The result is at least one correlation or coefficient between elements, is also negative. This is explained as a consequence of the Euclidean Foundation of classical statistics, where the scale is absolute and not relative.

In particular, for a D-part composition [x_1, \ldots, x_D] with the component sum x_1 + . . . + x_D = 1, since

$$cov(x_1, x_1 + \ldots + x_D) = 0$$

we have

$$cov(x_1, x_2) + ... + cov(x_1, x_D) = -var(x_1)$$
.

The right-hand side here is negative except for the trivial case where the first component is constant.

The fact that data are closed, induces invalid correlations and as a result, all methods based on the covariance or correlation matrix of vectors of observations, are inappropriate to examine and analyze Compositional Data in crude or raw form (e.g. as simple percentages) [92]. Conventional statistical methods present uncertainty in the analysis of compositional data, as far as the results are concerned. The main reason is because it is not possible to distinguish between the spurious effects caused by the constant sum constraint and the effects that would be attributed to natural processes. Rock [90] in his paper describes some of the problems: trends and clusters on petrological ternary and principal components diagrams can have little or no geological significance; dendrograms produced by cluster analysis can be severely biased; results from discriminant analysis are likely to be illusory; any correlation coefficient will be affected to an unknown degree by spurious effects induced by the constant sum constraint, etc. In general, problems appear with all methods based on regression and multivariate analysis, Discriminant Analysis or Principal Component Analysis and they seem to perform better on unconstrained random variables.

6.2 Approaches in the Statistical Analysis of CoDa

In the early 1980's the analysis of Compositional Data began to obtain a more efficient form. The key to such analysis is the relative magnitudes and variations of the parts in a D-part composition, rather than their absolute values. Thus, the information provided is essentially about ratios and any meaningful function (scale-invariant) of a composition should be expressed under such terms. The principal justification for using ratios of components is the Sub-Compositional coherence, which is a fundamental property of Aitchison's approach to compositional data analysis. Ratios are unaltered in the process of forming sub-compositions ($s_i/s_j=u_i/u_j$) which should mean that there exists some form of covariance structure based upon them.

However, mathematically and statistically speaking, ratios are somewhat difficult to handle. For example, between $var(u_i/u_i)$ and $var(u_i/u_i)$ there does not exist any simple relationship. Therefore, in order to overcome this difficulty, Aitchison was the first to introduce the logratio method, because of the simplicity of relationships such as

$$var\{log(x_i/x_j)\}=var\{log(x_j/x_i)\}.$$

Since there is also a one-to-one correspondence between compositions and a full set of logratios, for example,

$$[y_1...y_{D-1}] = [\log(x_1/x_D)...\log(x_{D-1}/x_D)]$$

with inverse

$$[x_1 x_2 \dots x_D] = [\exp(y_1) \dots \exp(y_{D-1}) + 1] / \{\exp(y_1) + \dots + \exp(y_{D-1}) + 1\}$$

any problem or hypothesis concerning compositions can be fully expressed in terms of log ratios and vice versa.

The proposed methodology is simple; first transform each of the compositions (ui,...,ud) to their log-ratio vectors and then apply standard multivariate procedures upon them. The conclusions of the unconstrained multivariate analysis can then be translated back into conclusions about the compositions, and the analysis is complete.

The aforementioned methodology represents a transformation technique, widely utilized in statistics. Starting with McAlister [93] and his logarithmic transformation, the lognormal distribution and the significance of the geometric mean, the log-ratio transformation comes in line with a long tradition of statistical methodology.

6.3 The Simplex S^D – Fundamental Properties of CoDa Analysis

There has been much debate against transformation techniques over the scientific community [94, 95, 96, 97, 98, 99, 100, 101, 102, 103]. However, while most of them are still valid, new approaches have been developed towards the statistical analysis of compositional data. *Staying-in-the-simplex* approach, represents part of them, offering the advantage of keeping the analysis free of dependence upon transformations and results in unconstrained multivariate analysis. Therein, compositional data analysis is conducted within a simple algebraic-geometric structure on the simplex. At this point, the term simplex has to be defined.

One of the main differences between compositional and unconstrained data, is the sample space within which, each type lies. The natural sample space of CoDa is the (restricted) unit simplex S^d (while unconstrained data belong to the real space R). The simplex is a basic geometric element in a Euclidean space, and is defined as

$$S^{D} = \{x = [x_{1}, x_{2}, \dots, x_{D}] \in \mathbb{R}^{D} | x_{i} > 0, i = 1, 2, \dots, D; \sum_{i=1}^{D} x_{i} = n\}$$

The constant \varkappa simplex is positive and arbitrary. Frequent values for \varkappa are 1 (per unit), 100 (percent, %), 1000, etc. The simplex S^D is a line segment in one dimension (D=1), a triangle in two dimensions (D=2), a tetrahedron in three dimensions (D=3), and so on. As far the superscript in the S^D is concerned, it accounts for the effective dimension of D-part compositions and is often reduced to D-1, due to the unit-sum constraint. A unit-simplex is defined as

$$S^{D} = \{ [x_{1}, \dots, x_{D}] : x_{i} > 0 (i = 1, 2 \dots, D) | x_{1} + \dots + x_{D} = 1 \}$$

With this representation, scale invariance is an element to be ensured by formulating all statements concerning compositions in terms of ratios of components.

Scale invariance is one the fundamental principles governing the compositional data analysis according to Aitchison. What scale variance addresses, is that statistical inferences about compositional data should not depend upon the scale of the data.

More specifically, two vectors of D positive real components x, $y \in R D + (x_i, y_i \ge 0 \text{ for all } i = 1, 2, ..., D)$, are compositionally equivalent if there exists a positive scalar $\lambda \in R + \text{ such that } x = \lambda \cdot y$ and, equivalently, C(x) = C(y). It is highly reasonable to ask our analyses to yield the same result, independently of the value of λ . This is what Aitchison (1986) called scale invariance.

A function $f(\cdot)$ is scale-invariant if for any positive real value $\lambda \in R + and$ for any composition $x \in SD$, the function satisfies $f(\lambda x) = f(x)$, i.e. it yields the same result for all vectors compositionally equivalent. This can only be achieved if $f(\cdot)$ is a function only of log-ratios of the parts in x (equivalently, of ratios of parts) [102, 104]. According to Aitchison, apart from scale invariance, there are also two other conditions that should be satisfied in order for any statistical method to be performed on compositional data; permutation invariance and subcompositional coherence.

A function is permutation-invariant if it yields equivalent results when the order of parts of the composition is changed. Two examples might illustrate what "equivalent" means here. If we are computing the distance between our initial sandstone and our final sand compositions, this distance should be the same if we work with [Q, F, R] or if we work with [F, R, Q] (or any other permutation of the parts). On the other side, if we are interested in the change occurred from sandstone to sand, results should be equal after reordering. A classical way to get rid of the singularity of the classical covariance matrix of compositional data is to erase one component: this procedure is not permutation-invariant, as results will largely depend on which component is erased.

Before examining the topic of sub-compositional incoherence, the definition of subcomposition must be given. A composition only representing some of the possible components is called a sub-composition and most of real compositional data is actually representing a sub-composition, as we never analyze each and every possible component of our samples. Sub-compositions represent the marginals of compositional data analysis. Two compositions (a greater and a smaller one) sharing common parts (therefore, the smaller is the sub-composition) should produce common correlations for these parts, regardless of whether we analyzed only that sub-composition or a larger composition containing other parts. This is what coherence means. If this is not the case, then there is what is expressed as sub-compositional incoherence.

6.4 Perturbation and Powering

In any sample space there is, only certain operations can be performed. For example, in real space R^D translation and scalar multiplication are the most commonly used operations. However, the typical algebraic/geometric operations (addition/translation, product/scaling, scalar product/orthogonal projection, Euclidean distance) used to deal with conventional real vectors are neither sub-compositionally coherent nor scaling invariant. The simplex is a sample space characterized by a different, compositional geometry and such operations would not be adequate for any analysis within it. Two fundamental groups of operations for the simplex are the perturbation operations, analogous to translation in the real space, and power transformation, analogous to multiplication by a scalar in the real space. These operational sets were introduced by Aitchison [89], they underpin the complete algebraic – geometric structure of the simplex and both require in their definition the closure operation [104, 105]. Closure is nothing but the operation responsible for the constant sum constraint as it divides each component of a vector by the sum of the components and represents the projection of a vector with positive components onto the simplex.

For any two equivalent compositions x and X, in the same compositional class, there is a scale relationship (X_1 , ..., X_D)= (ax_1 , ..., ax_D) for some a >0, where each component of x is scaled by the same factor a to obtain the corresponding component of X. For any two compositions x and X in different compositional classes c and Ca similar, but differential, scaling relationship (x_1 , ..., X_D)= (p_1x_1 , ..., p_Dx_D) can always be found, simply by taking $p_i = X_i/x_i(i=1, ..., D)$.Denoting the operation between the positive perturbing vector $p = (p_1, ..., p_D)$ and the composition x by \oplus we have $p \oplus x = (p_1x_1, ..., p_Dx_D)$ and $X = p \oplus x$. Such a perturbation operator is then easily adapted to the simplex simply by defining $p \oplus u = (p_1u_1, ..., p_Du_D)/(p_1u_1 + ... + p_Du_D)$. Note that the roles of p and u are interchangeable in this definition and we can conveniently restrict p to lie in the simplex S^d. Perturbations thus defined form a group, with p⁻¹, the inverse of p, defined as (p_1^{-1} , ..., p_D^{-1})/($p_1^{-1}+... + p_D^{-1}$)and the identity perturbation $p \in S^d$ such that $U = p \oplus u$ and $u = p^{-1} \oplus U$, where $p = U \oplus u^{-1}$. Thus, the perturbation $U \oplus u^{-1}$, or equivalently X $\oplus x^{-1}$ characterizes the change from c to C; the change from X to x is simply the inverse perturbation $U \oplus u^{-1}$.

Powering or power transformation, as mentioned before, is the second fundamental operational group in the simplex. First, we define the power operation and then consider its relevance in compositional data analysis. For any real number $a \in R^1$ and any composition $x \in S^D$, we define:

$$X = \alpha \otimes x = C [x_1^{\alpha} \dots x_D^{\alpha}]$$

as the *a*-power transform of *x*. Such an operation arises in compositional data analysis in two distinct ways. First it may be of relevance directly because of the nature of the sampling process. More indirectly the power transformation can be useful in describing regression relations for compositions.

It is clear that powering \otimes and perturbation \bigoplus play a significant role as far as the geometry of S^{D} is concerned. Powering is an external operation whereas perturbation is an internal one, and it would be meaningless to establish that they define a vector or linear space structure on S^{D} . In particular, the \bigoplus operation defines an abelian group with identity e = [1, ..., 1] / D. Both operational groups are marked by certain properties, which will now be addressed.

$$x \oplus y = y \oplus x$$
, $(x \oplus y) \oplus z = x \oplus (y \oplus z)$, $a \otimes (x \oplus y) = (a \otimes x) \oplus (a \otimes y)$.

The operator Θ is the inverse of \oplus and is defined by:

$$\mathbf{x} \Theta \mathbf{y} = \mathbf{C} [\mathbf{x}_1 / \mathbf{y}_1 \dots \mathbf{x}_D / \mathbf{y}_D]$$

and plays an important role in the construction of compositional residuals.

The structure can be extended by the introduction of the simplicial metric

$$\Delta: S^D x S \xrightarrow{D} R \ge 0$$

Defined as follows:

$$\Delta(x, y) = \left[\sum_{i=1}^{D} \left\{ \log \frac{x_i}{g(x)} - \log \frac{y_i}{g(y)} \right\}^2 \right]^{1/2} = \left[\sum_{i$$

where g() is the geometric mean of the components of the composition. The metric Δ satisfies the usual metric axioms:

- *Positivity*: $\Delta(x, y) > 0 (x \neq y), \Delta(x, y) = 0 (x = y)$
- Symmetry: $\Delta(x, y) = \Delta(y, x)$
- *Power relationship:* $\Delta(a \otimes x, a \otimes y) = |a| \Delta(x, y)$
- Triangular inequality: $\Delta(x, z) + \Delta(z, y) \ge \Delta(x, y)$

The fact that this metric has also desirable properties relevant and logically necessary, such as scale, permutation and perturbation invariance and sub-compositional dominance, for meaningful statistical analysis of compositional data is now well established and the relevant properties are recorded briefly here:

- *Permutation invariance*: $\Delta(xP, yP) = \Delta(x, y)$, for any permutation matrix *P*.
- *Perturbation invariance*: $\Delta(x \oplus p, y \oplus p) = \Delta(x, y)$, where *p* is any perturbation.
- Sub-compositional dominance: if s_x and s_y are similar, say $(1, \ldots, C)$ -Subcompositions of x and y, then $\Delta_S^C(s_x, s_y) \leq \Delta_S^D(x, y)$.

6.5 The Log Ratio Methodology

The constant-sum constraint is a mathematical property embedded in any compositional data set, causing problems on the analysis of such a type of data. Aitchison [106, 107, 89] showed that the effects of this constraint on the covariance and correlation matrices disappear, if the raw percentage data are expressed as logarithms of ratios, where the denominator is the geometric mean of the percentages in each sample.

For applying statistical methods designed for the Euclidean geometry on compositional data, as wells as for representing them in the Aitchison geometry on the simplex, some kind of transformations are first necessary. The main idea that leads to such transformations is to find a basis (or a generating system) and to express compositions in coefficients of such a basis (coordinate system). This class of mappings is widely known under the term log ratio transformations. There are three types to be presented in the upcoming paragraphs: a) the additive log ratio transformation (alr) and inverse b) the centered log ratio transformation (clr), and finally, c) the isometric log-ratio transformation to the usual vector addition and scalar multiplication. However, only the latter two transformations move the whole Aitchison geometry to the Euclidean one, i.e. including the Aitchison inner product. As the proposed transformations are one-to-one transformations, the obtained results are usually back-transformed to the simplex in order to simplify the interpretation.

6.5.1 Additive Log Ratio Transformation (alr)

The additive log ratio (alr) transformation transforms raw compositional data from simplex to real (Euclidean) space. Alr transformation is also capable of performing its inverse

transformation (from real space to simplex) with its inverse *ALR*-1 (Aitchison, 2003). *ALR* differs from other transformations in that it maps a composition in the D-part simplex none isometrically to a D-1, dimensional Euclidean vector. As it maps, the last part is treated as a common denominator to the others, which means that in case the denominator changes, then the *ALR* transformations obtained, would be different. The additive log ratio transformation follows the idea to construct a (non-orthonormal) basis which is very easy to interpret, since the relation to the original D-1 first parts is preserved. Thus, for a composition x, a special case of the additive log ratio (alr) transformation [89] to R^{D-1} , is defined as:

$$alr(x) = \left(ln\frac{x_1}{x_D}, \dots, ln\frac{x_{D-1}}{x_D}\right)^{\prime}.$$

In this equation, there is a division of each of the first D-1 components by the final component. It is easy to see that also another part can be used as ratio part in the denominator. It is usually chosen in such a way that the interpretation of the result is facilitated. Note that different alr transformations are related by linear transformations (see, e.g., Filzmoser and Hron, 2008).

The inverse transformation ALR^{-1} : $R^{D-1} \rightarrow S^{D}$ is

$$x = alr^{-1}(x) = C[exp(y_1), exp(y_2)...exp(y_{D-1})1]$$

,where C is the closure operation. When data are in their transformed state, they can be analyzed by all those statistical methods not relying on a distance. The drawback of alr transformation is that it is not an isometric transformation from the simplex. It lacks symmetry and orthogonality dew to the use of a common numerator or denominator. This weakness could be solved by use of an appropriate metric with oblique coordinates in real ALRspace, but that is not a standard practice [91].

6.5.2 Centered Log Ratio Transformation (clr)

Taking a generating system on the simplex leads to the centered log ratio (clr) transformation (Aitchison, 1986) to R^{D} ,

$$clr(x) = \left[ln\frac{x}{g(x)} - \dots - ln\frac{x_D}{g(x)}\right]$$

,where g(x) is the geometric mean of the parts involved:

$$g(x) = \left(\prod_{i=1}^{D} x_i\right)^{1/D} = exp\left(\frac{1}{D}\sum_{i=1}^{D} lnx_i\right)$$

,or with the inverse transformation (clr⁻¹), from real space (clr coefficients) to the simplex (raw data) (Aitchison, 1986). The clr coordinates represent a generating system, not a basis, and therefore clr coordinates sum up to zero [108], i.e. we get a constrained transformed vector. As a result, correlations and covariances between clr parts are not sub-compositionally coherent.

6.5.3 Isometric Log Ratio Transformation (ilr)

The calculation of ilr coordinates is more complex and the generation of specific expressions is dominated by different rules. With ilr the data are transformed from the simplex to real space, as ilr coordinates, or conversely applying the inverse ilr⁻¹. Both features are defined by a sequential binary partition [108, 109]. The ilr transformation is defined as:

$$ilr(x)=(y1, y2, ..., yD-1) \in R^{D-1}$$
,

where $y_i = \sum_{j=1}^{D} y_{ij} ln x_j$, i = 1, 2, ..., D - 1 and

$$\psi_{i,j} = \sqrt{\frac{s_i}{r_i(s_i + r_i)}}$$
 if at step i the part j is + 1

or

$$\psi_{i,j} = -\sqrt{\frac{s_i}{r_i(s_i + r_i)}}$$
 if at step i the part j is -1

or

$$\psi_{i,i} = 0$$
 if at step i the part j is 0

with r_i the number of parts at step i as +1, and s_i the number of parts at step i as -1.

The ilr⁻¹ transformation is defined as:

X=ilr⁻¹(y)=(x₁, x₂, ..., x_D) \in S^D, where [x₁, x₂, ..., x_D]=Cexp[z₁, z₂, ..., z_D], $z_j = \sum_{j=1}^{D-1} \psi_{ij} y_j$, C stands for the closure operation [89].

6.6 The CoDaPack v2 Software Package

Over the last years, a new methodological approach has been developed for the statistical analysis of compositional data, based on the approach introduced in the early eighties by John Aitchison. This methodology is not straightforward to use with standard statistical packages. For this reason, in this project, we examine a new freeware software, The Compositional Data Package, which implements at this moment the most elementary of mentioned statistical methods. The features of this new software are very wide:

- Transformations between the real space to the simplex or vice versa such as the alr, clr and ilr transformations.
- Operations inside the simplex like centering, perturbation, power transformation, amalgamation, subcomposition (closure) or rounded zero replacement.
- 2-D and 3D graphical outputs like ternary diagrams, alr plots, clr plots, biplots, plots of principal components.

• Compositional Descriptive Statistics.

The software has been developed by members of the Research Group on Compositional Data Analysis at the Dept. Informàtica, Matemàtica Aplicada i Estadística (IMAE-UdG) under the projects Compositional Data Analysis and Related methods (CODA-RETOS) and Compositional and Spatial Data Analysis (COSDA). The core of the group belongs to the University of Girona (UdG), and includes members from the Technical University of Catalonia (UPC), and Biomathematics & Statistics Scotland (BioSS).

6.6.1 Interface of the CoDaPack software

This time the analysis will be conducted only on a small part of the data set, in order to examine briefly, how a different treatment approach would impact on the data. There will be a comparison of the results between the classical statistical analysis and the compositional statistical approach. For this attempt, the Saturate Fraction Compositional Model (SFCM) was selected, and in the next paragraphs there will be a presentation of the interface of the software package.

Data could be imported from Excel files or recovered from previous sessions. The observations are organized in rows and the variables in columns. CoDaPack v2 main window (Fig. 38) has four parts. On the very top there are the menus, on the left the active data frame and the name of its variables. The bigger part is the right side. On top of this part there is the place where alphanumerical results are placed, and on bottom there is the data.

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	15 16 17 18 19 20	9535.00 8193.00 15159.00 10441.00 10955.00 29309.00	9092.00 9372.00 12212.00 11081.00 11762.00 31584.00	9244.00 9393.00 11261.00 11348.00 12202.00 39173.00	8369.00 8411.00 9174.00 9887.00 10051.00 37710.00	10504.00 10220.00 11056.00 11760.00 13694.00 49958.00	313.00 313.00 308.00 815.00 1021.00 3433.00	3271.00 3258.00 3448.00 3152.00 12418.00	438.00 268.00 529.00 666.00 3642.00	4007.00 4401.00 4731.00 6304.00 1803.00	20 21 18 60
	15 16 17 18 19 20 21	9535.00 8193.00 15159.00 10441.00 10955.00 29309.00 21385.00	9092.00 9372.00 12212.00 11081.00 11762.00 31584.00 26845.00	9244.00 9393.00 11261.00 11348.00 12202.00 39173.00 35457.00	8369.00 8411.00 9174.00 9887.00 10051.00 37710.00 34940.00	10504.00 10220.00 11056.00 11760.00 13694.00 49958.00 46438.00	313.00 308.00 815.00 1021.00 3433.00 3160.00	3271.00 3258.00 3448.00 3152.00 12418.00 11426.00	438.00 268.00 529.00 666.00 3642.00 3028.00	4007.00 4401.00 4731.00 6304.00 1803.00 20486.00	20 21 18 60 57
	15 16 17 18 19 20 21 22	9535.00 8193.00 15159.00 10441.00 10955.00 29309.00 21385.00 34368.00	9092.00 9372.00 12212.00 11081.00 11762.00 31584.00 26845.00 32780.00	9244.00 9393.00 11261.00 11348.00 12202.00 39173.00 35457.00 37356.00	8369.00 8411.00 9174.00 9887.00 10051.00 37710.00 34940.00 34190.00	10504.00 10220.00 11056.00 11760.00 13694.00 49958.00 46438.00 44769.00	313.00 313.00 308.00 815.00 1021.00 3433.00 3160.00 3530.00	3271.00 3258.00 3448.00 3152.00 12418.00 11426.00 11521.00	438.00 268.00 529.00 666.00 3642.00 3028.00 2936.00	4007.00 4401.00 4731.00 6304.00 1803.00 20486.00 21383.00	19: 20: 21: 18: 60: 57: 57:

Fig. 38 CoDaPack v2 main window.

In order to run a CoDaPack routine we first import the data. The software stores a set of data on Data Frames or Tables. It is possible to have opened more than one Data frame. A set of

Data frames could be saved as a Workspace and also it could be recovered by means of the item button Open Workspace (Fig. 39).

Each Data frame contains the name of variables and its numerical values. As far as the missing values are concerned, there are two kinds; non-detected or non-available data and there is a specific symbol to distinguish them. Non-detected data should begin with a character prefix, for example <, followed by the value of low detection limit while Non-Available data should use a symbol, for example "NA".



Fig. 39 Menu File

Data frames may be imported and exported from Excel files. After data are imported, (Fig. 40) we must indicate in which row starts the data, if there are labels, non-available symbol and non-detected prefix. At any time, we may can delete a Data Frame from the active workspace. The exportation saves the names of the variables into the first row of an Excel file and the data in rows below variable names.

Data Frame Name: SAts.xlsx Path: C:\Users\user\Desktop\SAts.xlsx	File
Path: C:\Users\user\Desktop\SAts.xlsx	File
Spreadsheet options	
Start reading at row: 1 vith H	headers
Non-available data: NA	
Prefix for non-detected data: <	

Fig. 40 Importing Data

Another part of the menu which is utilized in this project, is the Data menu (Fig. 41). In general, this menu manages three kinds of routines: 1) transformations of the data from the simplex

to the real space and vice versa, 2) operations inside the simplex and 3) management of variables.

Data	Transformations	3	ALR						
C13 C14 C15	Centering Subcomposition/Closure		CLR ILR Version 2.02.04 re is being developed by the EAD g						
C16	Amalgamation								
C17	Perturbation								
Pr	Power transformation								
C18	Set detection limit								
Ph	Devended even and evene								
C19	Kounded zero replaceme	nτ							
C20	Numeric to categorical								
C22									
C23	Categorical to Numeric		C13	C14		C15	C16		
C24	A dal Nicoreania Vaniabian		815.0	JU] .	2400.00	5279.00	00.6000		
C25	Add Numeric variables		5854.0	00 00	5289.00	8055.00	6005.00		
C26	Delete variables		3672.0	00 3	3861.00	4965.00	3634.00		
C27		100	1865.0	00 2	2858.00	3872.00	3127.00		
C28		101	2858.0	00 4	4003.00	5313.00	3970.00		
C29		102	1879.0	00 3	2440.00	3128.00	2325.00		
		103	5060 (00 0	3455 00	0124.00	5001 00		

Fig. 41 Menu: Data

The software package offers various options as far as the data analysis is concerned (Fig. 41). Beginning with the Data Menu, *Centering* is a feature with which the data are centered, that is, they are perturbed by the center or closed geometric mean of the data (Fig. 42).

Center Data Menu		×
Selected		Options
Available data:	Selected data:	Show Center
C13 ^ C14 C14 C15 C16 C17 Pr C18 Ph C19 C20 C21 C22 C23 C24 C25 C26 C27 V	> < Reset	
		Accept Cancel

Fig. 42 Data: Centering

This routine centers the data set, that is, it returns the data set Y formed by the D-part compositions $y = gN(X)^{-1} \otimes X$, where

$$gN(X) = C\left[\left(\prod_{k=1}^{N} x_{k1}\right)^{1/N}, \dots, \left(\prod_{k=1}^{N} x_{kD}\right)^{1/N}\right]$$

is the closed geometric mean of the data set X. The center of the set Y is e, the barycenter of the simplex; e.g. for D = 3 the geometric center of a ternary diagram is [0:333; 0:333; 0:333]. If Show Center is activated this routine writes the center of the parts selected on the output window.

The feature *Subcompostion/Closure* the data is closed, i.e. data are converted into parts of some whole summing to a given constant, Y = C(X): This constant is, by default 1:0 but could be entered by the user by means of the Closure form. If S parts, S < D; are selected, a subcomposition with S-parts is obtained (Fig. 43).

Closure Data Menu		×
Selected		Options
Available data:	Selected data:	Closure 1.0
C13 A C14 C15 C16 C17 Pr C18 Ph C19 C20 C21 C22 C23 C24 C25 C26 C27 V	> < Reset	
		Accept Cancel

Fig. 43 Data : Subcomposition/Closure

The *Amalgamation* feature amalgamates some columns of the data (Fig. 44). The result of amalgamation of some of the parts of a D-composition selected by the user is the sum of those parts. Amalgamation should be used only as a first step in preparing the data for further analysis, as this operation is non-linear in the Aitchison geometry and might lead to inconsistent results if compared to analysis made without amalgamation.



Fig. 44 Data: Amalgamation

With the *Perturbation* feature a vector perturbs the data. The output is a matrix of D-part compositions

$$y = p \quad x = C[p_1 x_1 p_2 x_2, ..., p_D x_D],$$

where C stands for the closure operation, and p is a given D-part composition. The user has to indicate on Perturbation box the vector p, which has to be the same length as the compositions x.



Fig. 45 Data : Perturbation

The *Power Transformation* feature applies a power transformation to the data. For $a \in R$; the power transformation returns

$$\mathsf{a} \otimes \mathsf{x}=\mathcal{C} \left[x_1^a, x_2^a, \ldots, x_D^a \right]$$

In this option, we have to indicate the constant of the operation on the *Power* box.

The *Rounded Zero Replacement* applies a transformation to the data to avoid zeros (Fig. 46). This transformation involves substituting an observation x, with zeros in some parts, by an observation y using the expression:

$$y_i = \begin{cases} \delta_i, & \text{if } x_i = 0\\ x_i \left(1 - \frac{\Sigma x_j = 0^{\delta_j}}{C_x} \right), & \text{if } x_i > 0 \end{cases}$$

where δ i is the replacement value for the i-th part defined by the user and Cx the components sum of observation x. This routine applies to non-detected data (the software distinguishes between non-available and non-detected data). There is an individual constant δ i for each non-detected value, that is stored on the data frame.



Fig. 46 Data : Rounded Zero Replacement

The *Numeric to Categorical* feature transforms the selected variables into strings and overwrites the results on the same variables.



Fig. 47 Data : Numeric to Categorical

The *Numeric to Categorical* feature, on the other hand, transforms the selected variables coded with a string into numerical ones, and overwrites the result on the same variables.

The *Add Numeric Variables feature*, imports date to the data set by a simple copy-paste action (Fig. 48).

Add numeric variables	×
Variable names	
Data	
Accept	

Fig. 48 Data : Add numeric variables

Finally, the *Delete Variables* routine deletes the variables the user selects from the workspace (Fig. 49).

🛦 Co	Da	Pack v2.02.04										-	- ×
ile Da	ıta	Statistics Graphs Help											
Data 113 114 115 116 117 r 118 h 119 120 121 122 123 124 124 13 14 15 16 17 r 18 h 129 120 121 121 121 121 121 121 121		Transformations Centering Subcomposition/Closure Amalgamation Petrubation Power transformation Set detection limit Rounded zero replaceme Numeric to categorical Categorical to Numeric Add Numeric Variables	nt	DaPack - Version 2.02.04 is software is being developed by the <u>EAD</u> group (Grup d'Estadistica i Anàlisi de Dades).									
25		Delete variables											
27 28 29 30 31 32			1 2	C13 227.00	C14 1161.00 5027.00	C15 88.00 12950.00	C16 4085.00 22813.00	C17 7506.00 39446.00	Pr 593.00 791.00	C18 1782.00 16639.00	Ph 553.00 1570.00	C19 4562.0 24556.0	C20 0 10: ^

Fig. 49 Data : Delete Variables

The CoDaPack software includes a *Statistics* Menu. The first option is the Compositional Statistics Summary (Fig. 50). This menu produces two types of descriptive statistics: the first related to logratios (Variation Array, CLR variance and Total Variance) and the second related

to compositional descriptive statistics (Centre, Min, Max and quartiles). This routine is utilized and the results are presented in the next chapter.



Fig. 50 Statistics : Compositional Statistics Summary

1. Variation Array: Returns a matrix where the upper diagonal contains the logratio variances and the lower diagonal contains the logratio means. That is, the ij-th component of the upper diagonal is var $[ln(X_i=X_j)]$; and the ij-th component of the lower diagonal is $E[ln(X_i=X_j)]$, where i, j= 1,2, ..., D.

2. CLR Variances: Returns, for each part, the sum of logratio variances that involve it. Thus, for the i-th clr component ξ i we have

$$var(\xi_i) = \frac{1}{2D} \sum_{i=1, j\neq 1}^{D} var[ln(X_i/X_j)].$$

3. Total Variance: The sum of all clr Variances is the Total Variance totvar.

4. Centre: Returns the center of the data set, that is, $\hat{\xi} = C[g_1g_2, ..., g_D]$, where $g_i = (\prod_{k=1}^N x_{ki})^{1/N}$ stands for the geometric mean of part Xi in data set X. The data set X has been previously closed.

5. Minimum and Maximum: For each part of the data set X it returns the maximum and the minimum of the closed data set.

6. Quartiles: For each part of the data set X it returns the first quartile Q1, the median Q2 and the third quartile Q3 of the closed data set. The user has to select the columns to close and where to put the results. There are two buttons in this routine:

The output of the routine is placed on the output part. It includes a color classification of the logratio variances (elements of the upper diagonal of Variation Array). It is assumed that the logarithm of the logratio variances follow a t-student distribution, then dark blue colores those elements below percentile 5, light blue from percentile 5 to 25, light red form percentiles 75 to 95 and dark red up to percentile 95.

The menu *Classical Statistics Summary* produces standard descriptive statistics, including mean (arithmetic), standard deviation, covariance matrix, Min, Max and quartiles). The output of the routine is placed on the output part.

The Additive-Logistic normality test feature allows the user to perform a test for logistic normality of a D-part composition (Fig. 51). It includes all marginal, univariate distributions (with a total of (D - 1) tests); all bivariate angle distributions (with a total of D(D-1)/2 tests); and the (D-1)-dimensional radius distribution. For each kind of test the Anderson-Darling, Cramer-von Misses and Watson statistics are computed and their significance is given.



Fig. 51 Statistics: Logistic Normality tests

The *Atypicality Indices* feature obtains the atypical observations and their indices under the assumption of Additive Logistic Normal distribution of the selected parts (Fig. 52). The user has to select the columns to calculate its atypical observations and the threshold of atypicality (usually 0:95) has to be given.



Fig. 52 Statistics : Atypicality indices

The last part in the Menu section is the *Graphs* Section (Fig. 53). The options this software offers, enable the user to create graphs in independent windows. The can customize the appearance of each graph and, in some cases, plot the observations in the graph according to a previous classification. These graphs can be zoomed and, in 3D, rotated.

🔺 CoDaPack v2.02.04								
File Data St	tatistics	Graphs	Help					
Data Frames	FinalCho	Te	ernary plot					
C13		Te	ernary plot [Empty]					
C14		Te	ernary Principal Compone	ents	12	>0.		
C15		Pr	redictive Region					
C16		0	enter Confidence Region		43	>0.		
Pr		-	chief confidence negion					
C18		A	LR plot		20	>0.		
Ph		C	LR plot					
C19		IL	R plot					
C20					85	[0.15,		
C22		C	LR biplot					
C23		IL	R/CLR plot		91	>0.		
C24		Ba	alance dendrogram					
C25 C26								
C27				43.6	632	<0.		
C28								
IC29								

Fig. 53 Graphs Menu

To perform a zoom in a graph it is possible to use the slider scroll at the bottom of the graph or just using the scroll wheel of the mouse. It is also possible to rotate a figure by means of the left button of the mouse. Holding the left mouse button and moving it the graph rotates following the direction of the mouse. If the graph is 2D then the figure just moves inside the windows without rotation. To move the graph inside the window holding the left mouse button and simultaneously holding the ALT key. Furthermore, the graphs can be saved by means of snapshots of what windows have each moment. This can be done with the menu File-Snapshot and the files produced could be in jpeg, eps, png and bitmap formats. The same menu File includes a submenu Configuration that allows to customize the elements of the graph like lines and labels by means of changing size and colors.

The Graphs menu will not be further presented here, as many of the options will be used straight on the data set, and the outcome will be discussed.

6.6.2 Application of the CoDaPack's routine on the Saturates' fraction

To examine how compositional data behave when treated according to Aitchison, only a part of the whole data was used; the Saturates' fraction (see Appendix). Components with zero values were removed from this data set, as they would cause problems to the transformation operations. In particular, samples A549, A1711,A1724, A2268, A2283, A2284, A2468, A2469, B515, B554, B014, B1279, B2121, B2122, C540, C1465 and finally, C1473 were removed.

The first step is to use the *Amalgamation* option. As mentioned before, amalgamation should be applied on the data to prepare them before further analysis. Amalgamation is equal to addition in R. The results are presented in the following table (Table 7).

Table 7 Amalgamation of all variables for each component

	A549	A550	A920	A1140	A1710	A1711	A1712	A1723	A1724	A1725	A2268	A2269	A2270	A2283	A2284	A2313	A2362	A2363	A2364	A2424
amalg	24574.0	164351.0	95894.0	118237.0	84688.0	96204.0	130134.0	58253.0	65913.0	66642.0	39317.0	24519.0	41602.0	53986.0	62386.0	65691.0	79984.0	76814.0	78769.0	239513.0
	A2425	A2426	A2427	A2428	A2429	A2430	A2431	A2432	A2433	A2434	A2435	A2436	A2468	A2469	A2470	A2611	A2627	A2706	A2884	A2892
amalg	229793.0	252557.0	255369.0	288556.0	147814.0	212470.0	152544.0	103844.0	142061.0	202393.0	284439.0	171464.0	94280.0	255283.0	348520.0	51185.0	115613.0	64207.0	137398.0	121527.0
	A2895	A2896	A2897	A2898	B515	B554	B1014	B1279	B1393	B1443	B2121	B2122	B2887	B1873	B1874	C495	C499	C503	C511	C513
amalg	239946.0	141455.0	191406.0	104755.0	63637.0	85889.0	37508.0	24213.0	83605.0	63678.0	18410.0	22286.0	37639.0	39058.0	40661.0	73354.0	80288.0	124331.0	300295.0	200714.0
	C529	C540	C548	C553	C557	C566	C574	C575	C579	C582	C589	C596	C711	C714	C721	C722	C725	C1386	C1387	C1388
amalg	286154.0	74231.0	152070.0	253089.0	268887.0	181418.0	133703.0	233839.0	124065.0	124764.0	49701.0	70429.0	69213.0	95710.0	201581.0	69094.0	53147.0	124722.0	162462.0	224475.0
	C1389	C1390	C1465	C1466	C1467	C1468	C1469	C1470	C1471	C1472	C1473	C1705	C1715	D756	D800	D801	D802	D841	D842	D924
amalg	93596.0	135161.0	33821.0	26273.0	16547.0	20901.0	77669.0	18958.0	22813.0	19675.0	15327.0	101468.0	80685.0	101365.0	164064.0	206299.0	87376.0	89224.0	56791.0	45470.0
	D1173	D1273	D1274	D1275	D1276	D1288	D1289	D1290	D1291	D1312	D1313	D1335	D1364	D1365	D1385	D2471	D2472	D2595	D2626	D2885
amalg	54842.0	33217.0	85176.0	47918.0	49561.0	52781.0	52667.0	41060.0	31760.0	82846.0	66946.0	45664.0	79499.0	117301.0	244337.0	403272.0	93609.0	83301.0	79110.0	139966.0

The next options utilized are the Compositional Statistics Summary and the Classical Statistics Summary (Table 8, Table 9).

Table 8 Compositional Statistics Summary

											Comp	ositional												
											N	IA's:												
												25												
											Samp	ole size:												
												95												
									_		Sta	tistics	_											
											Ce	enter												
									C13	C14	C15	C16	C17	Pr										
									0.0653	0.0869	0.1131	0.1035	0.1227	0.0265										
									0.0676	0.0205	0.0771	0.0462	0.0275	0.0255										
									C22	C24	0.0771	0.0405	0.0575	0.0555										
									0.0300	0.0290	0.0266	0.0233	0.0200	0.0168										
									C29	C30	C31	C32	C31	C32										
Variatio	on array:								0.0145	0.0116	0.0086	0.0072	0.0086	0.0072										
	Variance	e In(Xi/Xj)																						
vi\vi	C12	C14	C15	C16	C17	Dr	C18	Dh	C10	C20	C21	C22	C73	C24	C25	C26	C 27	C78	C29	C30	C21	C32	dr	1
7.1 (A)	015	014	015	010	C1/		C10		015	020	C21	C22	C25	024	025	020	(2)	020	625	0.50	0.51	0.52	variances	
C13	_	0.2668	0.2975	0.4900	0.5675	15.260	0.9115	20.073	0.9388	12.839	15,900	14.883	16.916	16.127	16.077	16.570	16.381	17.318	15.840	16.146	15,135	18,519	10.099	1
C14	0.2860		0.1872	0.2844	0.3442	11.968	0.5762	15.872	0.5804	0.8622	11.456	10.586	12.674	12.097	12.433	12.637	12.676	13.312	12.123	12.002	11.353	14.289	0.6816	
C15	0.5492	0.2632		0.0518	0.0812	0.8739	0.2740	12.048	0.2701	0.5247	0.7707	0.6867	0.8709	0.8133	0.8331	0.8605	0.8596	0.9301	0.8117	0.8116	0.7528	10.275	0.3700	
C16	0.4608	0.1748	-0.0884		0.0372	0.9882	0.2583	12.921	0.1937	0.4852	0.7392	0.6621	0.8605	0.8143	0.8593	0.8682	0.8736	0.9294	0.8137	0.7888	0.7329	10.108	0.3809	
C17	0.6304	0.3444	0.0812	0.1696		11.495	0.3356	14.719	0.2185	0.5932	0.8582	0.7788	0.9760	0.9261	0.9645	0.9806	0.9811	10.527	0.9213	0.8987	0.8330	11.264	0.4747	
Pr	-0.9007	-11.866	-14.498	-13.615	-15.310		0.4274	0.0982	0.8334	0.4168	0.4618	0.3732	0.4566	0.3901	0.3871	0.4090	0.4426	0.4735	0.4574	0.4543	0.4949	0.5272	0.3265	
C18	0.0353	-0.2507	-0.5139	-0.4255	-0.5951	0.9360		0.5457	0.1672	0.0679	0.2097	0.1376	0.2684	0.2278	0.2878	0.2702	0.2959	0.3095	0.2770	0.2563	0.2596	0.4260	0.0516	
Ph	-0.7596	-10.456	-13.088	-12.205	-13.900	0.1410	-0.7950		10.365	0.4776	0.4970	0.3924	0.4639	0.3790	0.3856	0.3866	0.4261	0.4401	0.4556	0.4437	0.5201	0.5050	0.4256	
C19	0.1660	-0.1200	-0.3831	-0.2948	-0.4644	10.667	0.1307	0.9257		0.2502	0.4559	0.3874	0.5627	0.5349	0.6184	0.5950	0.6189	0.6455	0.5700	0.5345	0.5160	0.7796	0.2570	
C20	-0.3448	-0.6308	-0.8940	-0.8056	-0.9752	0.5558	-0.3801	0.4148	-0.5109		0.1028	0.0385	0.1521	0.1318	0.2198	0.1831	0.2251	0.2148	0.2254	0.1878	0.2187	0.3730	0.0719	
C21	-0.5536	-0.8396	-11.028	-10.144	-11.840	0.3471	-0.5889	0.2060	-0.7196	-0.2088		0.0698	0.0337	0.1423	0.2067	0.1793	0.2167	0.2159	0.2446	0.2074	0.2532	0.3730	0.1509	
C22	-0.6104	-0.8964	-11.596	-10.712	-12.408	0.2902	-0.6457	0.1492	-0.7765	-0.2656	-0.0568		0.0699	0.0444	0.1110	0.0794	0.1145	0.1138	0.1347	0.1079	0.1522	0.2674	0.0734	
C23	-0.7775	-10.635	-13.26/	-12.383	-14.079	0.1231	-0.8128	-0.0179	-0.9436	-0.4327	-0.2239	-0.16/1	0.0346	0.0706	0.1153	0.0865	0.1093	0.1201	0.1534	0.1427	0.1841	0.2731	0.1489	
C24	-0.8121	-10.981	-13.013	-12.729	-14.425	0.0880	-0.8474	-0.0524	-0.9781	-0.4673	-0.2585	-0.2017	-0.0340	0.0005	0.0437	0.0122	0.0282	0.0406	0.0005	0.0742	0.1147	0.1903	0.1006	
C25	-0.8980	-11.840	-14.478	-13.594	-15.290	-0.1299	-0.9339	-0.1390	-10.040	-0.5538	-0.3450	-0.2882	-0.1211	-0.0805	-0 1208	0.0409	0.0469	0.0761	0.0853	0.1047	0.1423	0.2210	0.1339	
C20	-11 951	-14 711	-17 2/2	-16.450	-19 155	-0.1200	-12 204	-0.2058	-12 511	-0.8403	-0.4736	-0.5747	-0.2319	-0.2175	-0.1308	-0 1557	0.0075	0.0160	0.0331	0.0313	0.0845	0.1335	0.1105	
C28	-13.571	-16.431	-19.063	-18.179	-19.875	-0.4564	-13.924	-0.5974	-15.231	-10.123	-0.8035	-0.7467	-0.5796	-0.5450	-0.4585	-0.3276	-0.1720	0.0137	0.0238	0.0424	0.0631	0.1343	0.1485	
C29	-15.070	-17.930	-20.562	-19.678	-21.374	-0.6063	-15.423	-0.7473	-16.730	-11.622	-0.9534	-0.8966	-0.7295	-0.6949	-0.6084	-0.4776	-0.3219	-0.1499	0.0LL1	0.0297	0.0334	0.1249	0.1195	
C30	-17.317	-20.176	-22.808	-21.925	-23.620	-0.8310	-17.670	-0.9720	-18.977	-13.868	-11.781	-11.212	-0.9541	-0.9196	-0.8331	-0.7022	-0.5466	-0.3746	-0.2247		0.0322	0.1415	0.1145	
C31	-20.328	-23.188	-25.820	-24.936	-26.632	-11.321	-20.681	-12.731	-21.988	-16.880	-14.792	-14.224	-12.553	-12.207	-11.342	-10.034	-0.8477	-0.6757	-0.5258	-0.3011		0.1263	0.1173	
C32	-22.047	-24.906	-27.538	-26.655	-28.350	-13.040	-22.400	-14.450	-23.707	-18.598	-16.511	-15.942	-14.271	-13.926	-13.061	-11.752	-10.196	-0.8476	-0.6977	-0.4730	-0.1719		0.2524	
	Mean In(xi/xj)									-												56.537	Total Variance

The Menu Compositional Statistics Summary, as mentioned before, includes two types of descriptive statistics. On Table 8 we observe the Variation Array, CLR variance and Total Variance as well as the Center, Min, Max and quartiles. The sample size is 95 in this case, due to the fact that in the data set, there exist zero values. The inadvertency introduced by a logratio variance (here clr variance) is that the logarithm of zeros does not exist, so if there are such observations in the data

On the other hand, the Menu Classical Statistics Summary includes the arithmetic mean, standard deviation, covariance matrix, Min, Max and quartiles (Table 9). The first step in analyzing multivariate data is computing the mean vector and the variance-covariance matrix. The mean vector consists of the means of each variable and the variance-covariance matrix consists of the variances of the variables along the main diagonal and the covariances between each pair of variables in the other matrix positions. The variance and the standard deviation are important in data analysis because of their relationships to correlation and the

normal curve. Correlation between a pair of variables measures to what extent their values co-vary. The term covariance is undoubtedly associatively prompted immediately. There are numerous models for describing the behavioral nature of a simultaneous change in values, such as linear, exponential and more. Observing Table 9, it is evident that all variables are correlated positively. The strongest positive correlation forms between C14 and C13 (0.9434). What is interesting here, is that in the classical statistics summary, the sample size remains at each original form of 120 samples. This is contrast to the Compositional Statistics summary, where sample size reduces, due to the exclusion of zero values.

Table 9 Classical Statistics Summary

										Clasica	l statistics sur	nmary:										
											NA's:											
											0											
											Sample											
											size:											
											120											
							Statistics															
							Statistics	Mean	Std.Dev	0	25	50	75	100								
							C13	82.765.083	88.639.087	0.0000	28.380.000	54.460.000	104.410.000	528.180.000								
							C14	97.541.417	84.559.280	0.0000	42.280.000	76.800.000	133.040.000	421.600.000								
							C15	121.133.500	101.799.595	0.0000	53.530.000	92.440.000	144.660.000	461.170.000								
							C16	113.407.583	99.571.357	7.800.000	46.660.000	81.820.000	135.770.000	449.950.000								
							Pr	31 463 083	32 061 154	2 340 000	11 090 000	21 010 000	41 090 000	232 300 000								
							C18	69.584.167	51.639.032	10.080.000	30.240.000	56.290.000	96.550.000	226.860.000								
							Ph	39.796.000	45.425.844	1.410.000	11.890.000	24.180.000	53.310.000	332.640.000								
							C19	84.766.167	70.033.547	9.580.000	29.900.000	59.420.000	114.730.000	275.920.000								
							C20	50.742.083	46.542.324	0.0000	18.190.000	39.300.000	61.150.000	239.940.000								
							C21	42.336.167	40.446.243	1.070.000	15.320.000	30.870.000	52.330.000	208.640.000								
							C22	39.475.083	38.984.371	2.720.000	14.510.000	27.610.000	49.230.000 4F.240.000	206.220.000								
							C25	32 257 667	32 915 131	2 770 000	10.930.000	24.740.000	40.650.000	184 340 000								
							C25	29.748.917	28.974.933	2.290.000	9.380.000	21.880.000	39,380,000	176,900,000								
							C26	25.860.417	26.808.898	1.040.000	8.090.000	18.480.000	33.200.000	158.100.000								
							C27	22.054.167	22.940.668	1.750.000	7.380.000	15.070.000	28.450.000	143.090.000								
							C28	18.874.667	19.582.024	1.440.000	6.410.000	12.350.000	23.310.000	97.930.000								
							C29	15.620.417	15.579.848	0.0000	5.260.000	11.190.000	19.470.000	79.610.000								
							C30	13.196.333	13.438.383	730.000	4.390.000	9.790.000	16.330.000	71.050.000								
							(31	9.331.583	9.881.062	0.0000	3.300.000	5 420 000	9,000,000	124 920 000								
Correlation:							0.52	5.400.055	10.420.070	0.0000	2.750.000	5.450.000	5.000.000	104.020.000								
	C13	C14	C15	C16	C17	Pr	C18	Ph	C19	C20	C21	C22	C23	C24	C25	C26	C27	C28	C29	C30	C31	C32
C13	10.000	0.9434	0.8505	0.7750	0.7316	0.1084	0.3769	0.0310	0.5192	0.1427	0.0897	0.0689	0.0309	0.0319	0.0439	0.0321	0.0402	0.0024	0.0281	0.0054	0.0418	0.0148
C14	0.9434	10.000	0.9604	0.9007	0.8646	0.2542	0.5544	0.1844	0.6985	0.3212	0.2635	0.2370	0.1840	0.1874	0.1920	0.1798	0.1789	0.1345	0.1618	0.1328	0.1713	0.1180
C15 C16	0.8505	0.9604	10.000	10.000	0.9485	0.3125	0.6306	0.2415	0.7903	0.3870	0.3303	0.3023	0.2688	0.2505	0.2820	0.2426	0.2433	0.1980	0.2303	0.2289	0.2413	0.1689
C16 C17	0.7750	0.9007	0.9599	0.9789	10,000	0.1943	0.5905	0.1555	0.8272	0.3911	0.3277	0.2994	0.2445	0.2410	0.2455	0.2340	0.2323	0.2016	0.2303	0.2100	0.2640	0.1388
Pr	0.1084	0.2542	0.3125	0.1943	0.1779	10.000	0.6878	0.9485	0.4244	0.6665	0.7037	0.6977	0.7359	0.7382	0.7940	0.7458	0.7500	0.6854	0.6873	0.6799	0.6575	0.7282
C18	0.3769	0.5544	0.6306	0.6044	0.5905	0.6878	10.000	0.7424	0.8479	0.9215	0.8922	0.8728	0.8494	0.8214	0.8012	0.7920	0.7658	0.7739	0.7466	0.7647	0.7445	0.5363
Ph	0.0310	0.1844	0.2415	0.1533	0.1267	0.9485	0.7424	10.000	0.4478	0.7715	0.8127	0.8167	0.8470	0.8572	0.8836	0.8576	0.8525	0.8048	0.7838	0.7748	0.7580	0.7944
C19	0.5192	0.6985	0.7903	0.8272	0.8324	0.4244	0.8479	0.4478	10.000	0.7398	0.6854	0.6606	0.6024	0.5925	0.5589	0.5687	0.5472	0.5455	0.5475	0.5295	0.5701	0.3413
C20	0.1427	0.3212	0.3870	0.3911	0.3590	0.6665	0.9215	0.7715	0.7398	10.000	0.9888	0.9802	0.9314	0.9259	0.8545	0.8921	0.8542	0.8896	0.8478	0.8336	0.8486	0.5529
C21	0.0897	0.2635	0.3303	0.3277	0.2900	0.7037	0.8922	0.8127	0.6854	0.9888	10.000	0.9965	0.9649	0.9660	0.9071	0.9386	0.9079	0.9307	0.8921	0.8750	0.8843	0.6171
(22	0.0889	0.23/0	0.3023	0.2994	0.2620	0.6977	0.8728	0.8167	0.6024	0.9802	0.9965	10.000	0.9714	0.9760	0.9203	0.9527	0.9238	0.9460	0.90/1	0.8887	0.9002	0.5351
C23	0.0309	0.1874	0.2505	0.2445	0.1995	0.7339	0.8454	0.8572	0.5925	0.9259	0.9660	0.9763	0.9760	10.000	0.9687	0.99918	0.9787	0.9736	0.9458	0.9131	0.9245	0.7296
C25	0.0439	0.1920	0.2820	0.2455	0.2183	0.7940	0.8012	0.8836	0.5589	0.8545	0.9071	0.9203	0.9675	0.9687	10.000	0.9735	0.9760	0.9500	0.9359	0.9367	0.9076	0.7775
C26	0.0321	0.1798	0.2426	0.2340	0.1905	0.7458	0.7920	0.8576	0.5687	0.8921	0.9386	0.9527	0.9599	0.9918	0.9735	10.000	0.9950	0.9843	0.9711	0.9245	0.9484	0.7566
C27	0.0402	0.1789	0.2433	0.2323	0.1885	0.7500	0.7658	0.8525	0.5472	0.8542	0.9079	0.9238	0.9443	0.9787	0.9760	0.9950	10.000	0.9802	0.9751	0.9280	0.9513	0.7832
C28	0.0024	0.1345	0.1980	0.2016	0.1584	0.6854	0.7739	0.8048	0.5455	0.8896	0.9307	0.9460	0.9514	0.9736	0.9500	0.9843	0.9802	10.000	0.9860	0.9512	0.9663	0.7007
C29	0.0281	0.1618	0.2303	0.2363	0.1922	0.6873	0.7466	0.7838	0.5475	0.8478	0.8921	0.9071	0.9130	0.9458	0.9359	0.9711	0.9751	0.9860	10.000	0.9438	0.9764	0.6980
C30	0.0054	0.1328	0.2289	0.2166	0.1913	0.6799	0.7647	0.7748	0.5295	0.8336	0.8750	0.8887	0.9400	0.9131	0.9367	0.9245	0.9280	0.9512	0.9438	10.000	0.9364	0.6922
(31	0.0418	0.1/13	0.2413	0.2640	0.2250	0.05/5	0.5262	0.7580	0.5/01	0.5520	0.6843	0.9002	0.9014	0.9245	0.90/6	0.9484	0.9513	0.9663	0.9764	0.9364	10.000	10.000
C32	0.0148	0.1180	0.1089	0.1388	0.1108	0.7282	0.5303	0.7944	0.3413	0.5529	0.01/1	0.0351	0./14/	0.7290	0.7775	0.7506	0.7832	0.7007	0.6980	0.0922	0.7392	10.000

The CoDaPack softaware offers the option of a Ternary Principal Component Graph. This feature calculates the two (or three) compositional principal components for a 3-part (or 4-part) composition and displays the result in a ternary diagram. What is more, it returns, as a numerical result, the Principal Components and the cumulative proportion explained with each component.

It would be meaningful if the Principal Components of this 3-part composition, is based on variables that present a bigger weight, related to the rest variables of the data set. For this reason, the command *var(X)* was utilized on Matlab to examine which are the three variables with the most impact. It was found that C13, C14 and phytane obtain the biggest weights (0.1034, 0.0796 and 0.924 respectively). Below we present both the ternary principal

component graph for all components produced by CoDaPack, as well as the PC plot that is produced by matlab (using raw compositional data).

• A • B • C • D - PC1 • PC2



Fig. 54 Ternary Principal Component Graph for C13, C14 and phytane.



Fig. 55 Plot of the first two Principal Components for C13, C14 and phytane.

By examining the ternary principal component graph (Fig. 54), we observe that Family A oils are distinctively separated from the rest, presenting a sub-parallel alignment to the first principal component axis (PC1). Samples from families B, C and D follow a linear trend along the PC2 axis overlapping each other. In Table 10 we observe the numerical representation of the principal components for each variable, as well as, the cumulative proportion explained with each PC. Both PC1 and PC2 are positively correlated to the three variables. C13 is the most important in explaining PC1, whereas C14 is the most important in explaining PC2

Table 10 Principal Components as Numerical results and the Cumulative proportions explained with each principal component.

	C13	C14	Ph	Cum. Prop. Exp.
PC1	0.4715	0.3999	0.1287	0.9074
PC2	0.1465	0.5970	0.2565	1.0000

On the other hand, Fig. 55 displays a completely different principal component analysis result. As far as the discrimination of the four family affiliations is concerned, it is evident that there is no clear distinction among them. All samples follow strictly linear gradients, overlapping
significantly, at the same time. PC1 scores for all samples are positively high, whereas for PC2, the majority obtains negative scores. The first Principal Component in this case explains 81% of the total variance, and PC2 follows with 15% of the total variance.

A simple ternary plot of C13, C14 and phytane is displayed on Fig. 56. As in the ternary principal component graph, in this plot there is a significant overlapping among oils B, C and D. Family A oils form a quite distinct group along the C13-C14 axis. Along the C13-phytane axis there is a sample (number) which displays a different behavior from the rest and it is D1338. Fig. 57 displays the centered version of the same plot. It offers a better understanding of how oil samples exist in the ternary plot's space.





The ALR plot represents a plot of three (four in 3D) alr-transformed parts (Fig. 58). The new variables obtained with the ALR transformation are displayed in an orthogonal coordinate system to visualize how the plot changes when permuting the components or initial columns. Nevertheless, care is required when interpreting the plot, as the axis are not really orthogonal, but at 60°.



Fig. 58 ALR plot of C13, C14 and phytane

What is observed in the ALR plot, is that oil samples form a positive gradient of 30° along the intersection of alr.C14_Ph and alr.C13_Ph axes. The additive logratio transformation seems to reveal a linearity embodied in oil families. Once more, the most distinct group is that of family A oils. The overlapping still holds among the other oil families.

The CLR plot feature represents a plot in an orthogonal coordinate system of the data, after the centred logratio transformation (clr) of two (three in 3D) selected parts. It has the same capabilities as the ALR Plot.

The ILR plot feature displays a plot in an orthogonal coordinate system of the data after the isometric logratio transformation (ilr) of three (four in 3D) selected parts according to a sequential binary partition. The way to select the partition is the same as in Transformation-ILR routine. The partition selected in our case is the default (Table 11).





Table 11 Binary partition for ILR transformation

• A • B • C • D



Fig. 60 ILR plot of C13, C14 and phytane

In the ILR plot there are two distinct positive gradients sub-parallel to and below the ilr.1 axis (Fig. 60). One of the two gradients, consists of oil samples solely from Family A and the other consists of oil samples from families B, C and D. The projections of sample points of family A oils do not overlap with any of the other, in contrast to the rest that overlap significantly.

The CLR biplot includes the selected variables C13, C14 and phytane. Once the graph is performed, we may choose 1) which 2D view we prefer (axes XY, YZ or XZ), 2) to display observations or not, and 3) which biplot display depending on the Form value; $\alpha = 0$ corresponds to a Covariance Biplot, $\alpha = 1$ Form Biplot, and $\alpha = 0.5$ Symmetric Scaling Biplot, which is the default value. In Fig. 61 the biplot is a Form Biplot.

What is more this routine returns, as a numerical result, the Principal Components and the cumulative proportion explained with each component (Table 12). Biplot consists on the decomposition of clr matrix, X = UDV'. If numerical output is desired the routine writes three matrices: UD, D and V. UD are the ilr coordinates of the original data.



Fig. 61 CLR biplot of C13, C14 and phytane

As far as the distinction of the families is concerned, more or less, the CLR Biplot presents the same results, as in the previous graphs.

Table 12 Principal Components explained by clr.13, clr.14 and phytane

	clr.C13	clr.C14	clr.Ph	Cum.Prop.Exp.
PC1	0.4878	0.3231	-0.8109	0.9074
PC2	-0.6548	0.7498	-0.0951	1.0000

Table 12 displays with which variable each principal component is explained along with the cumulative proportion explained. PC1 is positively correlated with clr.C13 and clr.C14, but negatively with clr.ph. PC2 is negatively correlated with clr.C13 and clr.ph, but positively with clr.C14.

Lastly, the Balance Dendrogram represents a dendrogram by means of a sequential binary partition of selected parts (Fig. 62). The way to select the partition is the same as in Transformation-ILR routine. Here the default partition is chosen (Table 14). As a numerical output, this routine returns on the output window the sequential binary partition used, the mean and the variance of each balance (Table 13). Also on the Data window are the ilr coordinates produced with this partition.



Fig. 62 Balance dendrogram of C13, C14 and phytane

 Table 13 Numerical output of Balance Dendrogram routine, including the mean and variance
 Including the mean and variance

Me	ean	Vari	ance
Balance 1	Balance 2	Balance 1	Balance 2
0.7370	-0.2022	1.1537	0.1334

Table 14 Default partition for the Balance Dendrogram routine

C13	C14	Ph
1	1	-1
1	-1	0

7. Conclusions

The aim of this project has been the examination of the way multivariate clustering methods perform on the classification of oil family affiliations. The methods implemented hereby include Hierarchical clustering, k-means clustering and Principal Component Analysis. The data set under study contained raw compositional information of four distinct oil families present at Williston Basin of Canada. For the needs of the study, four different models were developed out of the given geochemical information; the Saturates' Fraction Compositional Model, the Saturates' Fraction Ratios Model, the Gasoline Range Compositional Model. Focus was not placed on how the models would perform under the aforementioned statistical analysis, but the exact opposite. The effort was on the examination of the data set through a manifold manner.

Taking into consideration the performance of each method separately we conclude as follows:

- Hierarchical Clustering performed relatively well on all models. Family A oils were classified sufficiently and in some cases Family C oils appeared to form fair clusters. However, there was always considerable overlapping among families B, C and D.
- k-means failed in the task of classifying the given data set into distinct groups. In the SFCM and SFRM, it produced a two-cluster solution, with one cluster including mainly samples from Family A, and another cluster containing the rest. Judging, however, from the k-means plots, the clusters produced, did not present clear boundaries between them. In the GRCM and BCM, k-means produced a three-cluster solution, but significant overlapping among all families was observed. This was also evident from the respective k-means plots.
- Principal Component Analysis performed similarly to hierarchical clustering. It mainly distinguished Family A samples and presented significant overlapping among the rest oil samples. In BCB especially, there was an overlapping between families A and D, as well as with families C and D. Family B oil samples were dispersed in the plot.

All in all, the geochemical information under study, contains complex compositions of different oils. A blind application of multivariate data analysis methods on such data seems to be unable to classify them into distinct groups. Compositional data require probably different approaches concerning their analysis. Their special properties cause problems when analyzed with standard multivariate methods and a whole new chapter has been introduced by the scientific community on the way to examine them. The final chapter of this project deals with an alternative approach towards the analysis of compositional data, and results are compared to previous approaches. Principal Component Analysis in particular, presents a completely different picture when approached in a different manner. Further investigation, however, should be conducted on this type of data in order to understand their behavior and obtain meaningful information through their analysis.

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APPENDIX

Below we present the data set under study. The next tables include all raw data concerning the Biomarkers, the Gasoline range and the Saturated fraction. All models that were examined by multivariate statistical were derived from these three parts of the data set.

Biomarkers (Hopanes and Steranes) of the sample set

2425	A2426	A2427	A2428	A2429	A2430	A2431	A2432	A2433	A2434	A2435	A2436	A2468	A2469	A2470	A2611	A2627	A2706	A2884	A2892
0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	3650000	1110000	1930000	2990000	0
30000	2310000	1400000	0	872000	0	0	0	944000	1890000	0	0	0	0	0	3650000	1460000	1730000	2130000	1990000
30000	8780000	6560000	4610000	3360000	7500000	4590000	3350000	3010000	784000	3060000	8900000	37240000	34840000	19540000	17700000	6410000	9650000	12500000	7760000
1 00000	1300000	8590000	10200000	4280000	11700000	7860000	5270000	6190000	11500000	6500000	12800000	51820000	53980000	12650000	36500000	9500000	14400000	17300000	3930000
00000	28500000	19500000	22400000	11000000	26700000	15700000	12200000	13500000	25700000	12900000	22600000	135900000	116800000	25630000	74700000	21200000	31800000	38400000	8790000
00000	48000000	34700000	36900000	21900000	46600000	26900000	22600000	20800000	46400000	21000000	33300000	259700000	222300000	49010000	129000000	41400000	53400000	66900000	1500000
00000	19200000	11600000	16100000	8210000	19200000	10700000	8920000	8610000	18100000	8750000	10700000	00009966	92850000	16760000	54700000	17800000	22800000	25200000	5270000
1 00000	12400000	7550000	11100000	5770000	12400000	7120000	5470000	5740000	12000000	5680000	6530000	68410000	59630000	11480000	37800000	12200000	15000000	16600000	3850000
30000	1800000	1570000	1670000	1220000	2350000	1200000	1010000	965000	2300000	904000	849000	14840000	10560000	2731000	5890000	2050000	1870000	0	0
30000 1	11200000	7020000	10500000	5760000	12300000	6510000	5480000	5290000	11800000	5300000	4960000	70540000	58890000	11570000	36600000	12800000	15700000	16500000	3710000
00000	7330000	4700000	7080000	4090000	8270000	4420000	3520000	3430000	7850000	3500000	3150000	48970000	42700000	8838000	24000000	8950000	10500000	10700000	3560000
50000	5230000	3280000	5200000	3340000	6350000	3150000	2620000	2300000	5920000	2360000	1860000	38360000	32930000	6148000	19100000	7880000	9230000	7130000	2250000
00000	2760000	1870000	3150000	2060000	4010000	1770000	1520000	1350000	3690000	1450000	1020000	23290000	20630000	3837000	12000000	4980000	5500000	4770000	1260000
20000	4030000	2640000	3810000	3560000	6170000	2450000	2280000	1700000	5320000	1760000	1210000	65720000	28830000	5905000	16000000	9380000	8670000	6480000	1690000
00006	2280000	1430000	2400000	2210000	3880000	1400000	1280000	877000	3060000	1000000	597000	41110000	18250000	3576000	10100000	5830000	5520000	3560000	923000
1000	828000	501000	1550000	849000	1670000	618000	479000	448000	1410000	568000	20100	25600000	13250000	2297000	9570000	3820000	3630000	1630000	635000
0000	291000	200000	784000	408000	757000	228000	179000	186000	599000	227000	10000	14750000	6785000	1134000	5100000	2340000	2020000	761000	220000
3000	560000	519000	381000	368000	486000	311000	207000	298000	530000	401000	1030000	3841000	2877000	2492000	1600000	1050000	880000	1080000	1290000
40000	968000	710000	753000	423000	692000	551000	372000	424000	908000	614000	1300000	6462000	5582000	3957000	3840000	1490000	1780000	2170000	1950000
20000	1720000	1520000	1610000	000662	1430000	1110000	787000	810000	1840000	1140000	2370000	10820000	9838000	6672000	6200000	2740000	3250000	3810000	3330000
6000	235000	283000	439000	124000	348000	241000	155000	149000	413000	307000	475000	4360000	2580000	1530000	1570000	708000	727000	809000	333000
0006	184000	187000	240000	118000	213000	133000	109000	118000	243000	199000	245000	1760000	1040000	718000	642000	258000	408000	287000	236000
00000	1000000	908000	961000	517000	878000	650000	451000	514000	966000	738000	1210000	8594000	6405000	2794000	3220000	1550000	1820000	2010000	1180000
40000	1290000	1230000	1240000	683000	1280000	879000	630000	672000	1220000	944000	1530000	11220000	6871000	3647000	4170000	1840000	2170000	2350000	1650000
10000	1060000	926000	950000	538000	1030000	712000	485000	518000	898000	725000	1160000	8935000	5362000	2897000	3390000	1360000	1730000	1940000	1310000
00000	931000	894000	924000	489000	873000	628000	421000	492000	854000	672000	1010000	8251000	5234000	2464000	3290000	1460000	1540000	1730000	1150000
	2425 0 0 0 0 0 0 0 0 0 0 0 0 0	2425 A2426 0 0 80000 2310000 30000 8780000 30000 1300000 900000 1300000 30000 1300000 30000 12200000 30000 12200000 30000 1230000 30000 1230000 30000 1230000 50000 233000 30000 1230000 50000 2280000 90000 276000 50000 228000 90000 228000 90000 228000 90000 228000 90000 228000 90000 228000 90000 172000 90000 172000 90000 174000 90000 172000 90000 1290000 90000 1000000 90000 1000000 90000 1000000 931000 931000	2425 A2426 A2427 0 0 0 80000 2310000 878000 30000 878000 856000 90000 1300000 859000 90000 1300000 1400000 90000 1240000 850000 9120000 1240000 850000 90000 1240000 755000 90000 1240000 757000 90000 1230000 1470000 90000 733000 170000 90000 733000 187000 90000 238000 143000 90000 238000 143000 90000 238000 147000 90000 238000 147000 90000 238000 147000 90000 238000 147000 90000 238000 147000 90000 238000 147000 90000 140000 147000 90000 140000 1	2425 72426 72426 72428 0 0 0 0 80000 2310000 1400000 4610000 30000 8780000 8560000 1410000 00000 1300000 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GAM	7810000	3384000	3396000	1776000	662400	2399000	1325000	1194000	3544000	3882000	681700	4549000	3594000	2066000	0	72150000	65770000	65370000	0000696	90100000
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B2121	29221	50608	2040	0	18506	185932	127402	256485	6385	117179	27055	1544	786	3922	26411	221459	113878	33383	369586	240116	217023	565692	565128	350320	127114	88764	40886	67225	332321	498730	9649	438858	732212	199979	454999
B1443	482913	1458581	0	0	183375	1793018	1263633	3834118	0	2746982	86916	1775	118582	5911	416051	1529421	421966	48779	2748085	1704722	1586388	3223733	5642711	2227259	107622	1251303	164907	203149	796597	20651	159829	1950240	1985330	0	2894993
B1393	342289	833954	5551	0	171714	1044306	754058	1936610	0	1785196	66545	4914	190841	12801	630693	675349	218486	273058	1121140	1240709	1184249	2078723	2297995	2191021	602395	242485	68743	102817	638310	8336	123071	880252	1240921	238435	928244
B1279	45539	72005	775	0	8783	124546	80879	173698	0	121077	11860	68	923	1479	31570	129800	29128	17485	184399	118631	110451	271563	233694	199562	47500	45660	13216	22244	115388	88	37	121994	169080	47034	75260
B1014	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
B554	141883	240481	1767	0	30206	397225	265174	626552	0	518762	30646	1143	10869	4323	97039	351133	84183	65802	519764	458016	422281	925929	989254	651881	167291	169493	31901	49668	290791	5994	23844	369957	624614	106101	443095
B515	7599	13617	356	44	5955	80278	55498	127412	0	68978	12148	360	712	2531	14974	131890	44463	15780	203935	139146	127891	304098	377655	216008	66043	62425	23377	37215	183293	4166	10542	280639	166640	113010	529524
A2898	1539	10143	175	1683	850	7663	4678	75704	257	15461	906	0	5201	399	35024	11662	4468	1626	13461	4654	4404	13499	220821	94544	2558	8866	1394	1830	3042	20	25702	19029	10895	3329	246482
A2897	7371	24186	1212	3709	4228	27787	17731	130954	2272	38087	4444	588	8569	1666	44606	33144	13090	5485	34634	11688	10992	29221	282039	152989	8448	12006	4673	6106	8018	2	47320	39528	20019	3682	254510
A2896	4957	34169	317	4703	1638	16728	6966	173712	437	34839	1453	0	11041	616	53586	19591	7141	2606	22721	7899	7420	22332	386974	146371	3667	17052	1733	2414	4909	1	36694	26815	14667	7071	373185
A2895	4845	20384	373	4372	1883	16039	6866	102389	510	37599	1643	131	6698	869	38394	17063	8261	2648	20011	9478	8946	27455	234548	115546	4110	13861	1853	2364	5881	60	19591	20025	10816	3030	191893
	iC5	nC5	22DMC4	CYC5	23DMC4	2MC5	3MC5	nC6	22DMC5	MCYC5	24DMC5	223TMC4	Benzene	33DMC5	CYC6	2MC6	23DMC5	11DMCYC5	3MC6	c3DMCYC:	It3DMCYC	It2DMCYC:	nC7	MCYC6	22DMC6	ECYC5	25DMC6	24DMC6	223TMC5	234TMC5	Toluene	2MC7	3MC7	c4DMCYC	nC8

C1388	262484	460672	3180	159723	37364	703469	623535	1370787	0	1243485	42097	2502	1036419	5335	790381	658773	196525	80935	980124	644370	603365	1380585	1897831	1526450	165959	386730	74252	91227	414838	8289	721874	781250	178269	163993	1023797
C1387	243917	411995	3724	0	135115	499412	386393	932405	0	491547	22506	2031	829395	5051	659497	367819	110626	20218	503623	154043	142377	345509	1008065	772063	34624	152931	38504	41479	89837	3648	520902	322695	106038	39203	449886
C1386	514115	851472	4812	0	253968	974015	784000	1914212	0	905845	37644	2476	2314197	7079	1229279	699249	209709	42461	987340	255637	235480	579003	2136244	1553881	51627	344646	82959	87967	151326	6405	1607882	748001	831282	71684	1155940
C725	3360	5219	331	881	3458	24743	19929	38791	1100	30266	3361	262	38	1034	52203	41895	23627	8044	67169	25442	23253	57427	89743	156446	17016	19504	7536	10314	27331	763	453	36674	24616	18099	74981
C722	16159	23249	733	4055	6905	49197	38776	72728	1694	64299	5704	360	295	1406	108190	53828	31417	11615	87691	34743	32186	79027	109633	216382	19538	25730	7649	10733	29648	666	376	67170	2395	16382	60801
C721	1059	868	0	0	941	7860	6561	20762	448	9723	1324	0	1298	303	15577	22456	11736	2869	36750	12577	11140	29142	88918	72601	8567	12121	5066	6444	17169	476	1316	53854	1945	11053	79793
C714	8516	15920	339	2819	3629	31126	23246	73221	1049	47378	3256	378	27186	879	63779	38934	21722	8577	64372	29823	28001	70150	164883	154150	19371	20336	6077	9134	32067	0	43288	71118	105	18715	119322
C711	5277	11152	375	2211	3501	30507	23438	78933	1323	48819	4525	287	15416	1155	75545	47242	25042	11447	75393	32858	29685	72368	178477	173451	20996	20156	6361	9482	32372	921	40104	83120	16923	16699	145739
C596	108	31	0	P	268	739	937	1272	110	3768	281	0	511	120	4591	2879	3397	1100	5904	4671	4140	13753	6020	21671	4160	3874	1866	2375	9829	278	356	14725	8711	11661	13818
C589	7092	8229	245	1349	2437	18500	16015	16306	357	25262	1956	107	65	308	14389	14835	10771	3533	28587	20468	19189	50444	20320	46759	8002	10288	2157	3787	19282	261	29979	11705	21339	8360	3612
C582	109902	143764	1544	22071	29614	155447	128543	261384	1604	121982	6198	450	86216	1849	196536	108889	39889	4131	162350	31742	29789	84329	342723	256834	11809	54448	20051	20067	23220	1594	130594	158616	177234	7749	146919
C579	128450	230196	1816	44018	34074	272158	225116	456607	3848	443630	16552	668	408767	7216	250624	238474	73411	24427	343765	258874	245238	612454	661926	567970	105358	135793	39097	46232	246912	7160	561640	450567	135757	4694	141066
C575	7507	10869	207	2383	4362	24019	20146	44143	298	19257	1174	123	53034	623	32561	19463	10635	1019	29857	4572	4161	12514	60677	44179	1148	9304	3503	3325	2934	6421	22056	24922	7620	1992	40195
C574	5858	9939	92	1642	1555	12897	11214	23939	165	26351	1032	0	6773	147	12094	10430	6211	2936	19888	17457	16567	42131	34410	36827	5564	9012	1286	1878	11899	3446	4498	13044	16339	4989	13872
C566	42419	62352	1063	10735	19517	124051	108582	200914	1815	176850	10826	545	40088	2515	121497	124182	68055	21908	196295	130778	116841	306841	349575	317286	50797	69634	20810	25823	117732	2932	69802	209009	81291	64507	268618
C557	1051	1560	317	540	549	3236	3253	8060	93	6832	282	136	10800	0	9446	6651	3912	558	11713	4204	3936	11272	28836	24502	1365	6488	2018	2045	5018	3809	36387	10277	1424	3068	38773
C553	2646	9(9)	803	936	1520	20721	16263	42973	367	31060	2146	25	7736	257	9957	31618	17317	4679	61454	34390	30569	85044	117778	69888	9452	25006	7112	9081	35238	1109	42386	94308	69868	31247	168343
C548	131989	190458	3203	0	79737	397918	290090	696022	0	334499	26547	2214	110116	6143	539298	364706	121458	34969	485096	123305	111735	270261	1015137	812230	78720	107008	50284	61146	123671	5624	269444	478278	77483	86864	710465
C540	43712	74421	749	18392	9307	95337	77747	174752	0	160212	8062	309	187465	2498	110777	95411	38452	18268	151034	106027	99539	248773	322588	269413	45334	62001	17156	21013	94114	3474	355820	130435	118489	52757	280082
C529	8957	9343	336	3374	5511	21789	19579	29867	6581	33150	1943	191	320	7619	43051	21976	15155	5030	36571	: 16914	18707	45895	58627	47691	807	12492	4787	6175	20653	17139	3877	43531	20498	12270	56265
	iC5	nC5	22DMC4	CYC5	23DMC4	2MC5	3MC5	nC6	22DMC5	MCYC5	24DMC5	223TMC4	Benzene	33DMC5	CYC6	2MC6	23DMC5	11DMCYC	3MC6	c3DMCYC	It3DMCYC	It2DMCYC	nC7	MCYC6	22DMC6	ECYC5	25DMC6	24DMC6	223TMC5	234TMC5	Toluene	2MC7	3MC7	c4DMCYC	nC8

D924	8451	16851	255	3924	2439	24383	14715	44911	856	27710	2179	182	15580	514	20401	20139	14047	3251	25878	8396	7958	20488	58695	56494	4877	4240	3308	3807	5606	227	18940	27841	31146	5246	42449
D842	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
D841	31613	31714	689	9605	6919	56859	37328	68170	1498	77291	5993	605	47120	919	34413	55163	45105	6049	70166	32000	29353	88219	114553	144094	15915	21607	15634	15393	28822	77	48125	101967	116823	25549	121880
D802	13262	22587	622	4735	4961	49710	30886	78486	1587	54612	4472	23	33035	813	38647	49939	38493	7471	63884	27955	25190	68680	153370	152640	13209	18184	10672	11140	21282	569	24856	102987	112443	22354	153891
D801	2432	0069	88	703	1149	11545	7371	21435	565	12303	1288	155	17878	505	10136	13878	11777	2221	19504	7783	7191	18807	49575	46989	5692	4545	3757	3615	7431	174	20740	36040	38684	8152	55308
D800	4972	21438	142	2015	1556	18069	11124	36836	2696	16145	1808	201	22394	380	17503	23176	13105	3443	27968	9673	9316	22100	76658	76332	7571	4659	4347	5299	8167	67	23701	41323	48696	10203	72413
D756	5402	10667	511	1942	3420	31000	19927	91203	2414	37035	5781	634	34619	2391	40390	66370	34620	9407	79042	22427	21220	52303	323343	228625	22221	15186	16749	20347	22805	349	52657	149927	191992	34791	439879
C1715	95602	250854	4149	46113	40748	486101	286124	947644	13900	481765	39319	4743	1138601	10157	960986	393230	203790	48473	461991	133757	121386	266351	1176033	1648796	69240	97776	53659	63173	82199	5708	880246	451412	154052	85255	877285
C1705	41027	208001	2599	30916	27275	331941	226386	791434	6682	363735	23174	4030	881955	8219	846008	847034	563603	30939	819019	199864	160532	407121	2144461	2091279	73195	271073	86762	95492	118049	11178	2643579	964572	242720	117265	2453420
C1473	3432	6660	22	4113	1891	27179	29544	88550	0	136769	6234	0	83027	1474	93343	149359	46756	21896	249257	197474	192158	438366	601919	519397	84951	146581	33571	42471	232521	4546	182539	444076	627319	117589	582878
C1472	513158	1078195	2780	246974	66299	1166211	915932	2359148	0	1543941	49437	2838	1414900	7765	1252318	805435	267776	64498	1222403	596603	556810	1181387	2495998	1705678	123825	437922	80953	91432	325875	7367	766321	786463	948323	120309	1056009
C1471	278050	742486	1322	176044	23824	973071	784602	2223910	323	2146024	58755	1916	784891	6875	734317	1010105	271156	121699	1613518	1408259	1316310	2553040	3340468	2231750	310754	67779	96924	131868	826541	12476	545930	1211391	1048553	285370	1270570
C1470	581452	1294519	2906	389408	77499	1188430	1061767	3060950	0	1648916	53496	3583	2847729	10320	2030298	1019452	344214	46881	1539088	427734	398164	866385	3647116	2252173	82551	559915	117264	121577	230357	9068	2227438	1118179	164334	103817	1852102
C1469	233635	379404	1244	108393	38563	482539	421690	841034	0	797751	23955	1314	530983	2889	573071	398076	143906	40746	627831	364721	343407	776015	1077448	1020456	86799	266275	48972	57372	229209	5336	248074	490874	391009	92342	573637
C1468	818401	1722929	5122	489570	106345	2052294	1655125	4462517	0	2408116	84290	4486	2494797	13834	2860137	1542819	501786	75101	2268744	701858	650558	1426052	4557135	3142540	127023	722786	152926	166759	359116	12463	1433743	1351206	371093	139798	1781692
C1467	474917	1358678	980	239276	61489	1284815	960820	3208362	0	1508511	43498	1325	4136692	8066	1075681	1051743	253285	31464	1546617	542555	494237	905517	4302291	1772435	89035	604422	140321	136275	286521	8568	4149725	1663098	216886	116174	3268254
C1466	971382	2686237	3380	786094	112187	2957983	2267937	8133985	0	3678022	103539	4037	9548080	25773	3895748	2567840	653686	64131	3756242	878381	810070	1529074	10000000	4504933	130251	1398972	282292	267855	394611	16079	8742651	3004334	387331	161702	5779488
C1465	1196558	2600008	3786	579471	142786	2267821	1728583	5217965	0	2458811	75573	3702	6092301	12822	2636983	1561413	462533	49109	2263386	543569	503221	1036890	5706334	2834614	90940	800801	176626	169698	257145	11004	4871332	1666263	322287	111321	3008788
C1390	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0
C1389	113082	210604	1538	0	46708	294803	229265	460116	0	429871	18592	839	120512	2537	161530	225563	70503	36234	347866	285590	261537	604780	492933	466706	84150	110914	23758	32721	193400	3079	62401	227620	362127	69913	165114
	iC5	nC5	22DMC4	CYC5	23DMC4	2MC5	3MC5	nC6	22DMC5	MCYC5	24DMC5	223TMC4	Benzene	33DMC5	CYC6	2MC6	23DMC5	11DMCYC	3MC6	c3DMCYC	It3DMCYC.	It2DMCYC.	nC7	MCYC6	22DMC6	ECYC5	25DMC6	24DMC6	223TMC5	234TMC5	Toluene	2MC7	3MC7	c4DMCYC	nC8

D2885	6862	386	2221	2761	18787	13183	52873	1004	27588	2980	429	22035	677	26200	25441	18820	4145	32172	13002	12391	35575	148223	104587	7216	11180	5291	5980	10558	8952	62556	46430	18220	5349	186021	0
D2626	7501	15365	483	2134	1790	12362	7415	39545	547	15779	1433	174	3783	358	12248	10183	4784	1494	11110	4955	4562	11735	68748	39098	2293	3785	1399	1828	3183	44	8275	11982	2475	2624	66224
D2595	36759	43322	2150	5611	8003	53887	33019	90242	2633	43886	5293	781	26046	1781	38833	36876	25265	7139	43139	16401	15320	41754	120856	111824	9005	8831	6103	6744	10089	13978	25514	44236	13579	9959	99851
D2472	165	180	13	64	51	417	294	841	30	746	60	25	453	31	751	571	473	113	762	339	331	851	2207	2635	225	218	106	116	0	0	0	230	39	0	0
D2471	3172	2195	186	1174	1280	6930	5441	7232	490	11762	1059	331	5042	307	13894	6588	10858	2714	10650	6350	6110	22170	16678	47789	4971	3430	1997	2231	6370	260	6110	13261	2827	6037	35353
D1385	595958	1055444	11516	0	222786	1291374	702321	2055890	0	1120450	79447	8098	297055	15251	740591	797576	300049	83181	824554	321661	302644	674999	1757874	1710729	111621	157453	75440	82997	132962	4649	86862	529579	572842	85925	554837
D1365	5797	12654	363	0	3494	44591	24816	83160	0	38365	5345	515	22240	1427	28979	63716	23898	7817	67533	21760	20524	49132	158996	143107	13468	12147	10991	11850	16886	79	16991	87462	93297	15773	100817
D1364	100508	217707	8503	0	53121	458063	258889	1049220	0	389869	48869	5080	213658	13219	319845	461418	135735	31760	451893	141932	135534	304431	1374752	1024496	73669	69919	50935	59325	76302	2346	147039	363326	415171	64581	582640
D1335	461452	800296	14151	0	182316	1004955	564294	l655343	0	949440	69578	8631	293092	15520	688593	679974	220472	83961	729612	296911	285277	660559	l614640	1679247	118203	147661	71207	80640	136469	4874	99678	505404	583929	94990	585710
D1313	351958	661085	20688	0	170147	1058997	587529	2544993	0	944988	101115	11342	441902	26625	770259	938656	277863	106732	928245	323577	311929	697897	3682525	2422692	185419	213665	111060	129160	206961	7665	1383005	830669	894599	124735	1662119
D1312	258406	146856	9360	0	89184	422297	277420	122451	0	270838	26002	5961	20455	8326	197081	85512	184987	29721	153759	82302	84163	309621	48925	333469	36277	13598	12806	13309	37278	277	17104	17944	52356	13193	18597
D1291	200217	421189	9028	0	78452	453538	258305	1171081	0	381911	45738	4571	120542	12336	342043	405652	102452	48640	383217	109551	105280	217112	1385452	962672	58899	51875	40978	51718	52603	1230	87322	290217	353295	51108	555092
D1290	104537	220380	7086	0	35707	299757	168276	829083	0	219692	36797	3673	94026	10634	216561	320913	66707	34699	291614	71568	68034	136985	1168100	727803	46875	32772	35036	44356	37410	140	81012	237813	294141	41632	508473
D1289	59498	146041	3653	0	32194	226835	130229	693981	0	193259	30263	2783	103313	8461	186537	292969	78598	32950	289237	76414	72990	146447	1343104	766847	52109	42712	41118	51022	46989	408	146084	343356	392986	54770	900782
D1288	96470	215553	6791	0	38533	313104	176607	877412	0	247979	36637	3730	127127	10187	237965	307062	59628	41615	300621	82200	78546	158404	1257250	790420	48816	40437	33889	43335	42391	106	121548	247708	305775	44760	578157
D1276	45704	93538	4378	0	30431	188824	111036	414372	0	141408	20405	1840	46494	5112	114367	132689	63430	22614	160804	49822	46334	100909	410079	361503	23745	19975	15461	18879	22458	482	18126	93286	117778	17892	114918
D1275	7000	16739	622	0	6876	71264	41884	210766	0	66332	14362	1253	55992	3635	57485	184262	48655	14598	179412	46609	43329	93617	729702	393576	39064	29034	36966	41296	42293	154	84057	332090	327854	45491	675552
D1274	45552	92009	2953	10559	25450	432984	254604	787223	0	429296	49526	6311	347830	9764	242082	648771	297520	45047	713919	301995	283042	684343	1739653	1315354	128525	201564	110217	106272	201611	6949	151332	941268	887111	148247	1192059
D1273	134276	235331	4568	0	60570	465151	259735	722921	0	424697	38661	5060	166134	8040	297777	433597	166379	37231	447217	175473	165827	387425	955606	959877	82710	93598	55007	60490	103959	3696	68240	419138	455596	75179	460709
D1173	94538	116792	2515	0	44669	265428	156943	345932	0	239458	28710	1956	64017	4509	131683	281712	121186	25677	294136	106089	99094	227195	497407	575866	66692	52465	55512	59170	83314	2699	88237	348425	429471	71211	339542
	iC5	nC5	22DMC4	CYC5	23DMC4	2MC5	3MC5	nC6	22DMC5	MCYC5	24DMC5	223TMC4	Benzene	33DMC5	CYC6	2MC6	23DMC5	11DMCYC	3MC6	c3DMCYC.	It3DMCYC	It2DMCYC:	nC7	MCYC6	22DMC6	ECYC5	25DMC6	24DMC6	223TMC5	234TMC5	Toluene	2MC7	3MC7	c4DMCYC	nC8

A2424	29309	31584	39173	37710	49958	3433	12418	3642	1803	6074	4446	4047	2792	2397	2580	1869	1496	1235	1059	1078	754	656
A2364	10955	11762	12202	10051	13694	1021	3152	999	6304	1800	1206	977	877	704	622	557	546	456	386	284	299	248
A2363	10441	11081	11348	9887	11760	815	3448	529	4731	2126	1679	1488	1363	1118	1025	901	799	677	523	416	337	322
A2362	15159	12212	11261	9174	11056	308	3258	268	4401	2078	1659	1451	1314	1216	1032	902	829	745	556	439	371	295
A2313	8193	9372	9393	8411	10220	313	3271	438	4007	1986	1620	1412	1281	1078	938	809	806	646	498	378	336	285
A2284	9535	9092	9244	8369	10504	315	2697	516	4291	1571	1118	955	831	701	617	548	477	417	341	247	0	0
A2283	9767	8105	7605	6518	7520	234	2313	342	3074	1438	1131	1028	927	795	720	660	576	515	401	317	0	0
A2270	7164	6038	5710	5645	6894	467	1723	355	3091	829	657	474	449	404	332	290	262	244	194	130	132	118
A2269	4758	3755	3530	2931	3655	247	1008	141	1423	579	440	361	323	277	229	201	175	144	117	73	71	81
A2268	667.7	5801	5204	4807	5687	352	1660	309	2246	917	759	603	581	510	441	399	343	305	248	194	152	0
A1725	11180	10356	9877	9147	11696	886	2549	748	4116	1247	878	712	614	475	423	401	360	305	297	177	198	0
A1724	7667	9345	10149	9239	13317	1109	2809	803	4708	1401	1028	748	649	537	461	432	389	320	347	228	227	0
A1723	10292	9377	8756	8118	10167	906	2032	610	3480	961	674	483	452	326	291	284	230	195	190	160	151	118
A1712	6716	10711	14121	13577	16470	1007	9630	561	11056	7612	6686	5750	5138	4583	3835	3210	2783	2303	1718	1135	885	647
A1711	14058	14260	14466	12329	17445	1292	3894	1189	7763	1898	1367	984	955	697	695	660	580	516	526	328	302	0
A1710	13293	12853	12896	10628	14397	1003	3610	929	5625	1849	1350	1001	875	608	663	533	543	472	476	332	308	243
A1140	21200	19138	18368	15706	19893	1477	4523	1473	7503	2159	107	1135	103	1086	793	757	641	575	591	404	332	273
A920	1942	4948	9226	12767	20697	1554	6171	1449	11512	3578	2769	2256	2455	2487	2182	2124	1916	1700	1429	1206	835	691
A550	1147	5027	12950	22813	39446	791	16639	1570	24556	9134	5426	4350	3440	3508	2759	2507	2027	2242	1526	995	860	638
A549	227	1161	88	4085	7506	593	1782	553	4562	1038	615	442	373	314	252	221	193	180	159	131	66	0
	C13	C14	C15	C16	C17	Pr	C18	ЧЧ	C19	C20	C21	C22	C23	C24	C25	C26	C27	C28	C29	C30	3	C32

A2892	8557	10251	11842	13362	18016	798	8026	1064	10412	5128	4431	4313	3974	3703	3485	3027	2718	2322	2076	1688	1332	1002
A2884	15745	16910	20934	19120	25405	1332	6341	1034	11907	3396	2466	2210	1769	1602	1611	1227	1105	833	761	738	561	391
A2706	2676	5827	9134	9707	14130	844	3325	802	6319	1659	1394	1291	1043	981	970	817	738	641	636	546	387	340
A2627	7088	8210	10740	10455	14091	1314	6211	899	9007	4920	4892	4969	4581	4524	4245	4029	3631	3264	2887	2444	1871	1341
A2611	2838	4145	6084	7143	10207	499	2607	674	5942	1397	1110	1097	846	798	775	798	788	704	842	775	608	508
A2470	47419	40905	46117	43993	53529	4258	16933	3697	25494	10747	8469	7239	6567	5697	5209	4732	4479	3869	3139	2384	2028	1616
A2469	31823	31541	36703	37679	47857	3300	12321	2748	22824	6064	3966	3047	2679	2202	2029	2030	1831	1592	1172	1017	858	0
A2468	16221	13680	13857	11684	13441	1120	4099	1058	6223	2428	1863	1620	1467	1270	1088	930	819	640	442	330	0	0
A2436	9057	15778	24112	25631	36278	2617	9456	2418	17473	5415	4028	3636	2634	2361	2531	1902	1462	1068	1169	1071	870	497
A2435	20542	30177	42761	44995	59261	3483	14602	4122	27592	7474	5233	4634	3237	2804	3175	2329	1978	1340	1582	1367	1106	645
A2434	10434	18065	29226	30383	44615	3189	11941	2932	22585	5759	4283	3825	2716	2313	2472	1895	1415	1029	1156	991	677	492
A2433	10133	13825	20096	20687	29076	1931	7554	2097	14686	4132	3087	2579	2020	1786	2057	1427	1128	892	992	861	602	413
A2432	16490	14926	13984	13655	17535	1393	4601	1103	7057	2381	1958	1730	1393	1241	1220	831	673	488	392	361	256	176
A2431	25091	22118	24593	21008	25776	2007	6161	1602	10809	2845	1959	1787	1314	1123	1043	746	639	453	561	427	278	204
A2430	14303	22309	32431	32758	45418	2771	11799	3154	20526	5700	4058	3594	2521	2154	2287	1418	1403	931	979	903	616	437
A2429	11053	14391	20169	20166	27760	1872	8217	1821	12850	4977	4004	3455	2860	2598	2701	2085	1642	1288	1379	1086	827	613
A2428	52818	42160	44866	38064	48215	3022	11241	3416	20587	4776	3367	2945	2139	1854	2012	1474	1351	944	1119	606	799	478
A2427	5446	18940	36897	40935	58831	4547	7084	4055	26590	9490	7384	6525	5294	4367	4386	3320	2772	2156	2233	1716	1544	857
A2426	34368	32780	37356	34190	44769	3530	11521	2936	21383	5784	4162	3880	2792	2450	2543	1918	1507	1142	1150	1071	743	582
A2425	21385	26845	35457	34940	46438	3160	11426	3028	20486	5701	4019	3490	2474	2098	2331	1560	1244	896	940	803	589	483
	C13	C14	C15	C16	C17	ŗ	C18	Ρh	C19	C20	C21	C22	C23	C24	C25	C26	C27	C28	C29	C30	<u>3</u>	C32

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C513	10151	14615	16419	16952	15975	4732	15396	0677	15915	13403	10905	10109	8411	7796	6358	6011	4712	4291	3414	2919	2258	2182
C511	3262	8157	13731	18209	20203	7274	22369	12715	25010	23780	20864	19452	17767	16119	12732	12969	10781	9793	7961	7105	5182	4860
C503	7574	10373	11459	10907	9813	3794	9207	6019	9212	7939	6306	5980	4907	4431	3569	3347	2475	2331	1669	1350	965	704
C499	5809	7358	7590	7253	6555	2101	6263	2371	5572	4939	4006	3597	2993	2632	2228	2036	1648	1566	1288	1100	817	566
C495	2676	4647	5769	6193	5948	1922	5865	2937	5585	5030	4176	3827	3179	2874	2394	2265	1863	1703	1416	1269	086	836
B1874	4134	4490	4543	2922	3126	3930	2421	2278	1920	1819	1572	1327	1137	947	837	730	591	551	503	327	297	259
B1873	4309	4058	4231	3115	2931	3588	2297	1967	2373	1783	1485	1236	1039	863	791	684	548	508	442	318	284	208
B2887	2572	3638	3464	3423	3665	1495	2845	1103	3269	2294	1940	1702	1364	1106	891	753	573	425	366	348	227	176
B2122	1404	1950	2315	2136	2330	1478	1863	780	1720	1297	1039	841	692	561	443	363	284	269	230	163	128	0
B2121	1970	2126	2250	1808	1584	930	1279	613	1400	924	777	615	502	406	319	268	188	166	129	89	67	0
B1443	2071	3944	5529	6195	5611	3747	5256	4228	5001	4199	3545	3089	2169	1915	1783	1531	993	996	655	578	342	331
B1393	10008	9573	9616	7796	6644	5395	5894	3900	5411	4053	3332	2542	2048	1602	1176	1088	917	745	671	517	349	328
B1279	2407	2969	3378	2371	2170	1628	1917	1205	1201	970	677	601	485	409	484	316	251	269	258	149	86	0
B1014	5330	5288	4820	3658	3443	1263	2624	860	2284	1783	1343	1041	854	675	501	423	318	300	301	209	190	0
B554	5682	7936	9558	8459	8189	5415	7101	3668	6584	5749	3870	3277	2605	1865	1569	1198	948	736	610	458	412	0
B515	5227	7161	8710	7545	6575	3919	5384	2590	4492	3393	2263	1691	1190	912	731	589	428	377	275	185	0	0
A2898	3980	7913	11991	15456	23413	1118	6444	1322	11473	3930	2998	2655	2199	2008	375	1650	1395	1082	1035	956	766	596
A2897	19295	18796	22973	22106	29212	1814	11347	1385	16462	6682	5587	5395	4524	4323	4197	3713	3192	2881	2285	2124	1808	1305
A2896	15904	15942	17093	18994	26529	1244	7277	1392	12440	4101	3068	2761	2308	2154	2202	1815	1552	1167	1113	1031	783	585
A2895	30812	28845	33834	31975	42396	2245	11566	2136	20784	6123	4424	4207	3329	3130	3137	2510	2250	1768	1715	1446	1138	176
	C13	C14	C15	C16	C17	P	C18	Ρh	C19	C20	C21	C22	C23	C24	C25	C26	C27	C28	C29	C30	<u>3</u>	C32

C1388	14228	17572	18983	17639	16661	7659	16890	12493	16908	13995	11770	10555	8814	8131	6474	5928	4991	4689	3516	3123	1877	1629
C1387	4307	8950	11552	12980	13429	4723	13485	8213	14424	12631	10856	9763	7901	7889	5667	4952	3519	2849	1787	1357	673	613
C1386	4082	7136	9145	9783	9493	3233	9056	6652	9351	8803	7408	6965	5620	5550	4456	4148	3392	3222	2375	2070	1182	1150
C7 25	2875	5102	6000	5652	4993	2286	4524	3430	3711	2664	2123	1788	1086	950	859	106	802	931	773	728	481	488
C122	4292	6995	7861	7154	6157	2743	5608	3999	5001	3230	2706	2281	1498	1333	1162	1266	1117	1290	1119	1046	614	602
C721	12007	18016	21681	20941	18963	5801	12020	7410	10649	6697	8797	1548	7181	7434	6016	1885	5065	4956	3745	3268	1850	1749
C714	5521	7952	8656	8340	<u> 9108</u>	24/2	7195	3302	7113	2380	4616	42.22	3696	3369	2652	2744	2332	21.87	1739	1610	1032	1011
C711	3760	5468	6360	6020	5798	1999	2330	2460	5510	43.63	3683	32,26	2704	2422	1953	1866	1458	1464	1171	1008	602	268
C596	4651	5 626	6133	58.88	5325	2560	4974	1924E	45.24	4315	3419	3250	2712	2533	2120	2032	1512	1505	1090	696	729	638
C589	536	1396	2191	2571	2044	5155	3059	5588	3731	3235	2895	3092	2193	2018	1821	1580	1368	1399	1109	1083	<u>866</u>	821
C582	4671	7680	9343	9948	9405	3271	9445	692.2	9565	8842	7185	7055	5679	5334	4185	4373	3268	2810	1985	1679	1115	1004
CS 79	12002	12711	12320	10727	6096	5406	8137	6759	7396	6945	5464	4923	3925	3520	2864	2797	2020	1878	1478	1326	1/6	887
C575	7219	13304	16893	13564	18072	6447	18606	13940	17560	17109	14065	13.755	11467	10422	8250	8087	6183	5746	4266	3715	2637	2532
C574	4465	8596	11191	10855	10625	7004	10148	8450	10043	8985	7391	7011	5603	5103	4019	3787	2715	2419	1913	1535	1006	698
C566	2 29	841	Z71B	5774	8937	3794	11701	8065	14583	15304	13708	14399	12812	11732	9841	9851	8277	8647	6446	5711	4198	3855
C567	5409	12127	17886	12442	2:1078	7980	22686	16509	23127	20831	16992	16573	13966	12360	10263	8096	7404	7174	4877	4210	2013	2572
C563	18484	24206	27607	81.82	23546	16204	20992	16374	20123	15810	12266	10769	81.79	6855	5779	4874	3505	3082	2358	1848	1207	839
C548	15825	16192	15105	13554	11548	3356	10865	5695	10018	8588	6765	6296	2200	4768	3802	3633	2845	2568	1926	1633	1146	842
C540	921	8	12546	1384	10799	6820	9665	7208	958	•	4	22	7775	532	6214	형	620	208	•	4071	•	3309
CS29	1740	5568	11039	15510	19461	7968	22578	13514	24071	23994	20800	20622	17080	15406	12536	11991	10086	9437	6981	6306	4949	4497
	댢	CI4	닪	CIE	C17	'n	쁞	H	6	ซื	ទ	3	ដ	3	3	8	2	8	ຮ	ខី	ទ	8

24	55	28	72	27	38	51	33	14	31	30	30	26	74	47	71	22	72	60	5	9	1	9
D92	186	285	387	312	380	226	283	331	238	205	198	195	187	207	217	172	15(12(77	84	62	36
D842	3672	3861	4965	3634	4320	4109	3068	5331	2527	2149	2161	2146	2121	2352	2677	1884	1703	1183	901	970	587	470
D841	5854	6289	8055	6005	7051	6993	5457	9051	3089	2977	2634	2702	2812	3148	3973	2974	2783	2088	1654	1794	1088	753
D802	815	2450	5279	5669	7879	4597	5709	6003	5804	5214	5550	5294	5209	5152	5213	3338	2655	1713	1177	979	386	1291
D801	5533	8059	14665	12789	13454	14000	12353	15474	10190	9710	9242	8780	8447	8820	10010	9280	8466	7368	7491	5359	3989	2820
D800	7853	9855	14084	11473	12114	8277	9670	10640	10900	7869	7818	7610	7205	7475	8039	6204	5575	4092	3125	2170	1287	729
D756	7269	8859	9905	7581	8711	3082	5998	4363	6351	4521	4419	4142	3962	4065	3938	3392	3143	2383	1846	1683	852	006
C1715	4960	6206	6556	6041	5938	1772	5629	2183	5668	5003	4595	4406	3912	3635	3189	2905	2376	2055	1607	867	639	543
C1705	5757	7751	7729	7756	7422	1872	6910	2878	6885	6094	5287	4964	4340	3931	3473	3326	3045	3074	2840	2251	1899	1984
C1473	0	274	787	1099	1428	959	1282	1140	1228	1168	977	958	771	662	537	463	364	364	269	276	168	153
C1472	0	263	1030	1361	1656	808	1529	1241	1695	1397	1141	1154	1003	894	744	697	592	615	490	523	392	450
C1471	137	382	1069	1659	2135	1474	1987	1763	1808	1848	1561	1489	995	970	845	684	559	515	366	355	212	0
C1470	0	0	0	780	1563	642	1717	1252	1958	1704	1458	1306	1233	1093	850	748	635	594	464	421	263	277
C1469	0	1016	3651	5360	5954	2710	6274	4981	6923	6036	4869	4511	3464	3664	2906	2846	2396	2493	1965	2119	1679	1852
C1468	0	132	767	1364	1718	713	1636	1238	1890	1583	1370	1384	1191	1069	827	791	666	724	513	550	377	398
C1467	0	311	1022	1349	1504	716	1348	1231	1470	1231	1049	1059	853	969	550	601	405	388	293	295	176	0
C1466	0	0	904	1643	2274	831	2218	1647	2381	2067	1773	1747	1510	1336	1096	983	807	822	608	637	451	538
C1465	0	0	1974	2451	3005	1147	2769	2117	2946	2551	2199	2151	1826	1637	1303	1174	1016	1063	737	758	500	497
C1390	2415	5903	8644	8962	10326	7873	11223	9276	9558	10100	8034	7039	5490	5195	4558	3621	3190	3228	2749	3028	2252	2497
C1389	4227	0269	8454	8145	7899	4883	7902	5928	7086	6115	4955	4145	3212	2967	2284	1967	1509	1386	1106	995	703	758
	C13	C14	C15	C16	C17	ŗ	C18	Ρh	C19	C20	C21	C22	C23	C24	C25	C26	C27	C28	C29	C30	3	C32

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D2885	7943	10509	13650	9843	12688	3723	8166	4704	8627	6337	6499	6756	6464	6794	6629	5472	4698	3029	2510	2437	1438	1050
D2626	4348	4567	5985	6210	9171	995	3802	893	6243	3123	3166	3432	3261	3425	3537	3285	3055	2700	2799	2222	1697	1194
D2595	1733	3447	4917	4598	5988	1952	4244	2296	5107	4139	4428	4580	4622	4959	4991	4495	4168	3441	3326	2575	1847	1448
D2472	4616	7563	9830	7426	8365	4582	5710	5076	5515	4306	4165	3943	3863	3642	3384	2852	2604	1974	1426	1015	987	765
D2471	15786	23301	28660	23549	27983	23230	22205	33264	22667	17432	17542	17295	17715	18434	17690	15810	14309	9392	7537	5787	5202	18482
D1385	12827	18449	22939	17488	21138	9384	15243	12882	16050	11586	11272	10650	10023	10618	9621	8263	7363	5833	4620	4586	1795	1707
D1365	6611	8740	10739	7959	9196	4712	6812	6919	6471	4821	5069	4992	4931	5237	5809	4380	4013	3000	2923	1409	1533	1025
D1364	4602	158	7798	6420	7556	2875	5308	3979	5095	3822	3744	3617	3632	3731	3926	3109	2919	2221	1947	1184	959	897
D1335	2733	3461	4427	3253	3881	2010	2733	2674	2492	1625	1753	1784	1848	1994	2188	1728	1571	1168	1082	563	414	282
D1313	6753	7186	7823	5991	6724	1500	4301	1794	3974	2848	1348	1529	1668	1909	2006	2005	1952	1652	1521	1018	662	645
D1312	4347	6303	8070	5827	7398	5126	4982	6815	1797	1553	1797	2182	2667	3251	4075	3242	3176	2271	2300	1172	1280	3215
D1291	2463	2878	3354	2514	3002	602	1854	805	1850	1454	1393	1306	1248	1233	1162	1051	936	779	691	465	346	267
D1290	3818	4228	4872	3488	4122	816	2419	879	2346	1753	1627	1491	1445	1369	1295	1151	1063	913	729	521	415	300
D1289	5499	5648	6244	4587	4912	1080	3024	1323	2892	2346	1435	1508	1609	1721	1689	1601	1526	1281	1078	714	551	399
D1288	4665	5287	6199	4666	5098	1070	3276	1426	2956	1838	1769	1786	1806	1835	1813	1655	1568	1299	1092	728	550	399
D1276	3989	5027	5386	4044	4750	1499	3223	1952	2990	1748	1538	1491	1502	1573	1640	1552	1367	1200	1188	761	630	511
D1275	4195	4335	5353	3866	4530	1565	3059	2011	2797	1864	1532	1452	1432	1531	1619	1518	1335	1124	1120	680	563	437
D1274	5069	6455	8124	5821	6677	4325	5578	6181	2892	2615	2689	2434	2679	3242	3790	3127	3037	2328	2306	1342	1147	2196
D1273	1879	2440	3128	2325	2853	1495	2029	2081	2156	1164	1107	1054	1149	1264	1391	1256	1131	901	883	501	324	706
D1173	2858	4003	5313	3970	4675	3877	3458	4730	2061	1784	1818	1760	1835	2185	2569	1848	1772	1296	1210	773	639	408
	C13	C14	C15	C16	C17	ŗ	C18	Ч	C19	C20	C21	C22	C23	C24	C25	C26	C27	C28	C29	C30	<u>3</u>	C32