

**CHEMOTAXONOMY: THE ROLE OF PHYTOCHEMICALS IN CHEMOTAXONOMIC DELINEATION OF TAXA**

**Abstract**

Chemotaxonomy is concerned with the systematic study of phytochemical variation between plants. This variation has been essentially used for classification purposes ever since 'folk taxonomies', based on certain obvious plant characteristics which were instinctively employed by mankind centuries ago which included characters such as edibility, taste, colour, smell and medicinal value were founded subjectively on such chemical properties. The growth in the knowledge of the chemical complexity of plants became high from the desires of Europeans for exotic spices and condiments which provoked investigations into their medicinal properties. This Knowledge about the subject was summarised in herbals, and concentrated on information about physiologically active secondary metabolite such as alkaloids and saponins. During the eighteenth and nineteenth centuries knowledge in the field increased, and some taxonomists made use of several chemical characteristics in attempts to delineate plants taxa and to demonstrate their phylogeny. Chemotaxonomy has undoubtedly made a big contribution to taxonomic work in the past and will most certainly continue to do so in future. The valuable information it offers is best used in conjunction with other sources of taxonomic evidence and thus a multidisciplinary approach is required in order to establish a system of classification which reflects natural relationships as accurately as possible.

**Keywords: Chemotaxonomy, Phytochemicals, Primary Metabolites, Secondary Metabolites.**

**INTRODUCTION**

The aspect of taxonomy/systematics which employs the application of chemistry is referred to as chemotaxonomy or chemical taxonomy [1]. Relying on the external morphology alone is not adequate for proper delineation of taxa, hence other branches of study are of considerable value in proper assessment of the systematic status of a taxon and its phylogeny [1]. The taxonomic contributions of chemotaxonomy have made an equally great help to support the ideas of classification and phylogeny [1]. The rise in the study of chemotaxonomy has ushered in the development of sophisticated techniques in chemical analysis which can detect even trace of chemical compounds [2].

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employed by mankind centuries ago which included characters such as edibility, taste, colour, smell and medicinal value were founded subjectively on such chemical properties [3]. As long ago as the first century after Christ the aromatic mints had been recognised and grouped together by Dioscorides [4].

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It is now generally accepted that certain compounds and related substances may be characteristic of certain taxonomic groups [3, 5]. However, the investigation of phytochemicals for chemotaxonomic purposes have been employed at all levels of the taxonomic hierarchy, from sub variety rank [3, 7, 8, 9]. It is thought that when the groups in question show differences in their content, their ability to form a chemical substance was retained by virtue of metabolic processes retained by the group or its ancestors. By implication the pathway of chemical evolution is seen to be established in order to offer

insight to the Evolutionary history of the group, as well as to the understanding of the present –day relationships within and between groups [3, 6]

### **Chemotaxonomic Classification**

The phenolics, alkaloids, terpenoids, and non-protein amino acids are among the important and widely exploited groups of compounds utilized for chemotaxonomic classification [7, 8, 9]. These groups of compounds exhibit a wide variation in chemical diversity, distribution and function [7, 9, 11]. The system of chemotaxonomic classification relies on the chemical similarity of taxon [11, 12].

However, [7] proposed three broad categories of compounds are used in chemotaxonomy:

1. Primary metabolites
2. Secondary metabolites and
3. Semantides

#### **1. PRIMARY METABOLITES**

Primary metabolites are the compounds that are involved in the fundamental metabolic pathways [7]. A greater number of the primary metabolites occur universally and are utilized by the plant for growth and development [14, 15]. These compounds are abundant in nature and are present almost in all parts of plants, hence they are less significant in chemotaxonomic classification [7]. These molecules are useful in chemotaxonomic delineation on the basis of their quantities. For example, carbohydrate sedoheptulose is abundantly present in genus *Sedum* hence, making the accumulation of sedoheptulose in the species of genus *Sedum* a useful chemotaxonomic parameter [7, 14]. Furthermore, [16] reported that the water soluble polysaccharides (WSP) were used as chemotaxonomic markers where the gas liquid chromatographic analysis on WSP from annatto tree (*Bixa orellana* L.) showed hemispherical type contained 38% rhamnose, while conical and ovate types contained 17% and 34% glucose, respectively. Thus, glucose and rhamnose content of WSP could be used to distinguish the three landraces of annatto trees [16].

#### **2. SECONDARY METABOLITES**

Secondary metabolites are the compounds that are used for protection and defence against predators and pathogens in plants and they usually perform non-essential functions in the plants [14]. These compounds are limited in occurrence and hence very useful in chemotaxonomic classification.

[7] Identified the following Secondary metabolites and they include: glycoside, alkaloid, volatile oil, flavonoid, plant phenols and terpenoids.

#### **a. Glycosides in Chemotaxonomy**

Glycosides are the compounds in which one or more sugars are combined with non-sugar molecules through glycosidic linkage [7]. These compounds are grouped as O-glycoside, C-glycoside, N-glycoside and S-glycoside based on the glycosidic linkage, [7]. Due to the common distribution of O-glycosides like rhein, its chemotaxonomic value is low. The *R. rugosa* flavonol glycosides were reported to be important chemotaxonomic parameters for the classification of species in Cinnamomeae [17]. The use of flavonol glycosides as chemotaxonomic parameters was reported to be useful for the identification of *Rosa* species belonging to sections [17]. The C-glycosides like aloin, cascaroside which retain a direct carbon linkage between sugar and non-sugar are reported to be less dominant in nature and they are found in some plants comprising anthraquinone derivatives [18, 19] such as aloin in *Aloe-Liliaceae* [20, 21] cascaroside in *Cascararhamnaceae* [22]. S-Glycosides Sinigrin are exemplified by those produce isothiocyanate on hydrolysis. These compounds serve as chemotaxonomic characteristic of the family Cruciferae, Moringaceae, Capparaceae [23].

#### **b. Cyanogenic Glycoside in Chemotaxonomy**

The cyanogenic glycosides are the compounds that providing defence to plants [14]. Plant species undergo cyanogenesis which is the ability to produce hydrogen cyanide (HCN) by enzymatic hydrolysis of cyanogenic glycosides [24]. Cyanogenesis was reported for the first time in the genera *Beilschmiedia*, *Cardwellia*, *Cleistanthus*, *Elaeocarpus*, *Embelia*, *Mischocarpus*, *Opisthiola*, *Parsonsia* and *Polyscias* [25]. [14] Reported that different amino acid like phenyl alanine, tyrosine, valine, leucine, and isoleucine are precursor for the biosynthesis of cyanogenic glycosides and their presence are restricted to particular families. For example, a cyanogenic glycoside synthesized from leucine commonly occurs in the subfamily amygdaloideae (almond) and maloideae (apple) of family rosaceae [14]. The

glycosides derived from tyrosine commonly occur in the families of the order mangnoliales and laurales [14].

### **c. Glucosinolates in Chemotaxonomy**

Glucosinolates are sulfur-nitrogen-containing plant secondary metabolites common in the order Capparales, [26]. They are mostly differentiated on the basis of alkyl component of glucosinolate compound. For example, *Brassica juncea* (mustard) from Indian subcontinent contain 3-butenyl glucosinolate and allylglucosinolate while those from Asiatic country contain only allyl compound [27, 28], hence querying the ancestry of Indian species, because that is the hybrid of *B. nigra* (allylglucosinate) and *B. compestris* (3-butenyl glucosinate).

### **d. Alkaloid in Chemotaxonomy**

Alkaloids are heterocyclic compounds containing Nitrogen [29, 30]. But, few non-heterocyclic alkaloids are also present [30]. The parent base compound in the alkaloid is the diagnostic character when it is considered for chemotaxonomic relevance. For instance, the indole alkaloids contain indole as the parent base and more than 2,500 indole alkaloids were isolated mainly from three plant families, Rubiaceae, Loganiaceae and Apocynaceae. The indole is formed from two building blocks secologanin and tryptamine or tryptophane through a single precursor, strictosidine [31]. This suggests a relationship between these families [31, 32]. Other indole alkaloids like physostigmine obtained from *Physostigma venenosum* (family Leguminosae) [33], yohimbine from *Rauwolfia serpentina* (family Apocyanaceae) [34] and *Corynanthe yohimbe* (family Rubiaceae) [35] and Vinblastine from *vincarosea* (family Apocyanaceae) [36]. The Pyridine and Piperidine alkaloids like Lobeline obtained from *Lobelia inflata* family Lobeliaceae [37]. Nicotine obtained from *Nicotiana tobaccum* family Solanaceae [38]. Anabasine obtained from *Nicotiana gluaca* Family Chenopodiaceae [39]. The presence of these alkaloids serve as chemotaxonomic diagnosis [7]. Anabasine occurs in tobacco, where it is formed from lysine and nicotinic acid, while in the legume and chenopod species this can be synthesized from two molecules of lysine [37]. Similarly, the alkaloids like isoquinoline alkaloids, tropane alkaloids, indole alkaloids etc have also been diagnostic in taxonomic delineation of taxa [39].

### **e. Plant Phenol in Chemotaxonomy**

Polyphenols are among the most widespread class of metabolites in nature [7]. It is estimated that 100,000 to 200,000 plant phenols exist [41] and some 20% of the carbon fixed by photosynthesis is channelled into the phenylpropanoid pathway, thus generating the majority of the natural occurring phenolics [42]. Flavonoids are largest group of phenolic compounds. They are mostly found in the vacuole of higher plant and absent in lower plant [7]. Different classes of plant phenols include flavones, flavanones, isoflavanones, isoflavonoids, anthocyanidins and chalcones. All flavonoids have common biosynthetic origin and therefore it possess the same basic structural element. For example, 2- phenylchromone skeleton. They may be present in many classes depending on degree of oxidation of pyran ring which may be open and cyclize into furan ring, e.g. 2-phenyl benzopyrylium: anthocyanin and 2-phenyl chromone: flavone, flavanol, isoflavone [42]. A chemotaxonomic study of practically all the species of the genus *Aloe* showed that flavonoids occur as major compounds in 31 out of a total of 380 species investigated [43].

#### **f. Terpenoids in Chemotaxonomy:**

Terpenoids are found mostly in higher plants belonging to the following families: *Myrtaceae*, *Lauraceae*, *Rutaceae*, *Lamiaceae*, *Asteraceae*, *Apiaceae*, *Poaceae*, and *Cupressaceae* [1]. The different types of terpenoids are discussed as follows:

**Monoterpenes:** These are acyclic or bicyclic, optically active hydrocarbons with boiling point ranging from 140° – 180°C which sometimes constitute 90% of essential oil as found in *Citrus* oils [1]. They are most abundantly present. The predominance of monoterpene (-)-enantiomers in the emission of some European *Pinus* and *Abies* species was explained by [34]. A monoterpene lactone nepetalactone has been reported to be the principal odour constituent of Catmint *Nepetacataria*, a plant which has a peculiar attraction for the domestic cat because of its odour [34].

**Sesquiterpenes:** These are common constituents of the essential oils of higher plants and contribute to the pharmacological properties of the plants [44]. Structurally, just like monoterpenes they are hydrocarbon alcohols and ketones. Examples are  $\beta$ - bisabolene, longifoline, farnesol, santalol, sinesals, cedryl acetate [1]. Recently by GC broad chemodenes were distinguished by the presence of carvone and presence of absence of dill apiole [45].

**Diterpenes:** They vary greatly in structure which depends strictly on their biogenesis [1]. Diterpene containing drugs have different applications such as anti-hypertensive, co-carcinogenic, anti-oxidant, hallucinogenic properties etc. Diterpenoids occur in the *Garryaceae* where they play diagnostic role in the delineation of taxa in this family due to the difficulty to classify them on the basis of morphological grounds [1]. The chemotaxonomy of *Sideritis* species was evaluated and its acetone extract was shown to possess insecticidal and ascaricidal activity. The extract was found to contain linearol, lineal, isolinearol and siderol [46].

**Triterpenes:** The taxonomic/diagnostic relevance of these compounds is feasible when they are combined with other constituents such as phenols and flavonoids [1]. They have therapeutic potential in many fields such as cystostatics, insecticides, antiinflammatory agent etc and they play role in confirming the relation of *Pittosporaceae*. This family has more affinity with *Araliaceae* than *Saxifragaceae*. The triterpenes of 5 *lithocarpus* species were examined and they were of friedo unrearranged oleanane group viz, friedelin, friedelan, 3- $\beta$ -taraxerol and  $\beta$ -amyrin. Glutinol was also present except in Lharlandi where frielan 2- $\alpha$ , 3- $\beta$  diol was found. In addition, 3 new cycloartane triterpene, lithocarpolone, lithocarpdiol and 24- methylene cycloartane 3- $\beta$ , 21 diol were found in *L. polystachya* [47].

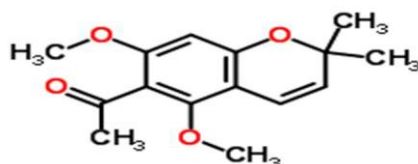
**Tetraterpenes:** These are carotenoids which interfere with photoxidation processes such as treatment of photosensitization linked to porphyria, also ingredient of tanning pills and food technology industry. A qualitative and quantitative examination of carotenoids of pure cultures of four marine micro algae including *Chroomonas salina*, *Vaucheria sassilis*, *Cacolithus* and *Huxleyi* was reported to have played diagnostic roles where the latex contained a new natural carotenoid and fucoxanthin [1, 47].

**Polyisoprenes:** In the delineation of the species *Erigeron bonariensis* (L.) which is a common weed which is traditionally used in urine problems (Asteraceae) [1] and belongs to the genus (*Erigeron*) with about 390 species of flowering plants [1]. Intercontinental plant inventions resulted in a number of taxonomic problems especially in distinguishing it from *Conyza*. From the investigation on the basis of chemotaxonomy it was concluded that the phenolic content and caffeol derivatives present in it has a closer relationship to *Erigeron* than species of *Conyza* [48].

#### **g. Essential oils in Chemotaxonomy:**

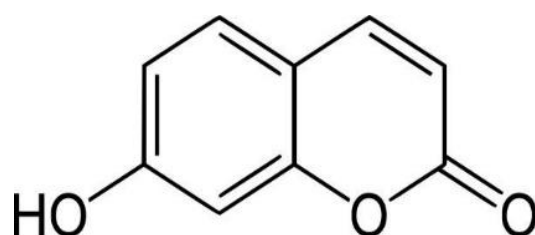
The impact of essential oils on scientific plant classification have been reported by [49] in the families discussed below:

**Rutaceae:** Rutaceae family plants are chemically characterized by the synthesis and accumulation of essential oils, furanocoumarins, anthranilic acid derived alkaloids and limonoids [1]. Cneoraceae have represented a taxon in certaesedis for a long time. Their oil cells and the chemical nature of their bitter principles and of their 2- methylchromones leave absolutely no doubts about their intimate rutaceous meliaceous- simaroubaceous affinity. Prenylation of aromatic compounds is common in this family; examples of this tendency are furano and dimethyl pyrano coumarins and a number of essential oil constituents evodionol [1, 7].



**Fig 1: Evodionol**

**Umbelliflorae:** *Umbelliferae* family constitute the plants with furano and dimethyl pyrano coumarins and essential oils which tend to contain phthalides; for example Ligustilide ferulol type monoterpenoids and acetylinic compounds like falcarinone [1]. [7] reported that essential oils in schizogenous ducts are highly characteristics of Araliaceae and umbelliferae together with other chemical characters they accentuate the overall similarities of these two families and the need for reclassification of other 5 families often included in umbelliflorae, cornaceae and allied families are iridoid producing taxa which seem to have affinities with *Dipsacales* rather than with *Araliaceae* and *Umbelliferae*. Prenylation of aromatic compounds is also common in this family such as Umbelliferone (coumarin).





## Fig 2: Umbelliferone

**Verbenaceae and Labiatae:** Many members of these two families are highly aromatic and yield essential oil [1]. The group of aromatic plants does not produce iridoid glycosides; at the most some non-glycosylated compounds like myodesertal and myodesertin may be present in essential oils; the non- aromatic group of Labiates is characterised by iridoid glycosides like ajugol, galiridoside, harpagide, lamiol and others. The main taxonomic importance is two chemical groups of *Labiatae* which coincide with classification proposed by [49] for this family. Iridoids are insecticidal and insect deterrant and well described for steam volatile iridoids.

**Piperales:** Overall presence of oil cells and isolation of aporphine type alkaloids from roots of *Piper auritum* and stems of *Piper sanctum* [48] confirms the affinity between Magnoliales and Piperales.

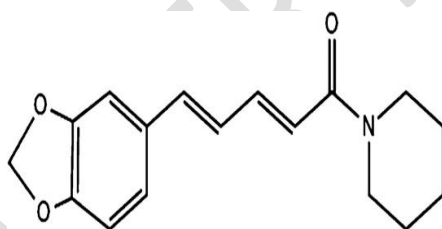


Fig 3: Piperine

**Malvales:** In this family there is a restriction of oleoresin and a widespread of mucilage cells and cavities [7].

**Myrtaceae:** Myrtaceae family consists of plants with essential oil that is rich in tannin [7]. Methylated, prenylated and acetylated phloroglucinol derivatives occurs frequently in their essential oils; production of this very characteristic of acetogenins (Torquatone, a phloroglucinol derivative; present in some species of *Eucalyptus* ) represent a chemical trend of this family [7].

**Dipsacales:** All members of *Dipsacaceae* produce iridoid or secoiridoid glucosides and lack essential oils except the genus *morina* whose species does not produce iridoids but have essential oil in oil cells [1].

### 3. SEMANTIDES

These comprise DNA, RNA and proteins. Because each of these is so intimately connected with genetic characteristics, many researchers consider them to be of immense taxonomic value, as they potentially are. However, the time equipment and know-how necessary for their effective analysis often limits their usefulness.

[9] Mentions three (4) main methods used in plant protein taxonomy. Viz;

- a. Electrophoresis
- b. Amino-acid sequencing
- c. Systematic serology

Electrophoretic techniques enable proteins to be 'fingerprinted' by establishing their relative size, charge and isoelectric point by separating them in variable gel mixtures across a voltage gradient [9]. Protein profiles produced via electrophoretic separation and subsequent staining have been used in various systematic studies investigating polyploidy taxa, as well as interspecific, intraspecific and population levels. Particular care and expertise are required in the use and interpretation of protein profiles [4].

It was reported by [4] that amino-acid sequencing attempts to establish the variation in the normal order of amino-acids in a single homologous protein through a variety of organisms. This analysis relies on the fact that a particular protein may vary to a certain extent without altering its essential function. One molecule used extensively for this purpose is cytochrome c, in which 79 out of the approximately 113 amino acids vary interspecifically, but alterations of even one of the other 34 terminates the functioning of the molecule. Generally, the number of differences parallels the relational distance between the organisms in traditional classification, but anomalies do present themselves, interfering that a measure of protein structure is not an infallible guide to degree of kinship.

Certain assumptions are made when analysing the results of this technique:

It is assumed that:

- The molecule has evolved via the minimum number of mutations

- No convergent evolution or back-mutation has occurred
- Different positions on the molecule are equally susceptible to substitution.

These assumptions weaken the evidence when the arrangement of a number of homologous proteins yields conflicting evidence [9, 51]. Some researchers have pointed out that to apply cytochrome patterns in chemo-systematics requires that one should take into account the quantitative and qualitative effects of the growth conditions on the cytochrome content [52, 53]. [9] Suggests that in order to prevent interpretative mistakes, results from a wide range of proteins, preferably studied by a number of different techniques, and should be pooled rather than placing total reliance on the screening of a single protein.

Synthetic serology is an immunological technique relying on the relative specificity of the immune reaction, and the fact that the degree of cross-reactivity is proportional to the level of relationship that exist in the organisms.

In plant serology, antisera to antigens from various taxa are raised in animals, using various plant extracts [4], and then the antisera can be used as a usual test alongside other plant extracts. The degree of coagulation that the other extracts cause in them is used as a degree of their resemblance to the original antigen. Refinements in the technique have made this method more specific than it was previously, and serology has been extensively used throughout the taxonomic levels from above family to below species, yielding many valuable data [9].

Nucleic acids have not yet been used very extensively in plant systematics due to the complexity of their analysis. Most techniques are of relatively recent origin, and so the data accumulated thus far are limited [4]. Theoretically these characters should be able to solve many phylogenetic problems, firstly because each organism has DNA with a unique base sequence, and secondly because the theory of evolution is based on the premise that related organisms should show similarities in their DNA which are not shown by unrelated species [49].

The most useful technique in this regard at present is DNA hybridisation in, which DNA double helices are induced to unwind. And then allowed to recombine with each other as well as similarly treated DNA from other species. This results in some hybrid double helices being

formed, the number and fidelity of recombination's theoretically depending on the compatibility of the two DNA base sequences. Some useful results have been obtained which shows the potential value of this method, but techniques have not yet been perfected. [4] Point out that variable results have been obtained depending on experimental conditions. Some evidence has suggested that *in vitro* replication of the DNA template is affected by factors such as temperature, and the absence of regulatory phenomena or specific factors that are present *in vivo* [51], and it seems well possible that this might just as well apply to recombination also.

Other techniques have been used to investigate DNA and RNA, but results, according to [6], are of limited application. He suggests that advances in gene cloning and genetic engineering may lead to more extensive use of nucleic acid characters in taxonomy, but a potential drawback to their extensive use in phylogeny is that live material is often a prerequisite.

Chemical variation is of considerable taxonomic value in several ways:

1. Confirmation or support of putative classifications derived from other sources of taxonomic characters, such as morphology.
2. Resolution of problems where relationships based on other evidence are ambiguous or conflicting.
3. Providing evidence to suggest more natural positioning of anomalous taxa, as well as to separate taxa. Often the presence of anomalous taxa in a group is accentuated by their chemical peculiarities.
4. Detection of confirmation of hybridization.
5. Providing additional on/off characters for numerical taxonomy by their presence or absence in taxa.

However, as with all other taxonomic characters, chemical variation must constantly be subject to critical appraisal of techniques and interpretations. Two major problems that appear to need addressing are the lack of standardization of the methodology and the inadequate sampling of groups.

## **Conclusion**

Chemotaxonomy has undoubtedly made a big contribution to taxonomic work in the past and will most certainly continue to do so in future. However, given the lack of fossil evidence and the need for live material in some analyses it seems that its contribution to phylogenetic classification must perforce remain limited. The valuable information it offers is best used in conjunction with other sources of taxonomic evidence and thus a multidisciplinary approach is required in order to establish a system of classification which reflects natural relationships as accurately as possible.

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