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- 9 DECEMBER 1966

# **Terpenoid Precursors of Hydrocarbons**

## from the Gasoline Range of Petroleum

Abstract. 2,6-Dimethyloctane and 2-methyl-3-ethylheptane were isolated from petroleum. These hydrocarbons which are present in relatively large amounts appear to be derived from the monoterpenoids.

The presence in petroleum of hydrocarbons that have skeletal structures or large fragments of the skeletal structures of certain compounds that are also in vegetable and animal products is well established (1, 2). Most, if not all, of these hydrocarbons have been isolated from petroleum distillates with a relatively high boiling point.

We have isolated two C10-branched paraffins that appear to be derived from the terpenoids. These hydrocarbons, which are present in relatively

large amounts, are 2,6-dimethyloctane and 2-methyl-3-ethylheptane. The finding of a large amount of 2,6-dimethyloctane was not unexpected since higher homologs of this series have previously been shown to be important constituents of petroleum (1). This hydrocarbon is probably formed from the acyclic monoterpenoids, though it could also be derived by splitting acyclic terpenoids which have a higher molecular weight. However, the finding, in large amount, of 2-methyl-3-ethylheptane

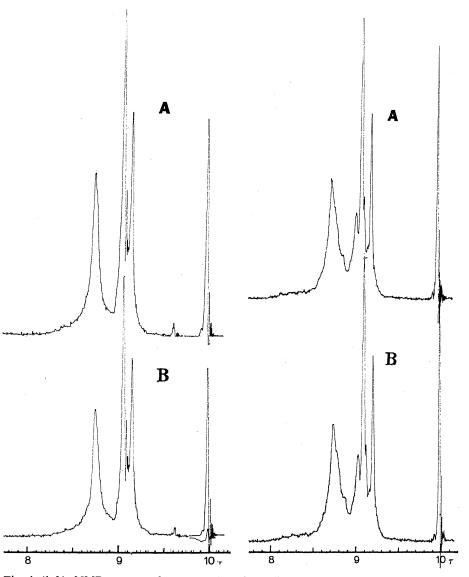
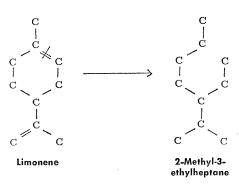


Fig. 1 (left). NMR spectra of two samples of 2,6-dimethyloctane. (A) Synthetic; (B) from petroleum. Fig. 2 (right). NMR spectra of two samples of 2-methyl-3-ethylheptane. (A) Synthetic; (B) from petroleum.

was unexpected since paraffins with this type of branching have not heretofore been found as major components of petroleum.

Limonene, which is abundant in many essential oils, such as in the rind of citrus fruits, in pine needles, and in turpentine, is the probable, although not necessarily exclusive, precursor. We suggest that 2-methyl-3-ethylheptane is derived from limonene by saturation of the double bond in the side chain and by rupture of the double bond in the ring:



These hydrocarbons were isolated by preparative scale gas-liquid chromatography of distillate fractions in the boiling range of 159.5° to 161.5°C of the branched paraffin-cycloparaffin portion of the reference petroleum of the API Research Project 6 (3). They were tentatively identified from their physical and spectrometric properties. These identifications were subsequently confirmed by direct comparison of their infrared and nuclear magnetic resonance spectra with those of synthetic samples (Figs. 1-4). The amounts were determined by analyzing the distillate fractions in the 159.5° to 161.5°C range by using both capillary and 0.25inch (0.63-cm) packed columns with polar partitioning liquids (silicon nitrile and diethylene glycol succinate). On these columns the two compounds are well separated, with 2,6-dimethyloctane having the lower retention time. The maximum concentration of 2,6-dimethyloctane occurred in fractions that boiled at 160.4°C; that of 2-methyl-3ethylheptane in fractions that boiled at 160,9°C. 2,6-dimethyloctane was found to constitute 0.50 volume percent and 2-methyl-3-ethylheptane 0.64 volume percent of the crude petroleum. For comparison, the corresponding value for all other branched paraffins that boiled in the range from 155° to 165°C (a total of 49 are theoretically possible) was 0.44 volume percent.

Because these two hydrocarbons appear to be derived from different precursors, the amounts present and their ratios may be expected to vary from crude to crude depending in part on the content of each precursor in the source materials from which the different crudes were generated. If subsequent investigations establish such a variation, the information may be valuable in several ways. For example, the petroleum industry frequently needs to

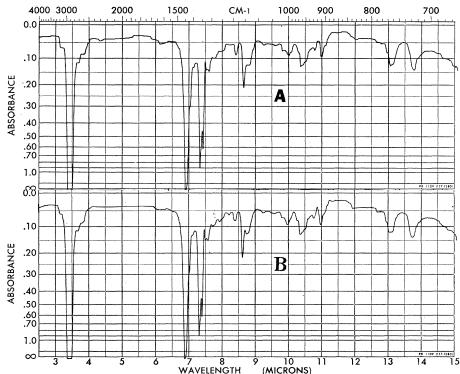


Fig. 3. Infrared spectra of two samples of 2,6-dimethyloctane. (A) Synthetic; (B) from petroleum.

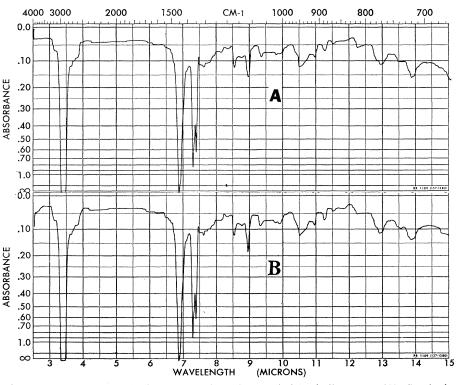


Fig. 4. Infrared spectra of two samples of 2-methyl-3-ethylheptane. (A) Synthetic; (B) from petroleum.

know whether petroleum from wells separated by several miles comes from the same or from different sources. A knowledge of the content of these two hydrocarbons in the given wells may aid in solving this problem. With the information now available an analysis for these two compounds can be made by gas-liquid chromatography, using only retention times for identification provided samples of the pure hydrocarbons are available for calibration.

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### Cystinuria: Genetic Heterogeneity and Allelism

Abstract. Studies of four stoneforming cystinuric subjects from three unrelated pedigrees indicated that each was heterozygous for two of the three described mutant genes producing cystinuria (I, II, III). Their genotypes were I-II, II-III, I-III, and I-III, respectively. These doubly heterozygous patients were phenotypically indistinguishable from cystinuric homozygotes of genotype I-I, II-II, or III-III. The data provide the first direct evidence that all of the known mutations responsible for the genetic heterogeneity in cystinuria are allelic.

Results of definitive studies (1, 2) on the genetics of cystinuria from 27 pedigrees indicated that the occurrence of cystinuria followed classical Mendelian laws for autosomal recessive inheritance. Homozygous affected subjects excreted large quantities of cystine, lysine, arginine, and ornithine in their urine. Renal, ureteral, and vesical cal-

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culi composed of cystine were formed because of the very limited solubility of cystine. The findings in heterozygous individuals were heterogeneous. In some families urinary dibasic amino acid excretion among heterozygous individuals (parents and children of affected subjects) was normal, whereas in other families, modest to marked excesses of cystine and lysine were excreted by all heterozygotes. Thus Harris suggested that cystinuria was, in fact, more than one disease entity. We have reported that there are three types of cystinuria (3). Type I was characterized by the absence of mediated intestinal transport mechanisms for dibasic amino acids in homozygotes and by normal urinary cystine values in heterozygotes. Type II differed most significantly in that heterozygotes excreted markedly increased quantities of cystine. In type III cystinuria, intestinal transport of all dibasic amino acids was retained by homozygotes, and heterozygotes excreted cystine in slight excess. Quantitative determinations of lysine, arginine, and ornithine in these heterozygotes have confirmed the differences noted in cystine excretion. Thus we have demonstrated statistically significant differences among type I, II, and III heterozygotes for each of the four dibasic amino acids measured (4).

In the first nine pedigrees studied, our results were in agreement with those of Harris (1), Crawhall (5), and their co-workers in that all heterozygotes within a single pedigree showed remarkably similar values for the aforementioned urinary amino acids. That is, there was no example in these nine pedigrees of more than a single type of heterozygote within a single family. Recently, however, we have found three unrelated families in which two types of heterozygotes have been demonstrated in a single pedigree. These results form the basis of this report.

Three individuals, G.F. (31-year-old male), J.P. (25-year-old male), and B.B. (35-year-old female), showed recurrent renal or ureteral cystine stone formation. Each excreted in the urine great excesses of cystine, lysine, ornithine, and arginine, the values being in the range noted for cystinuric homozygotes. However investigation of their families revealed (Figs. 1 and 2) that G.F.'s father (F.F.) showed a urinary amino acid pattern typical for a type III heterozygote (Fig. 1) while his mother (M.F.) and son showed a normal urine pattern characteristic of type I heterozygotes. These results were confirmed by analysis of three separate urine samples from each subject, and were strengthened by finding other type I and type III heterozygotes within the pedigree (Fig. 2). These data represent the first unequivocal demonstration of heterogeneity within a single cystinuric family. Similar, but not identical, results were encountered in the other two families (Fig. 2). J.P.'s father (H.P.) was a type II heterozygote (Fig. 1) while his mother (T.P.) was a type I heterozygote. Analyses for the four amino acids in urine samples from each of B.B.'s five children revealed that three were type II heterozygotes and two were type III heterozygotes (Fig. 2). Representative results from this pedigree also appear in Fig. 1. These findings suggested that J.P., B.B., and G.F. were not homozygous for one of the mutations producing cystinuria but, rather, were each heterozygous for two of the three mutant genes, their genotypes being I-II, II-III, and I-III, respectively.

These findings necessitated additional studies (Table 1) of intestinal amino acid transport in the affected probands -G.F., J.P., B.B., and E.M. (a paternal first cousin of G.F., once removed, with cystinuria and renal lithiasis). None of the patients from the three pedigrees were able to transport significant quantities of lysine, arginine, or cystine. The results were identical to those noted for type I homozygotes (3). However, oral tolerance studies with cystine did reveal subtle differences in both G.F. and E.M. in that there were small, but perceptible rises in plasma cystine (Fig. 3), results which were midway between those of type III homozygotes and type I homozygotes (3).

Such results indicate that the genetic heterogeneity in human cystinuria is even more complex than supposed. The pedigrees (Fig. 2) indicate that subjects heterozygous for two different mutations causing cystinuria are phenotypically indistinguishable from documented homozygotes, and thus the question of allelism must be considered.

Let us assume that cystinuria, like other disorders characterized by autosomal recessive inheritance, results from defective synthesis of a specific membrane polypeptide which catalyzes transport of the dibasic amino acids in gut and kidney. We can make certain pre-