

## APPLICABILITY OF INFRARED ALIPHATIC STRETCHING REGION FOR CHARACTERISATION OF OILS OF THE SAME GENETIC TYPE

## J. Stevanović<sup>1,2</sup>, A.R. Rakitin<sup>1</sup>, K. Stojanović<sup>2</sup>

<sup>1</sup>STC-NIS Naftagas, Upstream Laboratory, Serbia, <sup>2</sup>University of Belgrade, Serbia

The work focuses on the applicability of infrared (IR) aliphatic stretching region for characterisation of oils of the same genetic type underlining IR spectroscopy's obvious advantage as a robust and inexpensive analytical technique capable of simultaneously collecting information on all organic matter (OM) components, irrespective of their chemical nature, molecular weight or physical state. Paraffinic crude oils (56 samples) from the Turija-Sever oil field (southeastern part of the Pannonian Basin, Serbia) were studied. Our previous investigations based on detailed interpretation of biomarkers and aromatic hydrocarbons showed that the oils are similar, originate from mixed aquatic/terrestrial sources and were generated from Tertiary source rocks in an early stage of oil window (vitrinite reflectance  $\sim 0.65 \,$ %). However, certain slight differences among the oils were observed. They are reflected in a higher contribution of algal OM to the group I oils which were formed in reducing environment (western part of the field), compared to the oils from eastern part of the field (group II) with lower contribution of algal OM, formed under dysoxic conditions (Stevanović et al., 2019).

IR spectra of Turija-Sever oils are dominated by stretching, bending and rocking vibrations of aliphatic methyl and methylene groups. Stretching region 2800–3000 cm<sup>-1</sup> was decomposed into five Lorentzian bands and CH2/CH3 branching factor calculated from the ratio of deconvoluted asymmetric stretching peaks (Painter et al., 1981). Definitive 3–4 cm<sup>-1</sup> red shift between crudes with the least and most methyls per methylene group was found for both symmetric and asymmetric CH<sub>2</sub> stretching modes, whereas CH<sub>3</sub> asymmetric stretching mode only fluctuates by  $\pm 1$  cm<sup>-1</sup>. By analogy with chain-melting phase transitions in phospholipid membranes it was suggested that the observed peak behaviour is characteristic of the onset of gauche rotamer formation in mostly trans polymethylene fragments with increasing degree of methyl substitution (Mantsch, McElhaney, 1991). Interestingly, a cross-plot of the CH<sub>2</sub>/CH<sub>3</sub> ratios calculated from the corresponding symmetric and asymmetric peak absorbances after deconvolution exhibited an absence of correlation. Perusal of deconvoluted absorbances reveals that while both peaks of methylene are closely proportional in magnitude in all spectra, the methyl peaks are in disarray. To our knowledge, this fact has hitherto received little attention in the literature. Although in the original spectra of Turija-Sever oils CH<sub>3</sub> symmetric stretching is poorly resolved from the adjacent CH<sub>2</sub> feature, it is still possible to read the intensity of the former from the inflection at 2868–2870 cm<sup>-1</sup>. When symmetric vs. asymmetric CH<sub>2</sub>/CH<sub>3</sub> peak absorbances ratios from the original spectra were plotted, correlation becomes apparent. Test runs of the fitting program on synthetic spectra obtained from the sum of five ideal Lorentzians using different initial approximations for the fit parameters confirmed high algorithm sensitivity to the intensity and width of the symmetric methylene peak notwithstanding the fact that its contribution to the aliphatic stretching region is the lowest among all vibrational modes and in this study amounts to only 6-7 % in terms of the area. The findings allude to the necessity of introducing additional vibrational modes in order to accommodate for the diversity of steric and chemical environment of CH<sub>n</sub> groups in oil as noted earlier for coal.



Cross-plot between the normal to isoprenoid alkanes sum ratio,  $\Sigma(n-C_{10} - n-C_{40})/\Sigma(i-C_{13} - i-i)$ C<sub>20</sub>), derived from whole oil gas chromatography (GC-FID) and the IR CH<sub>2</sub>/CH<sub>3</sub> branching factor segregates Turija-Sever oils into two identical groups (I and II) previously established based on biomarkers and aromatic hydrocarbons. The group I oils showed values of the  $\Sigma(n-C_{10} - n-C_{40})/\Sigma(i-C_{13} - i-C_{20})$  ratio varying between 4.58 and 6.28, whereas CH<sub>2</sub>/CH<sub>3</sub> branching factor, ranged from 2.61 to 3.15. Group II is characterized by greater values of both parameters. The  $\Sigma(n-C_{10} - n-C_{40})/\Sigma(i-C_{13} - i-C_{20})$  ratio increases from 7.70 to 11.65, associated with  $CH_2/CH_3$  in range 3.06 – 3.88 (Fig. 1). Although higher branching factor is usually associated with lower OM maturity (Lis et al., 2005), enhanced CH<sub>2</sub>/CH<sub>3</sub> ratio for group II oils indicate that in the case of uniform OM maturity greater average length of polymethylene fragments results from a higher content of long chain *n*-alkanes signifying an increased impact of terrigenous OM (land plant waxes). Furthermore, generally high  $CH_2/CH_3$  ratios (>1.63) for both groups of oils are consistent with low maturity. More detailed inspection of the cross-plot reveals a markedly more pronounced correlation between the IR and GC-FID parameters for the second ( $R^2 = 0.69$ ) compared to the first group of oils  $(R^2 = 0.02)$ . This result is in agreement with a higher contribution of aquatic OM to group I oils enriched in methyl- and dimethylalkanes, originating mainly from cyanobacteria and Botryococcus braunii race A. Since the  $\Sigma(n-C_{10} - n-C_{40})/\Sigma(i-C_{13} - i-C_{20})$  ratio derived from GC-FID did not include methylated alkanes with the exception of regular isoprenoids, whereas IR spectra are sensitive to all methyl moieties, variation of the CH<sub>2</sub>/CH<sub>3</sub> branching factor for the group I oils supersedes the *n*-alkanes/isoprenoids/ ratio. For the group II oils, enriched in terrigenous OM, isoprenoids are the main source of aliphatic CH<sub>3</sub> groups, thus improving the correlation.



*Figure 1.* Ratio of normal to isoprenoid alkane sums from vs.  $CH_2/CH_3$  branching factor. Dotted lines represent linear fits.

## References

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