

Microstructure Characterization of Polyolefins by Automated Cross-Fractionation Chromatography (CFC).

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Introduction

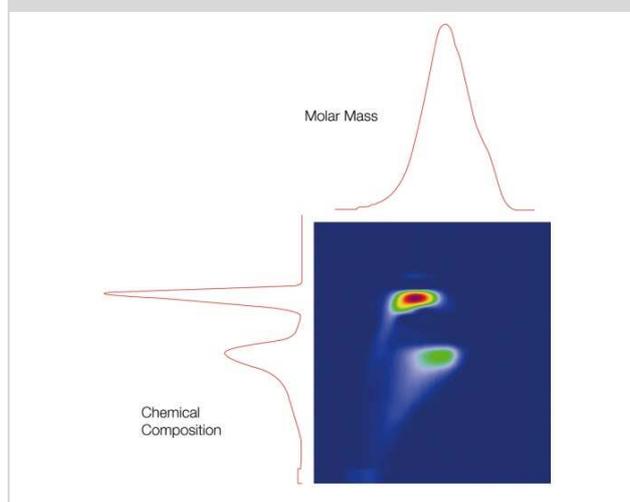
Polyolefin technology has made remarkable progress in the last decades through the development of new technologies which allow more control of the polymer structures by the product designers, so high value products are produced with varying and increasing microstructure complexity. Polymer characterization provides information about the molecular structure which builds a bridge between polymer properties and polymerization conditions.^{1,2}

In most industrial polyolefins there are essentially two molecular parameters of interest: the MMD (Molar Mass Distribution) and the CCD (Chemical Composition Distribution), which are manipulated to impart desired properties to the final product. The MMD tells how much material of a certain molar mass is present. The CCD tells how much material of a certain composition, or comonomer content, is present.

Separate information on MMD and CCD, which are routinely determined by SEC/GPC (Size Exclusion Chromatography, Gel Permeation Chromatography) and TREF (Temperature Rising Elution Fractionation) or CRYSTAF (Crystallization Analysis Fractionation)³ respectively, although important and in many cases sufficient, are not enough to fully characterize a Polyolefin resin and the full bivariate distribution is required.⁴

Hyphenated techniques, such as SEC-FTIR and TREF-LS that carry out the fractionation only along one molecular axis and determine some average of the other, have become popular in the last years,^{2,5} although they result in some information loss, the extent of which depends on the method employed and on the shape of the CCD x MMD surfaces of the materials (Figure 1).

Figure 1: Contour plot of the CCD x MMD and individual distributions as measured by TREF and GPC/SEC.



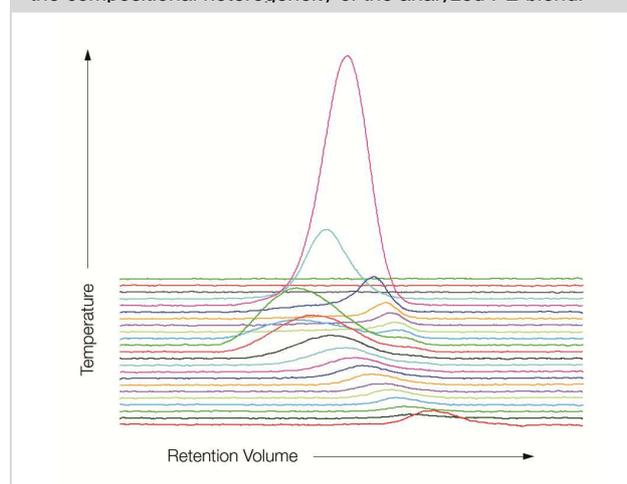
Thus, it is generally preferable to access the full CCD x MMD, which has been traditionally a challenging and time consuming task.⁶

With the introduction by Polymer Char of the CFC automated analytical-scale instrument for polyolefin cross-fractionation⁷ it is now possible to fully characterize complex comonomer / molar mass distributions on the order of hours instead of days and without the need of intermediate manual steps in a bench top instrument. Very high resolution can be achieved in both dimensions by using optimized analytical columns, analysis conditions and sample sizes.

Experimental

The instrument is based on a high resolution analytical TREF apparatus combined with a dedicated GPC/SEC columns oven and equipped with five vessels for sample preparation so up to five can be analyzed sequentially. Polymer detection is made through an IR4 infrared detector for maximum sensitivity in polyolefin applications, which also provides excellent long term baseline stability.

Figure 2: GPC/SEC chromatograms collected at 23 different fractionation temperatures, showing both the molar mass and the compositional heterogeneity of the analyzed PE blend.



The polymer sample is placed in solid form (40–200 mg) into a stainless steel vessel, where it is dissolved in a proper solvent, o-DCB or TCB, at temperatures of 140°C-160°C for 60-90 minutes under stirring. The instrument works by loading the polymer solution into the TREF column where it is crystallized by cooling down the TREF oven and then eluting the fractions in step-wise temperature increments towards the GPC columns.

A series of chromatograms are collected at the different fractionation temperatures (Figure 2). Those chromatograms describe the molar mass distribution of every compositional fraction. The relative area of the individual chromatograms reflects the compositional heterogeneity while their retention time and shape relates to the molar mass dimension.

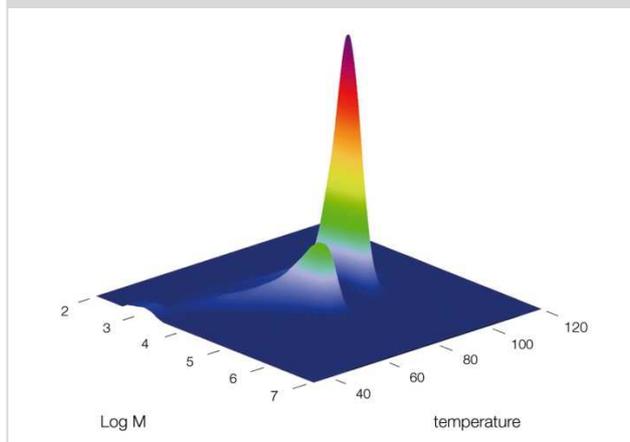
Experimental conditions may be varied to optimize resolution and signal quality depending on the chemical composition distribution, by setting any number and arrangement of temperature fractionation points as required: evenly distributed in all the range every 2°C or 3°C for a typical Z-N LLDPE sample having broad CCD, or concentrating more fractionation points every 1°C in a narrow range around 85°C - 95°C for HDPE which tends to have quite narrow composition distribution. Sample size is also optimized according to the fractionation conditions in order to prevent overloading of the SEC/GPC columns while keeping good enough signal from the IR detector.

An SEC/GPC columns calibration curve is established so that MMD and molar mass moments are calculated at every elution temperature. The relative weight fraction is obtained from the area under each chromatogram and it is combined with the individual fractions MMD to obtain the full CCD x MMD distribution for the sample. All the data processing and calibration is performed within a dedicated and comprehensive software package which additionally provides the TREF profile and MMD for the whole sample.

Results and discussion

A synthetic blend of two single site catalyst polyethylene-co-octene copolymers having different densities and molar masses was analyzed automatically using the CFC instrument. The full CCD x MMD is directly measured with high resolution in 12 hours by elution of 24 TREF fractions into the GPC columns.

Figure 3: MMD x CCD surface plot of a two-component PE blend.



The bivariate distribution is presented in a tridimensional surface plot (Figure 3), which can be processed in several ways within the software. The two components are observed in the graph, which also reveals clearly their distinct position in the molar mass / chemical composition plane. In this case more comonomer was added to the higher molar mass component of the blend, as

highlighted by the lower temperature peak which appears at a higher value in the log M scale.

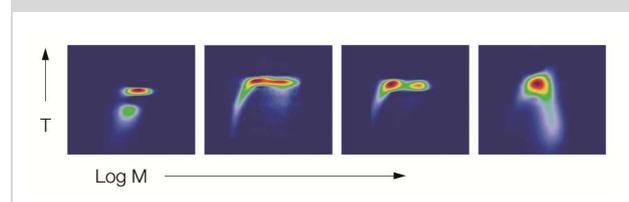
The TREF profile is recovered from the fractions relative weights using a smoothing fitting routine to obtain the differential weight fraction versus elution temperature.

The two dimensional data is also processed to obtain the molar mass distribution of the whole sample by integration over the temperature dimension providing all the distribution averages as well as overall polydispersity. Additionally, the software allows the integration of data across specific temperature ranges in order to study the MMD at different comonomer levels.

The microstructural characterization of different types of polyolefin products can benefit from the automation of this powerful technique:⁸ ZN-LLDPE, polypropylene impact copolymers, dual reactor products or polymer blends and also high density polyethylene with low comonomer levels.

The result of this cross-fractionation analysis can be presented as contour plots which map the molecular structure of polyolefins (in terms of comonomer and molar mass). Families of related or competitive products are easily compared in this way (Figure 4) helping the product designer identifying the relevant structural features and assisting to establish basic property-structure relationships.

Figure 4: Contour plots of different PE products showing distinct microstructural features.



Conclusions

A new integrated benchtop instrument is introduced, which enables the direct measurement of the CCD x MMD surface plot in polyolefins on the order of hours, and in a fully automated way. This method is capable of describing the resin microstructure without any information loss. Applications are found in all types of polyolefins when accurate knowledge of their microstructure is required.

References

- (1) Z. Zhang, *Macromol Symp.* 282 (2009), 111-127.
- (2) W. W. Yau, D. Gillespie, *Polymer* 42 (2001) 8947-8958.
- (3) B. Monrabal, in "Encyclopedia of Analytical Chemistry", R. A. Meyers, Ed., John Wiley & Sons Ltd., 2000.
- (4) S. Nakano, Y. Goto, *J. Appl. Polym. Sci.* (1981), 26, 4217.
- (5) W. W. Yau, *Macromol. Symp.* 2007, 257, 29-45.
- (6) A. Faldi, J.B.P. Soares, *Polymer* 42 (2001) 3057-3066.
- (7) A. Ortin, B. Monrabal, J. Sancho-Tello, *Macromol. Symp.* 257 (2007), 13-28.
- (8) C. Li Pi Shan, D. Gillespie, L. Hazlitt, *Ecorep* 2005. Lyon.