

Test Report

Molecular weight measurement of Polymer by GPC (VISCOPLEX® 12/10301)

manufactured by

Evonik Resource Efficiency GmbH
Oil Additives

Date: 2018-04-26

Tested by

Evonik Technology & Infrastructure GmbH

Product Line Analytik

1. Testing Laboratory

Adress:

Evonik Technology & Infrastructure GmbH

Kirschenallee

64293 Darmstadt

Germany

Person in Charge: Dr. Ania Niklaus

Analyzer: Jana Ernst

Signature:____

Date: 2018-04-26

Tel: +49 6151 18-6364; Fax: +49 6151 18-846364; mail: jana.ernst@evonik.com

2. Test substance:

VISCOPLEX®12/10301 supplied by Evonik Resource Efficiency GmbH; Oil Additives

CAS No: 2181814-52-6

3. Analytical conditions:

(1) Eluent: Tetrahydrofuran (eluent is continuously distilled and circulated by pump)

(2) GPC column:

5 SDV Columns 8 x 300 mm resp. 8 x 50 mm (company PSS at Mainz) 1 Solvent-Peak Separation Column 8 x100 mm (company Shodex)

No	Туре		Dimension	Serial no.
Precolumn	SDV	10μ	8 x 50 mm	7040721
1	SDV LinL	10μ	8 x 300 mm	2062903
2	SDV LinL	10μ	8 x 300 mm	6082302
3	SDV 100Å	10μ	8 x 300 mm	0071401
4	SDV 100Å	10μ	8 x 300 mm	0070508
5	KF-800D		8 x 100 mm	2007012

(3) Detector: RI: temperature 35 °C

(UV: Wavelength 239 nm, not used)

(4) Column temperature: 35 °C

(5) Flow rate: 1 ml/min

(6) Injection volume: 100 μl

(7) Sample concentration: 2 g/l

(8) Data system: PSS WinGPC-Software

4. GPC calibration

(1) Standard sample: PMMA, PSS (Mainz) or Polymer Laboratories

(2) Injection volume: 100 μl

(3) Standard concentration: 1 g/I (for Mw > 10^6 : 0.5 g/I, Mw > $2*10^6$: 0.25 g/I)

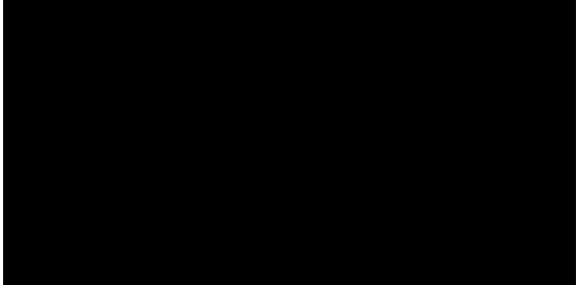
(4) Test date for calibration: 2017-12-01

(5) GPC Chromatogram of Standard Sample:



	No	Sample	Comment
	1	RI 1100er HP, PMMA Mischung I (2 8 12a 15)	
_	2	RI 1100er HP, PMMA Mischung II (4 10 13 16)	
_	3	RI 1100er HP, PMMA Mischung III (6 12 14 17)	
_	4	RI 1100er HP, PMMA Mischung 1-7	
_	5	RI 1100er HP, PMMA Mischung 3-9	
_	6	RI 1100er HP, PMMA Mischung 5-11	
_	7	RI 1100er HP, PMMA Mischung 21-23	
_	8	RI 1100er HP, PMMA Mischung 22	
	9	RI 1100er HP, PMMA18 mmg20113 660	
	10	RI 1100er HP, PMMA1 PS F-850 TS-31 11.200.000 0,25 g/l	
	11	RI 1100er HP, PMMA2 5911/7/1 4.960.000	
	12	RI 1100er HP, PMMA3 mm7086 3.250.000	

Figure 1a. GPC chromatograms of PAMA standard samples



Elutions Volumen [-]

Sample: PMMA18 mmg20113 660
Material: PMMA/THF/30°C at 633nm

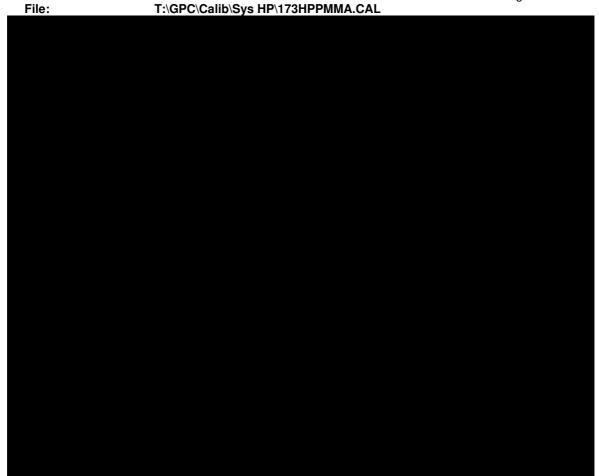
Figure 1b. Raw chromatogram of the lowest standard sample including system peaks and Internal Standard (The blue-marked line is the start marker of the next inject.)

(6) Calibration plot:



WINGPC Report Calibration

Page: 1



Calibration Parameters:

Fit Characteristics: Internal Standard:

Fit Type: PSS Poly 5 Name: o-Dichlorbenzol 0.2μ l / 100μ l (IP)

Date of Printing: 2018-04-26



Figure 2. Calibration plot



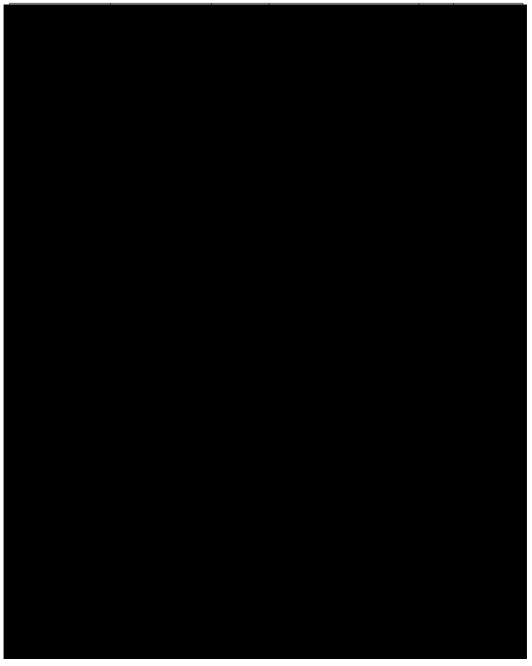
WINGPC Report Calibration

D:\AP\GPC\Calib\Sys HP\173HPPMMA.CAL

Polymer Standards: Poly(methyl methacrylate)

Calibration Table:

File:



Date of Printing: 2018-04-09



Page: 2



WINGPC Report Calibration

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Polymer Standards:

Calibration Table:

File:



5. Test method

The molecular weight results (number-average and weight-average molecular weight and distribution) were measured by GPC (Gel permeation chromatography) in dependence on DIN 55672-1

6. Test result

Table 2. Molecular weights, polydispersity, % content below 1000 g/mol and % content below 500 g/mol of VISCOPLEX ® 12/10301

Mn		Mw	Мр	Mw/Mn	content below	content below
[g/r	nol]	[g/mol]	[g/mol]		500 g/mol [%]	1000 g/mol [%]

Mn: number-average molecular weight

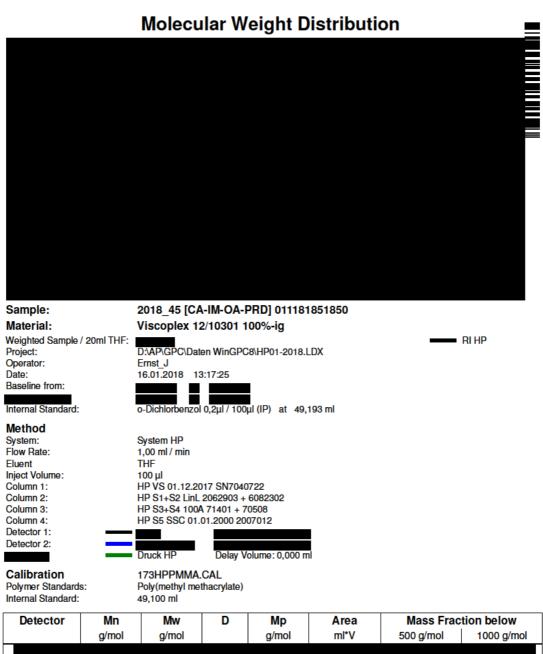
Mw: weight-average molecular weight

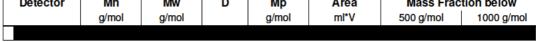
Mp: Peak average molecular weight

Mw/Mn: polydispersity

Sample test date: 16. January 2018 (see Date line on Fig. 3)

Page: 3





Date of Printing: 2018-04-10



Figure 3. GPC Chromatogram of VISCOPLEX® 12/10301



Sample: 2018_45 [CA-IM-OA-PRD] 011181851850

Material: Viscoplex 12/10301 100%-ig

Figure 4. Raw chromatogram of VISCOPLEX® 12/10301 including system peaks and Internal Standard (The blue-marked line is the start marker of the next inject.)

Table 3. GPC slice table

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^{*} The used GPC software produce two slice tables, one with elution volume, molar mass, detector signal (height) and integral distribution, the other with molar mass, differential distribution (area), and integral distribution. The scaling of both tables is different. The molar mass scaling is logarithmical and the elution table is normal. Even though the number of data points in both tables is

similar, the pairs of elution volume / molar mass correlate not exactly. Therefore we used the first table (see above) and we added the column "100-Integral [%]", which is manually calculated on base of Integral [%].