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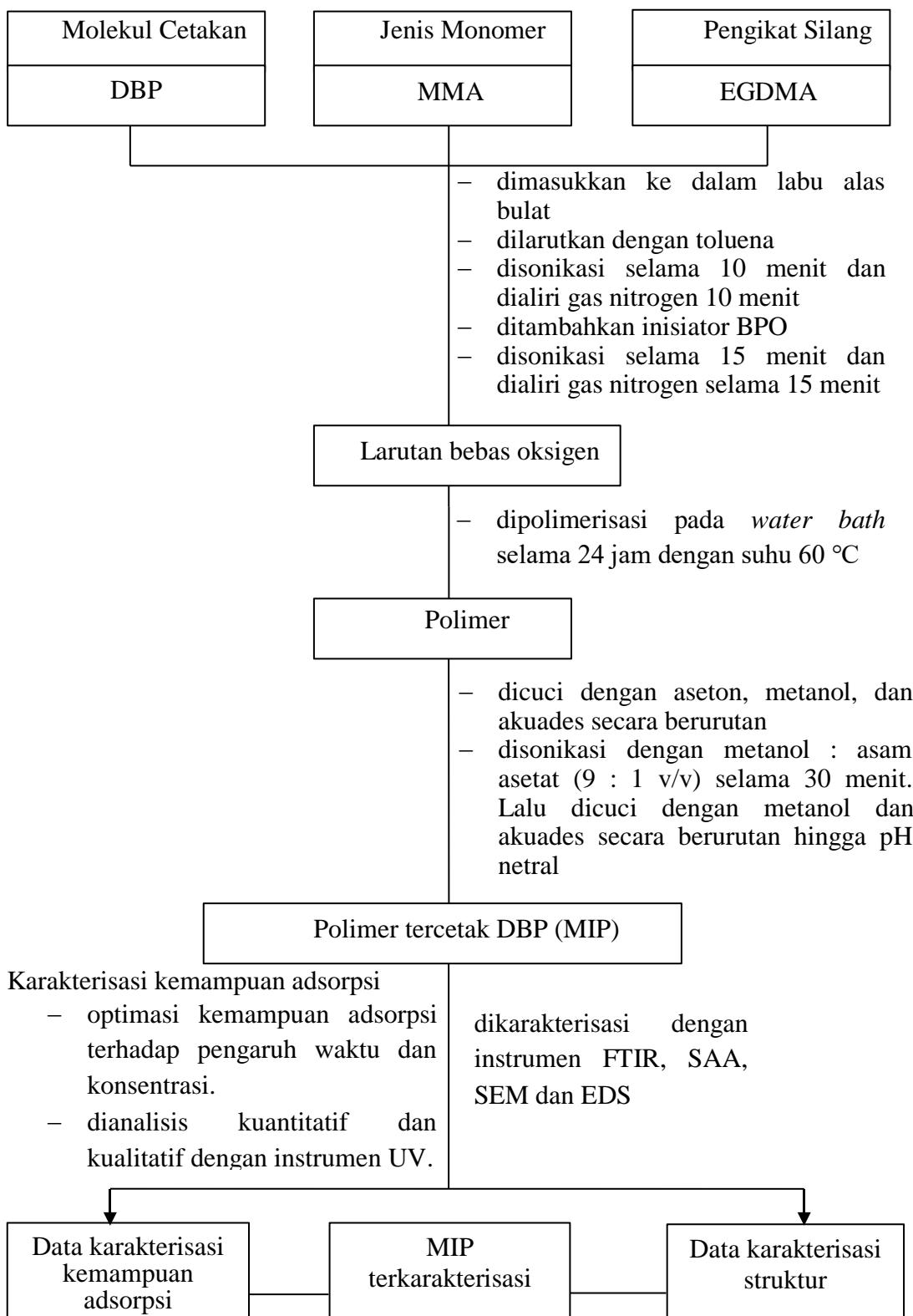
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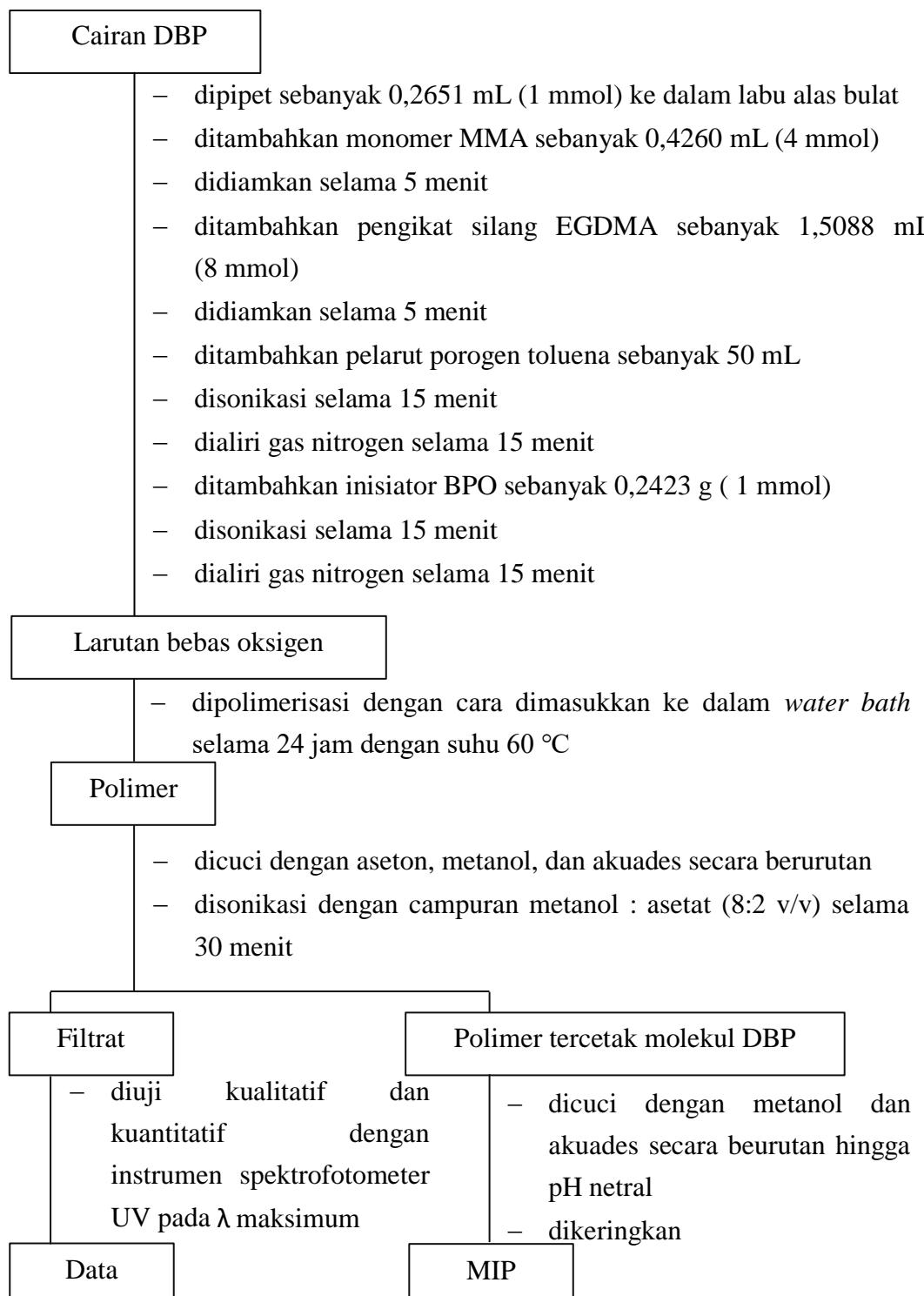
Lampiran 1. Skema sintesis Poolimer Bercetakan Molekul DBP menggunakan Metode Polimerisasi Presipitasi



Catatan: Sintesis NIP dibuat dengan metode yang sama dengan MIP, tapi tanpa DBP dan proses ekstraksi

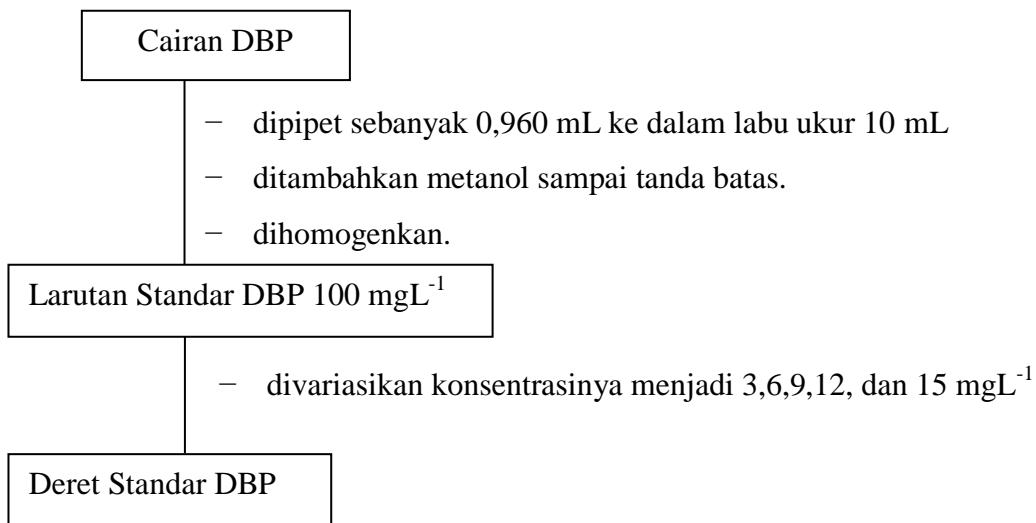
Lampiran 2. Bagan Alir Prosedur Kerja

1. Sintesis MIP DBP dan NIP

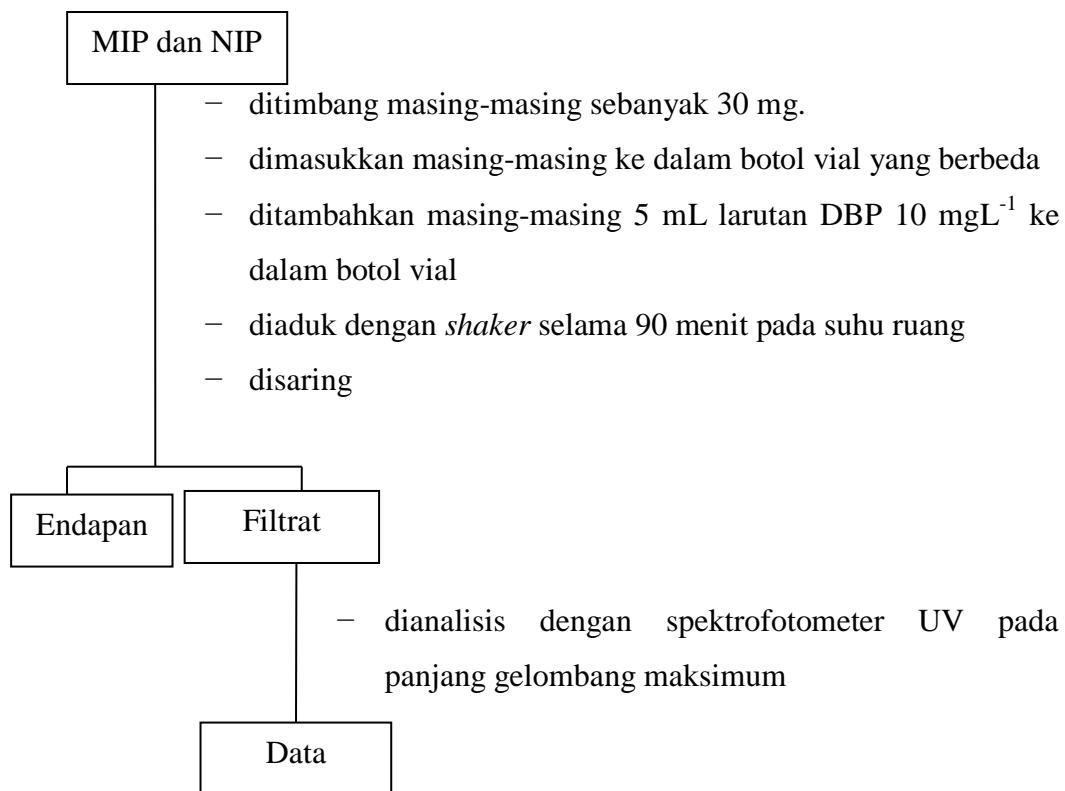


Catatan: Sintesis NIP dibuat dengan metode yang sama dengan MIP, tetapi tanpa menggunakan molekul cetakan DBP

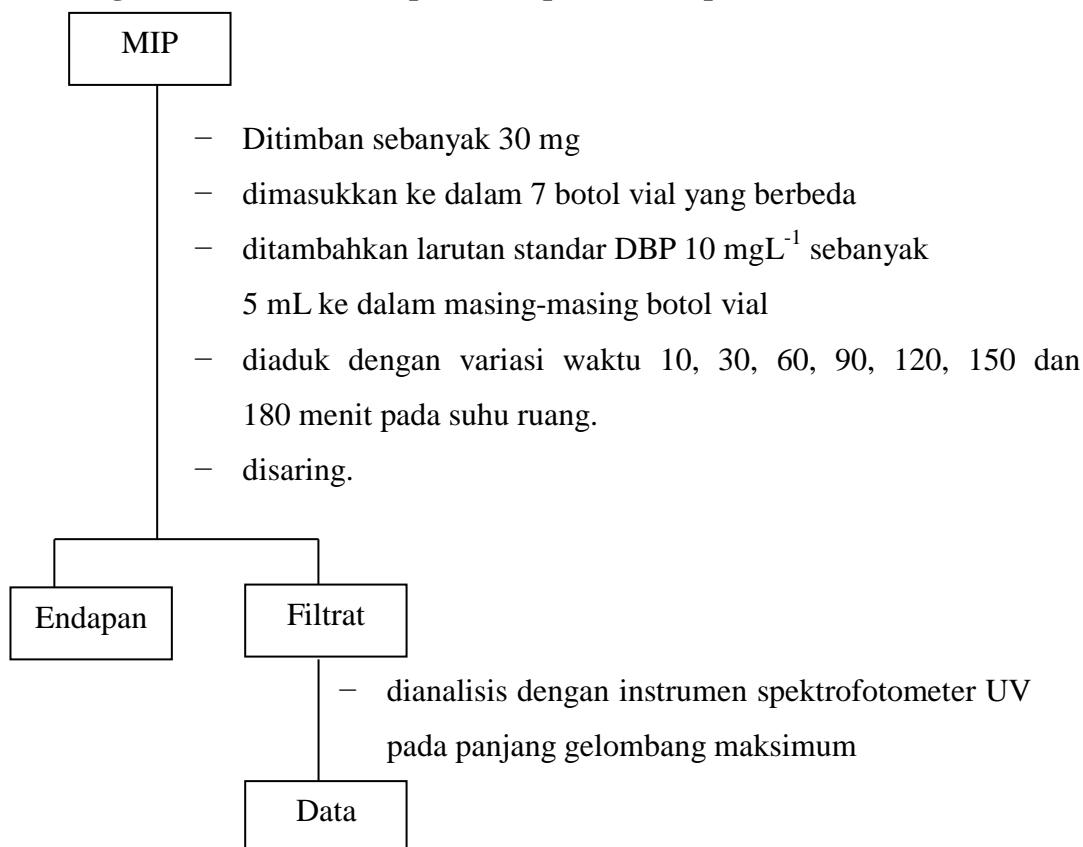
2. Pembuatan Larutan Standar DBP 100 mgL^{-1}



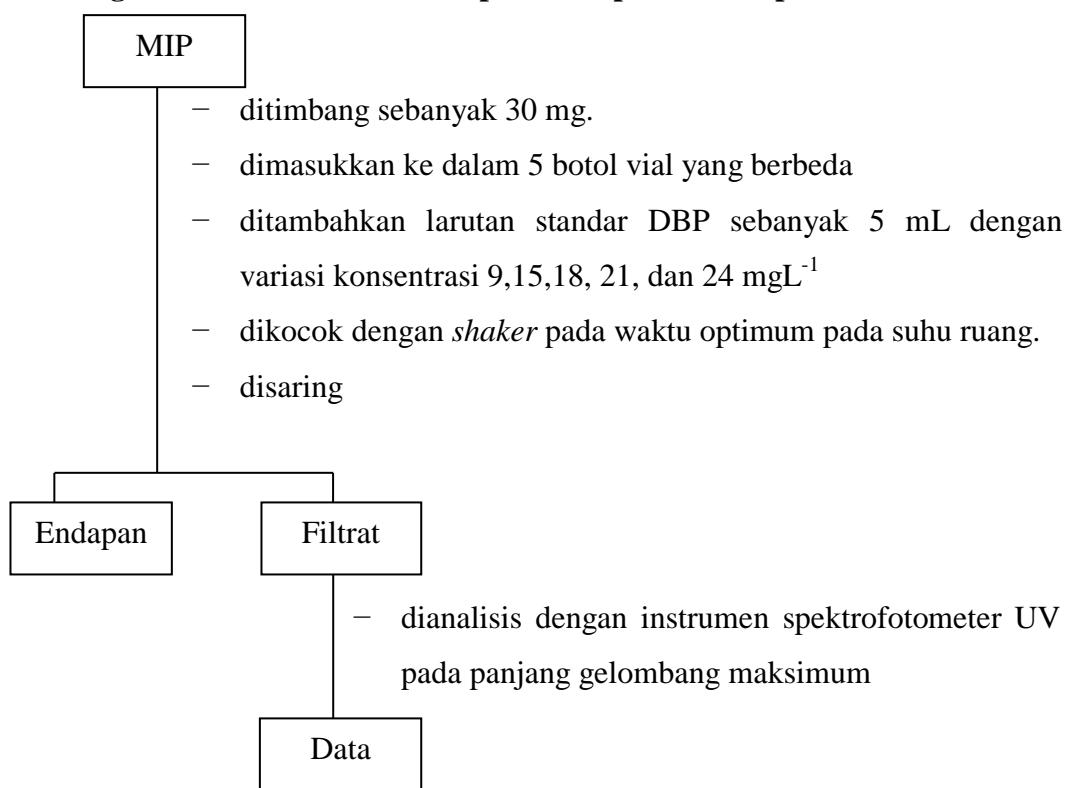
3. Uji Kemampuan Adsorpsi MIP DBP dan NIP



4. Pengaruh Waktu terhadap Kemampuan Adsorpsi MIP DBP



5. Pengaruh Konsentrasi terhadap Kemampuan Adsorpsi MIP DBP

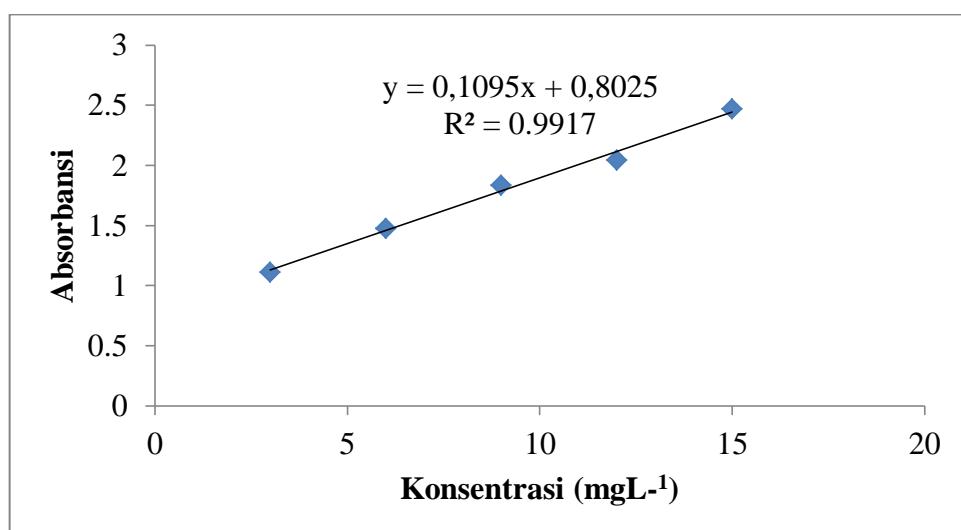


Lampiran 3. Data Spektrofotometer UV-Vis

1. Data absorbansi larutan standar DBP

No.	Sampel	Konsentrasi (mgL^{-1})	Absorbansi
1	DBP 1	3	1,112
2	DBP 2	6	1,478
3	DBP 3	9	1,835
4	DBP 4	12	2,043
5	DBP 5	15	2,472

2. Kurva hubungan antara absorbansi Vs konsentrasi larutan standar DBP



3. Data absorbansi kemampuan adsorpsi MIP dan NIP

No	Sampel	Absorbansi	$q_e(\text{mg/g})$	$\Delta q_e(\text{mg/g})$
1	Adsorpsi DBP oleh MIP	1,575	0,4909	0,2512
2	Adsorpsi DBP oleh NIP	1,740	0,2397	

4. Data absorbansi adsorpsi DBP oleh MIP terhadap pengaruh waktu

No.	Sampel	Waktu (menit)	Absorbansi
1	DBP	10	1,695
2	DBP	30	1,671
3	DBP	60	1,66
4	DBP	90	1,634
5	DBP	120	1,577
6	DBP	150	1,55
7	DBP	180	1,596

5. Data penentuan kinetika adsorpsi orde satu semu dan orde dua semu

No.	Waktu	$C_e \text{ (mgL}^{-1}\text{)}$	$q_t \text{ (mg/g)}$	$q_e - q_t$	$\log (q_e - q_t)$	t/q_t
1	0	0	0	0,52892	-0,27661	0
2	10	8,151	0,30822	0,2207	-0,6562	32,4444
3	30	7,932	0,34475	0,18417	-0,73478	87,0199
4	60	7,831	0,36149	0,16743	-0,77617	165,979
5	90	7,594	0,40107	0,12785	-0,89329	224,402
6	120	7,073	0,48782	0,0411	-1,3862	245,991
7	150	6,826	0,52892	0	0	283,597
8	180	7,247	0,4589	0,07002	-1,15481	392,239

Catatan:

q_t adalah q_e pada waktu t

q_e adalah q_t pada waktu optimum

6. Data absorbansi adsorpsi DBP oleh MIP terhadap pengaruh konsentrasi

No.	Sampel	Konsentrasi (mgL^{-1})	Absorbansi
1	DBP	9	1,005
2	DBP	15	1,385
3	DBP	18	1,592
4	DBP	21	1,796
5	DBP	24	1,969

7. Data persamaan isothermal Langmuir dan Freundlich

No.	Sampel	Konsen trasi (mgL ⁻¹)	C _e (mgL ⁻¹)	q _e (mg/g)	log C _e	log q _e	1/C _e	1/q _e
1	MIP_DBP	9	1, 8493	1,192	0,26701	0,0762	0,54074	0,83908
2	MIP_DBP	15	5,3196	1,613	0,72588	0,20774	0,18798	0,661981
3	MIP_DBP	18	7,21	1,798	0,85794	0,25487	0,1387	0,55607
4	MIP_DBP	21	9,0731	1,988	0,95775	0,29838	0,11022	0,50306
5	MIP_DBP	24	10,653	2,225	1,02747	0,34723	0,09387	0,44954

Lampiran 4. Perhitungan

1. Nilai konsentrasi adsorpsi DBP oleh MIP dan NIP

$$y = 0,1095x + 0,8025$$

a. Adsorpsi DBP oleh MIP

$$y = 1,575$$

$$y = 0,1095x + 0,8025$$

$$1,575 = 0,1095x + 0,8025$$

$$x = \frac{1,575 - 0,8025}{0,1095}$$

$$x = 7,0548 \text{ mgL}^{-1}$$

b. Adsorpsi DBP oleh NIP

$$y = 1,74$$

$$y = 0,1095x + 0,8025$$

$$1,74 = 0,1095x + 0,8025$$

$$x = \frac{1,74 - 0,8025}{0,1095}$$

$$x = 8,5616 \text{ mgL}^{-1}$$

2. Nilai Kemampuan Adsorpsi DBP oleh MIP dan NIP

$$q_e = \frac{(C_o - C_e) V}{W}$$

$$\text{Diketahui: } C_o = 10 \text{ mgL}^{-1} \quad W = 0,03 \text{ g}$$

$$V = 0,005 \text{ L} \quad C_e = \text{Konsentrasi setelah adsorpsi}$$

a. Kemampuan Adsorpsi DBP oleh MIP

$$q_e = \frac{(10 - 7,0548) 0,005}{0,03}$$

$$q_e = \frac{0,014726}{0,03}$$

$$q_e = 0,4909 \text{ mg/g}$$

b. Kemampuan Adsorpsi DBP oleh NIP

$$q_e = \frac{(10 - 8,5616) 0,005}{0,03}$$

$$q_e = \frac{0,007192}{0,03}$$

$$q_e = 0,2387 \text{ mg/g}$$

3. Nilai konsentrasi adsorpsi dan kemampuan adsorpsi DBP oleh MIP terhadap pengaruh waktu

Waktu (menit)	y (absorbansi)	x (konsentrasi) (mgL^{-1})	q_e (mg/g)
10	1,695	8,1507	0,3082
30	1,671	7,9315	0,3448
60	1,66	7,8311	0,3615
90	1,634	7,5936	0,4011
120	1,577	7,0731	0,4878
150	1,55	6,8265	0,5289
180	1,596	7,2465	0,4589

Contoh perhitungan konsentrasi adsorpsi dan kemampuan adsorpsi DBP oleh MIP terhadap pengaruh waktu:

a. Konsentrasi Adsorpsi DBP oleh MIP 10 menit

$$y = 0,1095x + 0,8025$$

$$y = 1,695$$

$$y = 0,1095x + 0,8025$$

$$1,695 = 0,1095x + 0,8025$$

$$x = \frac{1,695 - 0,8025}{0,1095}$$

$$x = 8,1507 \text{ mgL}^{-1}$$

b. Kemampuan Adsorpsi DBP oleh MIP 10 menit

$$q_e = \frac{(C_o - C_e) V}{W}$$

$$\text{Diketahui: } C_o = 10 \text{ mgL}^{-1} \quad W = 0,03 \text{ g}$$

$$C_e = \text{Konsentrasi setelah adsorpsi} \quad V = 0,005 \text{ L}$$

$$q_e = \frac{(10 - 8,1507) 0,005}{0,03}$$

$$q_e = \frac{0,0092465}{0,03}$$

$$q_e = 0,3082 \text{ mg/g}$$

4. Nilai konsentrasi adsorpsi dan kemampuan adsorpsi DBP oleh MIP terhadap pengaruh konsentrasi

Konsentrasi awal (mgL^{-1})	y (absorbansi)	x (konsentrasi) (mgL^{-1})	q_e (mg/g)
9	1,005	1,8493	1,1918
15	1,385	5,3196	1,6134
18	1,592	7,2100	1,7983
21	1,796	9,0731	1,9878
24	1,969	10,6530	2,2245

Contoh perhitungan konsentrasi adsorpsi dan kemampuan adsorpsi DBP oleh MIP terhadap pengaruh konsentrasi:

a. Adsorpsi DBP oleh MIP 9 mgL^{-1}

$$y = 0,1095x + 0,8025$$

$$y = 1,005$$

$$y = 0,1095x + 0,8025$$

$$1,005 = 0,1095x + 0,8025$$

$$x = \frac{1,005 - 0,8025}{0,1095}$$

$$x = 1,8493 \text{ mgL}^{-1}$$

b. Kemampuan Adsorpsi DEHP oleh MIP 9 mgL^{-1}

$$q_e = \frac{(C_o - C_e) V}{W}$$

Diketahui: C_o = konsentrasi awal $W = 0,03 \text{ g}$

C_e = Konsentrasi setelah adsorpsi $V = 0,005 \text{ L}$

$$q_e = \frac{(9 - 1,8493) \cdot 0,005}{0,03}$$

$$q_e = \frac{0,0357535}{0,03}$$

$$q_e = 1,1918 \text{ mg/g}$$

Lampiran 5. Foto Hasil Penelitian



Proses persiapan alat dan bahan



Proses pencampuran bahan dan prapolimerisasi



Sonikasi



Pengaliran gas nitrogen untuk menghilangkan gas oksigen



Polimerisasi dalam *waterbath*



Polimer terbentuk berwarna putih



Pengeringan polimer



Proses pencucian polimer dengan aseton, metanol, dan akuades agar bersih dari pengotor



Proses ekstraksi (sonikasi)



Pencucian polimer dengan akuades



Penentuan pH



Penimbangan polimer hasil sintesis



Pembuatan deret standar DBP



Uji kemampuan adsorpsi MIP dan NIP



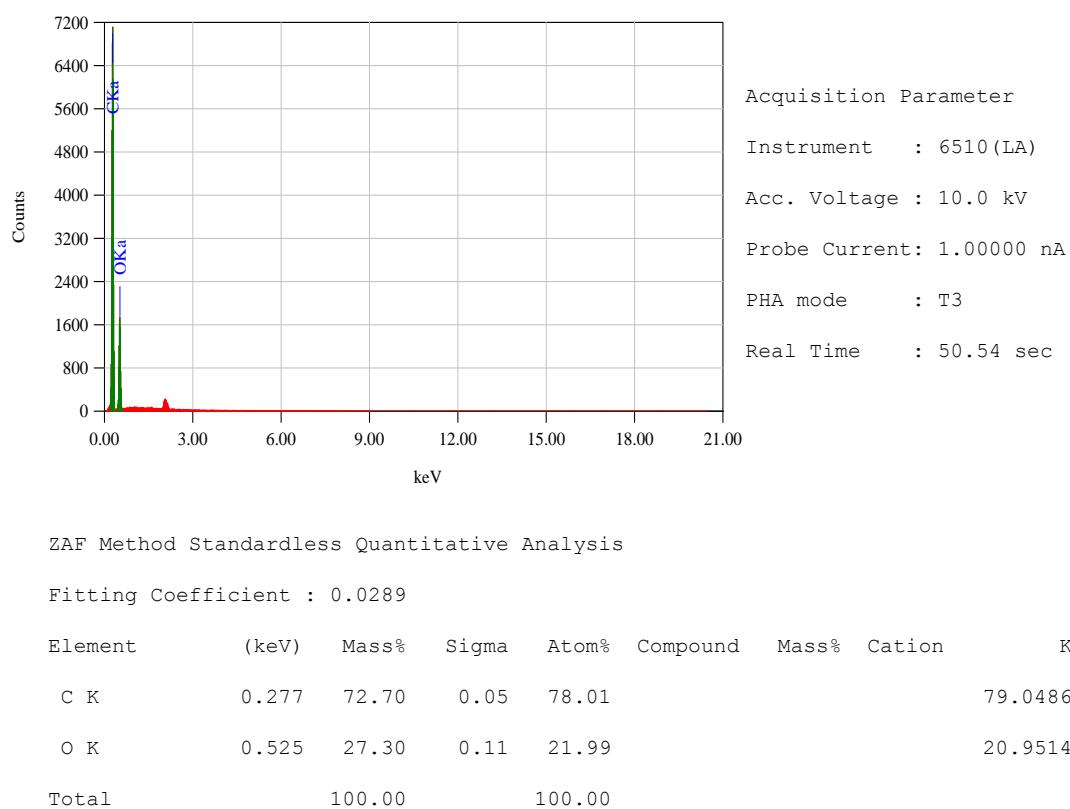
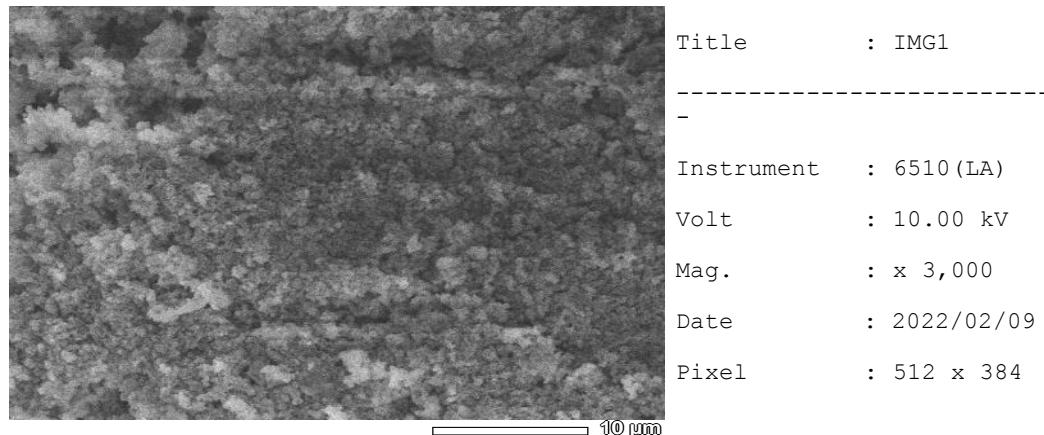
Pengocokan dengan alat *shaker* untuk pengaruh waktu dan konsentrasi terhadap adsorpsi DEHP



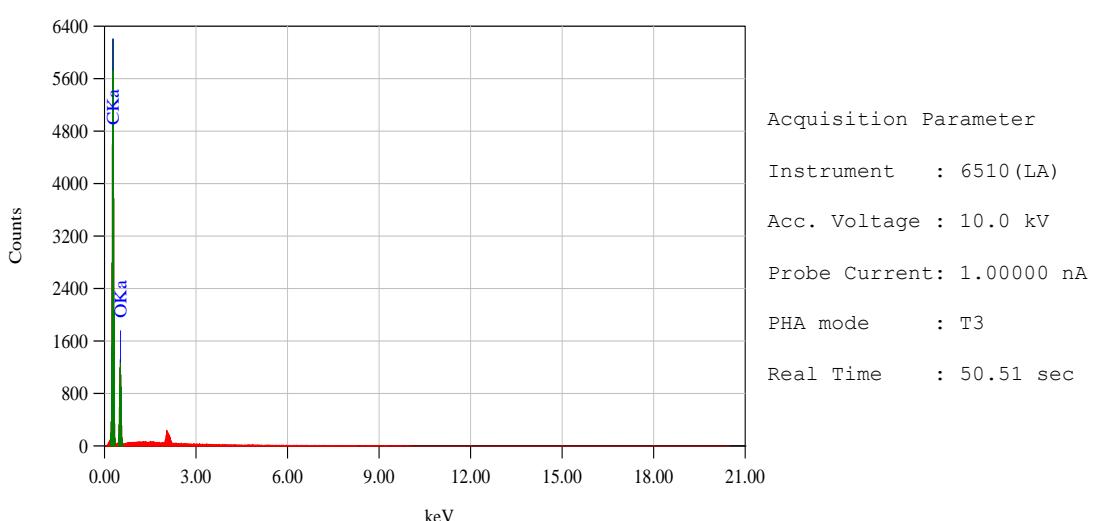
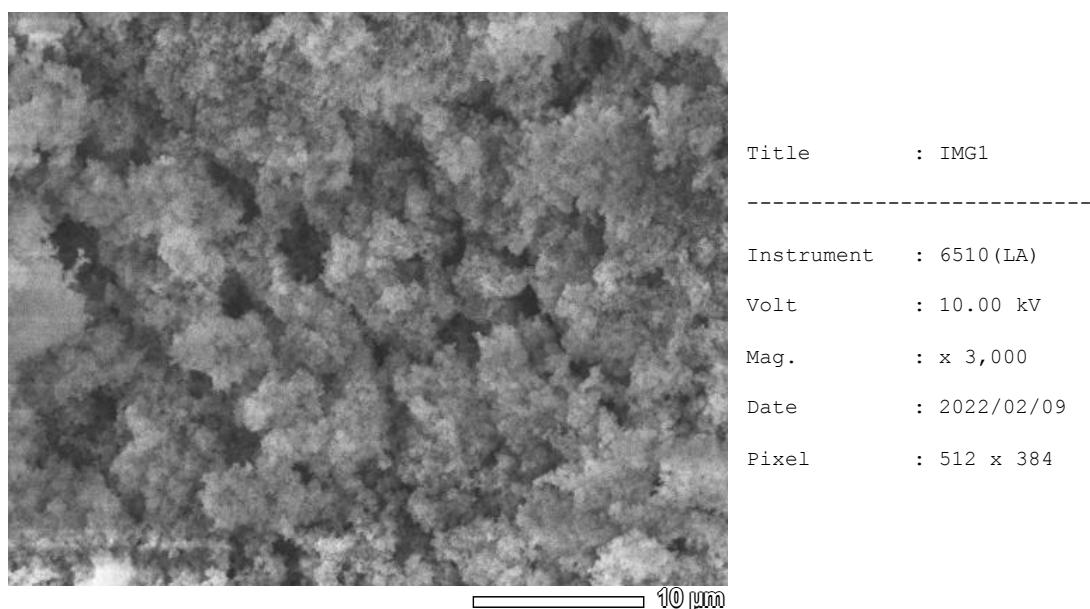
Polimer hasil sintesis
MIP_DBP_MMA-co-EGDMA dan
NIP_MMA-co-EGDMA

Lampiran 6. Karakterisasi EDS

1. NIP_MMA-co-EGDMA



2. MIP_DBP_MMA-co-EGDMA(BE)

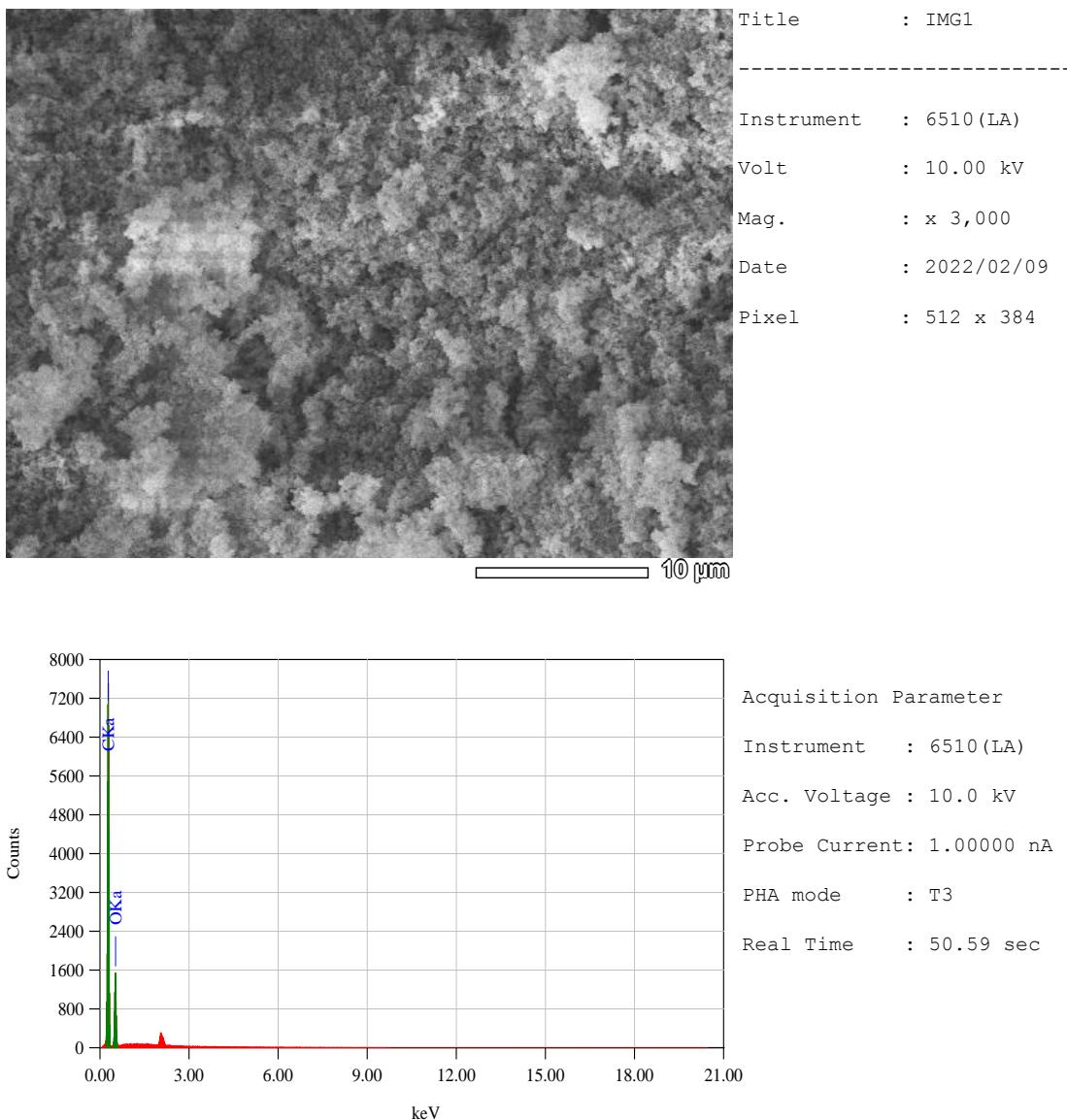


ZAF Method Standardless Quantitative Analysis

Fitting Coefficient : 0.0317

Element	(keV)	Mass%	Sigma	Atom%	Compound	Mass%	Cation	K
C K	0.277	75.26	0.22	80.21				81.8204
O K	0.525	24.74	0.28	19.79				18.1796
Total		100.00		100.00				

3. MIP_DBP_MMA-co-EGDMA(TE)



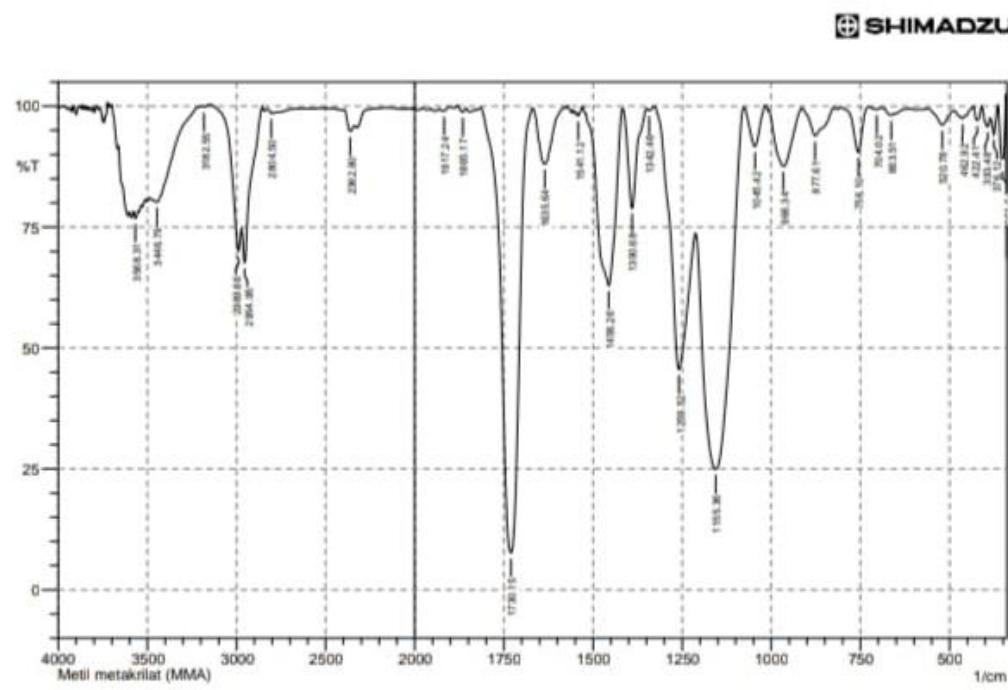
ZAF Method Standardless Quantitative Analysis

Fitting Coefficient : 0.0324

Element	(keV)	Mass%	Sigma	Atom%	Compound	Mass%	Cation	K
C K	0.277	72.15	0.24	77.11				81.6018
O K	0.525	27.85	0.31	22.89				18.4982
Total		100.00		100.00				

Lampiran 7. Karakterisasi FTIR

1. Spektrum Monomer MMA



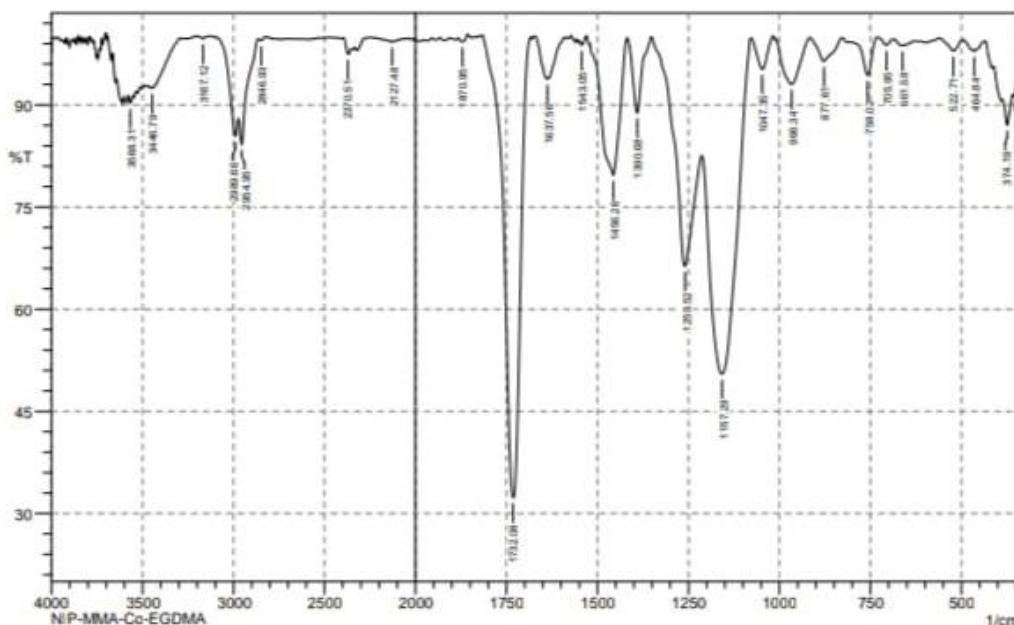
No.	Peak	Intensity	Corr. Intensity	Base (H)	Base (L)	Area	Corr. Area
1	351.04	87.56	13.926	364.55	341.4	0.722	0.646
2	376.12	94.238	4.408	385.76	364.55	0.313	0.197
3	393.48	95.78	2.501	410.84	385.76	0.298	0.159
4	422.41	97.112	2.808	432.05	410.84	0.184	0.176
5	462.92	97.471	0.81	487.99	455.2	0.274	0.072
6	520.78	96.086	3.256	563.21	487.99	0.683	0.475
7	663.51	98.138	0.666	682.8	655.8	0.17	0.056
8	704.02	99.303	0.304	715.59	690.52	0.058	0.016
9	756.1	90.517	9.28	802.39	727.16	1.18	1.111
10	877.81	93.878	5.746	912.33	825.53	1.313	1.182
11	966.34	87.56	11.789	1006.84	914.26	2.999	2.751
12	1045.42	91.69	8.284	1076.28	1018.41	1.126	1.117
13	1155.36	24.936	59.596	1211.3	1078.21	45.218	36.375
14	1259.52	45.581	38.795	1327.03	1213.23	18.989	11.339
15	1342.46	99.092	0.566	1348.24	1327.03	0.051	0.03
16	1390.68	78.95	20.546	1415.75	1350.17	2.781	2.638
17	1456.26	62.906	36.253	1519.91	1417.68	10.702	10.268
18	1541.12	98.051	1.011	1546.91	1529.55	0.097	0.043
19	1635.64	88.049	11.528	1666.5	1581.63	2.275	2.134
20	1730.15	7.664	91.911	1815.02	1670.35	43.294	43.039
21	1865.17	98.8	0.976	1880.6	1857.45	0.074	0.062
22	1917.24	98.993	0.644	1930.74	1903.74	0.084	0.042
23	2362.8	94.789	2.675	2393.66	2335.8	0.989	0.355
24	2804.5	98.483	0.943	2827.64	2659.84	0.706	0.31
25	2954.95	67.618	10.245	2970.38	2856.58	8.43	1.328
26	2989.66	70.012	7.133	3145.9	2972.31	8.205	0.95
27	3182.55	99.826	0.296	3192.19	3147.83	-0.018	0.02
28	3446.79	80.207	1.226	3460.3	3194.12	10.241	0.301
29	3568.31	76.803	1.373	3577.95	3554.81	2.577	0.103

Comment:
Metil metakrilat (MMA)

Date/Time: 2/14/2022 3:11:17 PM
No. of Scans:
Resolution:
Apodization:

2. Spektrum NIP-MMA-co-EGDMA

 SHIMADZU



No.	Peak	Intensity	Corr. Intensity	Base (H)	Base (L)	Area	Corr. Area
1	352.97	91.28	3.465	358.76	341.4	0.471	0.181
2	374.19	87.108	3.934	387.69	360.69	1.354	0.255
3	464.84	97.888	0.726	495.71	455.2	0.258	0.072
4	522.71	97.936	1.757	561.29	495.71	0.315	0.233
5	661.58	98.71	0.33	682.8	655.8	0.12	0.033
6	705.95	98.787	0.972	731.02	688.59	0.121	0.082
7	758.02	94.371	5.281	800.46	731.02	0.742	0.612
8	877.61	96.423	3.468	916.19	829.39	0.737	0.695
9	966.34	93.032	6.706	1008.77	916.19	1.654	1.555
10	1047.35	95.252	4.726	1078.21	1016.41	0.643	0.636
11	1157.29	50.495	39.228	1213.23	1078.21	23.346	17.766
12	1259.52	66.346	21.862	1350.17	1215.15	11.172	5.433
13	1390.68	88.884	11.03	1417.68	1352.1	1.373	1.352
14	1456.26	79.727	19.646	1519.91	1419.61	5.29	4.961
15	1543.05	98.796	0.678	1546.91	1529.55	0.049	0.027
16	1637.56	93.783	6.222	1670.35	1591.27	1.099	1.096
17	1732.06	32.362	67.692	1815.02	1683.86	20.115	20.164
18	1870.95	99.299	0.789	1882.52	1857.45	0.045	0.057
19	2127.48	99.271	0.5	2250.93	2077.33	0.306	0.162
20	2370.51	97.45	1.347	2395.59	2355.08	0.287	0.109
21	2846.93	99.458	0.362	2862.36	2821.86	0.054	0.028
22	2954.95	84.264	5.579	2972.31	2864.29	3.607	0.622
23	2989.66	85.424	4.015	3093.82	2974.23	3.398	0.526
24	3167.12	99.753	0.358	3205.69	3136.25	0.008	0.042
25	3446.79	92.476	0.77	3460.3	3296.35	3.018	0.306
26	3568.31	90.283	1.067	3579.88	3654.81	1.056	0.07

Comment:
NIP-MMA-Co-EGDMA

Date/Time: 2/14/2022 2:34:22 PM

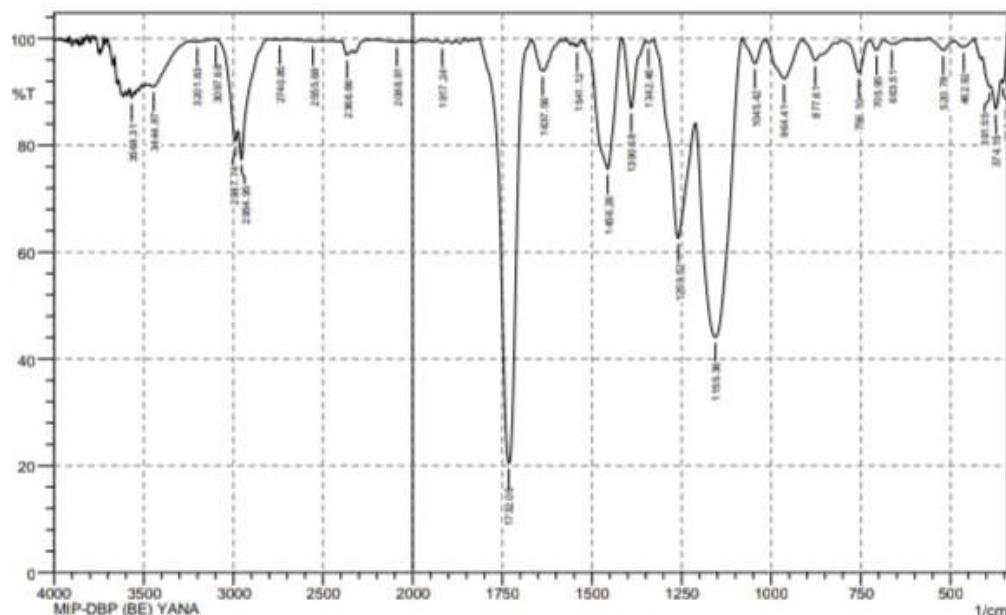
No. of Scans:

Resolution:

Apodization:

3. Spektrum MIP_DBP_MMA-co-EGDMA(BE)

 SHIMADZU



No.	Peak	Intensity	Corr. Intensity	Base (H)	Base (L)	Area	Corr. Area
1	352.97	91.773	3.31	358.76	341.4	0.464	0.189
2	374.19	86.819	4.794	387.69	358.76	1.432	0.341
3	391.55	90.956	1.013	410.84	387.69	0.738	0.085
4	462.92	98.375	0.512	470.63	432.05	0.167	0.043
5	520.78	97.821	1.932	563.21	489.92	0.33	0.264
6	663.51	98.969	0.337	682.8	655.8	0.092	0.025
7	705.95	97.845	1.858	725.23	682.8	0.207	0.153
8	756.1	93.586	6.099	802.39	725.23	0.858	0.753
9	877.61	95.878	3.679	912.33	823.6	0.885	0.686
10	964.41	92.482	7.373	1016.48	912.33	1.89	1.824
11	1045.42	95.328	4.657	1082.07	1018.41	0.635	0.633
12	1155.36	44.046	46.979	1211.3	1082.07	26.533	21.772
13	1259.52	62.65	27.805	1327.03	1213.23	11.811	7.384
14	1342.46	99.237	0.46	1350.17	1327.03	0.055	0.026
15	1390.68	87.054	12.85	1415.75	1362.1	1.61	1.563
16	1456.26	75.572	24.266	1519.91	1417.68	6.298	6.183
17	1541.12	98.506	0.68	1544.98	1529.55	0.067	0.028
18	1637.56	93.913	5.857	1666.5	1589.34	1.106	1.025
19	1732.08	20.448	79.447	1815.02	1670.35	26.347	26.296
20	1917.24	99.189	0.428	1930.74	1907.6	0.064	0.027
21	2088.91	95.42	0.063	2102.41	2077.33	0.061	0.005
22	2366.66	96.954	1.736	2395.59	2333.87	0.6	0.231
23	2555.68	99.552	0.028	2700.34	2546.04	0.195	0.007
24	2740.85	99.745	0.121	2791	2700.34	0.072	0.019
25	2954.95	77.419	7.022	2972.31	2804.5	6.325	0.809
26	2987.74	80.828	3.942	3088.03	2974.23	4.543	0.503
27	3097.68	99.788	0.084	3113.11	3088.03	0.018	0.004
28	3201.83	99.422	0.042	3207.62	3184.48	0.053	0.002
29	3444.87	90.978	0.868	3460.3	3257.77	4.122	0.203
30	3568.31	88.96	1.247	3577.95	3554.81	1.121	0.079

Comment:
MIP-DBP (BE) YANA

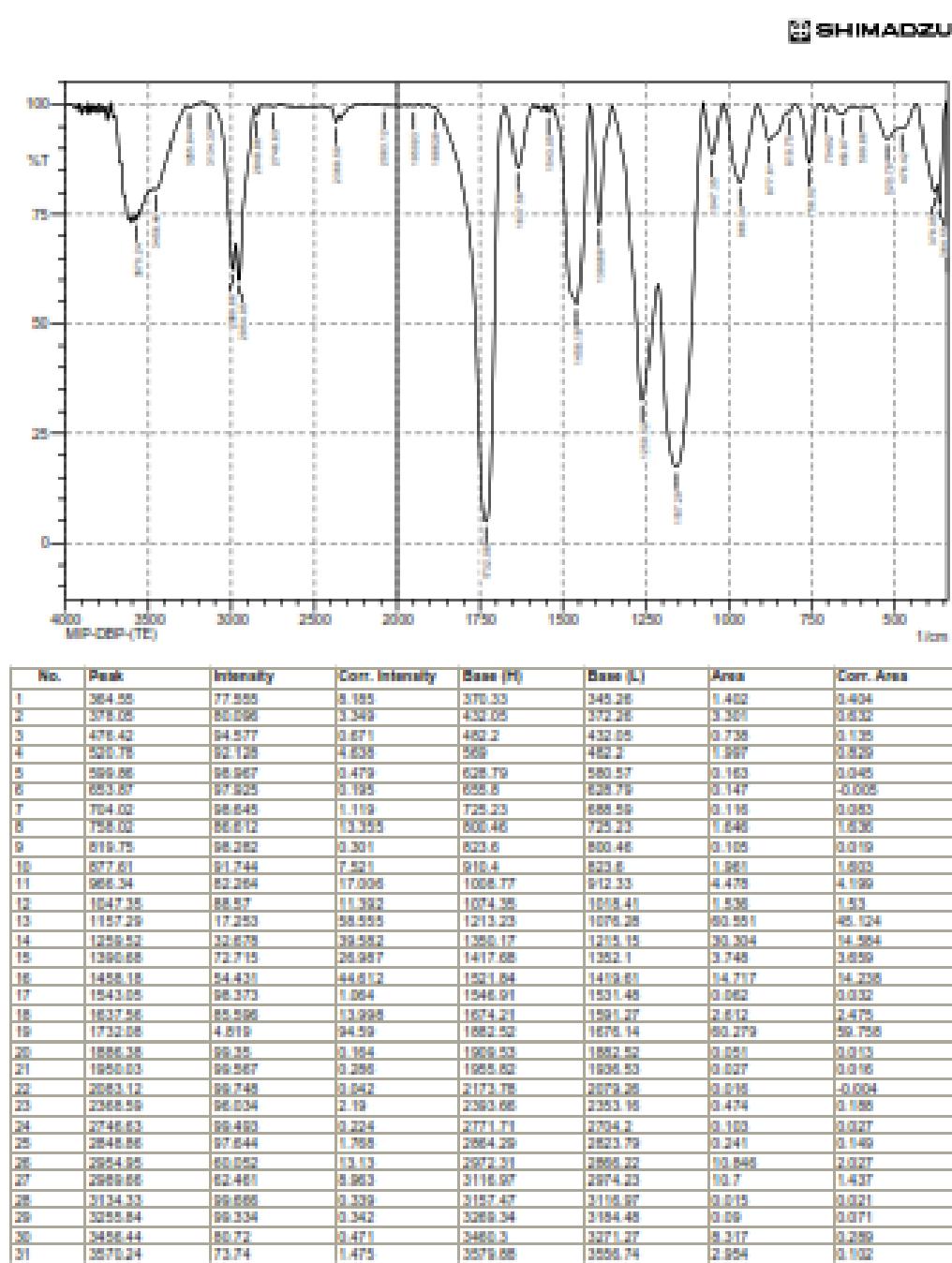
Date/Time: 2/14/2022 2:40:32 PM

No. of Scans:

Resolution:

Apodization:

4. Spektrum MIP_DBP_MMA-co-EGDMA(TE)



Lampiran 8. Karakterisasi SAA



TriStar II 3020 2.00 TriStar II 3020 Version 2.00 Unit
1 Port 1 Serial #: 1108 Page 1

Sample: MIP_DBP_MMA-Co-EGDMA (TE)
Operator: Sarah
Submitter: 30391
File: C:\TriStar II 3020\data\SAMPE...\\MIP_DBP_MMA-Co-EGDMA-TE.SMP

Started: 3/9/2022 7:18:50 AM Analysis Adsorptive: N2
Completed: 3/9/2022 3:44:43 PM Analysis Bath Temp.: -195.860 °C
Report Time: 3/11/2022 8:09:22 AM Thermal Correction: No
Sample Mass: 0.2273 g Warm Free Space: 11.2384 cm³ Measured
Cold Free Space: 32.1591 cm³ Equilibration Interval: 5 s
Low Pressure Dose: None Sample Density: 1.000 g/cm³
Automatic Degas: No

Summary Report

Surface Area

Single point surface area at P/P₀ = 0.304038660: 154.4254 m²/g

BET Surface Area: 157.1970 m²/g

t-Plot Micropore Area: 44.7217 m²/g

t-Plot External Surface Area: 112.4754 m²/g

BJH Adsorption cumulative surface area of pores
between 1.7000 nm and 300.0000 nm diameter: 76.812 m²/g

BJH Desorption cumulative surface area of pores
between 1.7000 nm and 300.0000 nm diameter: 83.8432 m²/g

D-H Adsorption cumulative surface area of pores
between 1.7000 nm and 300.0000 nm diameter: 88.596 m²/g

D-H Desorption cumulative surface area of pores
between 1.7000 nm and 300.0000 nm diameter: 75.4948 m²/g

Pore Volume

Single point adsorption total pore volume of pores
less than 171.1684 nm diameter at P/P₀ = 0.988677175: 0.244900 cm³/g

t-Plot micropore volume: 0.023223 cm³/g

BJH Adsorption cumulative volume of pores
between 1.7000 nm and 300.0000 nm diameter: 0.197929 cm³/g

BJH Desorption cumulative volume of pores
between 1.7000 nm and 300.0000 nm diameter: 0.192406 cm³/g

Pore Size

Adsorption average pore width (4V/A by BET): 6.23167 nm

BJH Adsorption average pore diameter (4V/A): 10.3071 nm

BJH Desorption average pore diameter (4V/A): 9.1793 nm

D-H Adsorption average pore diameter (4V/A): 9.2271 nm

D-H Desorption average pore diameter (4V/A): 9.7970 nm



TriStar II 3020 2.00

TriStar II 3020 Version 2.00 Unit
1 Port 1

Serial #: 1108

Page 3

Sample: MIP_DBP_MMA-Co-EGDMA (TE)
Operator: Sarah
Submitter: 30391
File: C:\TriStar II 3020\data\SAMPE...\\MIP_DBP_MMA-Co-EGDMA-TE.SMP

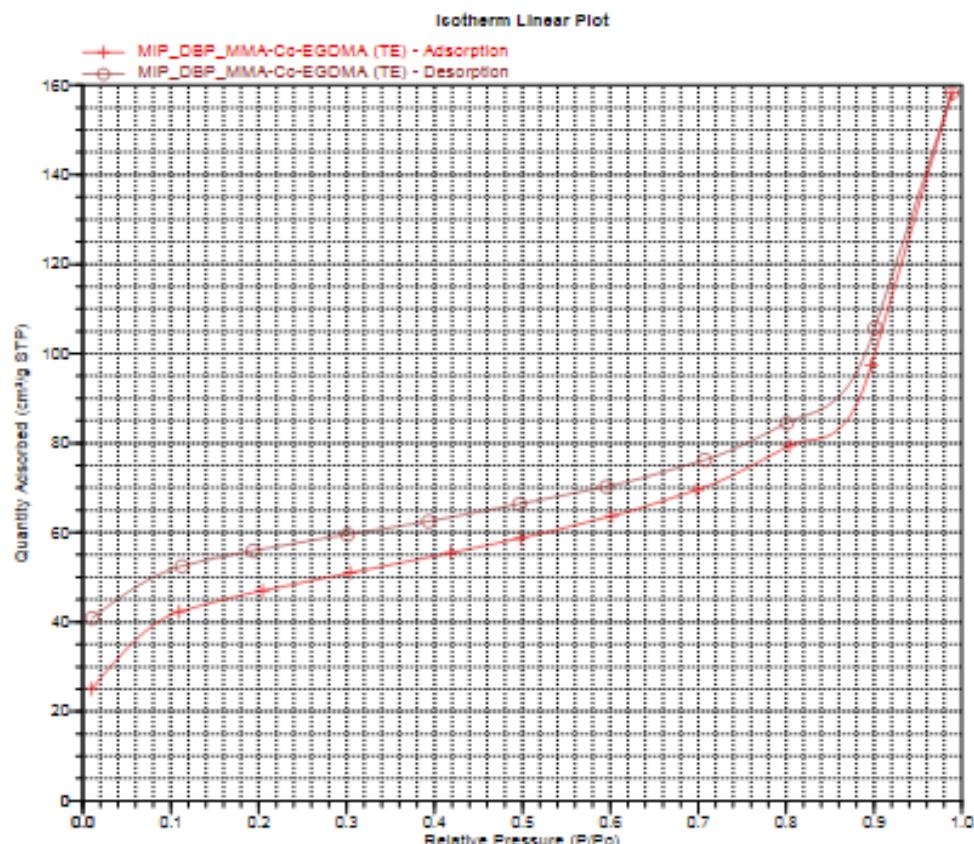
Started: 3/9/2022 7:18:50 AM Analysis Adsorptive: N2
Completed: 3/9/2022 3:44:43 PM Analysis Bath Temp.: -195.850 °C
Report Time: 3/11/2022 8:05:22 AM Thermal Correction: No
Sample Mass: 0.2273 g Warm Free Space: 11.2384 cm³ Measured
Cold Free Space: 32.1591 cm³ Equilibration Interval: 5 s
Low Pressure Dose: None Sample Density: 1.000 g/cm³
Automatic Degas: No

Isotherm Tabular Report

Relative Pressure (P/P ₀)	Absolute Pressure (mmHg)	Quantity Adsorbed (cm ³ /g STP)	Elapsed Time (h:min)	Saturation Pressure (mmHg)
	760.000000			
0.009928568	7.545712	25.0539	02:53	
0.109238082	83.020943	42.2710	05:08	
0.204374434	155.324570	46.9872	05:25	
0.304038660	231.069382	50.9712	05:36	
0.418241320	317.883403	55.4349	05:47	
0.498288408	378.699188	58.7549	05:54	
0.601668900	457.268433	63.6914	06:03	
0.698865067	531.137451	69.5482	06:11	
0.803898942	610.983196	79.3982	06:22	
0.897094807	681.792053	97.4385	06:33	
0.988677175	751.394653	158.3268	06:38	
0.901173883	684.892151	105.7902	06:53	
0.800705599	608.538255	84.4959	07:04	
0.707451509	537.663147	76.2484	07:10	
0.595522188	452.596883	70.2574	07:15	
0.498751685	377.531281	66.3865	07:20	
0.393303801	298.910889	62.5104	07:24	
0.301256903	228.955246	59.5691	07:28	
0.193523246	147.077667	55.9389	07:34	
0.112766145	85.702271	52.4791	07:42	
0.010461469	7.950716	40.8810	08:24	

Sample: MIP_DBP_MMA-Co-EGDMA (TE)
Operator: Sarah
Submitter: 30391
File: C:\TriStar II 3020\data\SAMPLES\MIP_DBP_MMA-Co-EGDMA-TE.SMP

Started: 3/9/2022 7:18:50 AM Analysis Adsorptive: N2
Completed: 3/9/2022 3:44:43 PM Analysis Bath Temp.: -195.850 °C
Report Time: 3/11/2022 8:05:22 AM Thermal Correction: No
Sample Mass: 0.2273 g Warm Free Space: 11.2384 cm³ Measured
Cold Free Space: 32.1591 cm³ Equilibration Interval: 5 s
Low Pressure Dose: None Sample Density: 1.000 g/cm³
Automatic Degas: No





TriStar II 3020 2.00

TriStar II 3020 Version 2.00 Unit
1 Port 1

Serial #: 1108

Page 12

Sample: MIP_DBP_MMA-Co-EGDMA (TE)
Operator: Sarah
Submitter: 30391
File: C:\TriStar II 3020\data\SAMPE...\\MIP_DBP_MMA-Co-EGDMA-TE.SMP

Started: 3/9/2022 7:18:50 AM Analysis Adsorptive: N2
Completed: 3/9/2022 3:44:43 PM Analysis Bath Temp.: -195.850 °C
Report Time: 3/11/2022 8:05:22 AM Thermal Correction: No
Sample Mass: 0.2273 g Warm Free Space: 11.2384 cm³ Measured
Cold Free Space: 32.1591 cm³ Equilibration Interval: 5 s
Low Pressure Dose: None Sample Density: 1.000 g/cm³
Automatic Degas: No

BJH Adsorption Pore Distribution Report

Faas Correction

Harkins and Jura

$$t = [13.99 / (0.034 - \log(P/P_0))] ^ {0.5}$$

Diameter Range: 1.7000 nm to 300.0000 nm

Adsorbate Property Factor: 0.95300 nm

Density Conversion Factor: 0.0015468

Fraction of Pores Open at Both Ends: 0.00

Pore Diameter Range (nm)	Average Diameter (nm)	Incremental Pore Volume (cm³/g)	Cumulative Pore Volume (cm³/g)	Incremental Pore Area (m²/g)	Cumulative Pore Area (m²/g)
171.0 - 20.0	21.7	0.119870	0.119870	22.080	22.080
20.0 - 10.7	12.5	0.031441	0.151311	10.022	32.102
10.7 - 8.9	7.9	0.015403	0.166714	7.782	39.884
8.9 - 5.1	5.7	0.008222	0.174936	5.780	45.673
5.1 - 3.9	4.3	0.006789	0.181724	6.328	52.001
3.9 - 3.2	3.5	0.004456	0.186181	5.156	57.157
3.2 - 2.4	2.7	0.006172	0.192353	9.102	66.259
2.4 - 1.9	2.1	0.005577	0.197929	10.553	76.812

Activate Windows

Lampiran 9. Contoh Perhitungan Nilai K_1 dan K_2 berdasarkan Persamaan Orde Satu Semu dan Orde Dua Semu

1. Penentuan K_1 dari persamaan orde satu semu

Persamaan orde satu semu:

$$\log (q_e - q_t) = \log q_e - K_1 t/2,303$$

- $\log q_e = \text{Intercept}$
 $q_e = \text{Inv. log Intercept}$

- $\log q_e = -0,5801$
 $q_e = 0,2629$

- $K_1/2,303 = \text{Slope}$
 $K_1 = \text{Slope} \times 2,303$
 $K_1 = -0,0019 \times 2,303$
 $K_1 = -0,0044$

2. Penentuan K_2 dari persamaan orde dua semu

Persamaan orde satu semu:

$$t/q_t = 1/K_2 q_e^2 + t/q_e$$

- $1/q_e = \text{Slope}$
 $q_e = 1/\text{Slope}$
 $q_e = 1/1,9876$
 $q_e = 0,5031$

- $1/K_2 q_e^2 = \text{Intercept}$
 $K_2 = 1/\text{Intercept} \times q_e^2$
 $K_2 = 1/19,952 \times (0,5031)^2$
 $K_2 = 0,0127$

Lampiran 10. Perhitungan Nilai Kapasitas Adsorpsi berdasarkan Model Persamaan Isotermal Langmuir dan Isotermal Freundlich

1. Isotermal Adsorpsi Langmuir

Persamaan:

$$\frac{1}{q_e} = \frac{1}{q_m K_L} x \frac{1}{C_e} + \frac{1}{K_L}$$

Keterangan:

- C_e = Konsentrasi saat kesetimbangan (mg/L)
 q_e = Jumlah zat teradsorpsi saat kesetimbangan (mg/g)
 X_m = Kapasitas adsorpsi maksimum monolayer (mg/g)
 K_L = Konstanta afinitas adsorpsi atau konstanta kesetimbangan (L/mg)

Berdasarkan model isotermal Langmuir diperoleh persamaan garis:

$$y = 0,7885x + 0,4245$$

$$\frac{1}{q_m} = 0,4245 \quad \text{maka,} \quad q_m = \frac{1}{0,4245} = 2,3557$$
$$\frac{1}{q_m K_L} = 0,7885 \quad \text{maka,} \quad K_L = \frac{1}{0,7885 \times 2,3557} = 0,5384$$

2. Isotermal Adsorpsi Freundlich

Persamaan:

$$\log q_e = \frac{1}{n} \log C_e + \log K_F$$

Keterangan:

- C_e = Konsentrasi saat kesetimbangan (mg/L)
 q_e = Jumlah zat teradsorpsi saat kesetimbangan (mg/g)
 K_F = Konstanta Freundlich menyatakan tingkat adsorpsi (mg/g)
 $\frac{1}{n}$ = Konstanta Freundlich menyatakan faktor heterogenitas
 n = Intensitas adsorpsi

Berdasarkan model isotermal Freundlich diperoleh persamaan garis:

$$y = 0,3395x - 0,0236$$
$$\frac{1}{n} = 0,3395 \quad \text{maka,} \quad n = \frac{1}{0,3395} = 2,9455$$

$$\log K_F = -0,0236$$
$$K_F = \text{Inv. log } (-0,0489)$$
$$K_F = 0,9479 \text{ mg/g}$$