

APPENDICES

APPENDIX A

METHODOLOGY OF ANALYSIS

Appendix A1 Chemical Oxygen Demand (COD) analysis (Closed reflux, Titration method)

Apparatus

- Digestion vessel 16 x 100 mm
- Hot air oven
- Heating block

Reagents

- Potassium standard solution. The following reagents were added into 500 ml distilled water. 4.913 g $K_2Cr_2O_7$ (Primary standard grade) previously dried at 103 °C for 2 h, then added 167 ml conc. sulfuric and 33.3 g $HgSO_4$. The mixture was done until complete dissociation, thrown away to be cooled in room temperature, and finally diluted to 1000 ml.
- Sulfuric acid reagent. Added 1% W/V into conc. H_2SO_4 . The mixture was let for 1-2 days for completely dissolved Ag_2SO_4 .
- Ferroine Indicator
- 0.1 M Ferrous Ammonium Sulfate standard solution (FAS) 39.2 g $Fe(NH_4)_2(SO_4)_2$ dissolved into 500 ml distilled water, then added 20 ml conc. H_2SO_4 , and finally diluted to 1000 ml.

Note: 0.1 M Ferrous Ammonium Sulfate standard solutions compare with Potassium Dichromate standard solution. Using 2.5 ml distilled water, 1.5 ml $K_2Cr_2O_7$, 3.5 ml H_2SO_4 and 1-2 drops Ferroine Indicator into flask titrate with 0.1 Ferrous Ammonium Sulfate standard solution end point green blue to red brown.

$$\text{Concentration of FAS (M)} = \frac{\text{ml } K_2Cr_2O_7 \times 0.1}{\text{ml FAS}}$$

Procedure

- 2.5 ml of Sample was added to digestion vessel
- Add 1.5 ml digestion reagent, and then 3.5 ml sulfuric acid was slowly into the vessel.
- Seal was tightly with cork tube. Afterward, the vessel was gently inverted several times. (As mixing should put on glove and mask), and the vessel was then placed in the hot air oven.
- The vessel was heated at 150 °C for 2 h, and then left to be cooled in room temperature.
- Afterward, Ferroine Indicator was dropped for 1-2 drop, and finally titrate with 0.1 M 0.1 M Ferrous Ammonium Sulfate standard solutions. (End point green blue change to red brown)

Calculation

$$\text{COD (mg/l)} = \frac{(A-B) \times M \times 8000}{\text{ml of Sample}}$$

- where
- A = ml of FAS titrate with blank
 - B = ml of FAS titrate with sample
 - M = Molarity of FAS

Note: Distilled water was used to blank

Appendix A2 Total Suspended Solids (TSS) Analysis

Apparatus

- Desiccators
- Drying oven
- Glass-fiber filter disk (47 mm ID)
- buchner funnel
- Suction flask

Procedure

- The glass-filter disk was dried in a hot air oven at 105 °C for 1 h. The disk was cooled in desiccators to balance temperature and weighed.
- The Sample was chosen to yield between 10 and 200 mg dried residue.
- The glass-filter was placed in buchner funnel which connect vacuum machine.
- The filter was wet with a bit volume of distilled water for fix in buchner funnel
- The sample volume was sieved in the Buchner funnel, after complete filtration also suction and wash filter for 3 min.
- The filter was carefully removed from Buchner funnel.
- The filter was dried at least 1 h at 105 °C in an oven, afterward was cooled in desiccators in room temperature, and weight.

Calculation

$$\text{mg Total Suspended Solids/l} = \frac{(A-B) \times 1000 (\text{mg})(\text{ml/l})}{\text{sample volume}(\text{ml})}$$

A = Weight of filter + dried residue (mg)

B = Weight of filter (mg)

Appendix A3 Volatile Suspended Solids (VSS) Analysis**Procedure**

- The residue produced by TSS method was ignited in a furnace at 550 ± 50 °C.
- A furnace was heated up to temperature after inserting sample.
- Usually, 15 to 20 min ignition is required for 200 mg residue.
- The filter disk was left to partially cool in air until most of the heat was dissipated.
- The dist was transferred to desiccators, and weight as soon as if was cooled in room temperature.

Calculation

$$\text{mg Volatile Suspended solid/L} = \frac{(A-B) \times 1000}{\text{sample volume, ml}}$$

A = Weight of residue + disk before ignition (mg)

B = Weight of residue + disk after ignition (mg)

Appendix A4 Total Solid (TS) Analysis**Apparatus**

- 50 ml crucible
- Water bath
- Analytical balance
- Muffle furnace
- Desiccator
- Drying oven 103 to 105°C

Procedure

- Preparation of evaporation crucible : If volatile solids are to be measured ignite clean evaporating crucible at 550°C for 1 h in a muffle furnace. Store and cool crucible in desiccator until needed. Weigh immediately before use.
- 25 ml sample , that homogeneous samples and then evaporate to dryness on water bath.
- Dry evaporate sample for 1 h in an oven at 103 to 105°C, and then cool crucible in desiccators for 30 min or until a constant weight.
- Determine weight in a weight balance.

Calculation

$$\text{mg Total solid/l} = \frac{(A-B) \times 1000}{\text{sample volume, ml}}$$

A = Weight of residue + crucible before ignition (mg)

B = Weight of residue + crucible after ignition (mg)

Appendix A5 Volatile Solids Analysis**Apparatus**

- The residue from Total Solids
- Muffle furnace
- Weight balance

Procedure

- The residue from Total Solids is ignited in the muffle furnace at 550°C for 1 h.
- Cool in desiccators to a constant temperature and weigh.
- Weigh Analysis in weigh balance.

Calculation

$$\text{mg Volatile solid/l} = \frac{(A-B) \times 1000}{\text{sample volume, ml}}$$

A = Weight of residue from total solids + crucible before ignition

B = Weight of residue + crucible after ignition (ml)

Appendix A6 Total Nitrogen Analysis (Kjeldahl Methods)**Apparatus**

- Weigh balance
- Hood
- Digestor
- Distilling unit
- 250 ml Digestor tube
- 250 ml Erlenmyer flask
- 100, 500 and 1000 ml of volumetric flasks
- 250 ml beaker
- 50 ml burette

- 5, 10, 15 and 20 ml of pipettes
- 25 and 100 ml of Cylinder
- Stir glass and rubber pump

Reagents

- conc.H₂SO₄
- 30% H₂O₂
- Sodium hydroxide Solution. The following reagents were added 1 kg sodium hydroxide (commercial grade NaOH) in to 1000 ml of distilled water
- Mixed catalyst, that consists of K₂SO₄ and CuSO₄·5H₂O in 100:10 ratio.
- Indicator (collected in a light brown bottle)

0.2 g of methyl red dissolve into 100 ml of 95% ethyl alcohol then collect in bottle

0.1 g of methyl blou dissolve into 100 ml of 95% ethyl alcohol and then collect in bottle

- 4% Boric indicator. Add 40 g of boric acid into 1 l of volumetric flak, that have a bit distilled water in it and add 10 ml of indicator (mixed 10 ml methyl red and 5 ml methyl blou) and then diluted to 1000 l.
- 0.1 N Sulfuric acid standard solution: pipette 1.5 ml conc.H₂SO₄ transfer to a 500 ml volumetric flask, fill flask to mark with distilled water. Titration certainly determine concentration of sulfuric acid standard solution with 40 ml of 0.05 N sodium carbonate. The following calculation express:

$$\text{Normality of sulfuric acid} = \frac{A \times B}{53 \times C}$$

where: A = g Na₂CO₃ into 1 l flask
 B = ml of Na₂CO₃ solution taken for titration,
 C = ml acid used

Procedure

Digestor

- Add 100 ml sample in 250 ml digestor tube (blank use distilled water) and add 10.5 g mixed catalyst.
- Add 15 ml conc. Sulfuric acid and 2 ml hydrogen peroxide.
- Initial in digestor machine at temperature about 250°C, when temperature increase to 420°C count time for 2 hours.

Distillation

- Sample is distilled by auto Distillation unit (Kjeltectm2200). The program is fixed by Add 50 ml water and 30 ml sodium hydroxide distill for 4 min until get 150 ml and then was analyzed by titration.
- Add 25 ml boric indicator in 250 ml erlenmyer flask, which is supported from auto distillation.

Titration

- Titration with 0.1 sulfuric acid standard solution. The end-point change green to purple.

Calculation

$$\%N = \frac{(A-B) \times N \times 1400}{\text{ml sample}}$$

where: A = ml acid used titrate sample
B = ml acid used titrate Blank
N = Normality of standard acid

Appendix A7 Grease and oil Analysis via Soxhlet (Ruibal-mendieta et al., 2001)

Apparatus

- 150 ml beaker
- Soxhlet & distill bottle
- Extraction thimbles
- Glass beads

- Electric heating mantle
- Vacuum pump
- buchner funnel
- Water bath
- Clean cotton
- Desiccator
- Muslin remnant of cloth

Reagent

- Conc. HCl
- n-hexane

Procedure

- Add muslin remnant of cloth and filter paper, respectively in the Buchner funnel, that connect with vacuum pump and then add 100 mL water into funnel.
- Add 25 ml sample pass filter in funnel until dry.
- Collect muslin cloth and filter paper in extraction thimble. The residue of oil is wiped by n-hexane. Dry in oven for 30 min at 103 to 105°C.
- Extration thimble is add in soxhlet, that connect with distill bottle and soxhlet extraction for 4 h. n-hexane is Solvent in distill bottle. Extraction ratio 20 rounds per h.
- Afterward, distill extraction bottle until dry (n-hexane is distilled). Dry constant weight in oven at 103 to 105°C and place in desiccators until weigh constant.
- Determine weight by weigh blance.

Calculation

$$\text{mg/l oil and grease} = \frac{\text{mg increase in weight of flask} \times 1,000}{\text{ml sample}}$$

Appendix A8 Carbohydrate Analysis (Colorimetric method)**Apparatus**

- Tube
- Water bath
- Spectro photometer
- Micro pipette

Reagent

- 80% (w/w) phenol: 20 g phenol dissolve in 80 ml deionize water at 25°C.
- conc. Sulfuric acid
- 100 ppm of D-glucose standard solution:

Procedure

Standard curve

- Series of sugar containing 2, 4, 6, 8 and 10 μl of sugar solution is pipetted into colorimetric tube, 50 μl of 80% phenol is added. Then 5 ml of concentrated sulfuric acid is added rapidly.
- The tubes are allowed to stand 10 min, and then they are shaken and placed for 20 min in a water bath at 30°C.
- Characteristic of solution is yellow-orange color can be measured at 480, 484 and 490 nm with spectro photometer. Blank are prepared by substituting deionize water for the sugar solution

Sample Analysis

- Add 2 ml sample, that it is diluted 1000 times by deionize water in tube
- Add 50 μl of 80% phenol and then add 5 mL conc. Sulfuric acid is rapidly.
- The tubes are allowed to stand 10 min, then they are shaken and placed for 20 min in a water bath at 30°C. The same previously method.
- The sample is analyzed by spectro photometer at 480, 484 and 490 nm. Plot graph in standard curve and calculate concentration.

APPENDIX B
GAS CHROMATOGRAPH'S STANDARD GAS

Table B1 Gas chromatograph's calibration on standard curve (60% Hydrogen and 10% Methane)

Order	Hydrogen		Methane	
	% H ₂	Area	% CH ₄	Area
1	60	1810012	10	59149
2	60	1669967	10	61181
3	60	1568000	10	62757
4	60	1693790	10	61549
5	60	1894693	10	72672
6	60	1940139	10	73603
7	60	1912454	10	72642
8	60	1956328	10	72083
9	60	1874962	10	71123
10	60	1899974	10	70349
Average		1822031.90		67710.80

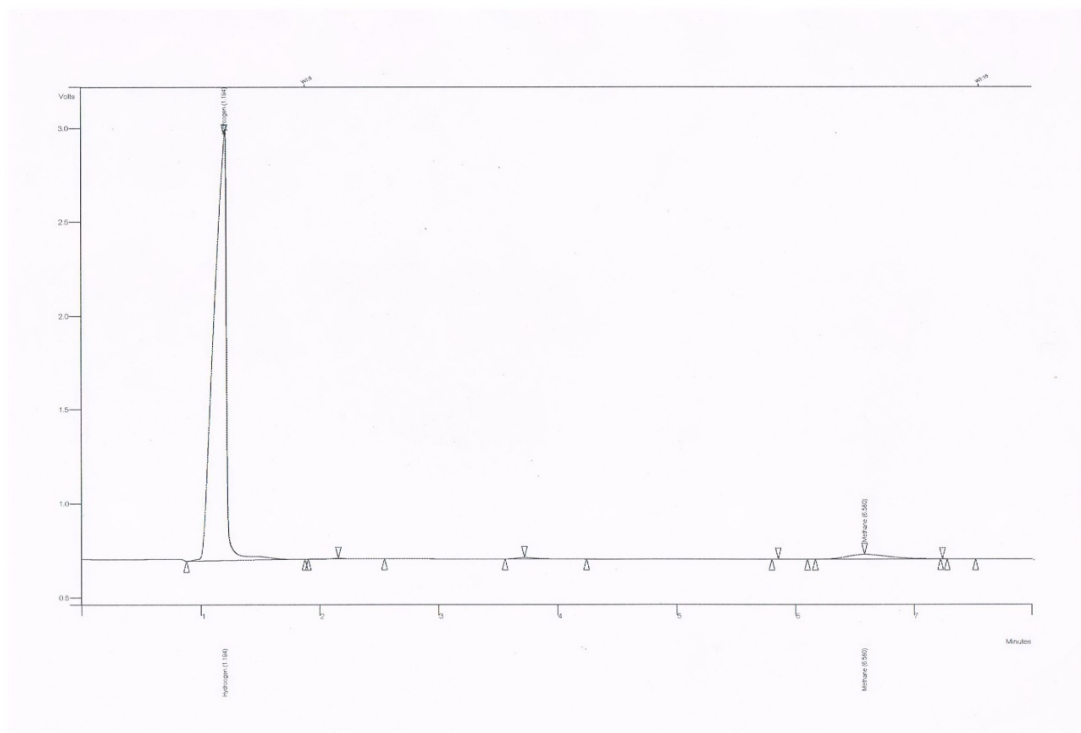


Figure B1 Graph of standard gas (Hydrogen and Methane)

Table B2 Gas chromatograph’s calibration on standard curve (30% Carbon dioxide)

Order	% CO ₂	Area
1	30	28325
2	30	27869
3	30	28738
4	30	28279
5	30	29111
6	30	29176
7	30	27006
8	30	31940
9	30	29337
10	30	27830
Average		28761.10

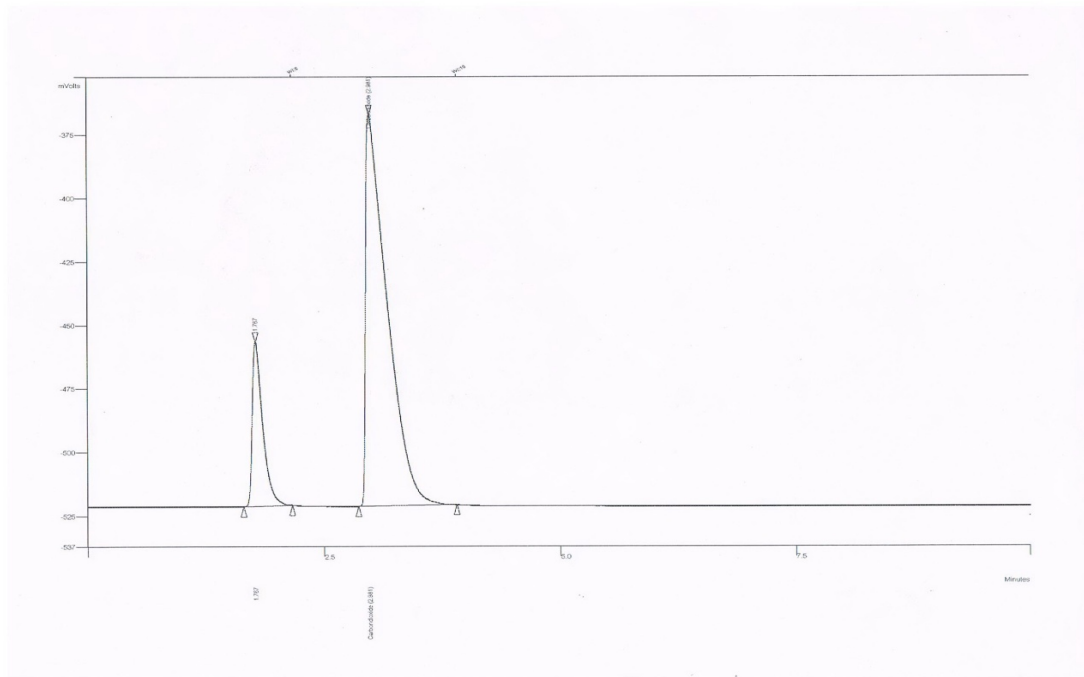


Figure B2 Graph of standard gas (Carbon dioxide)

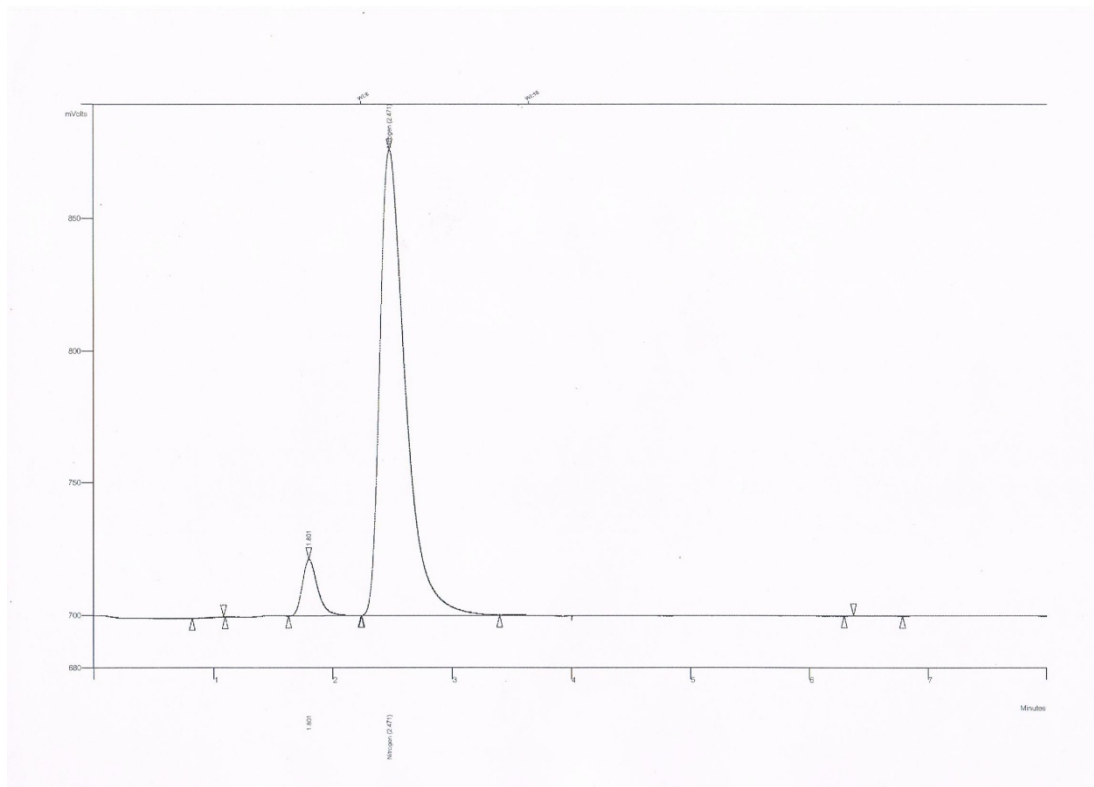


Figure B3 Graph of standard gas (Nitrogen)

Table B – 3 Gas chromatograph's calibration on standard curve (99.995% Nitrogen)

Order	% N ₂	Area
1	99.995	317564
2	99.995	180162
3	99.995	196075
4	99.995	227666
5	99.995	260304
6	99.995	193392
7	99.995	221885
8	99.995	90977
9	99.995	84379
10	99.995	197031
Average		196,954.50

APPENDIX C
VOLATILE FATTY ACID (VFA)

Table C1 Gas chromatograph's calibration on standard curve for acetate

Concentration of acetate (mg/l)	Peak area
4	7004
6	7537
8	9071
10	9604
40	34604
60	45937
80	63847
100	78604

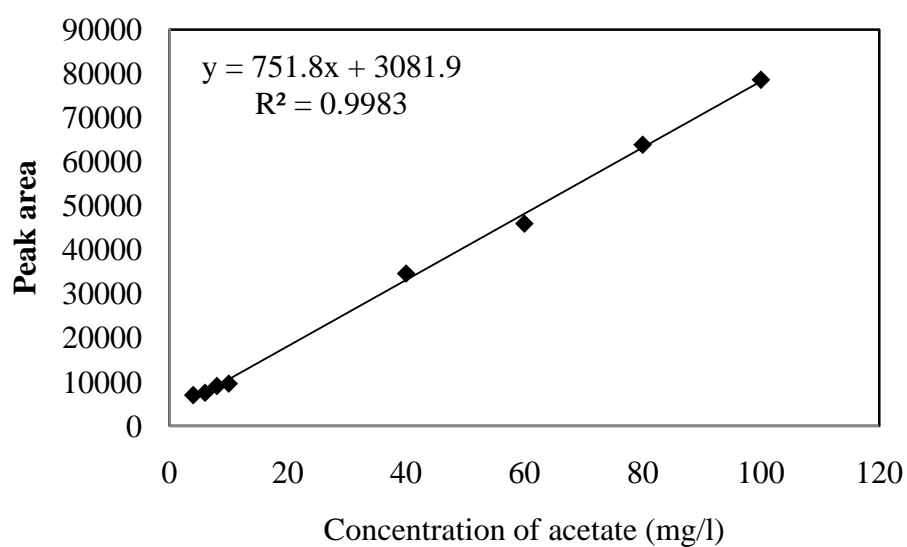


Figure C1 The relationship between peak area and different amount of standard acetic

Table C2 Gas chromatograph's calibration on standard curve for propionate

Concentration of propionate(mg/l)	Peak area
4	13853
6	16359
8	19780
10	20744
40	65198
60	94834
80	122836
100	145118

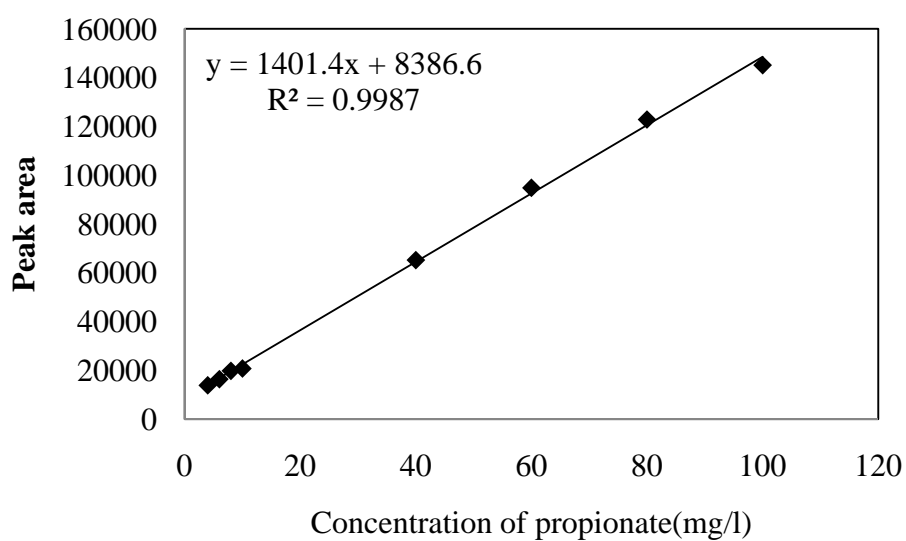
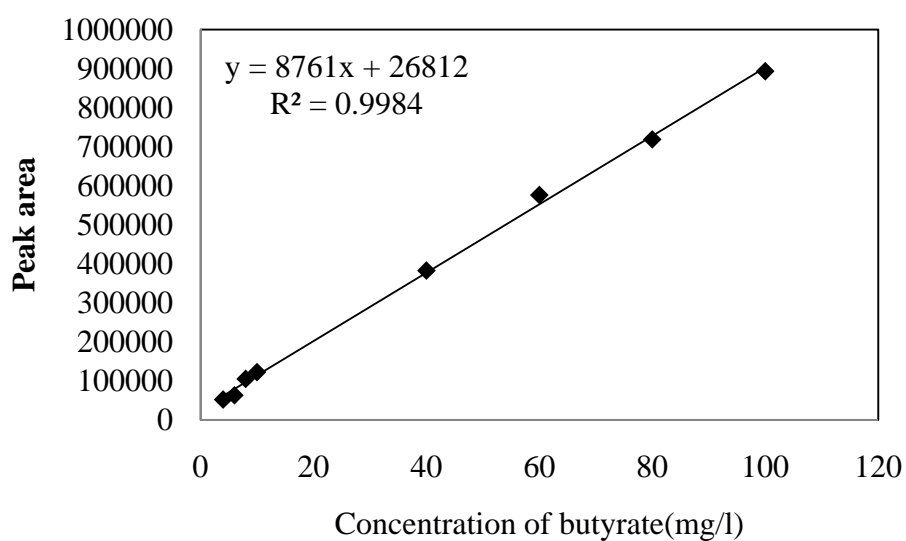
**Figure C2** The relationship between peak area and different amount of standard propionate

Table C3 Gas chromatograph's calibration on standard curve for butyrate

Concentration of butyrate(mg/l)	Peak area
4	51890
6	62763
8	104789
10	122153
40	382619
60	576262
80	718848
100	893549

**Figure C3** The relationship between peak area and different amount of standard butyrate

APPENDIX D
DATA OF BIOGAS

Appendix D 1 Raw data of hydrogen content in experiment batch reactor under mesophilic condition

Initial pH	Total biogas (ml)	H₂ (ml)	N₂ (ml)	CO₂ (ml)	% H₂	% N₂	% CO₂
pH4	116.83	0.00	49.72	67.11	0.00	42.56	57.44
PH5	167.83	0.10	53.79	113.94	0.06	32.05	67.89
pH6	561.33	166.22	107.16	287.95	29.61	19.09	51.30
pH7	357.42	122.68	76.34	158.39	34.32	21.36	44.32
pH8	1531.17	733.49	71.63	726.05	47.90	4.68	47.42
PH9	854.83	337.19	77.96	439.68	39.45	9.12	51.43
pH10	271.67	14.57	81.01	176.08	5.36	29.82	64.82
pH11	253.17	5.19	62.94	185.04	2.05	24.86	73.09
pH12	240.00	21.61	71.02	147.38	9.00	29.59	61.41

Appendix D2 Raw data of hydrogen content in experiment batch reactor under thermophilic condition

Initial pH	Total biogas (ml)	H₂ (ml)	N₂ (ml)	CO₂ (ml)	% H₂	% N₂	% CO₂
pH4	140.00	0.00	34.80	105.20	0.00	24.86	75.14
PH5	167.50	0.00	15.84	151.66	0.00	9.46	90.54
pH6	797.75	233.42	61.65	502.68	29.26	7.73	63.01
pH7	1118.00	356.58	61.34	700.08	31.89	5.49	62.62
pH8	1550.33	767.64	34.16	748.53	49.51	2.20	48.28
PH9	1339.67	423.55	36.98	879.14	31.62	2.76	65.62
pH10	1388.67	413.55	141.10	834.02	29.78	10.16	60.06
pH11	1832.17	280.62	163.12	1388.43	15.32	8.90	75.78
pH12	1079.00	320.23	112.56	646.21	29.68	10.43	59.89

Appendix D3 Raw data of hydrogen content in experiment batch reactor in different initial F/M ratios

F/M ratios	Total biogas (ml)	H₂ (ml)	N₂ (ml)	CO₂ (ml)	% H₂	% N₂	% CO₂
4	1571.67	648.43	87.54	835.69	41.26	5.57	53.17
6	1285.83	533.33	69.43	683.07	41.48	5.40	53.12
8	1164.50	477.94	60.79	625.77	41.04	5.22	53.74
10	1353.67	457.43	67.41	828.83	33.79	4.98	61.23
12	1272.00	426.30	68.43	777.27	33.51	5.38	61.11

Appendix D4 Raw data of hydrogen content in experiment batch reactor in different initial iron concentration

Initial Fe²⁺ concentration (mg FeSO₄/l)	Total biogas (ml)	H₂ (ml)	N₂ (ml)	CO₂ (ml)	% H₂	% N₂	% CO₂
0	1391.83	445.68	89.36	856.79	32.02	6.42	61.56
100	1532.67	588.25	89.66	854.76	38.38	5.85	55.77
200	1431.00	511.47	71.26	848.27	35.74	4.98	59.28
300	1313.00	405.90	75.23	831.87	30.91	5.73	63.36
400	861.00	367.88	44.86	448.26	42.73	5.21	52.06

Appendix D 5 Raw data of hydrogen content in experiment batch reactor in pretreatment method

Pretreatment methods	Total biogas (ml)	H₂ (ml)	N₂ (ml)	CO₂ (ml)	% H₂	% N₂	% CO₂
Heat	1435.50	588.65	588.65	103.93	41.01	7.24	51.75
Re Heat	1468.50	656.57	656.57	100.89	44.71	6.87	48.42
Chloroform	121.67	0.47	0.47	57.28	0.39	47.08	52.53
Ultrasonic	1537.50	443.36	443.36	153.44	28.84	9.98	61.18

APPENDIX E
ANOVA STATISTIC OF HYDROGEN ANALYSIS

Appendix E 1 Comparison of initial pH under mesophilic condition

	Sum of squares	df	Mean square	F	Sig.
Between groups	954955.034	8	119369.379	45727.510	.000
Within groups	23.494	9	2.610		
Total	954978.528	17			

(I) Initial pH	(J) Initial pH	95% Confidence interval				
		Mean difference (I-J)	Std. error	Sig.	Lower bound	Upper bound
4	5	.00000	1.61569	1.000	-3.6549	3.6549
	6	-166.22000*	1.61569	.000	-169.8749	-162.5651
	7	-122.68000*	1.61569	.000	-126.3349	-119.0251
	8	-733.89000*	1.61569	.000	-737.5449	-730.2351
	9	-337.18500*	1.61569	.000	-340.8399	-333.5301
	10	-14.57000*	1.61569	.000	-18.2249	-10.9151
	11	-5.19000*	1.61569	.011	-8.8449	-1.5351
	12	-21.61000*	1.61569	.000	-25.2649	-17.9551

(I) Initial pH	(J) Initial pH				95% Confidence interval	
		Mean difference (I-J)	Std. error	Sig.	Lower bound	Upper bound
5	4	.00000	1.61569	1.000	-3.6549	3.6549
	6	-166.22000*	1.61569	.000	-169.8749	-162.5651
	7	-122.68000*	1.61569	.000	-126.3349	-119.0251
	8	-733.89000*	1.61569	.000	-737.5449	-730.2351
	9	-337.18500*	1.61569	.000	-340.8399	-333.5301
	10	-14.57000*	1.61569	.000	-18.2249	-10.9151
	11	-5.19000*	1.61569	.011	-8.8449	-1.5351
	12	-21.61000*	1.61569	.000	-25.2649	-17.9551
6	4	166.22000*	1.61569	.000	162.5651	169.8749
	5	166.22000*	1.61569	.000	162.5651	169.8749
	7	43.54000*	1.61569	.000	39.8851	47.1949
	8	-567.67000*	1.61569	.000	-571.3249	-564.0151
	9	-170.96500*	1.61569	.000	-174.6199	-167.3101
	10	151.65000*	1.61569	.000	147.9951	155.3049
	11	161.03000*	1.61569	.000	157.3751	164.6849
	12	144.61000*	1.61569	.000	140.9551	148.2649
7	4	122.68000*	1.61569	.000	119.0251	126.3349
	5	122.68000*	1.61569	.000	119.0251	126.3349

(I) Initial pH	(J) Initial pH				95% Confidence interval	
		Mean difference (I-J)	Std. error	Sig.	Lower bound	Upper bound
7	6	-43.54000 [*]	1.61569	.000	-47.1949	-39.8851
	8	-611.21000 [*]	1.61569	.000	-614.8649	-607.5551
	9	-214.50500 [*]	1.61569	.000	-218.1599	-210.8501
	10	108.11000 [*]	1.61569	.000	104.4551	111.7649
	11	117.49000 [*]	1.61569	.000	113.8351	121.1449
	12	101.07000 [*]	1.61569	.000	97.4151	104.7249
8	4	733.89000 [*]	1.61569	.000	730.2351	737.5449
	5	733.89000 [*]	1.61569	.000	730.2351	737.5449
	6	567.67000 [*]	1.61569	.000	564.0151	571.3249
	7	611.21000 [*]	1.61569	.000	607.5551	614.8649
	9	396.70500 [*]	1.61569	.000	393.0501	400.3599
	10	719.32000 [*]	1.61569	.000	715.6651	722.9749
	11	728.70000 [*]	1.61569	.000	725.0451	732.3549
	12	712.28000 [*]	1.61569	.000	708.6251	715.9349
9	4	337.18500 [*]	1.61569	.000	333.5301	340.8399
	5	337.18500 [*]	1.61569	.000	333.5301	340.8399
	6	170.96500 [*]	1.61569	.000	167.3101	174.6199
	7	214.50500 [*]	1.61569	.000	210.8501	218.1599

(I) Initial pH	(J) Initial pH				95% Confidence interval	
		Mean difference (I-J)	Std. error	Sig.	Lower bound	Upper bound
9	8	-396.70500*	1.61569	.000	-400.3599	-393.0501
	10	322.61500*	1.61569	.000	318.9601	326.2699
	11	331.99500*	1.61569	.000	328.3401	335.6499
	12	315.57500*	1.61569	.000	311.9201	319.2299
10	4	14.57000*	1.61569	.000	10.9151	18.2249
	5	14.57000*	1.61569	.000	10.9151	18.2249
	6	-151.65000*	1.61569	.000	-155.3049	-147.9951
	7	-108.11000*	1.61569	.000	-111.7649	-104.4551
	8	-719.32000*	1.61569	.000	-722.9749	-715.6651
	9	-322.61500*	1.61569	.000	-326.2699	-318.9601
	11	9.38000*	1.61569	.000	5.7251	13.0349
	12	-7.04000*	1.61569	.002	-10.6949	-3.3851
11	4	5.19000*	1.61569	.011	1.5351	8.8449
	5	5.19000*	1.61569	.011	1.5351	8.8449
	6	-161.03000*	1.61569	.000	-164.6849	-157.3751
	7	-117.49000*	1.61569	.000	-121.1449	-113.8351
	8	-728.70000*	1.61569	.000	-732.3549	-725.0451
	9	-331.99500*	1.61569	.000	-335.6499	-328.3401
	10	-9.38000*	1.61569	.000	-13.0349	-5.7251

(I) Initial pH	(J) Initial pH				95% Confidence Interval	
		Mean difference (I-J)	Std. error	Sig.	Lower bound	Upper bound
11	12	-16.42000*	1.61569	.000	-20.0749	-12.7651
12	4	21.61000*	1.61569	.000	17.9551	25.2649
	5	21.61000*	1.61569	.000	17.9551	25.2649
	6	-144.61000*	1.61569	.000	-148.2649	-140.9551
	7	-101.07000*	1.61569	.000	-104.7249	-97.4151
	8	-712.28000*	1.61569	.000	-715.9349	-708.6251
	9	-315.57500*	1.61569	.000	-319.2299	-311.9201
	10	7.04000*	1.61569	.002	3.3851	10.6949
	11	16.42000*	1.61569	.000	12.7651	20.0749

*. The mean difference is significant at the 0.05 level.

Appendix E 2 Comparison of initial pH under thermophilic condition

	Sum of squares	df	Mean square	F	Sig.
Between groups	868500.104	8	108562.513	14887.303	.000
Within groups	65.631	9	7.292		
Total	868565.735	17			

(I) Initial pH	(J) Initial pH				95% Confidence interval	
		Mean difference (I-J)	Std. error	Sig.	Lower bound	Upper bound
4	5	.00000	2.70042	1.000	-6.1088	6.1088
	6	-233.42000*	2.70042	.000	-239.5288	-227.3112
	7	-356.59000*	2.70042	.000	-362.6988	-350.4812
	8	-767.64000*	2.70042	.000	-773.7488	-761.5312
	9	-423.55000*	2.70042	.000	-429.6588	-417.4412
	10	-413.55000*	2.70042	.000	-419.6588	-407.4412
	11	-280.62000*	2.70042	.000	-286.7288	-274.5112
	12	-320.23000*	2.70042	.000	-326.3388	-314.1212
5	4	.00000	2.70042	1.000	-6.1088	6.1088
	6	-233.42000*	2.70042	.000	-239.5288	-227.3112
	7	-356.59000*	2.70042	.000	-362.6988	-350.4812
	8	-767.64000*	2.70042	.000	-773.7488	-761.5312
	9	-423.55000*	2.70042	.000	-429.6588	-417.4412
	10	-413.55000*	2.70042	.000	-419.6588	-407.4412
	11	-280.62000*	2.70042	.000	-286.7288	-274.5112
	12	-320.23000*	2.70042	.000	-326.3388	-314.1212
6	4	233.42000*	2.70042	.000	227.3112	239.5288
	5	233.42000*	2.70042	.000	227.3112	239.5288
	7	-123.17000*	2.70042	.000	-129.2788	-117.0612

(I) Initial pH	(J) Initial pH				95% Confidence Interval	
		Mean difference (I-J)	Std. error	Sig.	Lower bound	Upper bound
6	8	-534.22000 [*]	2.70042	.000	-540.3288	-528.1112
	9	-190.13000 [*]	2.70042	.000	-196.2388	-184.0212
	10	-180.13000 [*]	2.70042	.000	-186.2388	-174.0212
	11	-47.20000 [*]	2.70042	.000	-53.3088	-41.0912
	12	-86.81000 [*]	2.70042	.000	-92.9188	-80.7012
7	4	356.59000 [*]	2.70042	.000	350.4812	362.6988
	5	356.59000 [*]	2.70042	.000	350.4812	362.6988
	6	123.17000 [*]	2.70042	.000	117.0612	129.2788
	8	-411.05000 [*]	2.70042	.000	-417.1588	-404.9412
	9	-66.96000 [*]	2.70042	.000	-73.0688	-60.8512
	10	-56.96000 [*]	2.70042	.000	-63.0688	-50.8512
	11	75.97000 [*]	2.70042	.000	69.8612	82.0788
	12	36.36000 [*]	2.70042	.000	30.2512	42.4688
8	4	767.64000 [*]	2.70042	.000	761.5312	773.7488
	5	767.64000 [*]	2.70042	.000	761.5312	773.7488
	6	534.22000 [*]	2.70042	.000	528.1112	540.3288
	7	411.05000 [*]	2.70042	.000	404.9412	417.1588
	9	344.09000 [*]	2.70042	.000	337.9812	350.1988
	10	354.09000 [*]	2.70042	.000	347.9812	360.1988

(I) Initial pH	(J) Initial pH				95% Confidence interval	
		Mean difference (I-J)	Std. error	Sig.	Lower bound	Upper bound
8	11	487.02000 [*]	2.70042	.000	480.9112	493.1288
	12	447.41000 [*]	2.70042	.000	441.3012	453.5188
9	4	423.55000 [*]	2.70042	.000	417.4412	429.6588
	5	423.55000 [*]	2.70042	.000	417.4412	429.6588
	6	190.13000 [*]	2.70042	.000	184.0212	196.2388
	7	66.96000 [*]	2.70042	.000	60.8512	73.0688
	8	-344.09000 [*]	2.70042	.000	-350.1988	-337.9812
	10	10.00000 [*]	2.70042	.005	3.8912	16.1088
	11	142.93000 [*]	2.70042	.000	136.8212	149.0388
	12	103.32000 [*]	2.70042	.000	97.2112	109.4288
10	4	413.55000 [*]	2.70042	.000	407.4412	419.6588
	5	413.55000 [*]	2.70042	.000	407.4412	419.6588
	6	180.13000 [*]	2.70042	.000	174.0212	186.2388
	7	56.96000 [*]	2.70042	.000	50.8512	63.0688
	8	-354.09000 [*]	2.70042	.000	-360.1988	-347.9812
	9	-10.00000 [*]	2.70042	.005	-16.1088	-3.8912
	11	132.93000 [*]	2.70042	.000	126.8212	139.0388
	12	93.32000 [*]	2.70042	.000	87.2112	99.4288
11	4	280.62000 [*]	2.70042	.000	274.5112	286.7288

(I) Initial pH	(J) Initial pH				95% Confidence interval	
		Mean difference (I-J)	Std. error	Sig.	Lower bound	Upper bound
11	5	280.62000*	2.70042	.000	274.5112	286.7288
	6	47.20000*	2.70042	.000	41.0912	53.3088
	7	-75.97000*	2.70042	.000	-82.0788	-69.8612
	8	-487.02000*	2.70042	.000	-493.1288	-480.9112
	9	-142.93000*	2.70042	.000	-149.0388	-136.8212
	10	-132.93000*	2.70042	.000	-139.0388	-126.8212
	12	-39.61000*	2.70042	.000	-45.7188	-33.5012
12	4	320.23000*	2.70042	.000	314.1212	326.3388
	5	320.23000*	2.70042	.000	314.1212	326.3388
	6	86.81000*	2.70042	.000	80.7012	92.9188
	7	-36.36000*	2.70042	.000	-42.4688	-30.2512
	8	-447.41000*	2.70042	.000	-453.5188	-441.3012
	9	-103.32000*	2.70042	.000	-109.4288	-97.2112
	10	-93.32000*	2.70042	.000	-99.4288	-87.2112
	11	39.61000*	2.70042	.000	33.5012	45.7188

*. The mean difference is significant at the 0.05 level.

Appendix E 3 Comparison of initial pH 8 under between mesophilic with thermophilic condition

Independent Samples Test

		Levene's test for equality of variances		t-test for equality of means	
		F	Sig.	t	df
Hydrogen	Equal variances assumed	6.685E14	.000	-27.599	2
	Equal variances not assumed			-27.599	1.918

Appendix E 4 Comparison of initial F/M ratios

	Sum of squares	df	Mean square	F	Sig.
Between groups	61029.592	4	15257.398	452.040	.000
Within groups	168.761	5	33.752		
Total	61198.353	9			

(I) F/M ratio	(J) F/M ratio	95% Confidence interval				
		Mean difference (I-J)	Std. error	Sig.	Lower bound	Upper bound
4	6	115.09500 [†]	5.80967	.000	100.1608	130.0292
	8	170.48500 [†]	5.80967	.000	155.5508	185.4192
	10	191.19500 [†]	5.80967	.000	176.2608	206.1292
	12	222.12500 [†]	5.80967	.000	207.1908	237.0592
6	4	-115.09500 [*]	5.80967	.000	-130.0292	-100.1608
	8	55.39000 [*]	5.80967	.000	40.4558	70.3242
	10	76.10000 [*]	5.80967	.000	61.1658	91.0342
	12	107.03000 [*]	5.80967	.000	92.0958	121.9642

(I) F/M ratio	(J) F/M ratio				95% Confidence interval	
		Mean difference (I-J)	Std. error	Sig.	Lower bound	Upper bound
8	4	-170.48500*	5.80967	.000	-185.4192	-155.5508
	6	-55.39000*	5.80967	.000	-70.3242	-40.4558
	10	20.71000*	5.80967	.016	5.7758	35.6442
	12	51.64000*	5.80967	.000	36.7058	66.5742
10	4	-191.19500*	5.80967	.000	-206.1292	-176.2608
	6	-76.10000*	5.80967	.000	-91.0342	-61.1658
	8	-20.71000*	5.80967	.016	-35.6442	-5.7758
	12	30.93000*	5.80967	.003	15.9958	45.8642
12	4	-222.12500*	5.80967	.000	-237.0592	-207.1908
	6	-107.03000*	5.80967	.000	-121.9642	-92.0958
	8	-51.64000*	5.80967	.000	-66.5742	-36.7058
	10	-30.93000*	5.80967	.003	-45.8642	-15.9958

*. The mean difference is significant at the 0.05 level.

Appendix E 5 Comparison of initial iron concentration

	Sum of squares	df	Mean square	F	Sig.
Between groups	52404.908	4	13101.227	3262.445	.000
Within groups	20.079	5	4.016		
Total	52424.987	9			

(I) Iron concentration	(J) Iron concentration				95% Confidence Interval	
		Mean difference (I-J)	Std. error	Sig.	Lower bound	Upper bound
0	100	-131.84000*	2.00394	.000	0	100
	200	-60.84000*	2.00394	.000		200
	300	36.78000*	2.00394	.000		300
	400	71.94500*	2.00394	.000		400
100	0	131.84000*	2.00394	.000	100	0
	200	71.00000*	2.00394	.000		200
	300	168.62000*	2.00394	.000		300
	400	203.78500*	2.00394	.000		400
200	0	60.84000*	2.00394	.000	200	0
	100	-71.00000*	2.00394	.000		100
	300	97.62000*	2.00394	.000		300
	400	132.78500*	2.00394	.000		400
300	0	-36.78000*	2.00394	.000	300	0
	100	-168.62000*	2.00394	.000		100
	200	-97.62000*	2.00394	.000		200
	400	35.16500*	2.00394	.000		400
400	0	-71.94500*	2.00394	.000	400	0
	100	-203.78500*	2.00394	.000		100
	200	-132.78500*	2.00394	.000		200
	300	-35.16500*	2.00394	.000		300

*. The mean difference is significant at the 0.05 level.

Appendix E 6 Comparison of pre-treatment method

	Sum of Squares	df	Mean square	F	Sig.
Between groups	521872.958	3	173957.653	67527.196	.000
Within groups	10.304	4	2.576		
Total	521883.263	7			

(I) pretreatment method	(J) pretreatment method	95% Confidence Interval				
		Mean Difference (I-J)	Std. error	Sig.	Lower bound	Upper bound
Heat	Re-heat	-67.91500 [*]	1.60503	.000	-72.3713	-63.4587
	Chloroform	588.18000 [*]	1.60503	.000	583.7237	592.6363
	Ultrasonic	145.29000 [*]	1.60503	.000	140.8337	149.7463
Re-heat	Heat	67.91500 [*]	1.60503	.000	63.4587	72.3713
	Chloroform	656.09500 [*]	1.60503	.000	651.6387	660.5513
	Ultrasonic	213.20500 [*]	1.60503	.000	208.7487	217.6613
Chloroform	Heat	-588.18000 [*]	1.60503	.000	-592.6363	-583.7237
	Re-heat	-656.09500 [*]	1.60503	.000	-660.5513	-651.6387
	Ultrasonic	-442.89000 [*]	1.60503	.000	-447.3463	-438.4337
Ultrasonic	Heat	-145.29000 [*]	1.60503	.000	-149.7463	-140.8337
	Re-heat	-213.20500 [*]	1.60503	.000	-217.6613	-208.7487
	Chloroform	442.89000 [*]	1.60503	.000	438.4337	447.3463

*. The mean difference is significant at the 0.05 level.