

Half-year report: 'Insect-based ingredients in aquafeed'

Manon Eggink, 01.08.2020

February

76h

During February, I have worked on a time schedule for the three year project of the PhD and design of the different trials that will be performed during the PhD. A more detailed schedule of the PhD project can be found in Appendix I. One pre-trial and a larger scale trial will be performed at ENORM to investigate the influence of the rearing substrate on the nutritional composition of black soldier fly larvae. Furthermore, different inclusion levels of ENORMs black soldier fly meal will be tested in fish trials with rainbow trout and Nile tilapia. Lastly, the effects of the replacement of currently used protein sources with black soldier fly meal on fish health will be investigated.

Furthermore, a desk study has been performed on the use of black soldier fly larvae as fish feed ingredients. The most important findings were that black soldier fly meal can be used as replacement of fishmeal to a certain extent, however, it is likely that crystalline amino acids need to be added to sustain growth, although this needs to be confirmed with analysis of ENORM samples. Additionally, it will be difficult to include black soldier fly larvae meal high in fat to replace current protein sources, therefore appropriate defatting processes are required. Lastly, black soldier fly oil can only replace current lipid sources to a small extent, due to the high amount of saturated fatty acids, which can negatively affect fish health and performance.

March

155.4h

March was used to elaborate and discuss the time schedule for the three years with the supervisors Anne Johanne Tang Dalsgaard and Ivar Lund. Additionally, the PhD proposal has been handed in, and accepted by the PhD board.

The first work package of the PhD project is to determine the nutritional composition of black soldier fly larvae and to quantify the chitin. Chitin is found in the exoskeleton of insects, and provides structure and protection. It is a polymer consisting of N-acetylglucosamine subunits connected by β 1,4-linkages, whereas chitosan is its deacetylated derivative consisting of N-glucosamine units with β 1,4-linkages. Several methods have been previously described to determine or quantify chitin, including HPLC, NMR, and FT-IR. Due to the absence of most of these machines at the DTU location in Hirtshals, it was decided to start with a method based on weight before and after extraction as previously described by Liu et al. (2012). In this method, samples are treated with an acid (HCl) to remove catechols & minerals and a base (NaOH) to remove cuticle proteins. For more details, see the protocol used in Appendix 2.

There were some observed issues with this method: 1) large sample sizes were required (5g) which makes it difficult to use when investigating larvae samples, 2) large quantities of HCl and

NaOH are needed, and 3) the method is not very accurate because sample is lost during the filtration step and it can be that there are still other components in the sample left e.g. lipids that can lead to an overestimation of the chitin. It has therefore been decided to investigate the use of two different methods to quantify chitin: spectrophotometry and fluorescence. For the future, it would be interesting to investigate the use of HPLC for chitin quantification, although, Manon has currently no experience in HPLC and therefore needs to do a course in HPLC which was originally planned in June but because of corona, the course has been delayed to late October- beginning of November. The corona outbreak had also affected the possibilities to perform lab analysis in March.

April

140.6h

During April, different fractions of larvae samples send by ENORM have been analyzed for proximate and amino acid composition in our lab. The fractions that were sent: hele larver, fiber juicer, larvejuicer juicer, fibre baader, larvae juicer baader, fibre baader, larvae juicer baader. It was observed that large differences in proximate composition were found between these different fractions, mainly in DM, protein, and fat content. Also for the amino acid profile, differences in the fractions were observed (Appendix 3). It should be mentioned that total organic nitrogen was determined using the Kjeldahl method, with a conversion factor of 6,25 to approach the protein content. However, recent findings have showed that the use of 6,25 overestimates the protein content for black soldier fly larvae due to the presence of non-protein nitrogen (Janssen et al., 2017). Janssen et al. (2017) recommended the use of 4,67 as conversion factor, and calculations using this conversion factor have therefore been included.

Sample ID	Dry matter (DM)	Ash (%DM)	Protein (%DM)	Fat (%DM)
Hele larver 1.1 100%	30,8	9,8	37,0 ^a 27,7 ^b	38,0
Fiber juicer 1.1	58,7	10,3	67,8 ^a 50,7 ^b	7,2
Larvejuicer juicer 1.1 91.3%	27,4	9,9	29,6 ^a 22,1 ^b	44,9
Fiber baader 1.1 20%	39,9	8,7	54,4 ^a 40,6 ^b	21,3
Larvejuicer baader 1.1	27,6	9,9	32,2 ^a 24,1 ^b	42,4
Fiber baader 12%	46,0	10,1	59,8 ^a 44,7 ^b	8,7
Larvejuicer baader 88%	29,8	9,3	31,2 ^a 23,3 ^b	50,0

^a Conversion factor 6,25; ^b Conversion factor 4,67

May

133.2h

In May, a desk study was performed on the different probable rearing substrates for black soldier fly rearing. After the desk study, different substrates were selected for the insect trials: crushed mussels, crushed shrimp waste, and brewer's mash. These are by- and waste-products widely available in Denmark. For external comparison, one group will be fed chicken feed whilst for internal comparison, one group will be fed ENORM mix. These substrates will be tested first on small-scale, and later on larger scale when observed that the larvae can use the substrate for growth.

Furthermore, ENORM has sent samples of the grax and protein fraction, which is samples before and after the drying process, respectively. Samples 1/3-3/3 are obtained from different containers but are replicates. It can be seen that the dry matter content is much higher in the protein fraction, which is expected as the protein fraction was dried. On dry matter basis, ash, protein, and fat content are a bit higher in the grax fraction compared to the protein fraction, indicating that the drying has a minor effect on the proximate composition. The amino acid profile of the protein and grax fraction were overall similar, showing that the drying process did not have large effects on the amino acid profile (Appendix 4).

Sample ID	Dry matter (DM)	Ash (%DM)	Protein (%DM)	Fat (%DM)
Protein 1/3	94,9	7,5	55,6 ^a 41,6 ^b	20,4
Protein 2/3	94,9	7,5	55,2 ^a 41,3 ^b	20,9
Protein 3/3	95,1	7,5	55,1 ^a 41,2 ^b	20,8
Grax 1/3	31,9	7,9	59,6 ^a 44,5 ^b	22,5
Grax 2/3	31,8	8,0	59,1 ^a 44,2 ^b	22,3
Grax 3/3	31,9	8,0	59,6 ^a 44,5 ^b	22,3

^a Conversion factor 6,25; ^b Conversion factor 4,67

June

155.4h

In June, ENORM has sent us the samples that were mechanically separated on size (0-200 µm, 200-400 µm, and >400 µm) assuming that the largest fraction (>400 µm) has the highest quantity of chitin. Additionally, a proto-type of insect meal has been sent to us to investigate the nutritional composition. It was found that the mechanical separation seem to increase protein and fat content whilst decreasing ash content. However, the mechanical separation has to be performed several times to investigate whether this trend is also seen in different batches of insect samples.

Sample ID	Dry matter (DM)	Ash (%DM)	Protein (%DM)	Fat (%DM)
0-200 um	93,3	8,4	53,7 ^a 40,1 ^b	23,9
0-400 um	94,6	7,6	55,0 ^a 41,1 ^b	21,2
200-400 um	94,0	6,9	56,1 ^a 41,9 ^b	19,7
>400 um	94,4	6,5	57,6 ^a 43,1 ^b	17,2
Insect meal	99,8	14,6	58,7 ^a 43,9 ^b	13,6

^a Conversion factor 6,25; ^b Conversion factor 4,67

July

170.2h

In July, most of the focus was put on finding a method for chitin quantification using either fluorescence methods or spectrophotometry. Two spectrophotometry methods were tested (Lehmann and White, 1975; Han and Heinonen, 2020) and one fluorescence method (Henriques et al., 2020). The spectrophotometry method is based on the reaction of hexosamines that deminate into 2,5-anhydrohexoses that react with 3-methyl-2-benzothiazolone hydrazone hydrochloride (MBTH) and FeCl₃ that yield into an intense blue colour. Whereas the fluorescence method is based on staining the chitin with calcofluor.

For the methods described by Lehmann and White (1975) and Henriques et al. (2020), large variations were observed between replicates as seen by the large standard deviation (Appendix 6 & 7). These variations are likely due to the difficulties with homogenization of these samples in distilled water in early steps of the quantification. However, the spectrophotometry method of Han and Heinonen (2020), which is also based on the reaction of MBTH and FeCl₃ similar to that of Lehmann and White (1975), and showed potential. Therefore, the protocol of Han and Heinonen (2020) is currently optimized to enhance repeatability (Appendix 5).

Additionally, ENORM has performed another time of mechanical separation with a different batch. Samples 101-104 (0-200 um) and 121-124 (200-400 um) were separated once, whereas the largest fraction (>400 um) was again separated to sample 201 (0-200 um), 221 (200-400 um), and 241-244 (>400 um). It can be seen that the fat content seems to decrease with mechanical separation, as was seen previously in June. The difference in ash and protein content between the different fractions as was seen in June, was less pronounced this time.

Sample ID	Dry matter (DM)	Ash (%DM)	Protein (%DM)	Fat (%DM)
101-104 (0-200 um)	96,5	13,6	54,9 ^a 41,0 ^b	21,5
121-124 (200-400 um)	96,4	13,5	55,7 ^a 41,6 ^b	20,2
201 (0-200 um)	96,3	13,6	54,9 ^a 41,0 ^b	19,2
221 (200-400 um)	96,7	11,9	56,2 ^a 42,0 ^b	16,3
241-244 (>400 um)	98,1	10,6	56,1 ^a 41,9 ^b	13,3

^a Conversion factor 6,25; ^b Conversion factor 4,67

Han, X., & Heinonen, M. (2020). Development of ultra-high performance liquid chromatographic and fluorescent method for the analysis of insect chitin. *Food Chemistry*, 127577.

Henriques, B. S., Garcia, E. S., Azambuja, P., & Genta, F. A. (2020). Determination of Chitin Content in Insects: An Alternate Method Based on Calcofluor Staining. *Frontiers in Physiology*, 11, 117.

Janssen, R. H., Vincken, J. P., van den Broek, L. A., Fogliano, V., & Lakemond, C. M. (2017). Nitrogen-to-protein conversion factors for three edible insects: *Tenebrio molitor*, *Alphitobius diaperinus*, and *Hermetia illucens*. *Journal of Agricultural and Food Chemistry*, 65(11), 2275-2278.

Lehmann, P. F., & White, L. O. (1975). Chitin assay used to demonstrate renal localization and cortisone-enhanced growth of *Aspergillus fumigatus* mycelium in mice. *Infection and immunity*, 12(5), 987-992.

Purkayastha, D., & Sarkar, S. (2020). Physicochemical Structure Analysis of Chitin Extracted from Pupa Exuviae and Dead Imago of Wild Black Soldier Fly (*Hermetia illucens*). *Journal of Polymers and the Environment*, 28(2), 445-457.

Waśko, A., Bulak, P., Polak-Berecka, M., Nowak, K., Polakowski, C., & Bieganowski, A. (2016). The first report of the physicochemical structure of chitin isolated from *Hermetia illucens*. *International Journal of Biological Macromolecules*, 92, 316-320.

Appendix 2

Chitin quantification based on Liu et al. (2012)

1. Dry samples for 2 days at 50 degrees Celsius
2. Micronize samples if needed
3. Store samples at 4 degrees until analysis
4. Weigh 5g sample in an Erlenmeyer
5. Treat the sample with 250 mL 1 M HCl at 100 degrees for 30 minutes
6. Filter the sample
7. Wash the sample with distilled water until neutrality is reached
8. Treat the sample with 250 mL 1 M NaOH at 80 degrees for 24 hours
9. Filter the sample
10. Wash the sample with distilled water until neutrality is reached
11. Dry the sample at 50 degrees in the oven until constant weight is achieved

Appendix 3

Relative amino acid content, % of total amino acids	Hele larver 1.1 100%	Fiber juicer 1.1	Larvejuice juicer 1.1 91.3%	Fiber baader 1.1 20%	Larvejuice baader 1.1	Fiber baader 12%	Larvejuice baader 88%
Hydroxyproline (Hypro)	ND ¹	ND ¹	ND ¹	ND ¹	ND ¹	ND ¹	ND ¹
<i>Histidine (His)</i>	3,5	2,9	3,8	3,2	3,5	3,0	3,5
Taurine (Tau)	0,3	0,2	0,3	0,0	0,3	0,0	0,3
Serine (Ser)	4,9	5,7	4,3	5,7	4,4	6,2	4,6
<i>Arginine (Arg)</i>	5,6	4,6	6,2	4,6	6,3	4,1	6,0
Glycine (Gly)	5,7	7,3	4,7	7,3	4,6	8,5	4,9
Aspartate (+ asparagine) (Asp + Asn) (3)	10,7	7,9	12,5	8,2	12,6	6,9	11,3
Glutamate (+ glutamine) (Glu + Gln) (3)	13,7	8,9	15,9	10,7	15,5	9,9	16,2
<i>Threonine (Thr)</i>	4,6	4,3	4,8	4,2	4,8	3,9	4,6
Alanine (Ala)	7,1	10,3	5,2	10,1	5,2	11,1	5,9
Cysteine (Cys)	0,0	0,0	0,0	0,0	0,0	0,0	0,0
Proline (Pro)	7,0	9,2	5,4	9,2	5,2	10,4	6,2
Cystine (Csn)	0,2	0,1	0,3	0,0	0,3	0,1	0,4
<i>Lysine (Lys)</i>	6,7	5,1	8,3	4,9	8,0	3,8	7,4
Tyrosine (Tyr)	6,2	7,9	5,1	7,6	5,5	8,2	5,6
<i>Methionine (Met)</i>	1,5	0,8	1,8	0,7	2,0	0,5	1,9
<i>Valine (Val)</i>	6,2	8,5	5,0	8,2	5,0	8,5	5,4
<i>Isoleucine (Ile)</i>	4,7	4,7	4,8	4,5	4,8	4,4	4,6
<i>Leucine (Leu)</i>	7,4	8,2	7,1	7,8	7,3	8,0	7,0
<i>Phenylalanine (Phe)</i>	4,1	3,4	4,6	3,1	4,7	2,5	4,3
<i>Tryptophan (Trp)</i> (4)	ND ¹	ND ¹	ND ¹	ND ¹	ND ¹	ND ¹	ND ¹
Sum of amino acids	100	100	100	100	100	100	100

¹ND= not detected

Appendix 4

Relative amino acid content, % of total amino acids	Protein 1/3	Protein 2/3	Protein 3/3	Grax 1/3	Grax 2/3	Grax 3/3
Hydroxyproline (Hypro) (2)	ND ¹	ND ¹	ND ¹	ND ¹	ND ¹	ND ¹
<i>Histidine (His)</i>	3,0	3,0	3,0	3,0	3,0	3,0
Taurine (Tau)	0,0	0,2	0,0	0,0	0,2	0,0
Serine (Ser)	5,0	5,0	5,1	5,0	5,1	5,2
<i>Arginine (Arg)</i>	4,7	4,7	4,7	4,5	4,6	4,3
Glycine (Gly)	6,4	6,4	6,3	6,4	6,5	6,6
Aspartate (+ asparagine) (Asp + Asn) (3)	9,2	9,2	9,3	9,6	9,2	9,0
Glutamate (+ glutamine) (Glu + Gln) (3)	11,2	11,2	11,3	11,5	11,1	11,2
<i>Threonine (Thr)</i>	4,6	4,6	4,6	4,6	4,6	4,5
Alanine (Ala)	8,8	8,7	8,7	8,7	8,8	9,1
Cysteine (Cys)	0,0	0,0	0,0	0,0	0,0	0,0
Proline (Pro)	7,5	7,5	7,4	7,4	7,5	7,7
Cystine (Csn)	0,2	0,2	0,2	0,2	0,1	0,1
<i>Lysine (Lys)</i>	5,7	5,7	5,7	6,1	5,9	5,8
Tyrosine (Tyr)	7,4	7,4	7,4	7,3	7,5	7,5
<i>Methionine (Met)</i>	1,6	1,5	1,6	1,3	1,3	1,3
<i>Valine (Val)</i>	6,9	6,9	6,9	6,7	6,8	6,9
<i>Isoleucine (Ile)</i>	5,1	5,1	5,1	5,1	5,0	5,0
<i>Leucine (Leu)</i>	8,3	8,3	8,2	8,3	8,3	8,3
<i>Phenylalanine (Phe)</i>	4,4	4,5	4,5	4,5	4,5	4,3
<i>Tryptophan (Trp)</i> (4)	ND ¹	ND ¹	ND ¹	ND ¹	ND ¹	ND ¹
Sum of amino acids	100	100	100	100	100	100

¹ND= not detected

Appendix 5

Sample preparation

1. Freeze samples at -20 degrees
2. Freeze dry to remove moisture until constant weight
3. Grind into a powder using micronisation and seal in plastic bags
4. Keep in desiccator under -20 degrees to keep the original composition

Preparation

Make a 0.5M NaOH solution

Cool the centrifuge to 5 degrees

Removal of protein

1. Add 0.5g sample to 10 mL 0.5M NaOH solution in a centrifuge tube

To remove proteins

1. Agitate with a stirrer for 2 hours
2. Centrifuge at 12000 rpm, 5 degrees, 15 minutes
3. Discard the supernatant
4. Wash the pellet with distilled water
5. Centrifuge at 12000 rpm, 5 degrees, 15 minutes
6. Add again 10 mL 0.5M NaOH solution to the sample
7. Agitate with a stirrer for 2 hours
8. Centrifuge at 12000 rpm, 5 degrees, 15 minutes
9. Wash pellet with distilled water
10. Move the pellet in buckets able to use for freeze drying
11. Freeze dry the samples until constant weight is achieved (~2-5 days)
12. Store in a desiccator at room temperature for later use

Demineralization

1. Hydrolyze 10 mg freeze dried deproteinized insect sample in 3 mL w/v 6 M HCl for 24h in a heating block at 100 degrees
2. Take 3 mL hydrolysate and add ± 1.4 mL 12 M NaOH to adjust the pH to 6.0-6.5 by adding NaOH solution
3. Move the mixture into a centrifuge tube and add up to 10 mL with distilled water
4. Take 1 mL of the sample into a new centrifuge tube and for the reagent blank 1 mL distilled water

Measurement of glucosamine

1. Add 1 mL 5% NaNO_2 and 1 mL of 5% KHSO_4 to each 1 mL sample solution and to a reagent blank
2. Leave solution standing for 15 minutes at room temperature with occasional shaking

Preparation

Make 0.5% w/v MBTH solution in distilled water

1. Add 1 mL of 12.5% $\text{NH}_4\text{SO}_3\text{NH}_2$ (slowly, as this gets foamy) and shake constantly for 5 minutes
2. Add 1 mL of 0.5% fresh MBTH and let it stand for 60 minutes at room temperature
3. Add 1 mL of 0.5% fresh FeCl_3 and let it stand for minimally 30 minutes
4. Measure the absorbance at 650 nm against the reagent blank

MBTH and FeCl_3 need to be made every three days and stored in the fridge.

Standard curve

A standard curve was constructed to determine the glucosamine content in the samples as a measurement for chitin.

GlcN-HCl standard solution:

1. Standard solution I
30 mg/10 mL GlcN-HCl in distilled water
2. Standard solution II
1 mL of standard solution I with 9 mL distilled water (1:10 dilution)

Standard solution II was diluted according to the scheme below to obtain a final volume of 10 mL per standard solution, all concentrations were made in duplicate:

Standard solution concentration GlcN-HCl ($\mu\text{g/mL}$)	Pipetted volume standard solution II (μL)	Pipetted volume distilled water (μL)
0.3	10	9990
0.99	33	9977
6	200	9800
12	400	9600
18	600	9400
24	800	9200
30	1000	9000

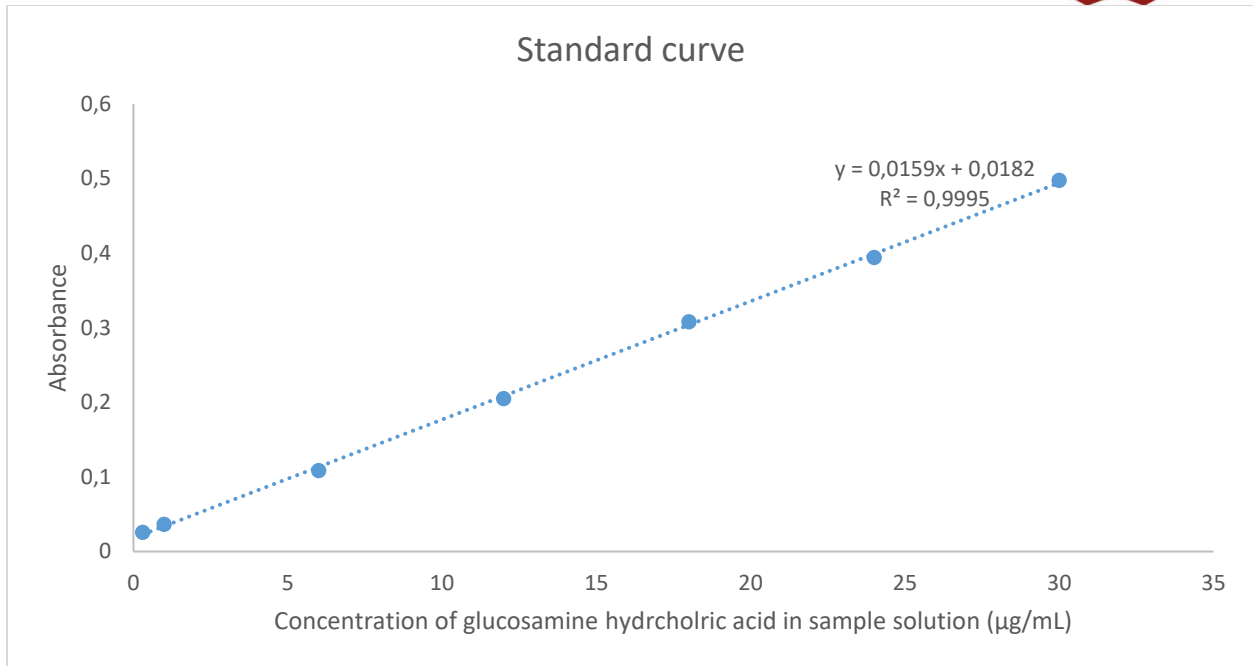
Measurement

1. Add 1 mL 5% NaNO_2 and 1 mL of 5% KHSO_4 to each 1 mL standard solution and to a reagent blank (1 mL distilled water)
2. Leave solution standing for 15 minutes at room temperature with occasional shaking
3. Add 1 mL of 12.5% $\text{NH}_4\text{SO}_3\text{NH}_2$ (slowly, as this gets foamy) and shake constantly for 5 minutes
4. Add 1 mL of 0.5% fresh MBTH and let it stand for 60 minutes at room temperature
5. Add 1 mL of 0.5% fresh FeCl_3 and let it stand for minimally 30 minutes
6. Measure the absorbance at 650 nm against the reagent blank

Standard curve calculations

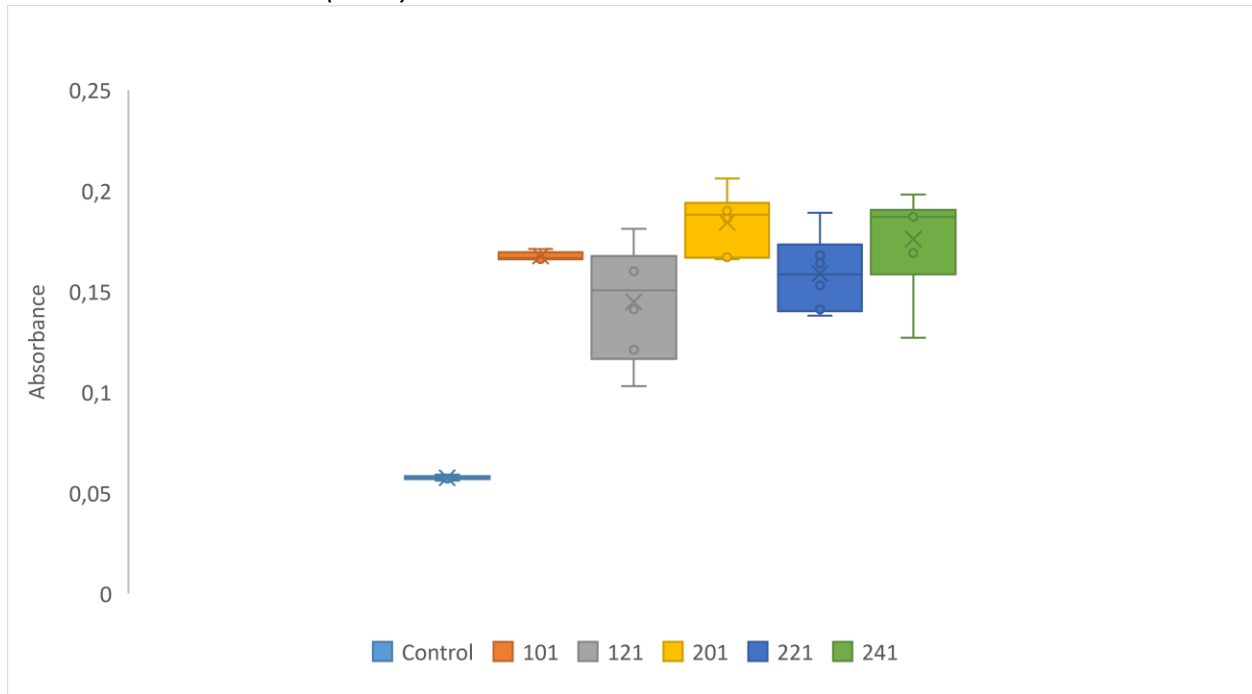
Y-axis: standard absorbance minus the absorbance of the reagent blank (background deduction)

X-axis: concentration of GlcN-HCl ($\mu\text{g/mL}$)



Appendix 6

Absorbance measured using spectrophotometry for internal control and samples 101-241 using the Lehmann and White (1975) method.



Appendix 7

Standard curve determined using fluorescence with chitin samples 1-50 mg initial weight

