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1	Extraction and characterization of protein fractions from five insect species
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19	ABSTRACT
20	Tenebrio molitor, Zophobas morio, Alphitobius diaperinus, Acheta domesticus and Blaptica
21	dubia were evaluated for their potential as a future protein source. Crude protein content
22	ranged from 19 - 22 % (Dumas analysis). Essential amino acid levels in all insect species
23	were comparable with soybean proteins, but lower than for casein. After aqueous extraction,
24	next to a fat fraction, a supernatant, pellet, and residue were obtained, containing $17 - 23 \%$,

25	33 - 39 %, $31 - 47$ % of total protein, respectively. At 3 % (w/v), supernatant fractions did
26	not form stable foams and gels at pH 3, 5, 7, and 10, except for gelation for A. domesticus at
27	pH 7. At 30 % w/v, gels at pH 7 and pH 10 were formed, but not at pH 3 and pH 5. In
28	conclusion, the insect species studied have potential to be used in foods due to: 1) absolute
29	protein levels; 2) protein quality; 3) ability to form gels.
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31	Keywords
32	Insect protein; Protein extraction; Protein characterization; Foaming; Gelation.
33	
34	1. Introduction
35	1.1. Insects as a source of food
36	In most developed countries, human consumption of insects is infrequent, or even culturally
37	inappropriate, although its nutritional value is comparable to conventional meat (van Huis,
38	2013). In many regions and countries of the world, insects form part of the human diet and it
39	is a misconception to believe that this is prompted by starvation (van Huis, 2013). About
40	1900 insect species are consumed globally as human food in the world
41	(<u>http://www.ent.wur.nl/UK/Edible+insects/Worldwide+species+list/</u>).
42	With an increase in the world population, increased consumer demand for protein, and the
43	amount of available agricultural land being constrained, the sustainable production of meat
44	will represent a serious challenge for the future. Insects can be considered as an alternative
45	protein source with less environmental impact (van Huis, 2013). Insects can be consumed as
46	a whole. However, they can also be processed in less recognizable forms, which may increase
47	consumer acceptability. Insects are already used as natural food ingredients, e.g. the red
48	colorant carmine (F120) used in vogurt is an extract of the female cochineal insect

49	1.2. Edible insects
50	Insects are consumed in different life stages like eggs, larvae, pupae or adults. The main
51	species consumed are, in order of importance: beetles (Coleoptera); caterpillars (Lepidoptera)
52	ants, bees and wasps (Hymenoptera); grasshoppers and locusts (Orthoptera); true bugs,
53	aphids and leafhoppers (Hemiptera); termites (Isoptera) and flies (Diptera) and some others.
54	Lepidoptera, Coleoptera, and Diptera (including flies) are commonly consumed in the larval
55	stage; while the Orthoptera, Hymenoptera, Hemiptera and Isoptera are mainly consumed in
56	the adult stage.
57	Cultivating edible insects for food consumption has several advantages: 1) Insects have a
58	high feed conversion efficiency compared with conventional livestock. For example, the feed
59	conversion ratio of house cricket (Acheta domesticus) can be calculated twice as efficient as
60	chickens, almost 4 times more efficient than pigs and over 12 times more than cattle (van
61	Huis, 2013); 2) Cultivating insects for protein has less environmental impact than cattle
62	ranching, due to the lower production of greenhouse gas and NH ₃ emissions (<u>van Huis, 2013</u>)
63	3) Besides the higher production yield and less environmental impact, insect feeds can be
64	obtained from a wider range of plants than that of conventional livestock, such as cattle or
65	swine (Durst & Shono, 2010). Overall, insect farming can be introduced in terms of a
66	sustainable form of agriculture.
67	1.3. Proteins of edible insects
68	As a food source, insects are potentially nutritious, rich in protein and fat, and providing a
69	certain amount of minerals and vitamins. Studies on protein quality, nutritional value, protein
70	content, and the amino acid composition of various insects are available (Ladrón de Guevara,
71	Padilla, García, Pino, & Ramos-Elorduy, 1995); (Renault, Bouchereau, Delettre, Hervant, &

Vernon, 2006); (Barker, Fitzpatrick, & Dierenfeld, 1998). The protein content of common

73 edible insects was around 9 - 25 % (Finke & Winn, 2004), and the Yellow mealworm beetle larvae (24 %) (Ghaly & Alkoaik, 2009), Zophobas morio larvae (19 %) (Finke, 2002), and 74 75 Acheta domesticus adult (19 %) (Finke & Winn, 2004), conventional meat protein sources 76 contain about 15 to 22 % protein (Ghaly & Alkoaik, 2009). In addition, some insects have not only protein content comparable to meat, but also to plant protein (up to 36.5 %). 77 People may consume insect food more easily when unrecognizable insect protein (extract) is 78 79 incorporated in food in comparison to consuming whole insects. (Del Valle, Mena, & 80 Bourges, 1982) also indicated that the extraction of proteins from insects for further use in food products is particularly relevant for countries that do not have the habit of consuming 81 82 insects, such as Europe and North America. In this study, there are five insect species selected based on their availability (species reared 83 84 by companies in the Netherlands): three species of Coleoptera considered edible, including 85 the Yellow mealworm (Tenebrio molitor), the Superworm (Zophobas morio), the Lesser 86 mealworm (Alphitobius diaperinus) and one species of Orthoptera; the House cricket (Acheta domesticus) considered edible and one of the Blattodea; the Dubia cockroach (Blaptica dubia) 87 not edible, but can be reared in large numbers and used for animal feed. 88 89 **Objective** 1.4. 90 Although researchers from entomological and zoo-biology science have studied intact edible insects, still very little information from a food science point of view is available on 91 92 characteristics and functionality of extracted insect proteins. 93 The aim of this study was to investigate if insects could be used as a future protein source in 94 food. Therefore, insect protein characteristics and functionality were determined and 95 evaluated for each of the five insect species. The specific objectives of this study were to: (a) 96 extract proteins and characterize obtained fractions; (b) evaluate protein purity and yield of

- 97 the obtained fractions; (c) establish some functional properties of the protein fractions
- 98 focused on foaming and gelation; (d) study protein quality by analysis of protein content and
- 99 amino acid composition.

100 **2.** Materials and methods

- 101 2.1. Insects used
- 102 Tenebrio molitor, Z. morio, A. diaperinus, A. domesticus and B. dubia were purchased from
- the commercial supplier Kreca V.O.F, Ermelo, the Netherlands. *Tenebrio molitor*, Z. morio,
- 104 A. diaperinus species were supplied in the larvae stage, A. domesticus and B. dubia in the
- adult stage. The feed for *T. molitor*, and *Z. morio* mainly consisted of wheat, wheat bran, oats,
- soy, rye, corn, carrot and beer yeast. The feed for A. diaperinus, A. domesticus and B. dubia
- mainly consisted of carrot and chicken mash obtained from Kreca V.O.F. All insects were
- sieved to get rid of feed and stored alive at 4 °C for about one day before processing.
- 109 2.2. Analysis of water content, protein, and fat content
- All fresh insects were frozen using liquid nitrogen and subsequently grinded using a blender
- 111 (Braun Multiquick 5 (600 Watt), Kronberg, Germany). Frozen grinded insects were freeze-
- dried (GRI Vriesdroger, GR Instruments B.V., Wijk bij Duurstede, the Netherlands) to
- determine moisture and dry matter content. The freeze-drying process was stopped at a stable
- sample weight. Next, the freeze-dried insects were used for protein content analysis. Crude
- protein content was determined by Dumas (Thermo Quest NA 2100 Nitrogen and Protein
- Analyser, Interscience, Breda, the Netherlands) using a protein-to-nitrogen conversion factor
- of 6.25. D-methionine (Sigma, CAS nr. 348-67-4) was used as a standard. Furthermore, fat
- 118 content was determined after hexane extraction (Biosolve, CAS nr. 110-54-3) in a Soxhlet
- apparatus for 6 hours. Afterwards, hexane was removed using a Rotary evaporator (R420,

- Buchi, Switzerland). Defatted insect meal was stored at 20 °C. All experiments were performed in two duplications of the same sample.
- 122 2.3. Determination of amino acid composition and protein quality
- 123 Amino acid composition of freeze-dried insect powder was analysed using ion exchange chromatography, following the International standard ISO 13903:2005. Tryptophan was 124 125 determined by reversed phase C₁₈ HPLC using fluorescence detection at 280 nm, according to 126 the procedure described by International standard ISO 13904:2005. The amino acid composition of the five insect species was compared to literature data of soybean protein and 127 casein, representing high quality proteins among vegetable and animal proteins (Sosulski & 128 Imafidon, 1990; Young & Pellett, 1994). Protein quality was evaluated by the essential amino 129 acid index (EAAI), which is based on the content of all essential amino acids compared to a 130 131 reference protein, being values for human requirements in this case (Smith & Nielsen, 2010). EAAI gives an estimate on the potential of using insects as a protein source for human 132 133 consumption without correcting for protein digestibility (Eq.1).
- 134 EAAI = (mg of lysine in 1 g of test protein mg of lysine in 1 g reference protein) × (etc. for the other 8 essential amino acids)
- 135 2.4. Protein extraction procedure

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For protein extraction, 400 g of N₂-frozen insects was used. After adding 1200 ml demineralized water, that was mixed with 2 g ascorbic acid beforehand, blending for one minute took place (Braun Multiquick 5 (600 Watt), Kronberg, Germany). Then the obtained insect suspension was sieved through a stainless steel filter sieve with a pore size of 500 μm. The filtrates and residues were collected. After centrifugation at 15,000 g for 30 min at 4 °C, three fractions were obtained from the filtrate: the supernatant, the pellet, and the fat fraction.

- The residue, the pellet and the supernatant fractions were freeze dried for further analysis.
- 143 The freeze-dried supernatant and pellet fractions of all insect species studied were
- 144 characterized in terms of colour, protein content and molecular weight distribution using
- SDS-PAGE. The extraction procedure was performed in duplicate starting twice with a new
- insect batch.
- 147 2.5. SDS-PAGE
- Sodium dodecyl sulphate polyacrylamide gel electrophoresis (SDS-PAGE) was used to
- determine the molecular weight distribution of the insect protein fractions. For the detection
- of the supernatant, pellet and residue fractions, 12.5 % acrylamide Phastgels (15 kDa to 250
- kDa) and 20 % acrylamide Phastgels (2 kDa to 150 kDa) (GE Healthcare Bio-Sciences AB,
- Uppsala, Sweden) were used. The applied markers were ordered from SigmaMarker (S8445,
- wide range, molecular weight 6.5 200 kDa SigmaMarker). The samples were dissolved in
- 154 20 mM Tris/HCl, 2 mM EDTA pH 8.0 buffers with protein concentration of 7 mg/ml and
- placed in an ultrasonic bath for 10 min. The protein concentration of the samples was
- 156 calculated based on protein content (Dumas) and amount of dry matter. Next, protein
- solutions were diluted with ratio 1:1 in a sample buffer, containing 20 mM Tris/HCl, 2 mM
- 158 EDTA pH 8.0 (Across Organics, Cas nr. 6381-92-6), 5 % (w/v) SDS (Sigma, Cas nr. 152-21-
- 3), 0.016 % (w/v) DTT (DL- Dithiothreitol, Sigma, Cas nr. 3483-12-4), 0.02 % Bromophenol
- 160 Blue (Merck, Cas nr. 115-39-9). Afterwards, the samples were heated at 100 °C for 5 min and
- 161 centrifuged for 2 min at 10,000 rpm before applying to the gel.
- 162 2.6. Foamability and foam stability
- The stability of foam stabilized by insect supernatant protein was determined using foam
- tubes with a diameter of 2.0 cm, and a glass grid at the bottom (Deak, Murphy & Johnson,
- 165 2007). The tubes were filled with 20 ml supernatant solution with a concentration of 3 % w/v,

166	at pH 3, 5, 7, and 10. The solutions were aerated from below with nitrogen gas, at a flow rate
167	of 10.0 ml/min. Some of the samples had insufficient foamability to form stable foam at these
168	concentrations. For those samples with sufficient foamability, the samples were aerated until
169	the foam level reached 30 cm. After stopping the flow of gas, the height of the foam was
170	determined as a function of time. From these curves, the half-time of the foam (the time in
171	which foam height is reduced by 50 %) was determined. All tests were performed in
172	duplicate.
173	duplicate. 2.7. Gel formation
174	2.7.1. Visual observation of gelation
175	Insect supernatant solutions were heated in a water bath (86 \pm 1 °C) for 10, 20 and 30 min.
176	The supernatant fractions were dissolved at concentrations of 3 % w/v and 30 % w/v at pH 3,
177	5, 7 and 10. Depending on the initial pH, the final pH was adjusted by slowly adding 1 and 5
178	M HCl/ NaOH solutions. Gel formation was determined through visual observation. If the
179	liquid was not moving upon turning the tube, it was considered a gel. This method was
180	previously used by (Beveridge, Jones, & Tung, 1984) for albumin gel formation. Experiments
181	were performed in duplicate.
182	2.7.2. Strain sweeps
183	Freeze-dried supernatant fractions from five insect species were used for this experiment.
184	Protein solutions were prepared as followed: freeze-dried supernatant fractions were
185	dissolved in demineralised water at a concentration of 15 % w/v, stirred for 30 minutes at
186	room temperature and adjusted to pH 7 using 1 M NaOH.
187	To determine the rheological properties of the supernatant protein solutions and gels made
188	from them, oscillatory strain tests were performed on a stress-controlled rheometer (Physica

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MCR 501, Anton Paar, Graz, Austria) with stainless steel and titanium CC-10 concentric cylinder geometry (diameter inner cylinder: 9.997 mm; diameter cup: 10.845 mm). After filling the geometry with supernatant solution, all samples were covered with a thin layer of silicone oil to prevent sample evaporation. Samples were first heated from 20 to 90 °C at a heating rate of 1 °C/min (phase 1), kept at 90 °C for 5 min (phase 2), and cooled to 20 °C at a rate of 3 °C/min (phase 3). During the temperature ramp, the storage modulus G' and loss modulus G" were determined by applying oscillatory deformations with a strain amplitude of 0.005 and a frequency of 0.1 Hz. The point at which G' started to increase and became greater than the background noise, was designated as the gelation temperature (Renkema, Knabben, & van Vliet, 2001). After formation of the gel, an oscillatory strain sweep was performed on the samples, with strains ranging from 10⁻⁴ to 10, and a frequency of 0.1 Hz. Strain sweeps were also performed to confirm whether this strain was in the linear response regime. All samples were tested at a supernatant fraction concentration of 15 % (protein content of around 8 % for five types of insects) w/v. Tenerio molitor was also tested at concentrations of 7 % (protein content of 4.1 %), and 30 % (protein content of 16.6 %) w/v. Values for G' for this fraction from the linear response regime were plotted against protein concentration C, and the exponent n, in the relation G'~Cⁿ, was determined using linear regression to obtain information on the structure of the gels. For all fractions the maximum linear strain, where G' starts to decrease as a function of increasing strain, was also determined. This was done by separately fitting the data points in the linear region and the fully nonlinear region, and extrapolating both curves to their point of intersection (see Figure 2C). This method of determining the maximum linear strain is only approximate, but since we are not interested in the absolute value of this strain, but rather in the differences in this strain for the various protein samples,

213	this approximation was considered sufficiently accurate. All tests were performed in
214	duplicate.
215	3. Results and discussion
216	3.1. Chemical composition of five insect species
217	The proximate composition of five insect species with regard to moisture, fat, protein was
218	determined on live weight basis (Table 1). The moisture content of the five insect species
219	ranged from 60 % to 71 %, fat content ranged from 3.6 % to 16 %, and crude protein from
220	19 % to 22 % (including chitin nitrogen). Other components, calculated by difference, ranged
221	from 3.4 % to 7.5 %.
222	The proximate composition of <i>T. molitor</i> was comparable to the results of (Barker,
223	Fitzpatrick, & Dierenfeld, 1998); (Finke, 2002); (Jones, Cooper, & Harding, 1972); (Ghaly &
224	Alkoaik, 2009). In addition, the crude protein content measured for A. domesticus and Z.
225	morio, 19.3 % and 20.6 % respectively, was comparable to the range described in literature,
226	namely 17.3 % to 20.5 % (Barker, Fitzpatrick, & Dierenfeld, 1998; Finke, 2002). For A.
227	diaperinus and B. dubia, no crude protein data are available in literature. The measured crude
228	protein contents of the five insect species might be relatively higher than their actual protein
229	content, since amounts of nitrogen are also bound in the exoskeletons as chitin. (Barker,
230	Fitzpatrick, & Dierenfeld, 1998) reported that 5 - 6 % of total nitrogen was measured as
231	chitin-bound nitrogen in <i>T. molitor</i> . This would lead to an overestimation in protein content
232	of $1.1 - 1.3$ % on a fresh weight basis. It is a reasonable estimate for true protein content in
233	most insect species. However, no detailed study on this issue is available.
234	The measured protein content of the tested insect species (around 20 %) in this study is
235	comparable with that of beef (18.4 %), chicken (22.0 %) and fish (18.3 %) (Ghaly, 2009b).
236	Further, measured insect protein content was higher than that of lamb (15.4 %), pork (14.6 %)

237 (Ghaly, 2009), eggs (13 %), and milk (3.5 %), but lower in comparison to soy (36.5%) 238 (Young & Pellett, 1994). 239 3.2. Amino acid composition and protein quality of five insect species 240 The insect protein quality of the insect species was estimated by the amino acid composition (Table 2). The larvae of A. diaperinus, T. molitor and Z. morio contained all the essential 241 242 amino acids in quantities that are necessary for humans (FAO/WHO/UNU, 1985). Also, the sum of the amount of total essential amino acids (EAA) for A. diaperinus, T. 243 molitor and Z. morio was comparable to that of soybean protein, but slightly lower than that 244 of casein, as reported by (Young & Pellett, 1991). Furthermore, the sum of EAA for A. 245 domesticus and B. dubia was lower than in casein and soybean protein, but EAA were 246 247 available in quantities that are necessary for human requirement (sum of 277 mg/g crude protein). The amino acid profiles found for T. molitor were similar to the profiles that were 248 reported by (Ghaly, 2009b); (Finke, 2002) and (Jones, Cooper, & Harding, 1972). The amino 249 acid profiles of Z. morio reported by (Finke, 2002) and those of A. domesticus reported by 250 251 (DeFoliart & Benevenga, 1989) were similar to ours. To our knowledge, no literature is 252 reported on the amino acid profiles for A. diaperinus and B. dubia before. The sum of total amount of amino acids (TAA) per g crude protein of A. diaperinus (927) 253 mg/g), T. molitor (910 mg/g) and Z. morio (931 mg/g) was higher than that in A. domesticus 254 255 (864 mg/g) and B. dubia (776 mg/g). The fact that the sum of the total amount of amino acids did not add up to 1000 mg/g crude protein is mainly explained by the presence of non-protein 256 257 nitrogen in the form of chitin. Acheta domesticus and B. dubia are used in adult form and are 258 known to contain a higher level of chitin as compared to T. molitor, A. diaperinus and Z. 259 morio.

The calculated essential amino acid index (EAAI) of A. diaperinus, 1. molitor and Z. morio
was somewhat higher than that of soybean, but lower than that of casein, also indicating that
the quality of the insect protein for these three insect species was comparable to conventional
food protein sources. The EAAI of A. domesticus and B. dubia was the lowest in comparison
to other insects, and lower than the EAAI for casein and soybean. For a more detailed insight
in insect protein quality, digestibility data need to be taken into account in future studies,
since digestibility is not included as a factor in determining EAAI. (Ramos-Elorduy, Moreno,
Prado, Perez, Otero, & De Guevara, 1997) found that protein digestibility, calculated from a
vitro study, ranged from 76 to 98 % for seventy-eight species of edible insects, representing
twenty-three insect families in Mexico. Their study indicated that insect proteins might have
a high nutritional value.
3.3. Protein distribution in obtained fractions and colour of supernatant fractions
A mass balance was built up based on protein content in the residue, pellet and supernatant
fractions (Fig. 1). The amount of protein in the fractions was calculated based on protein
content determined by Dumas, in combination with weight of the fractions (dry matter based)
The protein recoveries ranged from 86.5 % to 103 % (Fig. 1). The losses did occur during the
extraction procedure, especially for B. dubia. The pellet contained 32.6 % to 39.4 % of total
protein and the residue 31.4 % to 46.6 % of total protein (Fig. 1). The obtained pellet and
residue fractions were higher in protein content than that in the supernatant (17 % to 23.1 %)
for all five types of insects. The amount of proteins in the residue was higher than that in the
pellet, except for Z. morio (31.4 %).
In addition, the protein content on dry matter basis of each fraction ranged from 50 % to
61 % in the supernatant, from 65 % to 75 % in the pellet, from 58 % to 69 % in the residue
and around 0.1 % in the fat fraction. All chitin-bound nitrogen is expected to be present only

284	in the pellet and residue fractions, because chitin is insoluble in aqueous solvents (Goycoolea,
285	Argüelles-Monal, Peniche, Higuera-Ciapara, Doxastakis, & Kiosseoglou, 2000). Except for
286	the presence of chitin-bound nitrogen, there is also uncertainty in the protein-to-nitrogen
287	conversion factor of 6.25 leading to inaccuracy in the absolute protein content reported.
288	After aqueous extraction, the B. dubia had the lightest (light yellow), and the T. molitor the
289	darkest, colour (dark brown) among all insect supernatant solutions. The colour of A.
290	diaperinus, Z. morio and A. domesticus supernatant solutions was comparable. This visual
291	observation indicated that chemical reactions took place during processing. Preliminary
292	experiments showed that colour formation was most likely due to enzymatic browning
293	reactions. In addition, the colour of residue and pellet fractions was similar to that of the
294	supernatant fractions.
295	3.4. SDS-PAGE
296	The reduced SDS-PAGE using 12.5 % acrylamide gels results show a range of protein bands
297	of the supernatant fractions $<$ 95 kDa, and that of the pellet fractions $<$ 200 kDa for all five
298	insect species (Fig. 2). Five major groups of protein bands could be distinguished in Fig. 2,
299	namely bands \leq 14 kDa, 14 - 32 kDa, 32 - 95 kDa and $>$ 95 kDa. Due to insolubility in
300	sample buffer, protein bands of the residue fractions were absent on the gels used in this
301	experiment.
302	Based on intensity, the bands \leq 14 kDa were abundant, especially for <i>T. molitor</i> . SDS-PAGE
303	analysis using 20 % acrylamide gels showed that the band \leq 14 kDa consisted of a range of
304	protein bands from 6.5 kDa to 14 kDa for all insect species studied (results not shown). For <i>T</i> .
305	molitor, the bands ≤ 14 kDa could possibly originate from anti-freeze type of proteins
306	ranging from 8.5 - 13 kDa, including hemolymph proteins having a molecular weight ~12
307	kDa (Graham, Liou, Walker, & Davies, 1997); (Liou, Thibault, Walker, Davies, & Graham,

308	1999); (Graham, Tang, Baust, Liou, Reid, & Davies, 2001). For the other insect species
309	studied, no literature is available for specific proteins, not for those \leq 14 kDa but also not for
310	those > 14 kDa.
311	Next, the bands observed ranging from 14 to 32 kDa could possibly originate from <i>T. molitor</i>
312	cuticle proteins with molecular weights predominantly between 14 and 30 kDa (Andersen,
313	Rafn, Krogh, Hojrup, & Roepstorff, 1995), e.g. chymotrypsin-like proteinase (24 kDa)
314	(Elpidina, Tsybina, Dunaevsky, Belozersky, Zhuzhikov, & Oppert, 2005),.
315	The bands observed ranging from 32 to 95 kDa in the <i>T. molitor</i> supernatant fractions could
316	possibly be linked to enzymes and other proteins, e.g. melanization-inhibiting protein (43
317	kDa), β-glycosidase (59 kDa), trypsin-like proteinases (59 kDa), and melanization-engaging
318	types of protein (85 kDa) (Ferreira, Marana, Terra, & Ferreira, 2001); (Zhao, Soderhall, Park,
319	Ma, Osaki, Ha, et al., 2005); (Prabhakar, Chen, Elpidina, Vinokurov, Smith, Marshall, et al.,
320	2007); and (Cho, Choi, Moon, Kim, Kwon, Homma, et al., 1999).
321	Above 95 kDa, no bands were observed in the supernatant fractions of <i>T. molitor</i> . Compared
322	to T. molitor, the pattern of protein bands from supernatant fractions in A. diaperinus and A.
323	domesticus were similar, but not identical. For Z. morio and B. dubia, more bands were found
324	in the range of 30 to 95 kDa.
325	The observed bands with molecular weight > 95 kDa in the pellet fractions of T . molitor
326	possibly originate from vitellogenin-like protein with a molecular weight of 160 kDa (Lee,
327	Lee, Choi, Cho, Kwon, Kawabata, et al., 2000). No subunit structures of the proteins
328	mentioned were found using UniProt: Universal Protein Resource Knowledgebase (UniProt
329	ID: Q9H0H5), so that actual molecular weight reported in literature is similar to apparent
330	molecular weight on gel.

331	Besides the proteins mentioned before, proteins incorporated in the exoskeleton and muscle
332	proteins are present in the five types of insects and in the fractions obtained. For the adult
333	stage of A. domesticus and B. dubia muscle proteins include insect flight and leg muscles,
334	which mainly consist of large size proteins, e.g. M-line protein, (flight and leg muscle, 400
335	kDa); kettin (leg muscle isoform, 500 kDa); kettin (flight muscle isoform, 700 kDa) (Bullard
336	& Leonard, 1996); (Lakey, Ferguson, Labeit, Reedy, Larkins, Butcher, et al., 1990). For the
337	larval stage of T. molitor, A. diaperinus and Z. morio skeletal muscles, which likely consist of
338	large size proteins, are present.
339	3.5. Protein functionality measurements
340	Due to the insolubility of the pellet and residue fractions, only the supernatant fraction of the
341	protein was tested for its functionality with respect to foamability, foam stability, and
342	gelation.
343	3.5.1. Foamability and foam stability
344	As a reference for the foam stability measurements, albumin from chicken egg white was
345	used at a concentration of 1.5 % w/v. The reference sample is a good stabilizer for foam, and
346	was capable of producing foam with a half-time of 17 minutes. Zophobas morio formed foam
347	at pH 3, 7 and 10 with a half-time of 6 minutes, A. domesticus at pH 3 with a half-time of 4
348	minutes, and B. dubia produced foam at pH 5 with a half-time of 5 minutes. Foams with half-
349	time of < 6 minutes are not considered to be stable foams. All other supernatant fractions had
350	negligible foam ability at a concentration of 3 % w/v, at pH 3, 5, 7, and 10. This may be due

to the protein concentration in the supernatant fraction solution (around 1.7 % w/v) being too

low to generate stable foam. The stability of the foam can be influenced by protein structure,

protein concentration, and ionic strength. In addition, the stability of the foam can be also

influenced by presence of oil. As mentioned by (Lomakina & Mikova, 2006), the effect of oil

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355	at levels above $0.5\ \%$ reduced the volume of egg white foam. In our case, the supernatant
356	fractions obtained from five insect species also contained some amount of oil in
357	concentration of around 0.1 %, which may also influence foamability of proteins in
358	supernatant fractions.
359	3.5.2. Gelation
360	3.5.2.1 Visual observation of gelation
361	The visual appearance was determined of gels of five supernatant fraction solutions, with
362	fraction concentrations of 3 and 30 % w/v , at pH 3, 5, 7, and 10, after heating for 10 minutes
363	in a water bath at 86 \pm 1 ^{o}C (Table 3). A heating time of 20 and 30 minutes was also tested,
364	but no differences were seen in gel formation (not shown). Factors affecting the gel
365	properties in general are pH, protein concentration, and thermal treatment. The protein
366	concentrations selected for gelation are in the range from 0.5 to 25 % concentration that are
367	used in general to make gels. At a concentration of 3 % w/v, none of the protein fractions
368	showed gel formation, except for A. domesticus at pH 7. At pH 5 and pH 7, for all samples
369	(except A. domesticus at pH 7) heating induced the formation of visible large aggregates
370	rather than gel formation.
371	All 30 % w/v supernatant fractions formed a gel at pH 7 and 10, but not at pH 3. At pH 5,
372	very weak gels were formed, that yielded when turned upside down. In table 3, these samples
373	are designated as "V" (viscous fluid). All samples at pH 7 and 10 were turbid, indicating that
374	the characteristic size of the structures forming the gel was larger than the wavelength of
375	visible light. All gels were already formed after 10 minutes and longer heating times had no
376	influence on the appearance of the gel.
377	Some insect proteins have an isoelectric point of about 5. For instance, the pI of proteins from
378	silkworm (Bombyx mori) and spider (Nephila edulis) are 4.37 - 5.05, and 6.47, respectively

379	(Foo, Bini, Hensman, Knight, Lewis, & Kaplan, 2006). If our protein fractions also have a pI
380	of around pH 5, this may explain why all fractions at this pH formed aggregates at a
381	concentration of 3 % w/v, and very weak gels at concentrations of 30 % w/v. Close to the pI,
382	the electrostatic interactions between the proteins are very weak, which, upon denaturation,
383	tends to lead to the formation of dense aggregates. These dense aggregates have a much
384	higher gelling concentration than aggregates formed at a pH above or below the isoelectric
385	point. To form a firm gel at this pH, higher protein concentrations are needed.
386	Samples at pH 3 and 10 at 3 % w/v were more transparent than samples heated at pH 5 and 7.
387	The increased charge on the protein at pH 3 may prevent the proteins from aggregating, since
388	even at 30 % w/v these fractions did not form a gel or even a viscous fluid. The decrease in
389	turbidity observed at pH 10 suggests that the aggregates formed at this pH were less dense
390	and/or smaller than the ones formed at pH 5 and 7.
391	3.5.2.2. Rheological properties of gels
392	According to the visual observation of gelation, at a pH of 7 and a concentration of 3 % w/v a
393	weak gel was formed, and at 30 % w/v a strong gel was formed. Therefore, for studying gel
394	strength, fraction concentrations in between these two values (7.5 and 15 % w/v) were
395	chosen. For all five fractions, we determined the evolution of the storage modulus G' and loss
396	modulus G" during the temperature ramp at a concentration of 15 % w/v and a pH of 7. The
397	storage modulus is a measure for the elastic energy stored reversibly in a gel during
398	deformation, and characterizes its stiffness; the loss modulus is a measure for the energy
399	dissipated during deformation as a result of viscous friction. As an example, the results for
400	the mealworm supernatant fraction (the other fractions showed similar results) are provided
401	(Fig. 3A). G' gradually increased during the heating phase of the ramp. During the second
402	phase, when the temperature was kept constant at 90 °C, G' kept on increasing gradually.
403	This observation showed that the gel structure did not yet reach an equilibrium state. During

404	the cooling phase, both G' and G" increased sharply. This is typical for gels in which
405	hydrogen bonds are formed between structural elements (Ould Eleya, Ko, & Gunasekaran,
406	$\underline{2004}$). The gelation temperature observed ranged from about 51 °C to 63 °C (<i>T. molitor</i> 61.7
407	\pm 1.1°C , A. diaperinus 58.2 \pm 2.1 °C, Z. morio 51.2 \pm 1.5 °C, A. domesticus 56.2 \pm 0.7 °C, B.
408	dubia 63.2 \pm 0 °C, from which the lowest and the highest temperature were from Z. morio
409	and <i>B. dubia</i> supernatant fractions respectively (results not shown).
410	To obtain more information on the gel structure, the value of log G' of T. molitor supernatants
411	was determined as a function of log C (concentration) with fraction concentrations of 7.5 %
412	w/v, 15 % w/v and 30 % w/v (corresponding to actual protein concentrations of 4.1 %, 8.3 %
413	and 16.6 %) at 90 °C and 20 °C (Figure 3B). Values for G' at 90 °C were taken from end of
414	phase 2 from the ramp, and values at 20 °C were taken from the end of phase 3, which is
415	similar to the procedure of (Ould Eleya, Ko, & Gunasekaran, 2004). The values of the power-
416	law exponent n in the scaling relation $G^{\prime} \varpropto C^n$, were used for evaluation of gel structure
417	(Shih, Shih, Kim, Liu, & Aksay, 1990). The parameter n had a value equal to 3.0 ± 0.4 at the
418	end of the isothermal stage at 90 °C, and a value of at 2.8 ± 0.6 from the end of the cooling
419	stage at 20 °C. These two values are comparable, so there were no significant structural
420	rearrangements in the gel network upon cooling of the samples. An exponent n of about 2.8 is
421	typical for fractal protein gels and points to a fractal dimension d_f which is close to 2 (\underline{Ould}
422	Eleya, Ko, & Gunasekaran, 2004).
423	Fig. 3C shows G' at the end of phase 3 of the temperature ramp as a function of strain, for
424	insect supernatant gels at 20 °C and a concentration of 15 % w/v. The value for G' in the
425	linear response region of A. domesticus supernatant gels was around 2500 Pa, which was
426	almost 1.5 times stronger than that of B. dubia (around 1600 Pa), 6 times stronger than that of
427	Z. morio (around 390 Pa), and 25 times stronger than that of T. molitor (around 100 Pa) and
428	A. diaperinus (around 140 Pa). In interpreting these results, we must be careful, since the

429	actual protein concentrations in the fractions was lower than 15 % w/v, and differed slightly
430	from fraction to fraction. As seen before, the actual protein contents were for T. molitor 8.3
431	%; A. diaperinus 9.2 %; Z. morio 7.6 %; A. domesticus 9.2 % and B. dubia 7.4 %.
432	Several conclusions can be drawn from these results. Although the B. dubia supernatant
433	sample had the lowest actual protein content, it formed the strongest gels among all other
434	three insect species, except A. domesticus. Supernatants from A. diaperinus and A.
435	domesticus had similar protein concentration, but they showed significant differences in gel
436	strength. In addition, supernatants from B. dubia and A. domesticus that were in the adult
437	stage formed relatively stronger gels than the other three insect species that were in the larvae
438	stage. Apparently, the insect growth stage influences the body protein composition, and
439	different species differ in protein type and structure (Wilson, 2010).
440	All insect gels had a comparable maximum linear strain at supernatant fraction concentration
441	of 15 % w/v, with a value of around 50 %. An example is shown for Z. Morio (Fig. 3C). The
442	maximum linear strain is, of course, dependent on heating rate and protein concentration, and
443	it would therefore be interesting to investigate the concentration dependence of this property,
444	since it can provide additional information on the fractal dimension of the gels.
445	These detailed rheological results show that insect proteins can form gels that have similar
446	properties as those formed from conventional food proteins. It therefore shows that insect
447	proteins have indeed functionalities that are desirable for food application.
448	4. Conclusions
449	Proteins were extracted from five insect species and protein purity and yield of the obtained
450	fractions was evaluated: Around 20 % of total protein was found back in the supernatant, the
451	rest of the protein was divided about equally over the residue and the pellet fraction for all
452	five insect species after aqueous extraction. The extraction method is easy and feasible to

apply, but the yield of extracted supernatant fractions is relatively low. The purity of
measured protein content expressed as percentage of dry matter ranged from 50 % to 61 % of
supernatant fractions, from 65 % to 75 % of pellet fractions and from 58 % to 69 % of residue
fractions depending on the insect species.
We established some functional properties of the protein fractions, focusing on foaming and
gelation: The soluble protein fractions of all five types of insects had poor foaming capacity
at pH 3, 5, 7, and 10, but could form gels at a concentration of 30 % w/v. At a concentration
of 15 % w/v at pH 7 and 10, A. domesticus supernatant formed the strongest gels among all
insect species. The gelation temperature ranged from about 51 °C to 63 °C for all insect
species at pH 7. In addition, all insect gels had a comparable maximum linear strain at this
concentration, with a value of around 50 %.
We studied the protein quality of whole insects by analysis of protein content and amino acid
composition. The protein content of the five insect species was comparable to conventional
meat products in terms of protein quantification. The sum of EAA per g protein for all insect
species was comparable with the sum of EAA for soybean protein, lower than that for casein,
but higher than that for the daily protein requirement of an adult (<u>FAO/WHO/UNU</u> , 1985).
Differences in calculated EAAI were similar.
Although differences are observed in protein content, amino acid composition, protein
distribution of the fractions obtained, SDS-PAGE data, foaming and gelation properties, the
similarities between the insect species are more apparent than the differences. The fact that
gels could be formed for all five insect species, using the soluble fractions obtained by a
simple aqueous extraction procedure, is promising in terms of future food applications. More
research is needed for developing further extraction and purification procedures, and for more
detailed insight into functional properties.

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480 References

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Table 1. Proximate composition of five insect species on live weight basis (mean \pm S.D., n=2).

Table 2. Amino acid pattern of five insect species, casein, soybean protein, recommendation for adult and calculated essential amino acid index of five insect species and casein & soybean protein (<u>FAO/WHO/UNU</u>, 1985) and (<u>Young & Pellett</u>, 1991).

Table 3. Gel formation of supernatant fractions from five insect species (X: no gel formation; A: aggregation; V: viscous fluid; O: gel formation).

Table 1.

Insects	Moisture (%)	Fat (%)	Crude protein (%)	Other components (%)
			(including chitin	(e.g. carbohydrates,
			nitrogen)	minerals and vitamins)
T. molitor	63.5±1.8	9.9±1.0	19.1±1.3	7.5±2.2
A. diaperinus	64.5±1.0	8.5±0.2	20.6±0.1	6.4±1.0
Z. morio	59.9±5.4	16.0±0.7	20.7±0.3	3.4±5.5
A. domesticus	70.8±2.0	3.6±0.4	21.5±0.5	4.1±2.1
B. dubia	67.4±2.1	7.7±0.1	19.3±0.9	5.6±2.3

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Table 1. Proximate composition of five insect species on live weight basis (mean \pm S.D., n=2).

Table 2.

unit (mg/g	A.diaperinus	T.molitor	Z.morio	A.domesticus	B. dubia	Casein	Soybean	1985	
crude								FAO/WHO/U	
protein)								NU	
Essential amin	0								
acid (EAA)									
Histidine	34	29	31	21	23	32	25	15	
Isoleucine	43	43	46	36	31	54	47	30	
Leucine	66	73	71	66	56	95	85	59	
Lysine	61	54	54	53	43	85	63	45	
Methionine	26	26	24	25	23	35	24	22	
+Cysteine									
Phenyl-alanine	e 120	100	111	92	93	111	97	38	
+ tyrosine									
Threonine	39	39	40	35	32	42	38	23	
Tryptophan	12	12	14	9	8	14	11	6	
Valine	58	61	63	55	52	63	49	39	
Sum of EAA	459	437	454	392	361	531	439	277	
Non-essential									
amino acid									
Alanine	66	70	68	81	71				
Arginine	54	54	54	65	46				

Glutamic acid					67			
	123	109	127	110	96			
Glycine	46	50	48	51	53			
Proline	56	66	56	54	48			
Serine	40	44	42	38	34			
Sum of total AA	927	910	931	864	776			
EAAI	1.65	1.60	1.66	1.39	1.28	1.93	1.56	
				NA				

²Table 2. Amino acid pattern of five insect species, casein, soybean protein, recommendation for adult and calculated essential amino acid index of five insect species and casein & soybean protein (FAO/WHO/UNU, 1985) and (Young & Pellett, 1991).

Table 3.

	pH 3	pH 5	pH 7	pH 10
3 %				
T.molitor supernatant	X	A	A	X
A.diaperinus supernatant	X	A	A	X
Z.morio supernatant	X	A	A	X
A.domesticus supernatant	X	A	O	X
B.dubia supernatant	X	A	A	X
30 %				
T.molitor supernatant	X	V	O	0
A.diaperinus supernatant	X	V	O	O
Z.morio supernatant	X	v	0	O
A.domesticus supernatant	X	V	0	0
B.dubia supernatant	X	V	O	O

³ Table 3. Gel formation of supernatant fractions from five insect species (X: no gel formation; A: aggregation; V: viscous fluid; O: gel formation).

Figure Captions and Tables

- Fig. 1. Protein content of supernatant, pellet and residue fractions expressed as percentage of total protein and total recovery (n=2).
- Fig. 2. Molecular weight distribution of *T.molitor* protein fractions, determined by SDS-PAGE using 12.5% homogeneous phastgel and (Samples from left to right: supernatant, pellet and marker); marker is ranging from 6.5 kDa to 200 kDa. Mw is molecular weight.
- Fig. 3. A: Dynamic moduli G' and G" of T. molitor supernatant solution as a function of time. Heating and cooling phases are plotted as a secondary axis. B: Plots of the storage modulus G' as a function of protein concentration of mealworm supernatant fractions on a logarithmic scale at pH 7 (heating period 90 °C and cooling period 20 °C). C: Storage modulus G' (Pa) as a function of strain γ % for insect supernatant gelation at 20 °C at a supernatant fraction concentration of 15 % w/v.

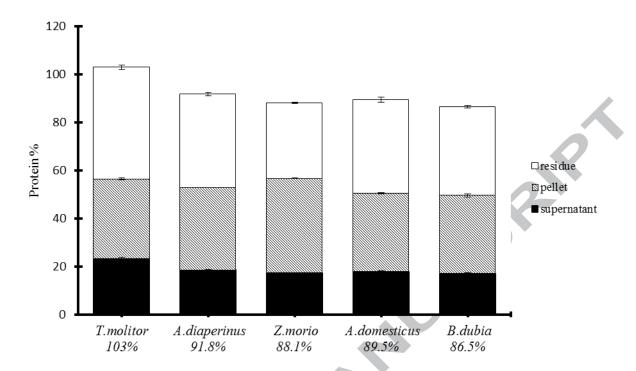


Fig. 1.

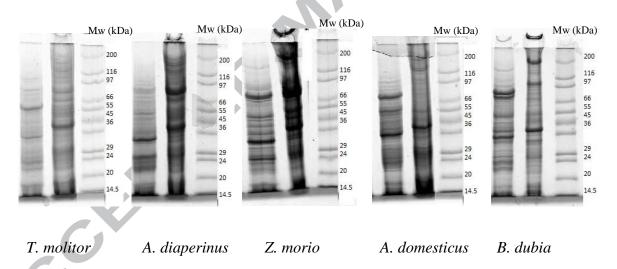
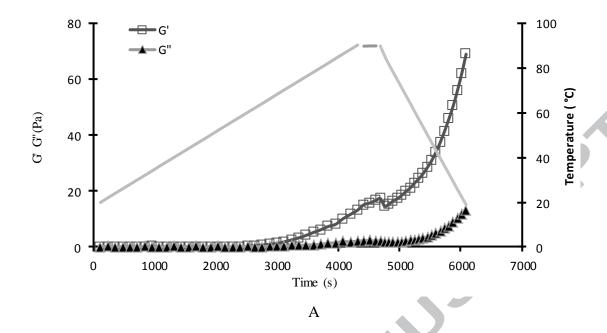
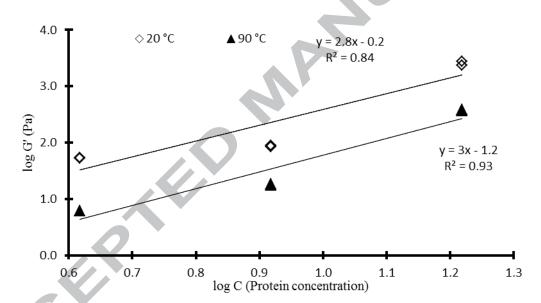
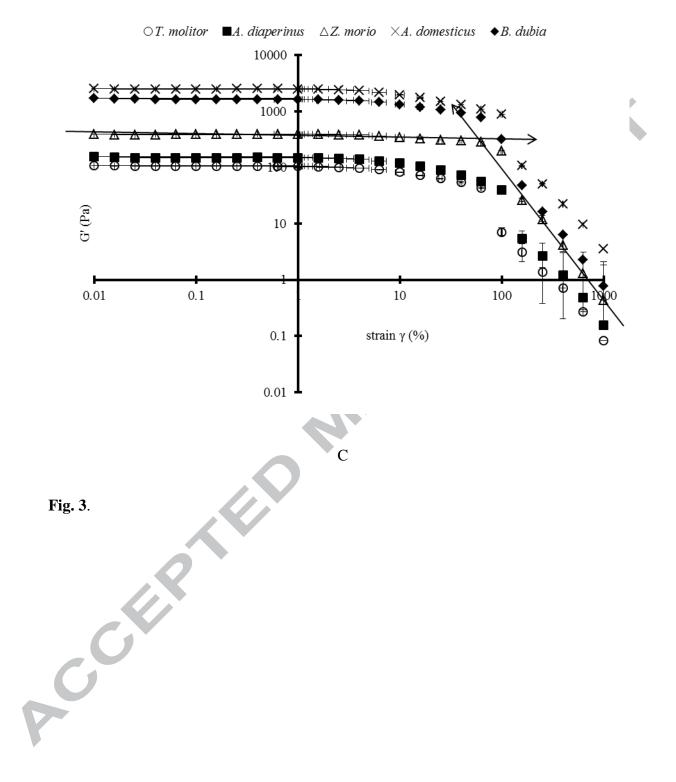


Fig. 2.







Highlights

Crude protein content of insects was similar to conventional meat products.

The amount of EAA of insects was higher than daily protein requirement of an adult.

The supernatant, pellet, fat and residue fractions were obtained after an aqueous extraction.

Protein bands were < 95 kDa for supernatant fractions and < 200 kDa for pellet fractions.

Most supernatant fractions did not foam, but could form gels depending on protein concentration and the pH.