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Article

# Operational Key for Microscopical Identification of Fragments from EU-Authorized Insect Species in Feed

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## Abstract

The microscopic identification of insect-derived processed animal proteins (PAPs) is essential for feed control within the European Union, where light microscopy remains the official method for detecting prohibited animal proteins. This study synthesizes current morphological knowledge for the insect species authorized as feed materials *Hermetia illucens*, *Musca domestica*, *Tenebrio molitor*, *Alphitobius diaperinus*, *Acheta domesticus*, *Gryllobates sigillatus*, *Gryllus assimilis* and *Bombyx mori* and provides a comprehensive framework for their recognition in processed feed. Diagnostic traits were compiled from reference laboratory material, the scientific literature and the EURL-AP micrograph collection, with emphasis on cuticular structures, setae, denticles, spiracles, tracheal elements, and species-specific features such as spine-like sensilla in *H. illucens*, gin-trap structures in *T. molitor*, and distinctive antennal or cercal fragments in Orthoptera. Dipteran PAPs are characterized by unsclerotized cuticle, denticle bands, and spiracular morphology, whereas Coleopteran fragments display stronger sclerotization, larger mouthparts, and urogomphi. Orthopteran meals exhibit the greatest structural diversity due to the use of nymphs and adults, yielding leg, wing, antennal, and cercal fragments. Across taxa, tracheal and muscle-fibre structures provide reliable confirmation of insect origin but lack species specificity. The study highlights persistent gaps in reference material, particularly for *Musca domestica*, and underscores the need for enhanced taxonomic training and the integration of automated image-recognition tools and complementary molecular methods for species-level confirmation. The identification key and morphological guidance presented here aim to support routine laboratory diagnostics and strengthen regulatory compliance in feed monitoring.:

**Keywords:** EU; feed; insects; microscopical identification

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## Introduction

Bovine spongiform encephalopathy (BSE), a prion disease transmitted through feed, is controlled in the European Union (EU) through banning of the use of processed animal proteins (PAPs) in ruminant diets. To strengthen feed safety, the EU implemented the General Food Law (Regulation (EC) No 178/2002) and the Hygiene Package (Regulations (EC) No 852/2004 and 183/2005). Within this regulatory framework, insect producers are responsible for ensuring product safety and full compliance with EU legislation.

Following recommendations from the European Food Safety Authority (EFSA), the use of insects in animal feed was introduced by Regulation (EU) 2017/893. This regulation amended Regulation (EC) No 999/2001 and Regulation (EU) No 142/2011, marking the first official step toward the inclusion of insect-derived PAPs in aquaculture feed. The authorization initially covered seven species: the black soldier fly (*Hermetia illucens*), the common housefly (*Musca domestica*), the yellow mealworm (*Tenebrio molitor*), the lesser mealworm (*Alphitobius diaperinus*), the house cricket (*Acheta domesticus*), the banded cricket (*Gryllobates sigillatus*), and the field cricket (*Gryllus assimilis*). In 2021, Commission Regulation (EU) 2021/1925 expanded the list of authorized species to include the silkworm (*Bombyx mori*). Later that year, Commission Regulation (EU) 2021/1372 partially lifted the

feed ban by permitting the use of insect PAPs, poultry PAPs, and pig PAPs in poultry and pig feed, while maintaining the prohibition on intra-species recycling.

Currently, two official analytical methods are recognized in the EU for detecting animal proteins in feed: light microscopy and real-time polymerase chain reaction (PCR) (European Commission 2009, 2013, 2020, 2022). Light microscopy, routinely applied since 2004, enables the detection of morphological markers of animal-derived PAPs, including insect material, after an extraction step, with a detection limit of approximately 0.1%. Its effectiveness is linked to its simplicity, the modest level of expertise required, and the absence of specialized equipment. When sufficient fragments are present, light microscopy can even support identification at higher taxonomic levels, such as order and, in some cases, family (Veys and Baeten, 2018).

The first work describing key diagnostic features of insect fragments under light microscopy was conducted by Ottoboni et al. (2017), focusing on *H. illucens*, *B. mori*, and *T. molitor*. Veys and Baeten (2018) expanded this knowledge by characterizing morphological traits of four legally authorized insect PAPs from three orders: Diptera (*H. illucens*), Coleoptera (*T. molitor* and *A. diaperinus*), and Orthoptera (*G. assimilis*). Weiner and Kwiatek (2022) further examined samples containing PAPs from *H. illucens* and *T. molitor*. More recently, Marien et al. (2024) proposed a specific real-time PCR method for the detection of *B. mori* and performed light microscopy observations on industrial feed samples containing *B. mori*, and pure insect meal samples of *H. illucens* and *T. molitor*.

The identification of fly larvae is extensively documented, mainly due to their medico-legal relevance rather than feed-science applications (Grzywacz et al., 2017; Barros et al., 2019; Walczak and Grzywacz, 2024). Grzywacz et al. (2017) developed species-level identification keys for Muscidae. Sultana et al. (2021) provided detailed descriptions, taxonomic keys, and illustrations for 17 Gryllidae species of Pakistan, bringing existing information up to date. Their work highlighted diagnostic differences between morphologically similar species and presented a taxonomic key for the species occurring in Sindh, including *A. domesticus*, *Acheta hispanicus*, *Gryllus (Gryllus) bimaculatus*, *Gryllus (Gryllus) campestris*, *Gryllus septentrionalis*, *G. sigillatus*, and *Gryllodes supplicans*. However, such tools are still lacking in the context of feed authentication and control. Their development would clearly support the accurate identification of PAPs from authorized species and help distinguish them from non-authorized ones, as previously emphasized by Veys and Baeten (2018).

In this work, we further investigated the morphological features of fragments from pure insect samples and, based on the available bibliography, provide a first draft of an identification key for the authorized species.

## Taxonomic Overview of Authorized Insects in Feed in EU

Currently, eight insect species belonging to four different taxonomic orders are authorized for use in feed within the European Union: Diptera (flies), Coleoptera (beetles), Orthoptera (crickets), and Lepidoptera (moths) (Table 1).

**Table 1.** The specific species and their respective taxonomic orders and families are as follows:

Common Name	Scientific Name	Order	Family
Black soldier fly	<i>Hermetia illucens</i>	Diptera	Stratiomyidae
Common housefly	<i>Musca domestica</i>	Diptera	Muscidae
Yellow mealworm	<i>Tenebrio molitor</i>	Coleoptera	Tenebrionidae
Lesser mealworm	<i>Alphitobius diaperinus</i>	Coleoptera	Tenebrionidae
House cricket	<i>Acheta domesticus</i>	Orthoptera	Gryllidae
Banded cricket	<i>Gryllodes sigillatus</i>	Orthoptera	Gryllidae
Field cricket	<i>Gryllus assimilis</i>	Orthoptera	Gryllidae
Silkworm	<i>Bombyx mori</i>	Lepidoptera	Bombycidae

## Diagnostic Morphological Features of Insects Authorized for Use in Feed

The most reliable features used by laboratories to identify insect material in processed feed are chitinous structures that withstand rendering and heat treatment. These include the cuticle, setae, tracheal elements, muscle fibres, spiracles, and mouthparts. Among these, cuticle structure is often the most informative diagnostic trait. All species share the presence of a tegument, although its degree of sclerotization varies markedly; from the soft cuticle of Diptera larvae to the more rigid and highly chitinized exoskeleton characteristic of the larval, pupal, nymphal, and adult stages of Coleoptera, Orthoptera, and Lepidoptera. Setae are present in all taxa, but their abundance, morphology, and function differ by species and developmental stage. Spine-like sensilla are a distinctive morphological feature of *H. illucens* larvae and are not consistently reported in the other authorized insect species used in feed; their presence therefore represents a species-specific trait characteristic of *H. illucens*. All insects considered in this study, across all active life stages (larvae, nymphs, pupae, adults), possess a tracheal system for respiration, with external openings (spiracles) that remain identifiable after processing. All developmental stages; larvae, pupae, nymphs, and adults, breathe through spiracles connected to the tracheal system, and mandibles are present in all these stages, making them common yet still useful diagnostic elements. Tracheoles appear as small, transparent, spiral-shaped tubes, and muscle fibres are universal identifiable features shared by all species, indicating only the presence of insect material but not allowing species-level discrimination. Muscle fragments typically appear as yellow or transparent rectangular pieces, sometimes showing visible zig-zag striation, which confirms the presence of animal tissue. The origin of muscle fibres from insects can be confirmed by the observation of associated tracheal elements.

The primary morphological distinctions among species include the presence or absence of legs, wings, and visible head structures such as antennae. Fly larvae, such as *H. illucens* and *M. domestica*, are typical maggots, lacking legs and possessing an indistinct head capsule or mouth hooks. While entirely absent in fly larvae, legs are present in all other life stages of the remaining species. In contrast, larvae of *T. molitor* and *A. diaperinus* have well-defined mouthparts (mandibles) and abdominal appendages. *A. diaperinus* is generally darker and more strongly sclerotized than *T. molitor*.

Crickets possess strongly developed hind legs for jumping in both nymph and adult stages. Sultana et al. (2021) described the legs of *A. domesticus* as yellowish with a few brown spots and numerous hairs, and noted that the posterior tibia bears eleven spines on its basal side; the cerci are well developed and pointed. In Gryllus species, the legs are blackish with brown spots, the posterior femora are relatively short and thick, and the posterior tibia is armed with six spines on each margin. *G. sigillatus* is characterized by reduced wings.

The following key is based on micrographs from the internal laboratory reference collection, the EURL-AP micrograph archive, and published literature (Table S1; Figures 1–7). Larvae of *M. domestica* were not available in our collection, and although fragment descriptions in processed feed exist for most authorized insect species, information for *M. domestica* larvae remains limited.

## Identification Key for EU—Authorized Insect PAPs

### 1. Initial separation based on presence of adult/nymph structures

1A. No wings, no compound eyes, no segmented antenna fragments, no cerci → Larval/Pupal pathway → go to 2

1B. Wings, compound eyes, segmented antenna fragments, cerci, articulated leg fragments present → Adult/Nymph pathway (Orthoptera) → go to 6

### 2. Larval / Pupal Pathway

2A. Absence of legs or claws

→ Diptera larvae → go to 3

2B. Thoracic legs, pseudolegs, mandibles, claws, urogomphi, or gin-traps present

→ Coleoptera or Lepidoptera larvae/pupae → go to 4

### 3. Diptera Larvae

3A. *Hermetia illucens* (Black soldier fly)

- Honeycomb-like reticulated cuticle (4–6-sided cells, thick walls, broad lumen)
- Dense small setae; long yellow-brown bristles
- **Spine-like sensilla present**
- Linear, unidirectional denticles (spinose bands)
- Dorsal posterior spiracles with multiple radial openings
- Anal opening bordered by short, thick spine-like setae
- Colour: grey-cream → dark brown

→ **Diagnostic conclusion: Honeycomb cuticle + spine-like sensilla + linear denticles = *H. illucens***

(Figure 1)

### 3B. *Musca domestica* (Housefly)

- Smooth cuticle
- Fine, sparse setae
- No spine-like sensilla, scale-like spine
- Posterior spiracles kidney-shaped with complete peritreme enclosing three M-shaped sinuous slits
- Pale, soft fragments

→ **Diagnostic conclusion: Smooth cuticle + M-shaped spiracles = *M. domestica***

## 4. Coleoptera and Lepidoptera Larvae / Pupae

### 4A. *Tenebrio molitor* (Yellow mealworm)

- Soft to amber, lightly sclerotized cuticle
- Greyish-yellow → greyish-amber
- Irregular light spots and scattered dark dots (some with very short bristles)
- Gin-trap fragments present (pupal origin)
- Mandibles, thoracic legs recognizable by their claws, urogomphi, anal spine

→ **Diagnostic conclusion: Gin traps + soft amber cuticle with spotted pattern = *T. molitor*** (Figure

3)

### 4B. *Alphitobius diaperinus* (Lesser mealworm)

- Moderately sclerotized, darker cuticle
- Uniform amber–brown pigmentation
- Sparse short setae
- Mandibles, thoracic legs recognizable by their claws, urogomphi, anal spine
- Gin-traps absent

→ **Diagnostic conclusion: Darker, more sclerotized cuticle + absence of gin traps = *A. diaperinus***

(Figure 4)

### 4C. *Bombyx mori* (Silkworm) – pupal fragments

- Yellow to dark brown cuticle
- Cuticular fragments display a pattern that partially resembles that of *H. illucens*
- Smooth surface with few setae
- Pupal leg fragments (leg sheaths)
- Spiracles present

→ **Diagnostic conclusion: Yellow→brown smooth cuticle + pupal leg fragments = *B. mori*** (Figure

2)

## 6. Adult / Nymph Pathway (Orthoptera)

(Identified by articulated legs, claws, antennae, cerci, wing fragments, compound eyes)

### Shared Orthopteran structures

- Pigmented, sclerotized cuticular fragments with setae attached or isolated
- Tibial spines, femoral and tarsal fragments
- Sclerotized mandibles
- Wing fragments (membranous or sclerotized)
- Compound eye fragments
- Cercal fragments
- Segmented antennal flagellomeres

- Claws (often reddish-tipped)
- Muscle tissue, tracheal system fragments, muscle with tracheae

#### 6A. *Acheta domesticus* (House cricket)

- Pale, lightly sclerotized cuticle
- Segmented antennal flagellomeres
- Cerci, Claws
- Leg fragments

Legs yellowish with a few brown spots; posterior tibia armed with eleven spines on the basal side.

→ **Diagnostic conclusion: Pale cuticle + typical cricket appendages = *A. domesticus* (Figure 5)**

#### 6B. *Grylloides sigillatus* (Banded cricket)

- Light cuticle (slightly darker than *A. domesticus*)
- Segmented antennal flagellomeres
- wings reduced → Less Wing fragments
- Cerci, Claws
- Leg fragments,

→ **Diagnostic conclusion: Light cuticle + cricket appendages = *G. sigillatus* (Figure 6)**

#### 6C. *Gryllus assimilis* (Field cricket)

- Darker, more strongly sclerotized cuticle
- Wing fragments
- Robust segmented antennal flagellomeres
- Cerci, Claws
- Leg fragments, mandibles, muscle + tracheae
- In *Gryllus* species, the legs are blackish with brown spots, the posterior femora are relatively short and thick, and the posterior tibia is armed with six spines on each margin.
- Honeycomb-like cuticle may occur

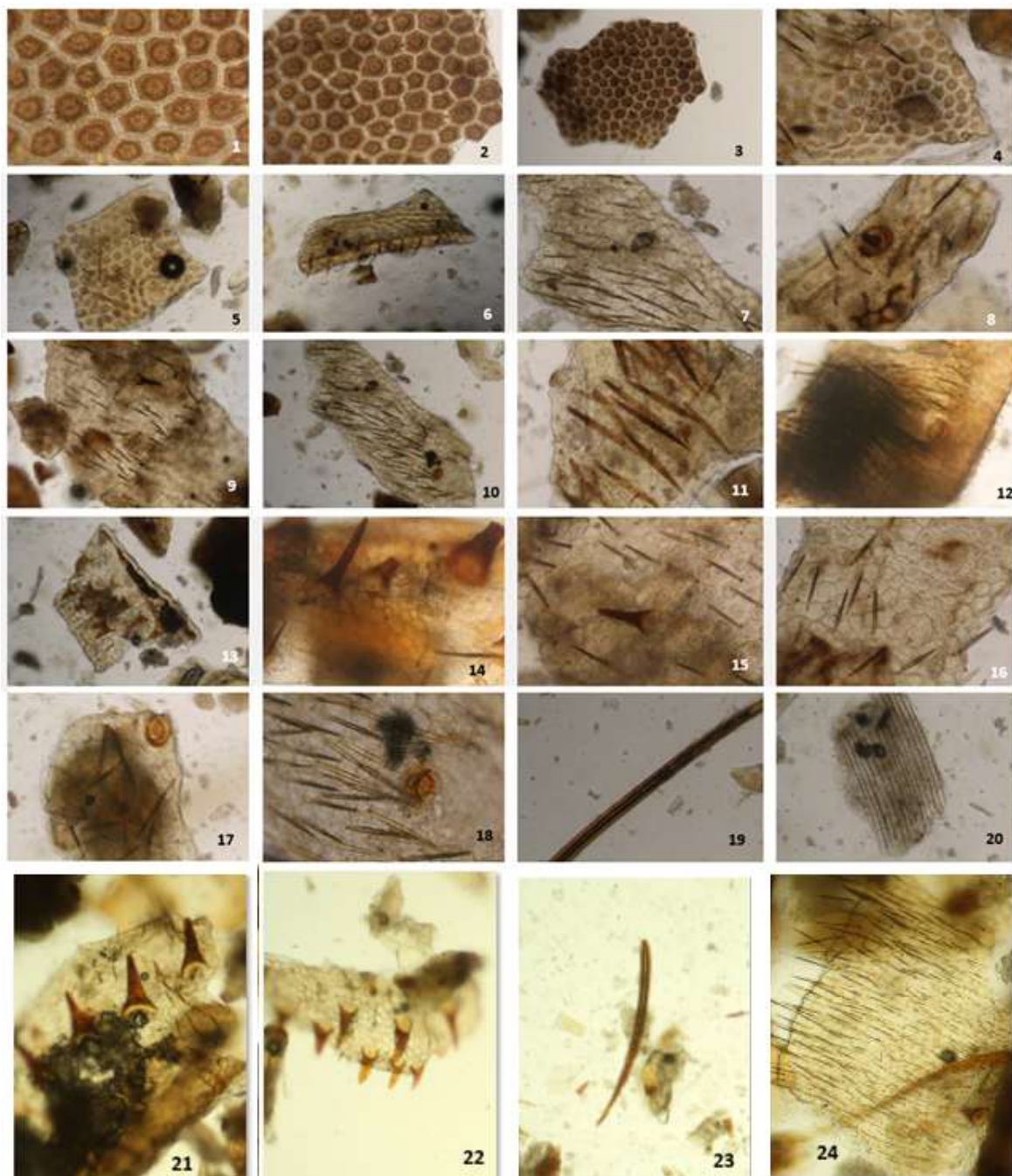
→ **Diagnostic conclusion: Dark cuticle + differentiated appendages + honeycombed fragments = *G. assimilis* (Figure 7).**

Diptera meal consists primarily of larval instars and pre-pupal stages (Veys and Baeten, 2018). The larvae are apodal (Barros et al., 2018). As a consequence of the holometabolous development of Diptera, these stages are only weakly differentiated, and therefore relatively few particles can be confidently identified as being of insect origin (Veys and Baeten, 2018). Nevertheless, the predominance of unsclerotized cuticle fragments, together with the presence of denticles arranged in spinose bands and long setae, provides reliable diagnostic criteria for recognizing Diptera larvae (Veys and Baeten, 2018; Szpila, 2009).

*H. illucens* material ranges in colour from grey-cream to dark brown (Barros et al., 2019; Weiner and Kwiatek, 2022; Marien et al., 2024). The cuticle of *H. illucens* is densely covered with setae (Oliveira et al., 2016; Ottoboni et al., 2017; Veys and Baeten, 2018), which under light microscopy may appear either attached to cuticular fragments or as isolated particles (Veys and Baeten, 2018).

Spine-like sensilla are distinctively present in *H. illucens* larvae and are not consistently reported in other authorized insect species used in feed, making them a species-specific trait of *H. illucens*. Long, yellow to yellow-brown bristles are also frequently observed (Ottoboni et al., 2017; Veys and Baeten, 2018; Weiner and Kwiatek, 2022). Small, linearly organized groups of denticles oriented in the same direction, corresponding to the spinose bands of the larval segments, commonly occur in *H. illucens* PAPs (Veys and Baeten, 2018). Under a light microscope, *H. illucens* can be recognized by its distinctive cuticular structures, which display irregular, cell-like patterns composed of four-, five-, or six-sided units with thick walls surrounding a lighter central lumen, giving the cuticle a characteristic honeycomb-like appearance; in some fragments, a central darker dot may also be visible (Ottoboni et al., 2017; Veys and Baeten, 2018; Weiner and Kwiatek, 2022; Rebora et al., 2023). Weiner and Kwiatek (2022) reported that some dark-brown cuticular fragments of *H. illucens* can resemble rape husks, complicating morphological classification. However, rape husks typically exhibit a darker central region, whereas the corresponding structures in *H. illucens* fragments show a less pronounced central

darkening (Weiner and Kwiatek, 2022). Ottoboni et al. (2017) also reported the presence of pyramidal cuticular structures, although their precise characterization in *H. illucens* remained challenging, and further investigation was recommended to better define these elements.



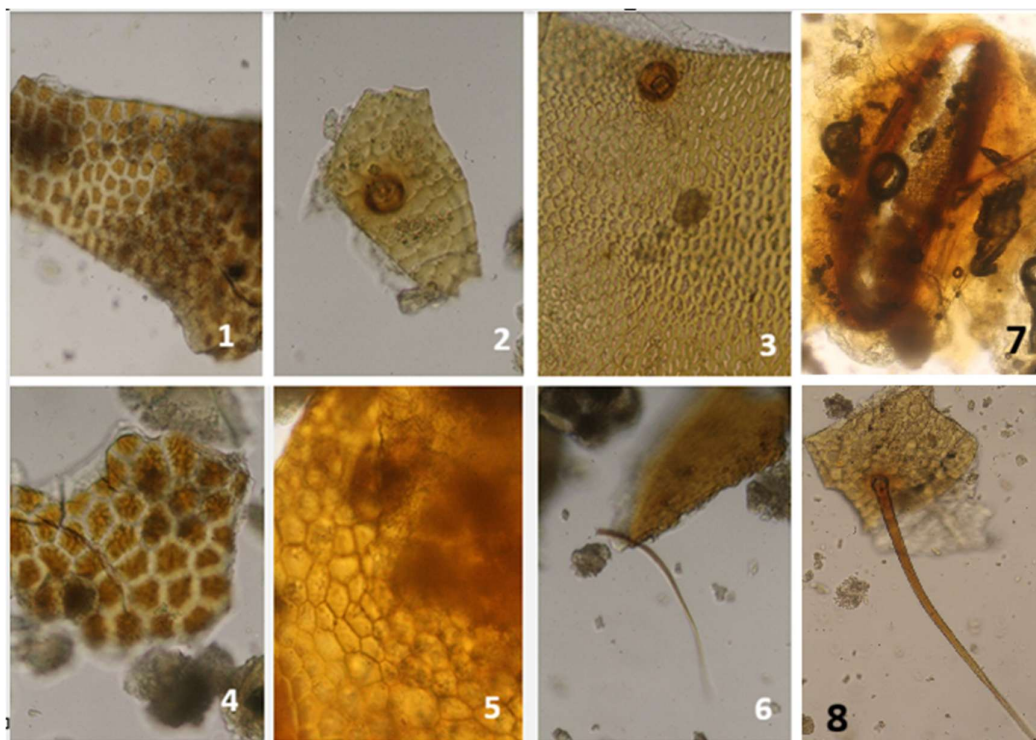
**Figure 1.** *H. illucens* under light microscopy: 1–3 Honeycomb-like cuticle; 4–11 Cuticular fragments with bristles; 12,15,17 Cuticular fragments with bristles and spiracle; 13–15 Spine-like sensilla; 19 Setae; 20 Fragments of the tracheal system; 21–24: Spine-like sensilla, bristles and setae.

The antennae of *H. illucens* are inserted anterolaterally on a moderately prominent ring-shaped socket with a distinct articular membrane (Fabian et al., 2025). Dorsal posterior spiracles, composed of multiple openings arranged radially on the ecdysial surface (Oliveira et al., 2016), may also be detected, as well as the anal opening, whose edges bear short, thick, spine-like setae (Oliveira et al., 2016; Barros et al., 2019; EURL-AP micrograph collection). Posterior spiracles are considered

the most reliable diagnostic structures for identifying fly larvae. Their morphology is highly distinctive, and the number and arrangement of spiracular openings can vary between species and even across developmental stages (Duncan et al., 2010; Raś et al., 2018).

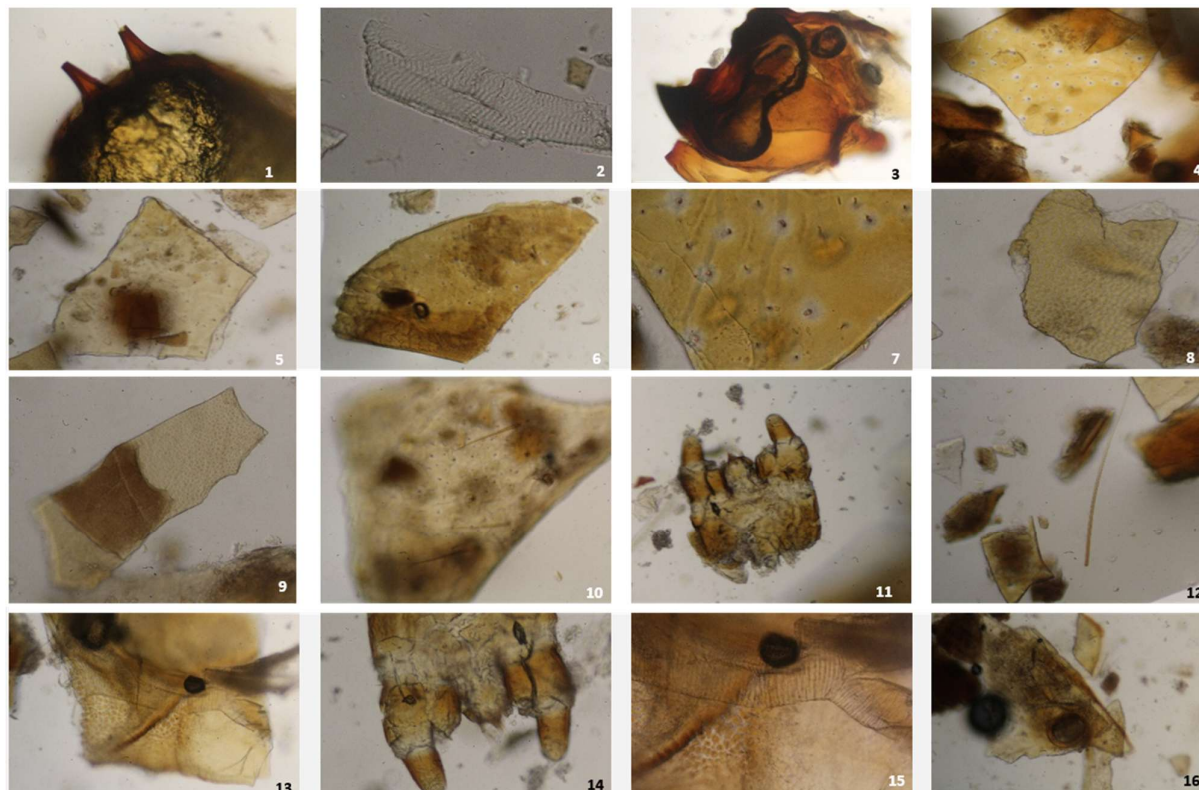
*M. domestica* larvae, characterized by their typical maggot-like form, lack the heavy sclerotization observed in *H. illucens*. For *M. domestica*, the morphology of the posterior spiracle is a key diagnostic feature (Apasrawirote et al., 2022). The spiracles are kidney-shaped, and a complete peritreme enclosing three distinctly sinuous, M-shaped slits is characteristic of this species (Grzywacz et al., 2017; Apasrawirote et al., 2022). Under high magnification, the cuticle may display minute bands of cuticular spines or spinules. Cortinhas et al. (2020) described the cephalic collar spines as composed of numerous scale-like structures, distinguishing them from those of other muscid species. They also noted spines on the anterior portion of the first thoracic segment. After segmentation between the cephalic region and the first thoracic segment, the ventrally concentrated spines become more slender (Cortinhas et al., 2020). These morphological traits may suggest the presence of similar spines in *M. domestica*. Despite these well-defined traits in intact larvae, the current literature provides very limited morphological descriptions of *M. domestica* fragments as they occur in processed feed. This gap is also reflected in the EURL-AP micrograph collection, where such fragmentary remains are not represented. As a result, the identification of house fly fragments remains challenging and is currently insufficiently supported by reference material.

*B. mori*, commercially processed and used at the pupal stage, exhibit cuticular fragments with a pattern that partially resembles that of *H. illucens*, with coloration ranging from yellow to dark brown (Figure 2; Harris, 1979; Ottoboni et al., 2017; Marien et al., 2024). Ottoboni et al. (2017) reported the absence of setae among PAP fragments derived from *B. mori*; however, Marien et al. (2024) documented their presence in this species. Similar structures are also observed in the laboratory reference sample (Figure 2). Kumar et al. (1999) described cremastral setae on the terminal abdominal segment of *B. mori*, which might explain those observations.



**Figure 2.** *B. mori* meal: 1–5: Cuticle fragments (specimens 2 and 3 showing oval spiracles); 6,8: Seta; 7: Spiracle.

Fragments of the exoskeleton of *T. molitor* are described as bright greyish-yellow to deep greyish-amber-brown in colour, often displaying irregular light spots and sporadic black dots (Ottoboni et al., 2017; Weiner and Kwiatek, 2022). Some of these dark dots bear very short, dark bristles (Weiner and Kwiatek, 2022). Rare dark-pigmented dots were also noted by Ottoboni et al. (2017), who described them as brownish; however, no bristles were observed in that study. The author further reported that *T. molitor* cuticular fragments can be difficult to recognize because these dots are not consistently present. Under light microscopy, the bases of sensilla trichoidea (hair-like) and sensilla chaetica (bristle-like) appear as small puncture-like features marking the points where setae were originally attached (Harris, 1979; Marien et al., 2024).

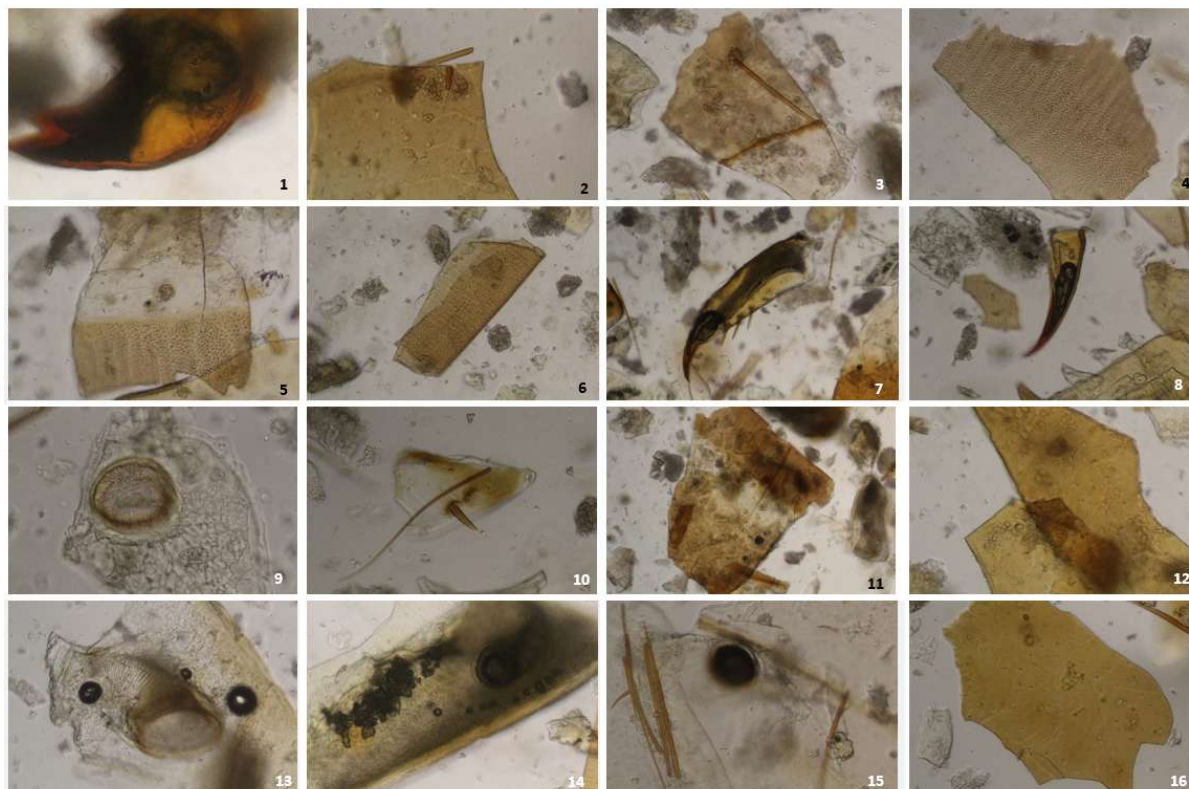


**Figure 3.** Fragments from *T. molitor*: 1 Urogomphi and anal spine fragments; 2 Muscle tissue; 3 Mandibular fragments; 4–10 Cuticular fragments; 11,14 Head capsule fragments; 12 Bristles; 13,15 Cuticle with tracheal structures; 16 Spiracle.

Pupae of *T. molitor* are characterized by the presence of gin-trap structures (EURL-AP micrograph collection; Hinton, 1946; Wilson, 1971). No published literature was found regarding the presence of gin-traps in *A. diaperinus*. Mouthpart fragments, including mandibles, urogomphi, and leg fragments identifiable by their claws, are frequently observed. *T. molitor* possesses ten pairs of spiracles throughout all developmental stages (Raś et al., 2018). Immediately behind the occluding apparatus of each spiracle lies the tracheal vestibule, a spherical three-dimensional structure from which multiple tracheae originate and extend throughout the insect body (Raś et al., 2018). Each spiracle consists of a peritreme that encircles the atrial orifice (Raś et al., 2018).

Fragments of *T. molitor* can be distinguished from those of *A. diaperinus* mainly by their larger overall size; structures such as head capsules and mandibles therefore appear proportionally larger within feed material. *T. molitor* cuticle fragments also tend to exhibit a smoother surface. In contrast to *A. diaperinus*, *T. molitor* fragments lack a clearly defined surface pattern or an organized arrangement of bristles (Ottoboni et al., 2017). Veys and Baeten (2018) reported no clear distinction between fragments of *T. molitor* and *A. diaperinus*. Fragments of urogomphi are expected in both

species, as documented in the EURL-AP micrograph collection for *T. molitor* and described for the larval stages of *A. diaperinus* in the literature (Figure 4; Chernaki et al., 2001). The presence of urogomphi in both sexes (Esquivel et al., 2012), which serve as reliable indicators of late-instar larvae, remains a definitive diagnostic feature for confirming Tenebrionid larval origin.



**Figure 4.** Fragments from *A. diaperinus*: 1 Mandibular fragment; 2–6 Moderately sclerotized cuticle and thoracic leg segments; 7–8 Claws; 9,14–15 Spiracles; 10 Cuticle with sparse short setae; 11,12,16 Additional cuticle fragments; 13 Spiracle with tracheal system fragments.

After the second instar, Coleopteran larvae and pupae show limited morphological variation (Park et al., 2014), yet they remain more morphologically differentiated than dipteran larvae, possessing developed mouthparts, legs, and a strongly sclerotized cuticle, which results in a higher proportion of rigid fragments in processed feed (Veys and Baeten, 2018). The larvae of both Coleopteran species lack denticles, in contrast to the denticle-bearing larvae of *H. illucens* (Veys and Baeten, 2018). The respiratory system of holometabolous insects consists of an internal network of cuticle-lined tracheae and tracheoles (Lowe et al., 2013; Iwan et al., 2015). During microscopic examination of insect meal, tracheolar fragments are more frequently observed in dipteran larvae such as *H. illucens* than in Coleopteran larvae (Veys and Baeten, 2018).

A wide diversity of particles can be observed under light microscopy in Orthopteran-derived PAP, as the meal is prepared from nymphs or imagos. These developmental stages possess legs, a well-developed head with eyes, mouthparts, and antennae, and, in the final nymphal instars, even wing pads are present (Veys and Baeten, 2018). The resulting particles include pigmented and sclerotized cuticular fragments; leg-derived structures such as tibial spines and femoral or tarsal pieces; sclerotized mandibular fragments; membranous or sclerotized wing fragments; eye material; cercal fragments; segmented antennal flagellomeres; honeycomb-like cuticle (notably in *G. assimilis*); cuticle bearing setae or isolated setae; claws; muscle tissue; tracheal system fragments; and muscle fibres associated with tracheae (Figure 5, 6, 7; EURL-AP micrograph collection; Veys and Baeten, 2018).



**Figure 5.** Fragments from *A. domesticus*. 1–3: cuticular fragments bearing setae; 4–5: segmented, cylindrical antennal flagellomeres; 6: tracheal structure; 7–8: tarsal claws; panel 9: isolated setae.

Veys and Baeten (2018) reported the presence of distinct short, segmented fragments originating from antennae or leg parts in *G. assimilis*, a feature not observed in PAPs produced from *H. illucens*, *T. molitor*, or *A. diaperinus*.

The hind tibiae of *G. assimilis* are characterized by highly sclerotized, robust spines accompanied by a dense covering of coarse setae. Wing fragments show dense, reticulated, leathery venation. The cerci are covered in long sensory hairs, each arising from a distinctive cup-shaped base (trichobothria) visible under high magnification. Adults of *G. assimilis* share many general traits with *A. domesticus*, but they are typically darker and exhibit a denser brown pubescence, giving them a noticeably “hairier” appearance under magnification. The mandibles of *G. assimilis* appear dark brown to black.

Small, non-functional wing fragments with simplified venation, together with sclerotized brownish mandibles, are highly diagnostic for *G. sigillatus*. The hind tibiae bear comb-like fixed spines that are noticeably smaller and more slender than those of the larger *G. assimilis*. The abdominal cerci are long and covered in sensory hairs. The cuticular hairs of *G. sigillatus* tend to be lighter in colour, in contrast to the dark brown to black bristles typical of *G. assimilis*. As a result, fragmented material from *G. assimilis* often contains a higher density of dark, coarse setae, whereas *A. domesticus* exhibits lighter and finer hairs.



**Figure 6.** Fragments from *G. sigillatus*. 1: cuticular fragment; 2: muscle tissue; 3–9: cuticular fragments with setae; 10–12: segmented antennal flagellomeres; 13: cuticle with isolated setae; 14–15: claws.



**Figure 7.** Fragments from *G. assimilis*: 1 segmented antennal flagellomere; 2–4: cuticular fragments; 5: muscle tissue with trachea; 6: segmented antennal flagellomere; 7: honeycomb-like cuticle; 8: antennal flagellomere; 9, 15–16: claws; 10–12: cuticle; 13: trachea; 14: isolated setae.

The EURL-AP micrograph collection covers only five insect species currently authorized as feed materials in the EU: *H. illucens*, *T. molitor*, *A. diaperinus*, *A. domesticus*, and *G. assimilis*. In *H. illucens*, the most frequently cited morphological structures include cuticle, anal segment, spine-like sensilla, tracheal system, muscle tissue, and spiracles. In *T. molitor*, reported fragments comprise cuticle, leg and claw pieces, tracheal system, urogomphi, antennae, anal spine, muscle, spiracle, gin trap, and labrum. For *A. diaperinus*, characteristic elements include cuticle, setae, legs, anal spine, tracheal system, muscle fibres, and mandibles. different diagnostic patterns were photographed for the cricket species; *G. assimilis* is characterized by cuticle, antennae, muscle fibres, mandibles, and wing fragments, whereas *A. domesticus* typically presents cuticle, cercus fragments, setae, claws, tracheal system, eye fragments, muscle fibres, and wing fragments. Despite its importance for laboratory analysis of insect processed-protein detection in feed in accordance with EU Regulation 152/2009, the collection still lacks several authorized species.

Structural details of cuticular fragments, setae or trichoid sensilla, and tracheolar structures, together with characteristic patterns of muscle fibres, have been shown to provide robust identification criteria for determining the insect origin of particles (Simpson and Douglas, 2013; Veys and Baeten, 2018; Weiner and Kwiatek, 2022). Trichoid sensilla, or setae, are common in all insect-derived PAPs. They vary in quantity, colour, shape, and size, and may appear either still attached to cuticular fragments or as isolated elements within the sample (Simpson and Douglas, 2013; Veys and Baeten, 2018). According to Veys and Baeten (2018), these setae are entirely or nearly unpolarized under polarized light microscopy, a feature that allows them to be distinguished from plant trichomes, commonly present in feed materials such as wheat bran or gluten, which exhibit strong polarization (Veys and Baeten, 2018).

Fragments of insect tracheae and tracheoles may occur on larger cuticular pieces as well as within muscle fibres, confirming the insect origin of the muscle tissue (Veys and Baeten, 2018; Weiner and Kwiatek, 2022). In the study by Veys and Baeten (2018), these structures were reported predominantly in PAP derived from *G. assimilis*, whereas they were far less frequently observed in material from *H. illucens*, *T. molitor*, or *A. diaperinus*. Weiner and Kwiatek (2022) also noted tracheolar fragments, together with muscle-fibre elements in *H. illucens* and *T. molitor*. At higher magnification, the authors observed a spiral, transverse thickening characteristic of tracheoles. Veys and Baeten (2018) recommended searching for tracheal structures in cases where the identification of exoskeletal remains is uncertain.

Insect muscle fibres often display a characteristic zig-zag sarcomeric striation, a feature that distinguishes them from the muscle fibres found in terrestrial or fish PAPs (Veys and Baeten, 2018; Weiner and Kwiatek, 2022). Although this striation reliably confirms that the material is of insect origin, it does not allow identification at lower taxonomic levels (Weiner and Kwiatek, 2022). These fragments therefore serve only as indicators of the presence of insect material and are not species-specific. In addition, the presence of tracheal structures within cuticular fragments and muscle fibres supports the classification of the material as insect-derived rather than originating from other animal taxa (Veys and Baeten, 2018).

Veys and Baeten (2018) also discussed the differences between insect PAPs originating from nearly identical instars, either larval or nymphal stages, and contamination by insect pests, in which imagoes are usually predominant. In such cases, remains of coleopteran elytra, membranous hindwings, compound eyes, antennae fragments, and other adult structures will inevitably be detected.

More specific methods for determining authorized insect species are recommended as a second step in feed monitoring, such as DNA-based approaches (van Raamsdonk et al., 2017; Weiner and Kwiatek, 2022). Molecular studies targeting sequences of *T. molitor*, *H. illucens*, and *B. mori* have been conducted by Debode et al. (2017) and Marien et al. (2018, 2025). It is important to emphasize, however, that DNA-based methods and other analytical techniques used in feed control should only be applied as complementary tools after light microscopy has confirmed the presence of PAP fragments. This approach continues the effective strategy that has been legally adopted for decades for the detection of PAPs in feed (Fumière et al., 2009; Veys et al., 2012; Lecrenier et al., 2016; Ottoboni et al., 2017; Veys and Baeten, 2018). Recently, Kaisin et al. (2025) proposed integrating automated recognition software with optical microscopy to reduce human error and accelerate identification.

## Conclusion

In the EU, light microscopy, currently the official method for detecting prohibited animal proteins, including insect processed animal proteins, remains a robust and accessible analytical tool. It offers high sensitivity and requires only basic laboratory skills and equipment. Light microscopy is effective in identifying insect PAPs, particularly cuticular fragments and intact structures such as mandibles, legs, or antennal segments, even after milling and drying. However, it does not provide species-level identification and cannot distinguish authorized from non-authorized insect species. For this reason, the continued integration of microscopy with complementary methods will be essential to meet future regulatory and analytical requirements.

There is now a widely recognized need for more specialized training to support taxonomic discrimination of insect species, beyond the standard training currently provided to control laboratories. Integrating automated recognition software with optical microscopy to reduce human error and accelerate identification is also crucial for addressing the limitations of light microscopy.

This work provides guidance for identifying particles of authorized insects in feed. Although key aspects of microscopic identification have been addressed, careful interpretation remains essential. The proposed identification key is useful when fragments are abundant and species-specific, but its diagnostic value decreases when only small, non-distinct structures such as tracheoles or muscle fibres are present. Overall, microscopic detection of insect particles is far more

effective when supported by a comprehensive photographic database to assist technicians in routine analysis.

## Methodology

Slides were prepared according to the specifications outlined in Section 2.1.2.1.4.4 of Annex VI to Regulation (EC) No 152/2009, as well as the EURL-AP Standard Operating Procedure (SOP) for staining reagents. Pure meals of *Hermetia illucens*, *Tenebrio molitor*, *Alphitobius diaperinus*, *Acheta domestica*, *Grylloides sigillatus*, and *Gryllus assimilis* were used for the preparation of reference slides. Laboratory reference material of *Bombyx mori* was acquired as an industrial ready-made meal. Material from *Musca domestica* remains difficult to obtain; therefore, morphological features for this species were extracted from the available scientific literature. Reference material observation and preparation were carried out in the Feed and Food Microscopy Laboratory at AGROLAB Alimentalia (Italy). Characteristic features of insects were determined by microscopic images using the Zeiss Axiovert 25 Inverted Phase Contrast Microscope. Observations were conducted at magnifications of 10X, 20X, and 32X, each paired with a 10X ocular lens. A 0.4 N.A. condenser equipped with a stage and phase slider was employed during examination. Photographic documentation of the slides included an additional 2.5X magnification.

Microsoft Copilot was used to assist with the structural refinement of the identification key; all scientific interpretations and taxonomic decisions were made by the authors.

**Supplementary Materials:** The following supporting information can be downloaded at the website of this paper posted on Preprints.org.

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