

Risk of exposure of a selected rural population in South Poland to allergenic mites. Part II: acarofauna of farm buildings

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Abstract

Exposure to mite allergens, especially from storage and dust mites, has been recognized as a risk factor for sensitization and allergy symptoms that could develop into asthma. The aim of this study was to investigate the occurrence of mites in debris and litter from selected farm buildings of the Małopolskie province, South Poland, with particular reference to allergenic and/or parasitic species as a potential risk factor of diseases among farmers. Sixty samples of various materials (organic dust, litter, debris and residues) from farm buildings (cowsheds, barns, chaff-cutter buildings, pigsties and poultry houses) were subjected to acarological examination. The samples were collected in Lachowice and Kurów (Suski district, Małopolskie). A total of 16,719 mites were isolated including specimens from the cohort Astigmatina (27 species) which comprised species considered as allergenic (e.g., Acarus siro complex, Tyrophagus putrescentiae, Lepidoglyphus destructor, Glycyphagus domesticus, Chortoglyphus arcuatus and Gymnoglyphus longior). Species of the families Acaridae (A. siro, A. farris and A. immobilis), Glycyphagidae (G. domesticus, L. destructor and L. michaeli) and Chortoglyphidae (C. arcuatus) have been found as numerically dominant among astigmatid mites. The majority of mites were found in cowsheds (approx. 32%) and in pigsties (25.9%). The remaining mites were found in barns (19.6%), chaff-cutter buildings (13.9%) and poultry houses (8.8%). The results suggest that the allergenic mites may constitute an occupational hazard for agricultural workers in all farming environments examined.

Keywords Storage mites · Acaridae · Glycyphagoidea · Pyroglyphidae · Litter

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Introduction

Storage mites, especially several species from the families Acaridae, Glycyphagidae and Chortoglyphidae (Acari: Sarcoptiformes, Astigmatina), are commonly found in stored food products, hay, straw, granaries, barns and other farming and occupational environments, in dust from railway freight wagons, as well as in samples of house dust. The most abundant mites are *Acarus siro, Acarus farris, Tyrophagus longior* and *Tyrophagus putrescentiae* from Acaridae, *Lepidoglyphus destructor* and *Glycyphagus domesticus* from Glycyphagidae and *Chortoglyphus arcuatus* from *Chortoglyphidae* (Boström et al. 1997; Franz et al. 1997; Mehl 1998; Sánchez-Ramos et al. 2004, 2007; Pike and Wickens 2008; Wong et al. 2011; Solarz 2012). These mites were identified as a source of clinically important allergens, causing occupational allergy (known as allergy to storage mites) among farmers, grain-storage workers and other agricultural workers (Fain et al. 1990; Revsbech and Dueholm 1990; van Hage-Hamsten and Johansson 1998). These mites cause IgE-mediated sensitization among the above-mentioned occupational populations, who develop asthma, rhinitis and conjunctivitis when exposed to organic dust containing mite allergens (Arlian 2002; Sánchez-Ramos et al. 2004; Berger et al. 2005; Solarz 2012).

The aim of this study was to investigate the occurrence of mites in debris and litter from selected farms of the Małopolskie province, South Poland, with particular reference to allergenic and/or parasitic species as a potential risk factor of diseases among farmers.

Materials and methods

The study was carried out from May 2011 to September 2012. A total of 60 samples of materials collected from four farms in two locations (Lachowice and Kurów) on the territory of the Suski district (Stryszawa and vicinity, Małopolskie) were examined as the potential sources of the allergenic and/or parasitic mites in the farming environments (Table 1). All samples were collected into plastic bags of 1 l capacity. These samples were sweepings containing organic dust, litter, debris, residues and other materials from certain farming environments. Generally 12 cowsheds, barns, pigsties, chaff-cutter buildings and poultry houses were analyzed for the occurrence of mites, including allergenic and/or parasitic taxa (Table 1). The mites were extracted using the Berlese method and preserved in 75% ethanol. For identification, all mites were mounted in Faure's medium on microscope slides, with the aid of the Olympus Europe Highlight 2100 stereomicroscope. All mite specimens collected were examined using differential interference contrast (Nomarski

Farming environments	Barns	Pigsties	Cowsheds	Poultry houses	Chaff- cutter buildings
No. samples examined	12	12	12	12	12
No. mites collected	3271	4321	5341	1465	2321
% of total count	19.56	25.85	31.95	8.76	13.88
No. mites per sample	109.03	144.03	198.03	48.83	77.37

 Table 1
 Summary statistics: number of samples collected from the particular farm buildings examined on the territory of Stryszawa and vicinity (Małopolskie province, South Poland)

DIC) (under a Zeiss Axioskope 2 plus light microscope) and phase contrast optics (under a Olympus CH 40 light microscope). Mites were identified using descriptions of taxa and the following publications: Hughes (1976), Baker (1999), Zhang and Fan (2005), Fan and Zhang (2007), Colloff (2009), Krantz and Walter (2009) and Solarz (2012). The results were expressed as the number of mites per g of sample (wet weight). Mite abundance was also calculated as the number of specimens per sample.

Data were analyzed using χ^2 tests in CSS-STATISTICA for Windows v.12 ($\alpha = 0.05$).

Results

Overall results

Species composition and dominance of the total mite fauna in samples from all types of farm buildings is listed in Table 2, whereas the abundance of allergenic acarofauna from particular types of farm buildings is given in Table 1. The weight of the samples ranged between 11.08 and 36.91 g (mean = 17.92 ± 6.86 g; median = 15.70 g). Mites were found in all samples examined – in total 16,719 mite specimens were isolated and 27 species were identified (Table 2). Most mites were found in cowsheds (approx. 32% of the total count) and in pigsties (25.9%), followed by barns (19.6%), chaff-cutter buildings (13.9%) and poultry houses (8.8%; Table 1). Samples were dominated by mites of the subfamilies Acaroidea and Glycyphagoidea. In addition, two species of pyroglyphid mites (Astigmatina) were collected (*Gymnoglyphus longior* and *Hirstia chelidonis*), as well as some members of Prostigmata (Cheyletidae, Tydeidae, Cunaxidae), Oribatida sensu lato and Mesostigmata (Table 2).

Species composition, frequency, dominance and abundance (numbers per g or per sample) of the acarofauna varied considerably between particular types of farm buildings examined (Tables 3, 4). Generally, the dominant species were A. *siro* (n = 5065; 30.3% of all mites), A. *farris* (n = 2009; 12.0%) and Acarus immobilis (n = 1265; 7.6%) of the family Acaridae (Astigmatina), and two species from the family Glycyphagidae (Astigmatina) – G. domesticus (n = 2051; 12.3%) and L. destructor (1790; 10.7%) (Table 2). Among the relatively numerous mites were cheyletids (Prostigmata) (1186; 7.1%) and some other astigmatid and potentially allergenic mites—*T. putrescentiae* (661; 3.95%) and Acarus *nidicolous* (534; 3.2%) from Acaridae, and Lepidoglyphus michaeli (562; 3.4%) and C. arcuatus (533; 3.2%) from Glycyphagoidea (Table 2). Other storage mites—*T. longior*, *Tyrolichus casei*, Mycetoglyphus fungivorus, Aleuroglyphus ovatus, Calogyphus berlesei, Rhizoglyphus spp. (all Acaridae), Lepidoglyphus fustifer, Glycyphagus privatus, Glycyphagus ornatus, Ctenoglyphus plumiger, Ctenoglyphus canestrinii and Diamesoglyphus intermedius (all Glycyphagidae)—were generally less abundant, as were the two pyroglyphids, *G. longior* and *H. chelidonis*.

Acarus siro, G. domesticus and L. destructor were also the most frequent mites in samples from the examined farm buildings with relative frequency 85.3, 66.7 and 66.0%, respectively (Table 2). The difference in frequency between A. siro and the two glycy-phagids was significant (χ^2 test: p < 0.005, in both cases). Acarus siro was the most frequent mite in almost all farm building types examined—only in poultry houses it was preceded by L. destructor; in cowsheds both species occurred with the same frequency (Table 3). The frequency of A. siro was highest in chaff-cutter buildings (100%) and in pigsties (93%). This species was significantly more frequent in chaff-cutter buildings than in cowsheds,

able 2 Species list, dominance,.bundance and frequency of	Mite taxa	Dominar	nce	Frequ	ency
nites found in samples from		N	D (%)	n	F (%)
he various types of farm puildings combined, situated on	Acarus siro	5065	30.29	128	85.33
he territory of Stryszawa and	Acarus farris	2009	12.02	73	48.66
vicinity (Małopolskie)	Acarus immobilis	1265	7.57	49	32.66
	Acarus nidicolous	534	3.19	11	7.33
	Acarus gracilis	8	0.05	3	2.00
	Tyrophagus putrescentiae	661	3.95	42	28.00
	Tyrophagus longior	55	0.33	17	11.33
	Tyrolichus casei	4	0.02	1	0.66
	Mycetoglyphus fungivorus	3	0.02	1	0.66
	Aleroglyphus ovatus	1	0.01	1	0.66
	Sancassania berlesei	4	0.02	1	0.66
	Rhizoglyphus echinopus	2	0.01	1	0.66
	Rhizoglyphus robinii	2	0.01	1	0.66
	Lepidoglyphus destructor	1790	10.71	99	66.00
	Lepidoglyphus michaeli	562	3.36	31	20.66
	Lepidoglyphus fustifer	259	1.55	18	12.00
	Glycyphagus domesticus	2051	12.27	100	66.66
	Glycyphagus ornatus	99	0.59	11	7.33
	Glycyphagus privatus	5	0.03	2	1.33
	Ctenoglyphus plumiger	212	1.27	16	10.66
	Ctenoglyphus canestrinii	14	0.08	5	3.33
	Diamesoglyphus intermedius	5	0.03	3	2.00
	Chortoglyphus arcuatus	533	3.19	30	20.00
	Gymnoglyphus longior	71	0.42	13	8.66
	Hirstia chelidonis	15	0.09	5	3.33
	Suidasia nesbitti	2	0.01	2	1.33
	<i>Calvolia</i> sp.	9	0.05	4	2.66
	Oribatida (sensu lato)	6	0.04	5	3.33
	Cheyletidae	1186	7.09	109	72.66
	Tydeidae	3	0.02	1	0.66
	Cunaxidae	32	0.19	7	4.66
	Heterostigmatina	11	0.07	4	2.66
	Prostigmata (other)	6	0.04	4	2.66
	Mesostigmata	235	1.41	47	31.33
	ACARI (total)	16,719	100	150	100

N number of mite specimens, *n* number of positive samples, *D* (%) percent of total count of mites collected, *F* (%) percent of the total number of samples positive for mites (n = 150)

pigsties, barns and poultry houses ($\chi^2 = 13.9$, 7.25, 22.22 and 40.96; $p \le 0.0071$). It was also significantly more frequent in pigsties than in poultry houses ($\chi^2 = 22.37$; $p \le 0.00001$) and barns ($\chi^2 = 7.24$; p = 0.0071), whereas differences in frequency of *A. siro* in pigsties and cowsheds were not significant (p = 0.16). Moreover, *A. siro* was significantly more

	Barns				Pigsties	es			Cowsheds	leds			Chaff	Chaff-cutter buildings	iplind	sgu	Poult	Poultry houses	s	
	Domi	Dominance	Freq	Frequency	Domi	Dominance	Freq	Frequency	Dominance	lance	Freq	Frequency	Domi	Dominance	Free	Frequency	Domi	Dominance	Freq	Frequency
	z	D (%)	u u	F (%)	z	D (%)	u	F (%)	z	D (%)	u u	F (%)	z	D (%)	u u	F (%)	z	D (%)	- u	F (%)
Acarus siro	945	28.89	24	80.00	732	16.94	28	93.33	2721	50.94	26	86.66	432	18.61	30	100.00	235	16.04	20	66.66
Acarus farris	956	29.23	15	50.00	495	11.46	14	46.66	271	5.07	17	56.66	270	11.63	20	99.99	17	1.16	٢	23.33
Acarus immobilis	241	7.37	8	26.66	311	7.20	13	43.33	89	1.67	13	43.33	624	26.88	10	33.33				
Acarus nidicolous					534	12.36	11	36.66												
Acarus gracilis	4	0.12	0	6.66					4	0.07	1	3.33								
Tyrophagus putrescentiae	13	0.40	5	16.66	479	11.08	10	33.33	17	0.32	4	13.33	116	5.00	15	50.00	36	2.46	×	26.66
Tyrophagus longior					8	0.18	б	10.00	26	0.49	5	16.66	21	0.90	6	30.00				
Rhizoglyphus echinopus									0	0.04	1	3.33								
Rhizoglyphus robinii									0	0.04	1	3.33								
Tyrolichus casei																	4	0.27	1	3.33
Mycetoglyphus fungivorus																	3	0.20	1	3.33
Aleuroglyphus ovatus																	1	0.07	1	3.33
Sancassania berlesei																	4	0.27	1	3.33
Lepidoglyphus destructor	78	2.38	21	70.00	1109	25.67	19	63.33	297	5.56	26	86.66	163	7.03	11	36.66	143	9.76	22	73.33
Lepidoglyphus fustifer					112	2.59	5	16.66	57	1.07	9	20.00	39	1.68	Э	10.00	51	3.48	4	13.33
Lepidoglyphus michaeli	103	3.15	6	30.00	5	0.12	1	3.33	411	7.70	12	40.00	31	1.34	5	16.66	12	0.82	4	13.33
Glycyphagus domesticus	373	11.40	19	63.33	281	6.50	16	53.33	721	13.50	24	80.00	145	6.25	25	83.33	531	36.25	16	53.33
Glycyphagus ornatus	1	0.03	1	3.33	4	0.09	1	3.33	94	1.76	6	30.00								
Glycyphagus privatus	3	0.09	1	3.33					0	0.04	1	3.33								
Ctenoglyphus plumiger	165	5.04	×	26.66	10	0.23	б	10.00	5	0.09	1	3.33	21	0.90	0	99.9	11	0.75	0	6.66
Ctenoglyphus canestrinii	13	0.40	4	13.33					1	0.02	1	3.33								
Diamesoglyphus intermedius					0	0.05	0	6.66					ю	0.13	1	3.33				
Chortoglyphus arcuatus	204	6.24	13	43.33	49	1.13	2	16.66	6	0.04	1	3.33	23	0.99	٢	23.33	255	17.41	2	16.66

Mite taxa	Barns				Pigsties	ies			Cowsheds	heds			Chaff	Chaff-cutter buildings	ipline	ngs	Poult	Poultry houses	SS	
	Dominance	lance	Freq	Frequency	Dom	Dominance	Free	Frequency	Domi	Dominance	Freq	Frequency	Dom	Dominance	Free	Frequency	Domi	Dominance	Free	Frequency
	z	D (%)	- u	F (%)	z	D (%)	u u	F (%)	z	D (%)	u u	F (%)	z	D (%)	- -	F (%)	z	D (%)	- -	F (%)
Gymnoglyphus longior	25	0.77	3	10.00					41	0.77	8	26.66	5	0.22	2	99.9				
Hirstia chelidonis	4	0.12	1	3.33					11	0.20	4	13.33								
Suidasia nesbitti	2	0.06	ы	6.66																
Calvolia sp.									7	0.13	б	10.00					7	0.14	1	3.33
Ornithonyssus sylviarum																	1	0.07	1	3.33
Oribatida (sensu lato)	5	0.15	4	13.33					1	0.02	1	3.33								
Cheyletidae	94	2.87	19	63.33	123	2.85	13	43.33	508	9.51	27	90.00	317	13.66	21	70.00	14	9.83	29	99.96
Tydeidae	3	0.09	1	3.33																
Cunaxidae	8	0.25	ŝ	10.00									6	0.39	ы	6.66	15	1.02	ы	6.66
Heterostigmatina													11	0.47	4	13.33				
Prostigmata (other)													9	0.26	4	13.33				
Mesostigmata	31	0.95	13	43.33	67	1.55	8	26.66	51	51 0.95	17	56.66	85	3.66	×	26.66				
ACARI (total)	3271	100	30	100	4321	100	30	100	5341	100	30	100	2321	100	30	100	1465	1465 100	30	100
N number of mite specimens, n number of positive samples, D (%) percent of total count of mites collected, F (%) percent of the total number of samples positive for mites (n = 30)	s, <i>n</i> numł	oer of p	ositive	e sample	ss, D (%) perce	nt of	total cou	nt of 1	mites col	lected	l, F (%)	percer	at of the	total	number c	of samp	les posi	tive fc	r mites

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Mites	Barns		Pigstie	s	Cowsh	eds	Chaff buildi	-cutter ings	Poult: house	-
	М	N/S	М	N/S	М	N/S	М	N/S	М	N/S
Acarus siro	10.78	39.38	8.44	26.14	30.15	104.65	4.07	14.4	2.58	11.75
Acarus farris	10.90	63.73	5.81	35.36	2.73	15.94	3.05	13.5	0.16	2.43
Acarus immobilis	2.74	30.13	3.23	23.92	0.89	6.85	7.06	62.4		
Tyrophagus longior			0.07	2.67	0.31	5.2	0.23	2.33		
Tyrophagus putrescentiae	0.68	2.6	7.03	47.9	1.01	4.25	1.30	7.73	0.38	4.5
Lepidoglyphus michaeli	0.82	11.4	0.05	5.0	3.78	34.25	0.34	6.2	0.66	3.0
Lepidoglyphus destructor	0.63	3.71	12.73	58.37	2.84	11.42	1.75	14.82	1.32	6.5
Lepidoglyphus fustifer			6.57	22.4	3.17	9.5	2.20	13.0	2.80	12.75
Glycyphagus domesticus	4.11	19.63	3.28	17.56	8.01	30.04	1.64	5.8	5.82	33.19
Glycyphagus ornatus			0.06	4.0	1.06	10.44				
Ctenoglyphus plumiger	1.88	20.62			0.05	5.0			0.11	5.5
Chortoglyphus arcuatus	2.05	15.69	0.53	9.8	0.11	2.0	0.06	3.29	2.80	51.0
Gymnoglyphus longior	0.27	8.33			2.23	5.13	0.05	2.5		
Hirstia chelidonis	0.05	4.0			0.61	2.75				
Cheyletidae	0.57	4.95	1.11	9.46	3.89	18.81	2.71	15.09	0.66	4.97

 Table 4
 Abundance of the most important allergenic mites in the particular farm buildings examined on the territory of Stryszawa and vicinity (Małopolskie)

M mean number of mites per g of sample, N/S number of mites per sample

frequent in cowsheds than in poultry houses ($\chi^2 = 12.3$; p = 0.0005), and more in barns than in poultry houses ($\chi^2 = 4.97$; p = 0.026), whereas the difference in frequency of *A. siro* in cowsheds vs. barns was not significant ($\chi^2 = 1.78$; p = 0.18) (Table 3).

Lepidopglyphus destructor occurred much less frequently in chaff-cutter buildings than in the remaining types of farm buildings examined ($p \le 0.00001$, in all cases). Moreover, it was more frequent in cowsheds (87%) than in poultry houses ($\chi^2 = 6.13$, p = 0.013), barns ($\chi^2 = 8.6$, p = 0.0034) and pigsties ($\chi^2 = 15.36$, p = 0.0001). It was also more frequent in poultry houses and barns than in pigsties but these differences were not significant ($\chi^2 = 2.3$ and 1.1, p = 0.13 and 0.29, respectively) (Tables 2, 3, 4).

Glycyphagus domesticus occurred most frequently in chaff-cutter buildings (83% positive samples) and cowsheds (80%) (Tables 2, 3, 4). It was more frequently found in chaff-cutter buildings and cowsheds than in the remaining types of farm buildings sampled (p < 0.01, in all cases). The difference in frequency in barns (63%), poultry houses (53%) and pigsties (53%) was not significant ($\chi^2 = 2.05$, p = 0.15) (Tables 3, 4).

Mite fauna in barns

In total 3271 mites were collected from barns, including 16 species from the cohort Astigmatina (Sarcoptiformes, Oribatida) (Tables 1, 3). Mite fauna from barns was dominated by two species of the genus *Acarus: A. farris* (29.2%) and *A. siro* (28.9%) (Astigmatina, Acaridae). Also numerous were *G. domesticus* (11.4%), *C. arcuatus* (6.2%) and *C. plumiger* (5.0%) from the superfamily Glycyphagoidea, and the acarid *A. immobilis* (7.4%) (Table 3). These mite species were also most abundant per g dust and per sample (Table 4). Among the less numerous remaining astigmatid mites were some allergenic taxa, as *L. michaeli* and *L. destructor* (Glycyphagidae), *T. putrescentiae* (Acaridae), *G. longior* and *H. chelidonis* (Pyroglyphidae), and *Suidasia nesbitti* (Suidasiidae) (Tables 3, 4). An important part of the non-astigmatid mites constituted predaceous and allergenic mites of the family Cheyletidae (Prostigmata) (2.3% of mites collected from barns). Mesostigmata constituted only 0.95% of mites collected.

Mite fauna in pigsties

In total 4321 mites were collected from the pigsties (Tables 1, 3). *Lepidopglyphus destructor* (Glycyphagidae) was predominant (approx. 25.7% of the total count), followed by the acarids *A. siro* (16.9%), *A. nidicolous* (12.4%), *A. farris* (11.5%), *T. putrescentiae* (11.1%) and *A. immobilis* (7.2%). The other eight species of astigmatid mites found were *T. longior* (Acaridae), *L. fustifer, L. michaeli, G. domesticus, G. ornatus, C. plumiger, D. intermedius* (Glycyphagidae) and *C. arcuatus* (Chortoglyphidae) (Table 3). Among the remaining mite taxa cheyletids were most numerous (2.85% of all mites collected from these pigsties).

Mite fauna in cowsheds

Acarofauna of cowsheds was also dominated by astigmatid mites: of the 5341 mites isolated from cowsheds, 4781 were of the cohort Astigmatina (89.5%) (Table 3). Twenty astigmatid species were found: A. siro, A. farris, A. immobilis, Acarus gracilis, T. longior, T. putrescentiae, Rhizoglyphus robini, R. echinopus (Acaridae), G. domesticus, G. ornatus, G. privatus, L. michaeli, L. destructor, L. fustifer, C. canestrinii (Glycyphagidae), C. arcuatus (Chortoglyphidae), G. longior, H. chelidonis (Pyroglyphidae) and Calvolia sp. (Winterschmidtiidae) (Table 3).

Acarus siro was the dominant species (50.9%) and the most abundant mite per sample and per g, whereas G. domesticus was the most numerous glycyphagid species (13.5%) (Tables 3, 4). Numerous among the astigmatid mites were A. farris (5.1%), L. michaeli (7.7%) and L. destructor (5.6%). Cheyletid mites constituted 9.5% of the total count of mites from the examined cowsheds (Table 3).

Mite fauna in chaff-cutter buildings

Acarofauna from chaff-cutter buildings was also dominated by the Acaridae: of the 2321 mites isolated (Tables 1, 3), 1893 specimens were from the cohort Astigmatina (81.6% of all mites collected from chaff-cutter buildings) and 1463 mites were acarids (63.0% of the total count). These mites were also most abundant both per sample and per g (Table 4). Among Astigmatina 13 species were found: *A. siro, A. farris, A. immobilis, T. putrescentiae, T. longior* (Acaridae), *L. fustifer, L. destructor, C. plumiger, D. intermedius* (Glycyphagidae), *C. arcuatus* (Chortoglyphidae) and *G. longior* (Pyroglyphidae) (Table 3). *Acarus immobilis* was the dominant and most abundant species, both per g and per sample (Tables 3, 4). Species of the *A. siro* complex constituted 57.1% of the total count of mites collected from the chaff-cutter buildings, including *A. siro* (18.6%) and *A. farris* (11.6%). Among the other numerous mites were *L. destructor* (7.0%), *G. domesticus* (6.3%), *T. putrescentiae* (5.0%), and members of the family Cheyletidae (13.7%) (Table 3).

Mite fauna in poultry houses

Among 1465 mites collected from poultry houses, *G. domesticus* was predominant (36.3% of the total count), followed by *C. arcuatus* (17.4%), *A. siro* (16.0%), cheyletids (9.8%) and *L. destructor* (9.8%) (Table 3). In total 14 species of astigmatid mites were found in poultry houses, including also *A. farris, T. putrescentiae, T. casei, M. fungivorus, A. ova-tus, Sancassania berlesei* (Acaridae), *L. fustifer, L. michaeli, C. plumiger* (Glycyphagidae) and *Calvolia* (Winterschmidtiidae) (Table 3). Members of the superfamily Glycyphagoidea were distinctly more abundant than the remaining astigmatid mites and constituted approximately 68.5% of all mites collected (Table 3). Among the mesostigmatid mites, one specimen of *Ornithonyssus sylviarum* was found (Table 3).

Abundance of mites in the examined farm buildings

The mean number of all mites collected per g of sample constituted 17.55. The number of mites per g sample was varied considerably among mite species and among particular types of farm buildings (Table 4). The most abundant species were *A. siro* (especially in cowsheds, barns and pigsties), *A. farris* (in barns) and *L. destructor* (mainly in pigsties) (Table 4). The total number of mites per sample constituted 278.65 (on average 111.5). The number of mites per sample was highest for cowsheds (198.0) and lowest for poultry houses (48.8) and chaff-cutter buildings (77.4) (Table 4). The most abundant species of mites per sample were *A. siro* (especially in cowsheds and barns), *A. farris* (in barns and pigsties), *L. destructor* (mainly in pigsties) and *G. domesticus* (mainly in barns). The remaining allergenic mites were distinctly less abundant both per sample and per g (Table 4).

Discussion

Agricultural work is considered to be a major risk factor for occupational diseases (Berger et al. 2005). All mites should be regarded as a potential source of mite allergens in environments associated with activity of man, not only in dwellings (Fain et al. 1990; Solarz 2006; Stejskal and Hubert 2008; Colloff 2009). So far, many groups of mites were not reported as occupational biohazards for humans (Arlian 2002). The greatest exposure to storage mites usually occurs in an agricultural setting where allergies to these mites are of major importance (van Hage-Hamsten and Johansson 1998; Arlian 2002; Cichecka et al. 2006). For the urban population, sensitivity to particular species of storage mites and their cross-reactivity with pyroglyphid dust mites is usually not important. Generally, the exposure of city people to storage mites is minimal (Solarz 2010). For farmers and other agricultural workers both storage and house dust mites may act as inhalant allergens. Therefore, for people living in (sub)agricultural settlements, sensitivity to various species of domestic and storage mites and cross-reactivity between species may be of clinical significance (Arlian 2002; Cichecka et al. 2006). On the other hand, sensitization to storage mites in urban dwellers has been reported in Spain, Denmark, Germany, Croatia, Poland and USA (Ebner et al. 1994; Garcia-Robaina et al. 1996; Macan et al. 1998; Gislason and Gislason 1999; Kanceljak-Macan et al. 2000; Arlian 2002; Solarz et al. 2008). Sensitization to T. putrescentiae and L. destructor is present in the urban population of Upper Silesia in similar proportions as to the *Dermatophagoides* spp. (Szilman et al. 2004; Cichecka et al. 2006; Solarz et al. 2008; Asman et al. 2009). Our previous studies suggest that allergenic mites belonging to the Acaridae, Glycyphagidae and Chortoglyphidae should be considered in Poland as potential occupational risk factors contributing to the occurrence of respiratory and dermal diseases among workers of zoos, coal-miners, bird breeders and farmers (Solarz and Solarz 1995; Solarz et al. 2004a, b, 1997; Cichecka et al. 2006).

The natural sources of allergenic mites in stores are still not well known. It has been suggested that the majority of the mite population is brought from the cultivated field into the stores, and that the open field is the main source of storage mites (Hallas and Iversen 1996), whereas bird nests are less important (Solarz et al. 2007).

The investigated farm buildings are typical for A. farris, T. longior, T. putrescentiae, Lepidoglyphus spp. and Glycyphagus spp. It is commonly known that A. siro, T. putrescentiae, T. casei, M. fungivorus, A. ovatus, Sancassania spp. and Rhizoglyphus spp. occur in various stored food products (Hughes 1976; Solarz 2012). Tyrophagus longior usually was the dominant species in haystacks and grainstacks standing in open fields (Hughes 1976). Moreover, T. longior, T. putrescentiae, L. destructor, Tarsonemus spp. and Cheyletidae were frequently found on farms in stored hay and straw (Hallas 1981; Terho et al. 1982, 1985; Hallas and Gudmundsson 1985; Korsgaard et al. 1985; Hallas and Solberg 1989; Hallas et al. 1991; Franz et al. 1997; Solarz et al. 1997; Boström et al. 1997; Mehl 1998; de Saint; Georges-Gridelet et al. 2003). All these mites are considered as allergenic (Revsbech and Andersen 1987, 1989; Luczynska et al. 1990; Armentia et al. 1994; Szilman et al. 2004; Stejskal and Hubert 2008; Asman et al. 2009). The species T. putrescentiae, A. siro complex, G. domesticus and L. destructor, which have actually been found in farming environments, are the most often studied allergenic mites of the cohort Astigmatina (van Hage-Hamsten et al. 1992; Kronqvist et al. 2000; Eriksson et al. 1998, 2001; Arlian 2002; Cichecka et al. 2006; Solarz et al. 2007; Stejskal and Hubert 2008). As mentioned above, these mites produce multiple allergens (Arlian 2002) and may cause occupational allergies. Several tarsonemid species are also considered as allergenic—some authors believe them to be a significant group of allergenic domestic mites (Korsgaard and Hallas 1979; Hallas and Korsgaard 1997).

The stated prevalence of cheyletids and glycyphagids in the farming environment is not in accordance with earlier data by Hallas (1981), Terho et al. (1982) and Solarz et al. (1997). The abundance of acarids was distinctly higher than that found in Solarnia (Lubliniec vicinity), in the vicinity of Kokotek and in Lesko (Krosno province) (Solarz et al. 1997), and it was similar as the abundance in dust samples from byres, hay and grain stores of 11 farms in Eastern Finland (Terho et al. 1982, 1985) and in samples of dust and organic debris from coal mines in Upper Silesia (Solarz and Solarz 1995) or in storage facilities in Iran (de Saint Georges-Gridelet et al. 2003). The number of astigmatid mite species found in this study was higher than in farming environments in Finland (Terho et al. 1982) and Iceland (Hallas 1981; Hallas and Gudmundsson 1985) or in coal mines in Poland (Solarz and Solarz 1995), and it was similar as in farm buildings examined earlier in Poland (Solarz et al. 1997; Solarz 2012) or in Iran (de Saint Georges-Gridelet et al. 2003). Some species of the family Pyroglyphidae (G. longior, H. chelidonis and Dermatophagoides farinae) have been found in farming environments in UK, Switzerland, Israel, Iran, USA, Canada (Hughes 1976; Fain et al. 1990; Mumcuoglu and Lutsky 1990; de Saint; Georges-Gridelet et al. 2003) and also in Poland (Solarz et al. 1997; Solarz 2012).

The present results confirm the occurrence of allergenic mites in cowsheds, barns, chaffcutter buildings and poultry houses (Hughes 1976; Terho et al. 1982; Fain et al. 1990; Mumcuoglu and Lutsky 1990; Solarz et al. 1997; Stejskal and Hubert 2008), and reveal the occurrence of these mites in pigsties. Thus, all these locations should be regarded as a potential source of storage mite allergens in the farming environment. As the occurrence and concentration of mites in samples from different farm buildings may vary to a considerable extent, further studies are highly desirable.

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