

Online Material

Appendix I. Characteristics of two honey types used for the studies of survival of *Pantoea agglomerans*.

	Blossom honey	Blossom honey with honey dew
Moisture	17.2%	17.4%
Electrical conductivity	230 μ S/cm	585 μ S/cm
pH	3.6	3.1
HMF*	<BG (21) mg/kg	<BG (21) mg/kg
Dominant pollen group	<i>Cardamine pratensis</i> <i>Tilia platyphyllos</i> <i>Allium</i> sp.	<i>Prunus</i> sp. <i>Malus</i> sp.
Subdominant pollen group	<i>Helianthus annuus</i> <i>Trifolium repens</i> <i>Betula pendula</i> <i>Taraxacum officinale</i> group <i>Acer campestre</i> <i>Ailanthus altissima</i> <i>Malus</i> sp. <i>Phacelia tanacetifolia</i> <i>Trifolium pratense</i> <i>Fragaria vesca</i> <i>Galium mollugo</i> <i>Vicia faba</i> <i>Robinia pseudacacia</i> <i>Melilotus albus</i> <i>Brassica napus</i> <i>Echium vulgare</i> 2 pollen types not identified	<i>Helianthus annuus</i> <i>Trifolium repens</i> <i>Betula pendula</i> <i>Taraxacum officinale</i> group <i>Acer campestre</i> <i>Campanula</i> sp. <i>Onobrychis viciifolia</i> <i>Hypericum perforatum</i> <i>Lotus corniculatus</i> <i>Plantago lanceolata</i> <i>Helianthemum nummularium</i> <i>Erica carnea</i> <i>Pinus nigra</i> <i>Cirsium arvense</i> <i>Achillea</i> sp. <i>Galium mollugo</i> <i>Salix</i> sp. 1 pollen type not identified fungi spores algae

* Hydroxymethylfurfural.

Appendix II. Origin of strains utilized in this work and frequency of isolation during sampling period.

No.*	Strain name	Origin	Origin and frequency of isolation			
			Honeysac	Pollenloads	Nectar	Plant
1	P21n	Pl ²⁰	2 ^{18,20}	9 ^{3,11,20,a,nk}		10 ^{1,2,13,17}
2	P31i	B ¹⁹	2 ^{19,20}	4 ^{19,20}	1 ¹⁹	16 ^{17,19}
3	P232b	Pl ¹¹	1 ¹¹			1 ¹¹
4	P273e	B ²				
5	P301f	B ²⁰	2 ^{11,20}	5 ^{11,20}	2 ¹¹	6 ^{11,20}
6	P361g	B ¹⁵		3 ¹⁵	1 ¹	2 ¹¹
7	P373n	Hs ^g	1 ^d	5 ^{a,c,h,nk}		2 ^{12,15}
8	P413o	Pl ^e	1 ^g	5 ^{15,a,e}		3 ¹⁵
9	P453b	B ¹¹	1 ¹⁰	3 ^{3,7,16}		1 ⁷
10	P484j	Pl ^g		1 ^c		1 ⁸
11	P533j	B ¹²		1 ⁱ		
12	P583k	B ¹⁵				1 ⁴
13	P715g	B ²⁰	1 ²⁰	1 ²⁰	1 ²⁰	2 ²⁰
14	P735b	B ⁴				
15	P745b	B ⁴	1 ⁵	2 ^{15,c}		3 ^{4,5,12}
16	P791g	B ¹⁵		3 ¹⁵		4 ¹⁵
17	P823l	B ⁸		2 ^{3,8}		
18	P862i	B ¹⁴	4 ^{3,14,np}	9 ^{3,5,14,nk,np}		11 ^{3,5,14}
19	P891f	B ²⁰	2 ²⁰			2 ²⁰
20	P975c	B ⁸				
21	P985b	B ⁴				
22	P1005t	Pl ³	1 ¹⁵	2 ⁵		3 ⁵
23	P1035f	B ³		2 ³		
24	P1137n	Hs ⁵				
25	P115bPa337	B ¹⁷		1 ¹⁷		2 ¹⁷
26	P1187p	Pl ⁵	1 ⁵	1 ⁵		5 ⁵
27	P1218j	B ¹⁶		2 ^{5,16}		1 ¹⁶
28	P1269j	Pl ^{nk}				
29	P1348j	B ¹⁶	3 ^{16,npa}	3 ^{16,npa}		6 ^{9,16}
30	P1378h	B ⁷	1 ⁷	1 ⁷		
31	P1416k	N ²¹	2 ^{16,20}	15 ^{3,10,16,19,20,b,f,g,nk,npa}		26 ^{9,10,14,16,17,19,20}
32	P16813f	Pl ^h				
33	P17113n	B ⁶	2 ^{nk}	1 ⁶		1 ¹⁹
34	P17813b	B ¹⁸	1 ¹⁸			

Appendix II. Continued.

No.*	Strain name	Origin	Origin and frequency of isolation			
			Honeysac	Pollenloads	Nectar	Plant
35	P18113b	B ¹⁸	1 ¹⁸	1 ¹⁸		
36	64b	B ¹⁹		3 ¹⁹	1 ¹⁹	4 ^{16,19}
37	68b	PI ^{nk}				
38	90b	B ²⁰				
39	91b	PI ⁶				
40	280b	B ²¹		1 ²¹	1 ²¹	
41	690b	PI ^{nk}				
42	700b	B ¹⁵		2 ¹⁵		1 ¹⁵
43	765b	B ¹⁴		2 ¹⁴		
44	1351b	PI ^b				
45	1376b	PI ¹⁷				1 ¹⁷
46	1410b	PI ^g		1 ³		
47	1436b	PI ³		1 ⁸		
48	1440b	B ¹⁵				
49	P190	B ¹⁸				4 ^{4,7,20,21}
50	P217	B ⁸				3 ^{8,9,16}

B: blossom; Hs: honey sac; PI: pollen loads; N: nectar.

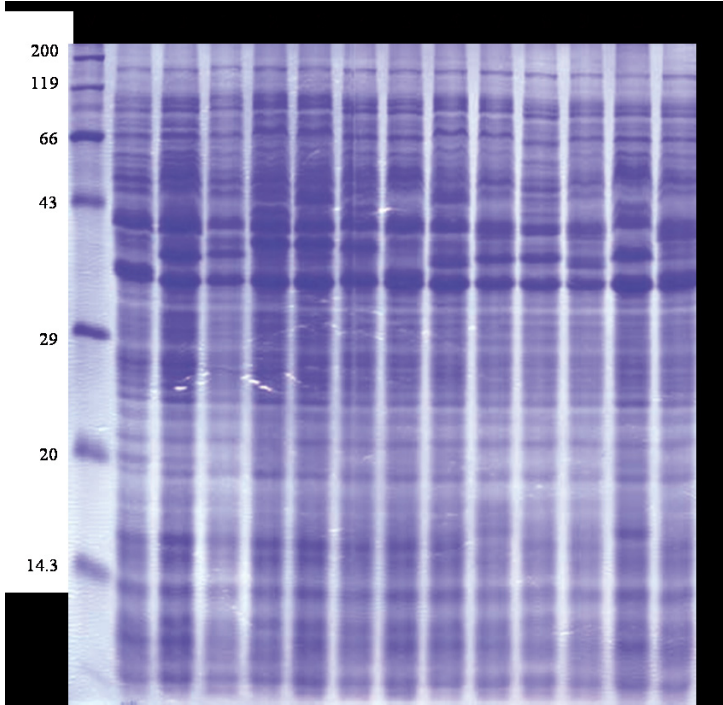
Superscript indicates plant or pollen origin, number indicates frequency of isolation.

Plants collected or their pollen was found in samples: ¹ *Agrimonia procera*, ² *Atropa belladonna nigra*, ³ *Carduus crispus*, ⁴ *Cichorium intybus*, ⁵ *Cirsium arvense*, ⁶ *Cuscuta europaea*, ⁷ *Dorycnium germanicum*, ⁸ *Echium vulgare*, ⁹ *Fagopyrum esculentum*, ¹⁰ *Helianthemum nummularium*, ¹¹ *Hemerocallis* sp., ¹² *Origanum vulgare*, ¹³ *Papaver rhoeas*, ¹⁴ *Parthenocissus triscupidata*, ¹⁵ *Phacelia tanacetifolia*, ¹⁶ *Picris hieracioides*, ¹⁷ *Plantago* sp., ¹⁸ *Solidago virgaurea*, ¹⁹ *Taraxacum officinale* group, ²⁰ *Tilia* sp., ²¹ *Trifolium repens*.

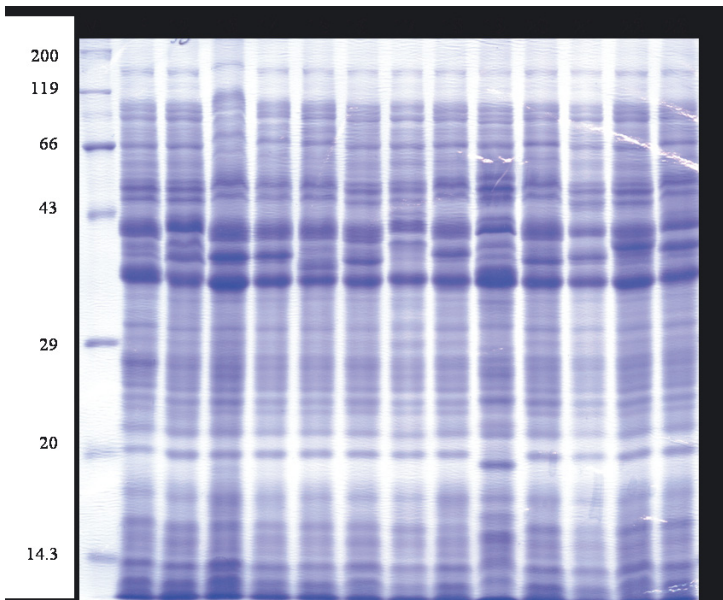
Plants which were not collected but their pollen was found in samples: ^a *Allium* sp., ^b *Artemisia santonicum*, ^c *Asparagus officinalis*, ^d *Brassica napus*, ^e *Fragaria ananassa*, ^f *Geum urbanum*, ^g *Helianthus annuus*, ^h *Odonites vulgaris*, ⁱ *Sinapis arvensis*.

^{nk} not identified; ^{np} no pollen; ^{npa} pollen analysis was not performed; * protein profile, RAPD profil.

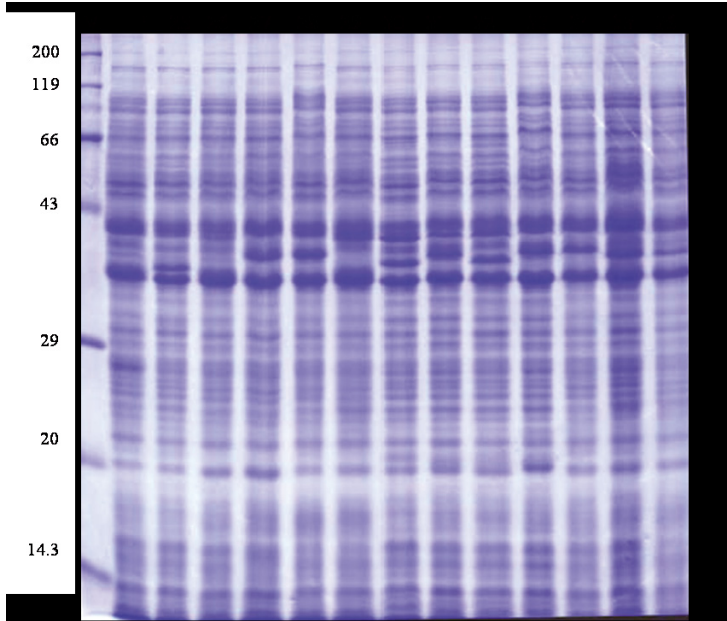
Appendix IIIa–e. SDS-PAGE of the whole-cell proteins of the *Pantoea agglomerans* strains. Lane M, molecular weight marker, numbers indicate molecular weight in kD, lane P, DSM 3493^T (a–e), lane D DSM 1619 (a). Appendix IIIa, b after Lane M and P numbers indicate strains listed in Appendix II (supplementary data).



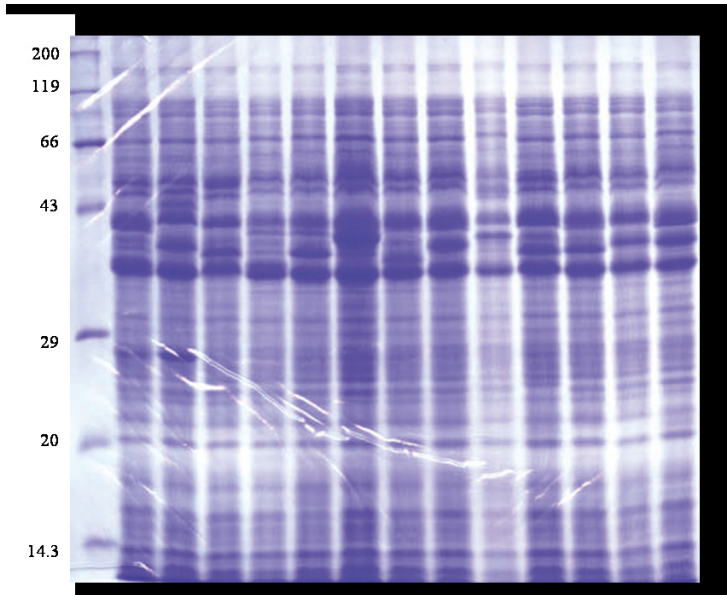
a



b

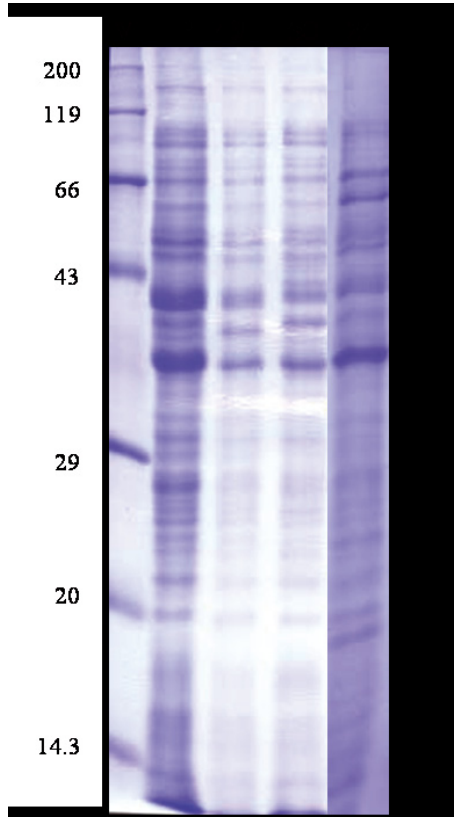


c



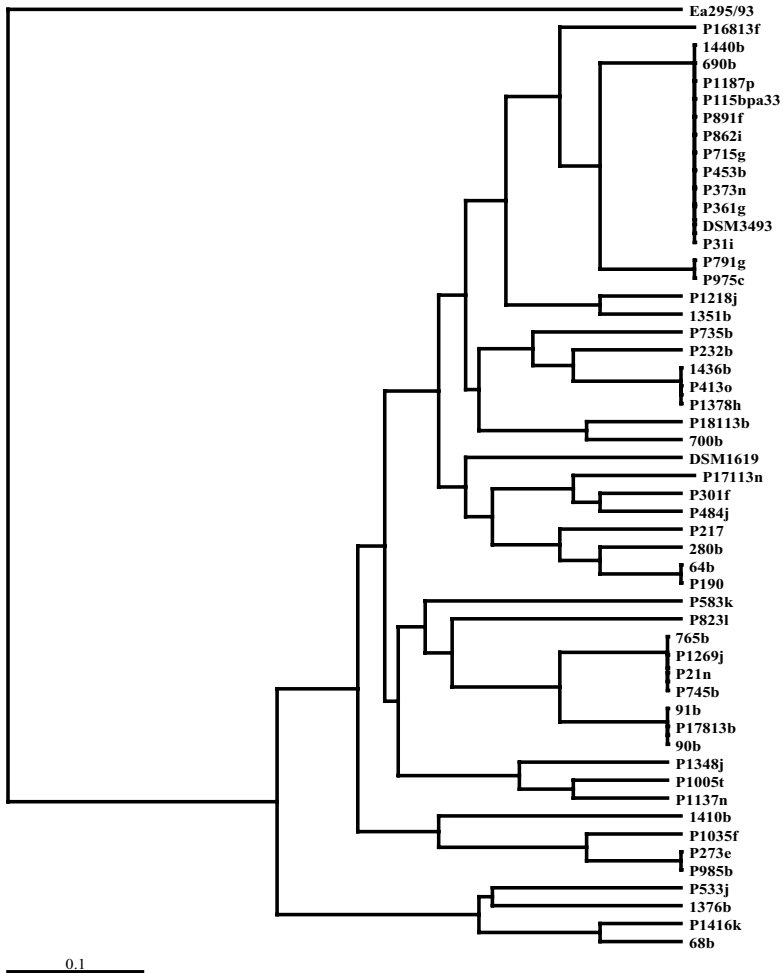
d

Appendix IIIa-e. Continued.



e

Appendix IIIa–e. Continued.



Appendix IV. UPGMA dendrogram derived by comparing the ERIC-PCR patterns obtained after ERIC-PCR from 52 *P. agglomerans* strains and one strain of *E. amylovora* 295/93. The scale indicates genetic distance measurements as calculated by UPGMA method.