

Zusätzliche Informationen zur Dissertation von Yvonne Gäbler, 2007

**Tabelle 7-47: Liste von Genen, deren Expression durch Pep13-Behandlung nach 4 h induziert wird.**  
*Macroarray*-Analyse in Petersilie.

Index	Klon	Organismus	AGI Code	Protein/Mögliche Funktion	Pep13		
					180204	080304	220304
3209	PC107-1-4	Arabidopsis thaliana	At5g05340	peroxidase (see 108-1-2)	3,67	5,63	17,24
851	PC116-3	Arabidopsis thaliana	At5g05340	peroxidase (see 107-1-4)	3,09	5,49	10,66
2236	PC137-1	Arabidopsis thaliana	At5g05340	peroxidase (see 107-1-4)	3,78	10,95	25,76
1852	PC137-3	Arabidopsis thaliana	At5g05340	peroxidase (see 107-1-4)	3,35	5,89	11,01
4181	PC137-3	Arabidopsis thaliana	At5g05340	peroxidase (see 107-1-4)	2,32	4,05	5,03
1660	PC137-4	Arabidopsis thaliana	At5g05340	peroxidase (see 107-1-4)	3,15	8,69	17,01
3989	PC137-4	Arabidopsis thaliana	At5g05340	peroxidase (see 107-1-4)	2,02	5,85	3,81
2242	PC145-1	Arabidopsis thaliana	At5g05340	peroxidase (see 107-1-4)	3,78	8,32	20,67
1666	PC145-4	Arabidopsis thaliana	At5g06060	peroxidase (see 107-1-4)	2,98	4,57	8,24
910	PC178-4	Petroselinum crispum		unnamed protein product (see 177-4)	2,46	3,47	3,99
4516	PC240-2	Petroselinum crispum		unnamed protein product (see 690-1)	2,05	6,64	8,73
4138	PC254-3	Danio rerio		PREDICTED: similar to zinc finger protein 91 (HPF7, HTF10)	2,99	4,4	3,56
682	PC258-2	Petroselinum crispum		Tyr-DC (Eli 5) (see 312-1)	2,23	2,55	2,73
3280	PC269-3	-		/	2,11	2,96	8,47
2806	PC277-2	Daucus carota		EDGP precursor (see 566-2)	2,12	3,62	2,43
3286	PC280-3	Petroselinum crispum		Tyr-DC (Eli 5) (see 312-1)	2,88	2,54	2,65
2902	PC283-1	-		/	4,39	6,4	11,57
2710	PC283-2	Arabidopsis thaliana	At3g26040	acyltransferase-like protein (see 774-3)	4,5	100	14,25
4066	PC295-3	-		/	7,29	3,64	2,95
3508	PC317-2	-		no significant similarity found	3,45	2,18	3,4
400	PC372-4	Nicotiana tabacum		peroxidase (see 107-1-4)	3,64	9,88	18,35
2312	PC399-4	Nicotiana tabacum		patatin-like protein I or 3, (see 396-3)	3,29	2,44	9,18
2510	PC405-1-3	-		/	2,7	6,55	10,52
3104	PC431-4	Arabidopsis thaliana	At1g19180	unknown protein	3,04	2,45	4,76
2462	PC478-4	Glycine max		homogentisate phytylprenyltransferase	2,64	2,31	3,13
3140	PC488-4	Arabidopsis thaliana	At1g22360	AT2 (see 362-1)	2,66	2,16	3,34
3932	PC521-1-4	Nicotiana tabacum		patatin-like protein 3 (see 399)	3,76	3	2,31
3961	PC575-4	Lycopersicon esculentum		Pirin	2,44	2,66	4,65

2875	PC667-2	-		/		7,72	2,18	13,86
2994	PC690-1	Petroselinum crispum		unnamed protein product (see 240-2)		2,1	2,61	5,23
2802	PC690-2	Petroselinum crispum		Tyr-DC (see 312-1)		2,04	2,09	2,7
3930	PC760-2-4	Daucus carota		cell attachment protein in somatic embryogenesis (see 758-2)		2,08	2,22	4,49
1643	PC774-4	Arabidopsis thaliana	At4g04410	putative polyprotein		2,55	2,39	3,32
1169	PC785-2	Lactuca sativa		carbohydrate oxidase		2,68	2,19	2,93

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**Tabelle 7-48: Liste von Genen, deren Expression durch HrpZ<sub>PspH</sub>-Behandlung nach 4 h induziert wird.**  
*Macroarray* -Analyse in Petersilie.

Index	Klon	Organismus	AGI Code	Protein/Mögliche Funktion	HrpZ <sub>PspH</sub>		
					121103	080304	220304
3869	PC070-4	Arabidopsis thaliana	At5g61790	calcium ion binding	2,23	2,1	2,69
2291	PC086-4	Arabidopsis thaliana	At1g23740	oxidoreductase/ zinc ion binding (see 162-2)	3,02	2,23	3,94
1241	PC162-2	Arabidopsis thaliana	At1g23740	oxidoreductase/ zinc ion binding (see 161-4)	2,86	2,33	3,7
1288	PC163-2	Arabidopsis thaliana	At1g23740	oxidoreductase/ zinc ion binding (see 162-2)	2,46	2,02	3,17
520	PC164-1-2	Arabidopsis thaliana	At1g23740	oxidoreductase/ zinc ion binding (see 162-2)	2,55	2,05	3,89
328	PC164-1-3	Arabidopsis thaliana	At1g23740	oxidoreductase/ zinc ion binding (see 162-2)	2,15	2,36	3,6
136	PC164-1-4	Arabidopsis thaliana	At1g23740	oxidoreductase/ zinc ion binding (see 162-2)	2,12	2,76	4,49
281	PC168-3	Arabidopsis thaliana	At1g23740	oxidoreductase/ zinc ion binding (see 162-2)	2,46	2,37	3,69
46	PC187-4	-		no significant similarity found	2,05	2,11	5,18
3059	PC216-1	Arabidopsis thaliana	At2g15220	unknown protein (see 177-1)	2,12	2,95	11,33
4409	PC222-2	Arabidopsis thaliana	At2g15220	unknown protein (see 216-1)	2,55	100	20,01
3856	PC268-4	-		/	2,99	2,52	4,24
3880	PC301-4	Daucus carota		cell attachment protein in somatic embryogenesis (see 372-3)	2,04	2,37	3,93
1744	PC360-3	Arabidopsis thaliana	At3g61980	serine-type endopeptidase inhibitor (see 318-1)	2,49	2,1	2,55
3832	PC366-1	Arabidopsis thaliana	At2g34790	electron carrier	2,26	2,52	17,1
1168	PC370-1	-		no significant similarity found	2,2	2,52	3,62
2906	PC417-1	Arabidopsis thaliana	At1g23740	oxidoreductase/ zinc ion binding (see 210-3)	2,12	2,12	3,33
2558	PC483-3	Petroselinum crispum		anionic peroxidase (see 261-1) (BUT no taq)	2,58	2,23	2,08
2366	PC483-4	Nicotiana tabacum		S-adenosylmethionine decarboxylase (BUT no apo)	2,58	2,28	3,96
4580	PC484-1	-		/	2,09	6,06	4,57
2948	PC489-1	Arabidopsis thaliana	At1g22360	AT2 (see 488-4)	2,69	2,38	3,14
2756	PC489-2	Daucus carota		EP4b	2,08	2,37	2,32
2605	PC547-1-3	Pyrus communis		isoflavone reductase related protein	2,96	2,55	2,19
3571	PC566-2	Daucus carota		EDGP precursor (see 763-4)	2,16	2,72	2,59
4063	PC584-3	Petroselinum crispum		anionic peroxidase (see 113-2)	2,34	2,5	2,43
2653	PC624-3	-		/	2,44	2,06	3,16
2659	PC632-3	Petroselinum crispum		Pathogenesis-related protein A (PR1-1)	2,52	2,73	2,77

2773	PC665-1-2	Linum usitatissimum		peroxidase (see 722-2)	3,86	3,31	3,23
3259	PC666-4	Linum usitatissimum		peroxidase (see 722-2)	3,87	3,11	4,01
4140	PC680-3	-		no significant similarity found	2,07	2,34	3,07
3186	PC689-4	Asparagus officinalis		cytochrome P450	4,29	2,51	4,13
3300	PC713-2-3	-		/	9,45	2,58	6,47
4362	PC716-2	Arabidopsis thaliana	At3g13610	oxidoreductase (see 685-1)	5,88	2,43	4,51
4170	PC716-3	Arabidopsis thaliana	At3g13610	oxidoreductase (see 685-1)	5,88	2,62	6,84
2346	PC721-4	Linum usitatissimum		peroxidase (see 672-3)	4,12	3,31	3,23
4278	PC739-2	Linum usitatissimum		peroxidase (see 722-2)	7,09	2,72	3,63
4086	PC739-3	Linum usitatissimum		peroxidase (see 722-2)	6,88	3,73	3,72
3894	PC739-4	-		/	3,44	3,23	3,19
2970	PC761-2-1	Daucus carota		cell attachment protein in somatic embryogenesis (see 758-2)	3,53	2,31	4,62
1835	PC774-3	Arabidopsis thaliana	At3g26040	acyltransferase-like protein (see 283-2)	2,82	2,36	4,78
1193	PC812-2	Nicotiana tabacum		PDR-like ABC-transporter (similar to 523-3)	2,86	3,39	4,93
1307	PC827-2	-		/	2,53	2,33	7,19
1409	PC837-1	Petroselinum crispum		omega-6 fatty acid desaturase (EC 1.14.99.-) (see 355-1)	2,24	2,81	3,47

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**Tabelle 7-49: Liste von Genen, deren Expression durch LPS-Behandlung nach 4 h induziert wird.**  
*Macroarray*-Analyse in Petersilie.

Index	Klon	Organismus	AGI Code	Protein/Mögliche Funktion	LPS		
					121103	080304	220304
4331	PC013-2	-		no significant similarity found	2,54	2,25	8,16
2705	PC055-2	Arabidopsis thaliana	At5g44100	ATP binding / casein kinase I	2,3	4,94	2,57
3533	PC208-2	-		no significant similarity found	2,89	2,05	3,56
1066	PC255-2-4	Papaver somniferum		tyrosin/dopa decarboxylase	2,15	2,24	2,25
4048	PC268-3	Arabidopsis thaliana	At1g44170	ALDH3H1	2,01	100	3,61
3752	PC390-1	Daucus carota		(S)-reticuline oxidase-like protein (see 397-1)	2,06	3,02	6,72
4064	PC427-3	-		no significant similarity found	2,16	2,03	2,98
4382	PC476-2	-		no significant similarity found	2,39	2,29	2,35
4039	PC540-3	-		no significant similarity found	2,05	100	3,47
3037	PC624-1	Oryza sativa		putative receptor-like protein kinase	2,37	3,45	5,19
3067	PC667-1	Medicago truncatula		E-class P450, group I (BUT no taq)	2,08	2,62	4,07
3354	PC761-1-3	Daucus carota		cell attachment protein in somatic embryogenesis (see 758-2)	3,57	2,51	5,53
3456	PC763-3	Arabidopsis thaliana	At4g28570	electron carrier	2,44	2,26	4,5
335	PC816-3	Arabidopsis thaliana	At1g13360	expressed protein	2,2	2,24	6,47

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**Tabelle 7-50: Liste von Genen, deren Expression durch Pep13- und HrpZ<sub>PspH</sub>-Behandlung nach 4 h induziert wird.**  
*Macroarray* -Analyse in Petersilie.

Index	Klon	Organismus	AGI Code	Protein/Mögliche Funktion	Pep13			HrpZ <sub>PspH</sub>		
					180204	080304	220304	121103	080304	220304
2339	PC100-1-4	-		no significant similarity found	7,04	2,8	5,86	6,48	3,3	5,53
4265	PC108-1-2	Arabidopsis thaliana	At5g05340	peroxidase (see 107-1-4)	4,86	8,54	18,59	2,34	7,71	11,05
3215	PC121-4	Petroselinum crispum		fatty acid desaturase/hydroxylase-like protein ELI7.1 (see 256-4)	3,75	7,28	16,77	3,72	4,08	12,18
479	PC202-2	Arabidopsis thaliana	At1g01800	oxidoreductase	2,7	2,63	3,92	2,07	2,32	2,83
2771	PC220-2	Arabidopsis thaliana	At2g15220	unknown protein (see 216-1)	3,02	2,95	8,82	3,45	2,32	10,41
4217	PC222-3	Arabidopsis thaliana	At2g15220	unknown protein (see 216-1)	2,53	4,56	27,2	2,36	2,13	28,34
4025	PC222-4	-		/	3,09	3,84	24,74	2,8	2,63	22,65
3724	PC355-1	Petroselinum crispum		omega-6 fatty acid desaturase (EC 1.14.99.-) (see 355-3)	3,78	4,18	5,65	4,4	2,54	4,47
3448	PC366-3	Arabidopsis thaliana	At2g34790	electron carrier (beste Sequenz)	2,33	2,86	6,6	2,84	3,29	7,81
3256	PC366-4	Arabidopsis thaliana	At2g34790	electron carrier	3,44	3,73	6,03	3,85	4,44	7,6
976	PC370-2	-		no significant similarity found	2,91	2,27	4,69	3,26	2,74	4,61
2696	PC399-2	Nicotiana tabacum		patatin-like protein I or 3, (see 396-3)	3,56	3,18	8,94	4,28	4,81	6,77
4562	PC458-1	-		no significant similarity found	2,82	2,89	3,76	2,64	2,64	2,72
4508	PC521-1-1	Nicotiana tabacum		patatin-like protein 3 (see 399)	3,81	2,84	10,86	2,5	2,42	9,81
3740	PC521-2-1	Nicotiana tabacum		patatin-like protein 3 (see 399)	3,31	2,89	7,88	2,85	3,07	7,05
3548	PC521-2-2	Nicotiana tabacum		patatin-like protein 3 (see 399)	3,88	2,16	8,55	4,46	2,18	6,01
3356	PC521-2-3	Nicotiana tabacum		patatin-like protein 3 (see 399)	3,86	4,01	14,82	4,59	4,08	13,55
3164	PC521-2-4	Nicotiana tabacum		patatin-like protein 3 (see 399)	3,48	2,13	8,28	3,48	2,17	5,03
2972	PC522-1-1	Nicotiana tabacum		patatin-like protein 3 (see 399)	5,64	2,64	10,36	4,88	2,85	8,58
2780	PC522-1-2	Nicotiana tabacum		patatin-like protein 3 (see 399)	4,74	2,49	7,42	3,53	2,29	7,02
2588	PC522-1-3	Nicotiana tabacum		patatin-like protein 3 (see 399)	3,79	3,18	13,14	3,78	2,67	11,18
4392	PC531-4	Nicotiana tabacum		patatin-like protein 3 (see 521-1-1)	4,78	3,41	8,16	2,87	2,94	7,51
2545	PC610-3	Lotus corniculatus var. Japonicus		Nod-factor receptor 1a	2,06	2,06	3,08	3	2,46	3,43
4219	PC665-2-3	Vigna angularis		peroxidase, cationic (ähnlich 665-1-2)	4,38	2,07	7,29	4,01	3,48	6,56
2418	PC690-4	Linum usitatissimum		peroxidase (see 672-3)	5,24	2,11	2,94	7,32	3,53	3,12
3684	PC713-2-1	Arabidopsis thaliana	At3g13610	oxidoreductase (see 685-1)	2,7	2,06	6,31	2,09	3,23	8,75
4554	PC716-1	Arabidopsis thaliana	At3g13610	oxidoreductase (see 685-1)	4,54	2,41	6,92	8,34	3,12	7,7

4560	PC722-1	Linum usitatissimum	peroxidase (see 672-3)	4,47	2,25	2,88	7,04	4,07	4,1
4470	PC739-1	Linum usitatissimum	peroxidase (see 722-2)	3,39	2,75	4	5,41	4,46	4,04
1979	PC829-2	-	/	4,24	3,62	8,4	4,64	4,21	5,67
1595	PC829-4	Nicotiana tabacum	patatin-like protein 3 (see 521-1-1)	2,92	2,82	9,53	3,39	2,94	9,15

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**Tabelle 7-51: Liste von Genen, deren Expression durch Pep13- und LPS-Behandlung nach 4 h induziert wird.**  
*Macroarray*-Analyse in Petersilie.

Index	Klon	Organismus	AGI Code	Protein/Mögliche Funktion	Pep13			LPS		
					180204	080304	220304	121103	080304	220304
4343	PC056-2	-		no significant similarity found	3,87	2,49	2,29	2,85	2,17	4,91
3923	PC217-4	Petroselinum crispum		Tyrosin-Decarboxylase (see 312-1)	2,71	5,28	9,71	2,29	3,06	3,81



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**Tabelle 7-52: Liste von Genen, deren Expression durch HrpZ<sub>PspH</sub>- und LPS-Behandlung nach 4 h induziert wird.**  
*Macroarray*-Analyse in Petersilie.

Index	Klon	Organismus	AGI Code	Protein/Mögliche Funktion	HrpZ <sub>PspH</sub>			LPS		
					121103	080304	220304	121103	080304	220304
3640	PC366-2	Arabidopsis thaliana	At2g34790	electron carrier	2,38	2,99	5,53	3,62	2,08	6,32
4340	PC405-2-2	Arabidopsis thaliana	At4g33850	catalytic	3,1	5,72	6,68	2,99	4,22	5,24
2420	PC407-4	-		/	2,34	2,49	2,71	2,17	3,85	3,06
4153	PC575-3	Solanum tuberosum		probable epoxide hydrolase (EC 3.3.2.3)	2,27	6,35	3,18	2,02	6	2,99
4332	PC680-2	-		no significant similarity found	2,08	2,25	2,05	2,03	2,1	3,79
2874	PC760-1-2	Daucus carota		cell attachment protein in somatic embryogenesis (see 758-2)	3	2,91	4,25	3,28	2,46	5,23
4506	PC760-2-1	Daucus carota		cell attachment protein in somatic embryogenesis (see 758-2)	2,16	2,81	10,97	2,64	2,4	10,55

**Tabelle 7-53: Liste von Genen, deren Expression durch Pep13-, HrpZ<sub>PspH</sub>- und LPS-Behandlung nach 4 h induziert wird.**  
Macroarray-Analyse in Petersilie.

Index	Klon	Organismus	AGI Code	Protein/Mögliche Funktion	Pep13			HrpZ <sub>PspH</sub>			LPS		
					180204	080304	220304	121103	080304	220304	121103	080304	220304
3881	PC108-1-4	Arabidopsis thaliana	At5g05340	peroxidase (see 107-1-4)	2,72	11,61	21,33	2,52	6,85	7,63	2,17	8,23	14,99
3791	PC121-1	Petroselinum crispum		fatty acid desaturase/hydroxylase-like protein ELI7.1 (see 256-4)	4,19	20,13	30,57	3,69	12,06	28,09	2,66	5,69	28,08
4463	PC126-1	Arabidopsis thaliana	At5g05340	peroxidase (see 107-1-4)	4,29	4,94	12,85	2,42	3,88	9,51	2,05	4,27	12,09
4589	PC190-1	Arabidopsis thaliana	At2g15220	unknown protein (see 216-1)	3,35	5,92	8,48	3,24	2,58	8,43	3,64	2,53	5,03
3821	PC191-1	Arabidopsis thaliana	At2g15220	unknown protein (see 216-1)	2,27	5,96	8,6	2,68	3,22	10,74	2,42	2,07	7,18
4115	PC217-3	Arabidopsis thaliana	At2g15220	unknown protein (see 216-1)	3,21	6,26	16,43	2,92	2,51	17,98	3,49	2,03	14,78
2963	PC220-1	Arabidopsis thaliana	At2g15220	unknown protein (see 216-1)	2,34	4,19	10,58	2,92	2,75	11,87	2,04	2,35	9,86
2410	PC256-4	Petroselinum crispum		fatty acid desaturase/hydroxylase-like protein ELI7.1 (see 289-3)	9,59	15,48	18,87	8,75	6,6	12,41	5,2	5,86	15,36
4432	PC268-1	-		no significant similarity found (BUT no apo)	2,82	4,15	14,39	2,24	3,31	14,11	2,39	3	17,78
3604	PC314-2	Petroselinum crispum		Eli 7.5 (see 352-3)	5,74	21,77	16,91	3,27	8,94	12,79	3,62	7,16	13,57
1840	PC326-3	Nicotiana tabacum		cytochrome P450 (see 454-3)	2,19	3,27	2,99	2,26	2,97	2,31	2,04	3,15	3,02
2668	PC352-3	Petroselinum crispum		Eli 7.5 (see 314-2)	5	19,64	25,04	4,74	6,95	18,79	3,23	7,03	14,73
3340	PC355-3	Petroselinum crispum		omega-6 fatty acid desaturase (EC 1.14.99.-) (see 355-1)	3,28	8,42	5,86	3,61	4,42	4,77	4,27	2,45	4
3730	PC362-1	Arabidopsis thaliana	At1g22360	AT2 (see 489-1)	2,18	3,32	11,53	2,33	2,98	8,35	2,51	2,01	8,61
592	PC372-3	Daucus carota		cell attachment protein in somatic embryogenesis (see 375-1)	2,43	3,58	4,51	3,27	3,87	4,73	2,61	3,12	4,76
2230	PC375-1	Daucus carota		cell attachment protein in somatic embryogenesis (see 638-3)	2,47	3,38	5,02	3,58	3,78	5,25	3,42	2,79	5,53
2038	PC375-2	Daucus carota		cell attachment protein in somatic embryogenesis (see 638-3)	2,39	3,82	4,79	3,32	3,52	5,06	3,21	3,01	6,44
1846	PC375-4	Daucus carota		cell attachment protein in somatic embryogenesis (see 638-3)	2,04	3,93	4,34	3,14	4,08	5,12	2,91	2,9	5,46
2318	PC405-1-4	-		/	2,92	2,32	4,53	3,06	2,65	4,9	2,67	2,5	4,67
4070	PC445-3	-		no significant similarity found	7,13	100	10,99	4,48	100	10,62	3,02	100	11,38
3649	PC529-1	Nicotiana tabacum		EIG-I24	2,45	4,6	7,32	3,64	4,5	7,54	3,41	3,21	9,04
4584	PC531-3	Daucus carota		elicitor induced glycoprotein iEP4	3,46	3,81	6,58	3,75	4,61	6,87	4,47	2,4	5,41
2815	PC583-2	-		no significant similarity found	2,45	3,1	9,38	2,52	2,66	7,3	2,83	2,17	7,76
4273	PC608-2	Daucus carota		probable cell-wall-bound EP4 protein, 45K, precursor	3,42	3,54	4,72	3,55	3,8	3,09	3,25	2,8	4,08
4375	PC611-2	Arabidopsis thaliana	At3g17840	RLK902 (see 492-1)	2,75	2,59	3,9	2,36	2,44	2,43	2,72	2,34	2,45
4201	PC638-3	Daucus carota		cell attachment protein in somatic embryogenesis (BUT no taq)	2,42	3,25	6,27	3,08	3,44	5,51	3,45	2,57	4,76
3654	PC677-1	Paracoccus denitrificans PD1222		conserved hypothetical protein (see 684-1)	3,72	4,53	7,31	5,02	6,12	6,99	5,67	2,87	12,63
3492	PC713-2-2	Arabidopsis thaliana	At3g13610	oxidoreductase (see 685-1)	2,22	3,35	5,42	2,89	4,55	5,16	3,56	2,04	6,69
3978	PC716-4	Arabidopsis thaliana	At3g13610	oxidoreductase (see 685-1)	4,35	2,96	7,05	5,34	3,54	8,53	5,45	2,12	13,9
3450	PC759-3	Daucus carota		cell attachment protein in somatic embryogenesis (see 758-2)	2,24	2,26	4,03	2,45	3,23	4,36	3,17	3,08	4,85
3066	PC760-1-1	Daucus carota		cell attachment protein in somatic embryogenesis (see 758-2)	2,61	2,19	5,01	2,72	3,12	5,62	3,85	2,75	6,71
4314	PC760-2-2	Daucus carota		cell attachment protein in somatic embryogenesis (see 758-2)	2,24	3,48	5,31	2,41	3,39	6,07	3,28	2,38	6,08
3546	PC761-1-2	Daucus carota		cell attachment protein in somatic embryogenesis (see 758-2)	2,09	2,15	4,72	2,42	2,39	4,89	3,19	2,14	6,68

293	PC770-3	-	/	3,84	9,2	6,19	2,33	3,39	8,13	2,66	3,38	5,89
1067	PC776-3	Petroselinum crispum	PR1-1	2,41	2,32	2,87	2,61	3,08	2,39	2,32	2,4	2,78
2177	PC836-1	Petroselinum crispum	fatty acid desaturase/hydroxylase-like protein Eli7.1 (see 256-4)	5,98	20,39	22,02	4,95	10,22	16,67	3,79	7,36	11,32
1601	PC836-4	Petroselinum crispum	fatty acid desaturase/hydroxylase-like protein Eli7.1 (see 256-4)	3,33	6,35	13,29	2,72	3,12	11,98	2,52	2,29	9,9
1511	daphs2-3	Petroselinum crispum	daphs2-3	5,18	4,44	9,35	4,08	2,89	8,13	3,64	2,33	6,3
1319	daphs2-5	Petroselinum crispum	daphs2-5	3,32	6,29	6,62	2,51	4,57	5,9	2,44	3,13	5,25
1127	daphs2-6	Petroselinum crispum	daphs2-6	3,23	4,35	4,99	2,48	3,22	4,1	2,28	2,01	3,52