The polygalacturonase gene *BcMF2* from *Brassica campestris* is associated with intine development

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Supplemental Material

Supplementary Table 1 Polygalacturonase amino acid sequences used in the molecular phylogenetic analysis.

Abbreviation	Species (gene name, expression)	SwissProt No.	GenBank No.
BcMF2	*Brassica campestris*	—	EU181170
Endopolygalacturonase			
Actde.pg	*Actinidia deliciosa* (leaf & bud)	P35336	L12019
Actde2.pg	*Actinidia deliciosa* (fruit)	—	AAF71160
Arath11.pg	*Arabidopsis thaliana* (root & seedling)	Q38958	X98130
Arath12.pg	*Arabidopsis thaliana*	O22699	AC002292
Brana1.pg	*Brassica napus* (*PG35-8*) (pod dehiscence)	Q42399	X95800
Brana2.pg	*Brassica napus* (*SAC66*) (pod development)	Q42636	Z49971
Brana3.pg	*Brassica napus* (*PGAZ*) (leaf abscission)	—	AJ250919
Cucme1.pg	*Cucumis melo* (fruit ripening)	O81244	AF062465
Cucme2.pg	*Cucumis melo* (*MPG 2*) (fruit ripening)	O81245	AF062466
Cucme3.pg	*Cucumis melo* (fruit ripening)	O81246	AF062467
Cucsa.pg	*Cucumis sativus* (stress induced)	—	AB035890
Glyma1.pg	*Glycine max* (roots, wound response)	Q9SWS3	AF128266
LycesA.pg	*Lycopersicon esculentum* (*PG2A*, fruit)	P05117	X04583
Lyces1.pg	*Lycopersicon esculentum* (*TAPG1*) (stigma, style)	O22311	AF001000
Lyces2.pg	*Lycopersicon esculentum* (*TAPG2*) (stigma, style)	Q96487	AF001001
Lyces3.pg	*Lycopersicon esculentum* (*TAPG3*) (abscission)	O22310	AF000999
Lyces4.pg	*Lycopersicon esculentum* (*TAPG4*) (stigma, style)	Q96488	U70481
Lyces5.pg	*Lycopersicon esculentum* (*TAPG5*) (abscission)	O22313	AF001003
Lyces6.pg	*Lycopersicon esculentum* (*TAPG6*)	O22610	AF029230

Lyces7.pg	*Lycopersicon esculentum* (stigma, style)	—	AF072732
Lyces8.pg	*Lycopersicon esculentum* (wound response)	—	AF118567
Lyces9.pg	*Lycopersicon esculentum* (seed)	—	AF138858
Maldo.pg	*Malus domestica* (fruit)	P48978	L27743
Orysa.pg	*Oryza sativa*	—	AP003140
Peram.pg	*Persea americana* (fruit)	Q02096	X66426
Pissa.pg	*Pisum sativum*	—	AF361321
Prupe1.pg	*Prunus persica* (*PRF5*)(fruit)	P48979	X76735
Prupe3.pg	*Prunus persica*	Q43063	X77231
Rubid.pg	*Rubus idaeus* (fruit)	O65886	AJ224147
TsPG	*Turnera subulata* (short style)	—	AY185765
Vitvi.pg	*Vitus vinifera* (fruit)	—	AY043233
Exopolygalacturonase			
Arath1.epg	*Arabidopsis thaliana* (flower)	P49063	X72292
Arath2.epg	*Arabidopsis thaliana* (*PGA2*) (flower)	O65401	X73222
Pollen Polygalacturonase			
Goshi.pp	*Gossypium hirsutum* (*G9*) (pollen)	Q39786	U09717
Medsa.pp	*Medicago sativa* (*P73*)(pollen)	Q40312	U20431
Nicta.pp	*Nicotiana tabacum* (*Npg1*) (pollen)	Q05967	X71020
Nicta2.pp	*Nicotiana tabacum* (sperm cell)	—	AF248538
Salgi2.pp	*Salix gilgiana* (male flower)	Q9MBB9	AB029458
TsPP	*Turnera subulata* (pollen)	—	AY185764
Brana.pep	*Brassica napus* (*Sta 44-4*)(pollen)	P35337	L19879
Oenor.pep	*Oenothera organensis* (pollen)	P24548	—
Phlpr.pep	*Phleum pratense* (pollen)	Q9XG86	AJ238848
Zeama1.pep	*Zea mays* (pollen)	P26216	X57627
Braol.pp	*Brassica oleracea* (*PG1*) (pollen)	—	AF518563

Arath1.pp	*Arabidopsis thaliana* (*PGA3*) (pollen)	—	AY096534
Arath2.pp	*Arabidopsis thaliana* (*PGA4*) (pollen)	—	NM_100158

Note: The terminations of abbreviations for the individual enzyme specificities following Markovič and Janeček (2001) are, pg (endopolygalacturonase), epg (exopolygalacturonase) pp (pollen polygalacturonase) pep (pollen exopolygalacturonase).