

## Marker -Trait Associations Known In Rosaceae Crops

(Table and references compiled by Dr. Nnadozie Oraguzie and Dr. Richard Bell)

Species	Trait	Trait locus (QTL, major locus, or known gene sequence) LOD score, map position, % variation	Source/Population	Linked markers	Reference	MAPS and/or MASS ? <sup>a</sup>
<b>APPLE</b>						
<i>Malus</i> × <i>domestica</i>	Resistance to apple scab	Major locus: <i>Va</i> ; LG1?	Antonovka PI172623, Fortune x PRI 1841-11; NY489 x PRI 1841-11	RAPD, SCAR	Hemmat et al., 2003	
<i>Malus</i> × <i>domestica</i>	Resistance to apple scab	Major locus: <i>Vb</i> ; LG1	Hansen's Baccata #2; Empiere x Hansen's Baccata #2	RAPD, SCAR	Hemmat et al., 2003	
<i>Malus</i> × <i>domestica</i>	Resistance to apple scab	Major locus: <i>Vb</i> ; LG12, 7.8	Golden Delicious x Hansens baccata#2	SSR	Erdin et al., 2006.	
<i>Malus</i> × <i>domestica</i>	Resistance to apple scab	Major locus: <i>Vd</i> ; LG 10, 2.0	Durello di Forli X Fiesta	RAPD	Tartarini et al., 2004.	
<i>Malus</i> × <i>domestica</i>	Resistance to apple scab	Major locus: <i>Vbj</i> ; LG2; 5.8	<i>Malus baccata jackii</i> ; A722-7 x Golden Delicious	SCAR	Gygax et al., 2004.	
<i>Malus</i> × <i>domestica</i>	Resistance to apple scab	Major locus: <i>Vd</i> ; LG10; 2.0	Durello di Forli	RAPD	Tartarini et al., 2004.	
<i>Malus</i> × <i>domestica</i>	Resistance to apple scab	Gene: <i>Vf</i> . LG1	Markers tested in 'Florina' x 'Nova Easygro' progeny and 55 cultivars or	SSR	Vinatzer et al. 2004	

<i>Malus</i> × <i>domestica</i>	Resistance to apple scab	Major locus: <i>Vfh</i>	selections <i>M. floribunda</i> 821		Bénaouf and Parisi, 2000.
<i>Malus</i> × <i>domestica</i>	Resistance to apple scab	Major locus: <i>Vg</i> ; LG12; 3.0	Golden Delicious; Prima X Fiesta	RFLP	Durel et al., 2000.
<i>Malus</i> × <i>domestica</i>	Resistance to apple scab	Major locus: <i>Vh8</i> ; LG2; 1.3, 4.3, 5.1 & 18.5, LODs5 & 3	<i>M. sieversii</i> W193B; Royal Gala x <i>M. Sieversii</i> W193B	SCAR & SSR	Bus et al., 2005a; New Phytologist 166:1035-1049
<i>Malus</i> × <i>domestica</i>	Resistance to apple scab	Major locus: <i>Vj</i>	Jonsib		Korban and Chen, 1992
<i>Malus</i> × <i>domestica</i>	Resistance to apple scab	Major locus: <i>Vm</i> , LG17, 0	<i>M. micromalus</i> ; Golden Delicious x Murray	SSR	Patocchi et al. 2005
<i>Malus</i> × <i>domestica</i>	Resistance to apple scab	Major locus: <i>Vh2</i> ; 2.5;	Russian apple R12740-7A; Royal Gala x TSR34T15	RAPD	Bus et al., 2005b
<i>Malus</i> × <i>domestica</i>	Resistance to apple scab	Major locus: <i>Vh2</i> ; LG2; 8.8 with SSR & 5.0 with RAPD-SCAR	Royal Gala x TSR34T15	SSR, RAPD	Bus et al., 2005b
<i>Malus</i> × <i>domestica</i>	Resistance to apple scab	Major locus: <i>Vr2</i> ; LG2, 0	Russian apple R12740-7A; GMAL 2473 x Idared	AFLP	Patocchi et al., 2004
<i>Malus</i> × <i>domestica</i>	Resistance to apple scab	Major locus: <i>Vh4</i> ; LG2; 4.0	Russian apple R12740-7A; Royal Gala x TSR33T239	SCAR	Bus et al., 2005b
<i>Malus</i> × <i>domestica</i>	Resistance to apple scab	Major locus: <i>VT57</i> ; 2.0	Sciglo x A68R03T057	SSR	Bus et al., 2005b

<i>Malus</i> <i>×domestica</i>	Resistance to apple scab	QTLs: LG 1-46.4 (isolate specific): LOD 8.2-8.9; 16.0-17.8%; LG 11-4.5: LOD 5.4-9.5; 16.5-22.8%; LG 15-20.3 (isolate specific): LOD 3.1-3.3; 5.6-6.0%; LG 17-19.1: LOD 3.6-5.9; 9.4-13.4%	Prima x Fiesta	RAPD, SSR, RFLP, SSR	Durel et al., 2003
<i>Malus</i> <i>×domestica</i>	Resistance to apple scab	Leaf scab: 6 QTLs: LG 6, 7, 10, 11, 12, 17: LOD 2.3-13.2; 4.0-23% Fruit scab: 2 QTLs: LG 15, 17: LOD 2.8-4.9; 7.0-9.0%	Fiesta x Discovery	SSRs	Liebhart et al. 2003b.
<i>Malus</i> <i>×domestica</i>	Resistance to apple scab	QTLs: LG 2 (co-locates with Vr and Vn8), 5, 12, 13, 15, 17; LG 1 (co-locates with Vf), 2; 3 major QTL for partial resistance to most isolates: LG 1, 2, 17: LOD 3.16-26.59; 5.1-51.1%, QTL for single isolate resistance, LG 5: LOD 5.4-12.57; 12.5-20.8%	Discovery x TN10-8	SSRs, AFLPs, Vg locus	Calenge et al., 2004.
<i>Malus</i> <i>×domestica</i>	Resistance to powdery mildew	Major locus: <i>Pl-1</i> ; <i>LG12</i> , 3-4	Idared x 78/18-4	CAPS	Lesemann & Dunemann, 2006.
<i>Malus</i> <i>×domestica</i>	Resistance to powdery mildew	Major locus: <i>Pl-2</i> , <i>LG11</i> ; 8.0	Royal Gala X A689-24	RAPD-SCAR	Gardiner et al., 2003.
<i>Malus</i> <i>×domestica</i>	Resistance to powdery mildew	Major locus: <i>Pl-d</i> ; <i>LG12</i> ; 8.0	Fiesta x A871-14	SSR	James et al. 2004.
<i>Malus</i> <i>×domestica</i>	Resistance to powdery mildew	Major locus: <i>Pl-m</i>	Fuji X MISop 93.051 G02-054	SCAR	Gardiner et al. 2003.
<i>Malus</i> <i>×domestica</i>	Resistance to powdery mildew	Major locus: <i>Pl-w</i> ; LG8	White Angel; Prima X Fiesta	Isozyme	Battle and Alston, 1996.

<i>Malus</i> <i>× domestica</i>	Resistance to powdery mildew	Major locus: <i>Pl-w</i> LG8;	D12 x White Angel	SRR, SCARS	Evans & James, 2003.
<i>Malus</i> <i>× domestica</i>	Resistance to powdery mildew	QTL; A679-2 map; LG 3, 5 & 16: Iduna map LG 2, 3, 5, 7, 8 & 9. % variance explained not shown	Iduna X A679-2	SSRs	Kellerhals et al., 2000.
<i>Malus</i> <i>× domestica</i>	Resistance to powdery mildew	QTL: U211 map: G2- LOD 12.0, 72.1% and LOD12.1, 71.5%; G3- LOD2.6, 37.5% and LOD 8.9; 72.4%; G4=LG12-LOD9.3, 64.9% & LOD8.8, 71.9% and LOD10.8; 72.0%. Idared map: G3=LG2- LOD3.3; 40.6% and LOD2.4; 39.4%, G5=LG15-LOD7.9; 61%	Idared x U211	AFLPs & SSR	Stankiewicz-Kosyl et al., 2005.
<i>Malus</i> <i>× domestica</i>	Resistance to powdery mildew	QTLs: LG 2: LOD 3.0-9.01; 7.4%-22.5%, LG 13: LOD 3.74-9.73; 7.5%-27.4%, LG 1: LOD 3.0, 7.4%, LG 8: LOD 5.27-8.45; 8.9%-19.5%, LG 10: LOD 3.99-4.02; 7.9%-8.3%, LG 14: LOD 3.45; 5.7%, LG 17: LOD 4.36-4.64; 8.8%-10.5%	Discovery x TN10-8	SSRs and AFLPs	Calenge and Durel, 2006.
<i>Malus</i> <i>× domestica</i>	Resistance to fireblight	QTL:26.82; LG7F-52.7; 42.6%	Prima x Fiesta Fiesta x Discovery	RAPD	Calenge et al., 2005.
<i>Malus</i> <i>× domestica</i>	Resistance to fireblight	QTL: 4.09; LG3P-53.8; 7.5%	Prima X Fiesta	SSR	Calenge et al., 2005.
<i>Malus</i> <i>× domestica</i>	Resistance to fireblight	QTL: 3.37; LG3F-0.0; 4.4-4.9%(depending on days of postinfection	Fiesta x Discovery	AFLP	Calenge et al., 2005.
<i>Malus</i> <i>× domestica</i>	Resistance to fireblight	QTL:3.53; LGD12-62.3; 5.4%	Fiesta x Discovery	SSR	Calenge et al., 2005.
<i>Malus</i> <i>× domestica</i>	Resistance to fireblight	QTL; 4.87; LG13D-10.2; 7.9%	Fiesta x Discovery	AFLP	Calenge et al., 2005

<i>Malus</i> × <i>domestica</i>	Resistance to fireblight	QTL: 7.5-8.1; LG7F-46.5-51.5	Fiesta x Discovery	AFLP	Khan et al., 2006
<i>Malus</i> × <i>domestica</i>	Resistance to fireblight	QTL: 9.54; LG3R-42.6	IdaRed X <i>M</i> × <i>.robusta</i>	SSR	Peil et al., 2007
<i>Malus</i> × <i>domestica</i>	Resistance to fireblight	QTL: 14.15; LG3R-48.2; 80% jointly with CH03e03	IdaRed X <i>M</i> × <i>.robusta</i>	SSR	Peil et al., 2007
<i>Malus</i> × <i>domestica</i>	Resistance to fireblight	QTL: 22.96; LG3R-58.9; 80% jointly with CH03g07	IdaRed X <i>M</i> × <i>.robusta</i>	SSR	Peil et al., 2007
<i>Malus</i> × <i>domestica</i>	Resistance to fireblight	QTL: LG7-46	Milwa x 1217	SCAR	Khan et al., 2007
<i>Malus</i> × <i>domestica</i>	Resistance to fireblight	QTL: LG7F-56	Milwa x 1217	SCAR	Khan et al., 2007
<i>Malus</i> × <i>domestica</i>	Resistance to fireblight	QTL: LG7F-46	Milwa x 1217	SSR	Khan et al., 2007.
<i>Malus</i> × <i>domestica</i>	Resistance to wooly apple aphid	Major locus: <i>Er-1</i> ; <i>LG8</i>	Discovery X TN10-8	SNP	Chagne et al. unpubl
<i>Malus</i> × <i>domestica</i>	Resistance to wooly apple aphid	Major locus <i>-Er-1</i> ; LG8NS-10.9	Sciglo x Northern Spy	SSR	Chagne et al. unpubl.; Bus et al., 2008.
<i>Malus</i> × <i>domestica</i> rootstock	Resistance to wooly apple aphid	Major locus: <i>Er-2</i> ; LG17R5-13.6	X3189 × <i>M.</i> × <i>robusta</i> 5	SSR	Bus et al., 2008.
<i>Malus</i> × <i>domestica</i> rootstock	Resistance to wooly apple aphid	Major locus: <i>Er-3</i> ; LG8, 12.0	M9 X Aotea	SSR	Chagne et al., unpubl.
<i>Malus</i> × <i>domestica</i>	Resistance to wooly apple aphid	Major locus: <i>Er-3</i> ; LG8A-10.9	Royal Gala × Aotea 1 F <sub>2</sub> derivative	SNP	Bus et al., 2008.
<i>Malus</i> × <i>domestica</i>	Resistance to rosy leaf	Major locus: <i>Sd-1</i>	Cox's Orange Pippin		Alston and Briggs, 1968.

<i>Malus</i> <i>×domestica</i>	curling Resistance to rosy leaf curling	Major locus: <i>Sd-1</i> , LG7, E6/M8R1 & 2B12 IcM away; and E6/M6R2, E6/M6R1	Fiesta X Discovery	BACs	Cevik & King, 2000
<i>Malus</i> <i>×domestica</i>	Resistance to rosy leaf curling	Major locus: <i>Sd-2</i> ; <i>LG7</i> ; 0	Double Red Northern Spy x Totem	SCAR, SSR	Cevik & King, 2002.
<i>Malus</i> <i>×domestica</i>	Resistance to rosy leaf curling	Major locus: <i>Sd-3</i>	<i>M. x robusta</i> OP MAL59/9		Alston and Briggs, 1977.
<i>Malus</i> <i>×domestica</i>	Resistance to rosy leaf curling	QTL: Fiesta map; LG17, locus 57.7, heritability 28.3%	Fiesta X Discovery	AFLP	Stoeckli et al., 2008.
<i>Malus</i> <i>×domestica</i>	Resistance to leaf curling midge	QTL: Fiesta LG7; locus4.5, heritability=50.2%	Fiesta X Discovery	AFLP	Stoeckli et al., 2008.
<i>Malus</i> <i>×domestica</i>	Tree habit - 7 components	QTLs: Height increment: LG 6, 7, 9, 10, 11, 12, 21 (3.9- 7.9%), Internode length: LG 5, 6, 9, 10 (4.6-23.1%), Internode no: LG 1, 5, 7, 10, 12, 21 (4.3-16.8%), Base diameter increment: LG 2, 7, 10, 14, 16, 21 (4.0-8.5%), Base diameter: LG 7, 9 (5.5-7.5%), Branch number: LG 7, 10 (7.1-24.3%), Leaf break: LG 3, 6, 7, 9, 11, 12, 15 (3.9-7.3%)	Wijcik McIntosh X NY75441-58	SSRs	Conner et al., 1998.
<i>Malus</i> <i>×domestica</i>	Tree habit in seedlings	QTLs: Stem diam: LG 2, 15, 17 (LOD 3.1-4.8; 6.0-10.0%); Leaf size: LG 9, 17 (LOD 3.0-4.2; 6.0-8.0%)	Fiesta x Discovery	SSRs	Liebhard et al., 2003a.
<i>Malus</i> <i>×domestica</i>	Tree habit in mature	QTLs: Height increment: LG 3, 5, 8, 11, 13, 17 (LOD 2.4-6.2; 5.0-11.0%)	Fiesta x Discovery	SSRs	Liebhard et al., 2003a.

	trees	Stem diam: LG 1, 2, 3, 8, 11, 13, 14, 15, 17 (LOD 1.7-6.5; 4.0-13%)			
<i>Malus</i> × <i>domestica</i>	Terminal bearing	Major locus: <i>Tb</i> , LG 6, possible QTL or masked by spurring	Rome Beauty X White Angel	RAPD	Lawson et al., 1995.
<i>Malus</i> × <i>domestica</i>	Tree habit	Many QTLs: LOD 1.0, 1.4 to 55.3% variation.	Braeburn x Telamon	AFLPs and SSRs	Kenis & Keulemans, 2004, 2007.
<i>Malus</i> × <i>domestica</i>	Tree columnar habit	Major locus : <i>Co</i> ; LG 10	Prima x Fiesta	SSR	Maliepaard et al. 1998.
<i>Malus</i> × <i>domestica</i>	Juvenile phase length	QTLs: LG 3, 15 (LOD 3.2-4.0; 6.0-8.0%)	Fiesta x Discovery	SSRs	Liebhard et al., 2003a.
<i>Malus</i> × <i>domestica</i> (rootstock)	Dwarfing	Major locus: <i>Dw1</i> , LG05; LODs 5&1	M9 x Robusta 5	SCAR & SSR	Rushholme-Pilcher et al. 2008
<i>Malus</i> × <i>domestica</i>	Bloom time	QTL: LG 1	Rome Beauty X White Angel	Isozyme	Lawson et al., 1995.
<i>Malus</i> × <i>domestica</i>	Blooming time	QTLs: LG 7, 10, 17 (LOD 2.5-3.6; 5-13%)	Fiesta x Discovery	SSRs	Liebhard et al., 2003a.
<i>Malus</i> × <i>domestica</i>	No. bunches	QTLs: LG 8, 15 (LOD 3.6-5.1; 7.0-10.0%)	Fiesta x Discovery	SSRs	Liebhard et al., 2003a.
<i>Malus</i> × <i>domestica</i>	Self-fertility	Gene : <i>SI</i> , LG17	Prima x Fiesta	Isozyme, RFLP	Maliepaard et al., 1998.
<i>Malus</i> × <i>domestica</i>	Fruit harvest date	QTLs: LG 3 (LOD 4.7; 16.0%)	Fiesta x Discovery	SSRs	Liebhard et al., 2003a.
<i>Malus</i> × <i>domestica</i>	Harvest date	QTL: 5.7; LG3D-75; 13%	Fiesta × Discovery	unspecified	Liebhard et al., 2003a.
<i>Malus</i> × <i>domestica</i>	Skin color	Major locus: <i>Rs</i> ; LG17	Rome Beauty x White Angel		Weeden et al., 1994.
<i>Malus</i> × <i>domestica</i>	Skin color	Major locus: <i>Rf</i>	Rome Beauty x White Angel	Isozyme	Weeden et al., 1994.

<i>Malus</i> <i>×domestica</i>	Skin color	Major locus: <i>Rf</i> ; LG 9	Prima x Fiesta	SCAR	Maliepaard et al., 1998.	
<i>Malus</i> <i>×domestica</i>	Fruit number	QTLs: LG 5, 15, 16 (LOD 3.2-4.5; 8.0-10.0%)	Fiesta x Discovery	unspecified	Liebhard et al., 2003a.	
<i>Malus</i> <i>×domestica</i>	Fruit weight	QTLs: LG 1, 3, 6, 8, 10, 12, 15, 16 (LOD 2.5-17.0; 7.0-31.0%)	Fiesta x Discovery	unspecified	Liebhard et al., 2003a.	
<i>Malus</i> <i>×domestica</i>	Fruit weight	LG 4 (LOD 4.53; 25.0%),	Prima x Fiesta	unspecified	King et al. 2001	
<i>Malus</i> <i>×domestica</i>	Fruit cortex color, red foliage	Gene: MdMYB10 ( <i>Rni</i> locus), LG9	Sciros x 91.136B6-77	SNP	Chagne et al., 2008.	
<i>Malus</i> <i>×domestica</i>	Fruit ripening	Gene: <i>Md-ACS1</i> ; <i>LG15</i>	Prima x Fiesta	SCAR	Costa et al., 2005.	MAPS MASS
<i>Malus</i> <i>×domestica</i>	Fruit ripening	Gene: <i>Md-AC01</i> ; <i>LG10</i>	Prima x Fiesta Fuji x Mondial Gala	SCAR	Costa et al., 2005.	MAPS
<i>Malus</i> <i>×domestica</i>	Flesh firmness	QTLs: LG 6, 11, 12, 14 (LOD 3.6-12.3; 6.0-27.0%)	Fiesta x Discovery	unspecified	Liebhard et al., 2003a.	
<i>Malus</i> <i>×domestica</i>	Fruit texture – 8 measures	QTLs: Fruit firmness: LG 1, 8, 10 (LOD 4.7-7.4; 16.0-22.0%), Resonant freq: LG 10 (LOD 4.6; 21.0%), Hardness: LG 10, Crispness: LG 1, 5, 10, 12, 13, 16 (LG 16, LOD 6.0; 24%), Juiciness: LG 1, 12, 16 (LG 16, LOD 14.8; 46%), Granularity: LG 2 (LOD 5.1; 24%), Slow breakdown: LG 1, Sponginess: LG 1, 5, 6, 16 (LG 16, LOD 7.7; 30.0%)	Prima x Fiesta	unspecified	King et al., 2000; Maliepaard et al 2001; HiDRAS (unpublished)	
<i>Malus</i> <i>×domestica</i>	Fruit texture – 7 measures	Compression: LG 1, 6, 8, 12, 15 (LOD 4.09-8.62; 16.0-27%), Wedge	Prima x Fiesta	unspecified	King et al. 2001	



		measures: LG 1, 7, 15, 16 (LOD 4.51-9.83; 15.0-32.0%), Specific gravity: LG 6 (LOD 7.99; 28.0%), 16 (LOD 4.54; 15.0%), Stress at first failure (compression): LG 13 (LOD 3.51), Work of fracture (wedge fracture): LG 7 (LOD 4.51), Circularity of cells: LG 3 (LOD 3.3)			
<i>Malus</i> <i>× domestica</i>	Sugar content	QTLs: LG 3, 6, 8, 9, 14 (LOD 3.1-5.1; 3.6-12%)	Fiesta x Discovery	unspecified	Liebhard et al., 2003a.
<i>Malus</i> <i>× domestica</i>	Fruit acidity	QTLs: LG 8, 16 (LOD 4.7-6.2; 42.0-46.0%)( <i>Ma</i> locus on LG 16)	Fiesta x Discovery	unspecified	Liebhard et al., 2003a.
<i>Malus</i> <i>× domestica</i>	Malic acid content	Major locus: <i>Ma; LG16; 0</i>	Prima x Fiesta	RAPD	Maliepaard et al., 1998.
<i>Malus</i> <i>× domestica</i>	Fruit taste: overall liking	QTLs: LG 12, 16 (LG 16, LOD 11.3; 38.0%)	Prima x Fiesta	unspecified	King et al., 2000
<i>Malus</i> <i>× domestica</i>	Volatile compounds based on PTR-MS	QTL: 10 QTL, LOD 2.5 associated with PTR_MS peaks at m/z= 28, 43, 57, 61, 103, 115 & 145	Fiesta x Discovery	SSR	Zini et al., 2005.
<i>Malus</i> <i>× domestica</i>	Allergens	Major locus: <i>Mal d 1.05</i> ; LG 6	Prima x Fiesta plus Jonathan x Prima	SNP, SSR	Gao et al., 2005a.
<i>Malus</i> <i>× domestica</i>	Allergens	Candidate genes: Family of seven <i>Mal d 1</i> genes ; LG 13	Prima x Fiesta plus Jonathan x Prima	SNP, SSR	Gao et al., 2005a.
<i>Malus</i> <i>× domestica</i>	Allergens	Candidate genes: Family of nine <i>Mal d 1</i> genes ; LG 16	Prima x Fiesta plus Jonathan x Prima	SNP, SSR	Gao et al., 2005a.
<i>Malus</i> <i>× domestica</i>	Allergens	Candidate genes: <i>Mal d 2.01A</i> ; LG 9	Prima x Fiesta	SNP	Gao et al., 2005b. Genet. 111:1087-1097.
<i>Malus</i>	Allergens	Candidate genes: <i>Mal d 3.01</i> ; LG	Jonathan x Prima	SNP	Gao et al., 2005c.

<i>Malus domestica</i>	Allergens	12 Candidate genes: <i>Mal d 3.02</i> ; LG 4	Prima x Fiesta plus Jonathan x Prima	SNP	Gao et al., 2005c.
<i>Malus domestica</i>	Allergens	Candidate genes: <i>Mal d 4.01</i> ; LG9	Prima x Fiesta	SNP	Gao et al., 2005b.
<i>Malus domestica</i>	Allergens	Candidate genes: 2 copies of <i>Mal d 4.01</i> ; LG9	plus Jonathan x Prima	SNP	van de Weg, pers. comm.
<i>Malus domestica</i>	Allergens	Candidate genes: <i>Mal d 4.2A</i> ; LG2	plus Jonathan x Prima	SNP	van de Weg, pers. comm.
<i>Malus domestica</i>	Allergens	Candidate genes: <i>Mal d 4.3A</i> ; LG8	plus Jonathan x Prima	SNP, SSR	van de Weg, pers. comm.
<b>PEAR</b>					
<i>Pyrus communis</i>	Red fruit skin and foliage	Major locus: unspecified	Bon Rouge x Packham's Triumph	SSR	Booi et al. 2005.
<i>Pyrus communis</i>	Leaf scab	2 QTLs: LG 3 & 4, both 88%; LOD>10	Abbe Fetel (AF) x Max Red Bartlett (MRB)	AFLPs & SSRs	Pierantoni et al., 2007.
<i>Pyrus communis</i>	Fire blight	QTL: LGHS2a-0.0, QTL : 3.25 ; LGHS2a-9.0 ; 16.4% QTL : 2.06 ; LFHS2b-10.7 ; 9.6% QTL : 2.64 ; LGHSA-9.8 ; 12.0% QTL : 1.85 ; LGHS-21.9 ; 8.5%	Passa Crassane x Harrow Sweet	SSRs, AFLPs, AFLP-RGAs	Dondini et al., 2004.
<i>Pyrus communis</i> and other <i>Pyrus</i>	Self-incompatibility	Gene : <i>S</i> locus (approx. 30 alleles within the genus)	Various cultivars	PCR of allelic sequences	Sanzol et al. 2008; Zhang et al., 2007
<i>Pyrus communis</i> × <i>P. bretschnei</i>	Dwarf	Major locus: <i>pcDw</i> (marker is 8.3 cM away from gene)	Aihuali x Chili	RAPD and SCAR	Jia et al. 2007.

<i>deri</i> <i>Pyrus</i> <i>pyrifolia</i>	Asian pear scab resistance	Major locus: <i>Vn</i> ; LG unspecified – Kin-61.9; marker at 44.5	Kinchaku (R) x Kosui (S)	RAPD	Iketani et al., 2001.
<i>Pyrus</i> <i>pyrifolia</i>	Asian pear scab resistance	Major locus: <i>Vnk</i> ; LG1; LOD 5.0, 2.4-12.4cM away	Shuurei x 314-32 (Kinchaku x Housui) Housui x 30-38 (Chikusui x Kinchaku)	AFLP-STS RAPD-STS	Terakami et al., 2004.
<i>Pyrus</i> <i>pyrifolia</i>	<i>Alternaria</i> black spot susceptibilit y	Major locus: <i>A</i> ; LG unspecified- Kin- 66.8; markers at 53.0 and 48.1	Kinchaku (S) x Kosui (R)	RAPD	Iketani et al., 2001.
<i>Pyrus</i> <i>pyrifolia</i>	<i>Alternaria</i> black spot susceptibilit y	Major locus: <i>A</i> ; marker is 3.1 cM from gene	Osa Nijisseiki open- pollinated	RAPD	Banno et al., 1999; Banno et al. 2002.
<i>Pyrus</i> <i>pyrifolia</i>	<i>Alternaria</i> black spot susceptibilit y	Major locus: <i>Ani</i> ; LG11Osa-3.4; SSRs at 0.0 and 12.1; LODs 24.5 and 18.1	Osa Nijisseiki x Okusankichi	SSRs	Terakami et al., 2007.
<i>Pyrus</i> <i>pyrifolia</i>	<i>Alternaria</i> black spot susceptibilit y	Major locus: <i>Ana</i> ; LG11N-2.5, SSRs at 2.5 and 10.1; LODs 12.0 and 7.4	Oushuu x Nansui	SSRs	Terakami et al., 2007.
<i>Pyrus</i> <i>pyrifolia</i>	Fruit ethylene biosynthesis	Genes: <i>PP-ACS1</i> (high ethylene) <i>and PP-ACS2</i> (moderate ethylene)	F <sub>2</sub> of OT-16 (F <sub>1</sub> of Osa Nijiseiki x Cili)	CAPS: A and B	Itai et al. 2003.
<i>Pyrus</i> <i>pyrifolia</i>	Fruit russet	Major locus: <i>I</i> (modifier of <i>R</i> ); not mapped; marker associated with unrusseted phenotype.	Kousui x Kinchaku; Niitaka x Chikusui	RAPD	Inoue et al. 2006.

<i>Pyrus pyrifolia</i> × <i>P. communis</i>	Red leaf color	Major locus: <i>Re</i> ; LG12O	Osa Nijisseiki x Oharabeni	RAPD	Banno et al. 2002.
<i>Pyrus pyrifolia</i> × <i>P. ussuriensis</i>	Fruit hardness	Not mapped	Niitaka x Suhyanri	RAPD, AFLP	Kim et al; 2005.
<i>Pyrus pyrifolia</i> × <i>P. ussuriensis</i>	Skin color	Not mapped	Niitaka x Suhyanri	RAPD, AFLPs	Kim et al; 2005.
<i>Pyrus pyrifolia</i> × <i>P. ussuriensis</i>	Fruit grit	Not mapped	Niitaka x Suhyanri	AFLPs	Kim et al; 2005.
<i>Pyrus pyrifolia</i> × <i>P. ussuriensis</i>	Fruit shape	Not mapped	Niitaka x Suhyanri	AFLPs	Kim et al; 2005.
<b>PEACH</b>					
<i>Prunus persica</i>	Leaf curl resistance	QTLs: 2.5-6.6, LG3D-28, 16-37%; 2.5-4.4, LG6D-28, 6-29%	Summergrande x <i>P. davidiana</i> P1908	RFLPs	Viruel et al., 1998.
<i>Prunus persica</i>	Powdery mildew resistance	QTL:2.2-4.3, LG7-21, 13.5-26.5%	( <i>P. ferganensis</i> x IF7310828) BC1	RFLP	Verde et al., 2002.
<i>Prunus persica</i> rootstock	Peach Tree Short Life (PTSL)	QTL: n/a; LG1 – 14; QTL: n/a; LG2 - 0-8; 34; 43; QTL: n/a; LG4 - 36; QTL: n/a; LG6 - 44.7	‘Guardian® 3-17-7’ x ‘Nemaguard’	SSRs	Liu et al., 2008.

<i>Prunus persica</i>	Powdery mildew resistance	QTLs: 2.4-3.2, LG1-86,7-11%; 2.6-13.0, LG6-31.85, 5-56% 17, LG8-19.9,32-57%	Summergrand x <i>P. davidiana</i> P1908 F <sub>1</sub> , F <sub>2</sub> , & BC <sub>2</sub> (crossed to Zéphir)	RFLPs	Foulongne et al., 2003
<i>Prunus persica</i> rootstock	Rootknot nematode resistance	Major loci: <i>Mi</i> and <i>Mij</i> , LG1	Lovell × Nemared	AFLPs	Abbott et al., 1998.
<i>Prunus persica</i>	Internode length	QTL: 4.1, LG1b, 26.2%	<i>P. ferganensis</i> x IF7310828) BC1		Verde et al., 2002
<i>Prunus persica</i>	Blooming time	QTL: 5.6-6.1, LG5-59,32.1-35.1% (co-segregates with F gene)	( <i>P. ferganensis</i> x IF7310828) BC1	B4C10	Verde et al., 2002.
<i>Prunus persica</i>	Fruit development period	QTL: 9; LG4 – 8; 76.2%	Ferjalou Jalousia x Fantasia	SSR	Etienne et al., 2002.
<i>Prunus persica</i>	Ripening date	QTL: 9; LG4 - 0; 77.7%	Ferjalou Jalousia x Fantasia	SSR	Etienne et al., 2002.
<i>Prunus persica</i>	Fruit maturity (harvest)	QTLs	Suncrest x Bailey	RAPD, AFLPs, RFLP	Abbott et al., 1998.
<i>Prunus persica</i>	Ripening time	QTL: 3.3-5.3, LG6-2, 22.4-30.6%; QTL: 2.6-3.3, LG2-6, 15.8-19.8% QTL: 4.2-4.6, LG6-2, 26.4-28.8%	( <i>P. ferganensis</i> x IF7310828) BC1	SSR, RFLP	Verde et al., 2002.
<i>Prunus persica</i>	Fruit skin color	QTL: 2.3-2.7, LG2-6, 14.1-17.5% QTL: 3.3, LG2-8, 19.9%	( <i>P. ferganensis</i> x IF7310828) BC1	SSR, RFLP	Verde et al., 2002.
<i>Prunus persica</i>	Fresh weight	QTL: 4.9; LG6 – 59.5; 52%	Ferjalou Jalousia x Fantasia	RFLP	Etienne et al., 2002.
<i>Prunus persica</i>	Fruit weight, fruit diameter	QTLs	Suncrest x Bailey	AFLPs RAPDs	Abbott et al., 1998.

<i>Prunus persica</i>	Fruit softening and flesh-stone adhesion	Gene: endoPG, LG4 ( <i>Freestone-melting flesh</i> locus)	'Dr. Davis' x 'Georgia Belle'	EST-SSR	Peace et al., 2005. MAPS
<i>Prunus persica</i>	Fruit fructose and glucose contents	QTLs	Suncrest x Bailey	RFLPs, AFLPs	Abbott et al., 1998.
<i>Prunus persica</i>	Fruit sugar (SSC) content	QTL: 4.1; LG4; 34% ; QTL: 2.5; LG6 ; 17%	Ferjalou Jalousia x Fantasia	SSR, RFLP	Etienne et al., 2002.
<i>Prunus persica</i>	Fruit sugar (SSC) content	Candidate gene: <i>RRUpe Vp2</i>	Texas (almond) x Earlygold F <sub>2</sub>	SSCP	Etienne et al., 2002.
<i>Prunus persica</i>	Fruit sugar (SSC) content	QTL: 2.0-2.9, LG6-0, 14.4-17.9 QTL: 3.3, LG2-4, 20.9%	( <i>P. ferganensis</i> x IF7310828) BC1	SSR, RFLP	Verde et al., 2002.
<i>Prunus persica</i>	Fruit sweetness: fructose, glucose, sucrose content	QTL: 4.5, LG4-10, 38% QTL: 4.6, LG4-8, 41% QTL: 4.5 LG6-63.8, 39% Major locus 4.1, LG5-23.8, 39%	Ferjalou Jalousia x Fantasia	RFLPS, <i>D</i> locus	Etienne et al., 2002.
<i>Prunus persica</i>	Fruit acidity: malic & citric acid content, TA, pH	Major locus: <i>D</i> (low acid): 10.7, LG5-31.4, 72%; 9.1, LG%-33.4, 77%; 10.8, LG5-33.4, 80%; 10.4, LG%-27.4, 86%	Ferjalou Jalousia x Fantasia		Etienne et al., 2002
<i>Prunus</i>	Fruit quinic	1 QTL on LG8	Ferjalou Jalousia x	RFLP	Etienne et al.,

<i>Persica</i>	acid content		Fantasia		2002.	
<i>Prunus</i>	Cold-	QTL: qP-Brn5.1 <sup>m</sup> , LG5-21; 39%	'Dr. Davis' x	EST-SSR &	Ogundiwin et al.,	
<i>Persica</i>	storage flesh browning	with candidate gene (PpLDOX)	'Georgia Belle'	SNP	2008.	
<b>ALMOND</b>						
<i>Prunus</i>	Self-fertility	Gene: S <sub>r</sub> -allele; LG6	'Tuono'	CEBASf	Sanchez-Perez et	MAPS
<i>dulcis</i>	& cross- compatibilit y	(same gene as for other <i>Prunus</i> )			al., 2004.	
<i>Prunus</i>	Anther color	Major locus: Ag; G3	Texas (almond) x	RFLP	Joobeur 1998	
<i>dulcis</i>			Earlygold (peach)			
<i>Prunus</i>	Blooming	Major locus: Lb; G4	D.3.5 (almond) x	RAPD	Ballester et al.,	
<i>dulcis</i>	time		Bertina (almond)		2001	
<i>Prunus</i>	Flower	Major locus: B; G1	Garfi (almond) x	RFLP	Jauregui 1998	
<i>dulcis</i>	color		Nemared (peach)			
<i>Prunus</i>	Kernel taste	Major locus: Sk; G5	Texas (almond) x	RFLP	Joobeur 1998	
<i>dulcis</i>			Earlygold (peach)			
<i>Prunus</i>	Nematode	Major locus: Mi; G2	Garfi (almond) x	RFLP	Jauregui 1998	
<i>dulcis</i>	resistance		Nemared (peach)			
<i>Prunus</i>	Shell	Major locus: D; G2	Ferragnés (almond)	RFLP	Arus et al., 1999	
<i>dulcis</i>	hardness		x Tuono (almond)			
<b>SWEET CHERRY</b>						
<i>Prunus</i>	Cross-	Gene: SFB at S locus	Various		Ikeda et al. 2004	MAPS
<i>avium</i>	compatibility					
<i>Prunus</i>	Self –	Gene: SFB4', LG6, same gene as	JI 2420	SFB4' CAP	Ikeda et al., 2004.	MAPS
<i>avium</i>	fertility	above (but specific allele)		marker		
<b>SOUR CHERRY</b>						
<i>Prunus</i>	Bloom	11 QTLs, LOD>2.4, 12.9%-	Rheinische	RFLP	Wang et al., 2000.	
<i>cerasus</i>	time, % pistil death, % pollen	25.9%; 3 QTL for bloom time, % pistil death and % pollen	Schattenmorelle (RS) x Erdi Boterno			

<i>Prunus cerasus</i>	germination Fruit weight	germination mapped to LG EB1 See above	(EB) RS x EB	RFLP	Wang et al., 2000.
<i>Prunus cerasus</i>	Fruit % Brix	See above	RS x EB	RFLP	Wang et al., 2000.
<i>Prunus cerasus</i>	Ripening time	See above	RS x EB	RFLP	Wang et al., 2000.

### APRICOT

<i>Prunus armeniaca</i>	Plum pox resistance	Major locus: LG1	Goldrich x Currot; Polonais x Stark Early Orange; Lito (Stark Early Orange x Tyrinthos) selfed	SSR and SSCP	Sicard et al. 2008.
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### STRAWBERRY

<i>Fragaria ×ananassa</i>	Day-neutrality	QTL: > 4.8, MI 05-1/MD05/MN 05-2, LG 28, 26.1 %; 5.2, 21.9 %, LG 28, 5.4 %; 5.8, CA05 LG 6, 14.4 %; 2.6, MN05-1, LG 6, 15 %; 2.4, MN05-3, LG 1, 11.5%; 2.4, MN05-4, LG 3, 13.0%	‘Tribute’ x ‘Honeoye’	AFLP, RAPD- SCAR	Weebadde et al., 2008	MASS
<i>Fragaria ×ananassa</i>	Fruit firmness	QTL: 3.7, f379, LG -1, 10 %; 3.3, f211, LG-6, 8 %; 3.3, f211, LG -6, 8 %; 3.9, f389, LG 35, 11 %	Capitola x CF1116	AFLP, SSR	Lerceteau-Köhler et al., 2004.	
<i>Fragaria ×ananassa</i>	Fruit maturity	QTL: 5.4, m574, LG 30, 15 %	Capitola x CF1116	AFLP, SSR	Lerceteau-Köhler et al., 2004	
<i>Fragaria ×ananassa</i>	Fruit fresh weight	QTL: 3.9, m97, LG 4, 9.3 %	Capitola x CF1116	AFLP, SSR	Lerceteau-Köhler et al., 2004.	
<i>Fragaria ×ananassa</i>	Fruit length, diameter	QTL:5.9, m425, LG 5, 12.9 %; 3.5, m563, LG 2b, 7 %; 7.5, m167, LG 5, 15 %; 3.1, m169, LG 7,	Capitola x CF1116	AFLP, SSR	Lerceteau-Köhler et al., 2004	



<i>Fragaria</i> × <i>ananassa</i>	Fruit citrate content	6.5 %; 3.3, m307, LG 7, 7 % QTL: 3.0, m407, LG 6, 7 %	Capitola x CF1116	AFLP, SSR	Lerceteau-Köhler et al., 2004	
<i>Fragaria</i> × <i>ananassa</i>	Fruit sucrose content	QTL: 3.5, f171, LG 37, 9 %; 3.7, f259, LG 7, 8 %	Capitola x CF1116	AFLP, SSR	Lerceteau-Köhler et al., 2004	
<i>Fragaria</i> × <i>ananassa</i>	Fruit sugar content (SSC)	QTL: 3.1, m577, LG 7, 7 %	Capitola x CF1116	AFLP, SSR	Lerceteau-Köhler et al., 2004	
<i>Fragaria</i> × <i>ananassa</i>	Resistance to <i>C. acutatum</i>	QTL: 2.3, f163, LG 6, 12%; 2.0, f219-f347, LG 11, 8%; 2.3, m193, LG 14, 10 %; m321, 2.8, LG 40, 10%	Capitola x CF1116	AFLP, SSR	Denoyes-Rothan et al., 2004	
<i>Fragaria</i> × <i>ananassa</i>	Resistance to <i>P. cactorum</i>	QTL: 2.1, f93, LG 1, 8 %; 2.4, f105-f291, LG 19, 10 %; 3.5, m243-m523, LG 2a, 12 %; 2.1, m671, LG 11, 6 %; 2.2, m153, LG 40, 10 %	Capitola x CF1116	AFLP, SSR	Denoyes-Rothan et al., 2004	
<i>Fragaria</i> × <i>ananassa</i>	Resistance to <i>P. fragariae</i>	Major locus?	Md683 x Senga Sengana and Pedigreed set of cultivars and breeding lines	SCAR	Haymes et al. 1997	MAPS MASS in 09
<i>Fragaria</i> × <i>ananassa</i>	Resistance to <i>P. fragariae</i>	Major locus? <i>MMAB</i> , 1 cM	Redgauntled x Senga Sengana and Pedigreed set of cultivars and breeding lines	SCAR	Van de Weg et al. ms in prep	MAPS MASS in 09
<b>RASPBERRY</b>						
<i>Rubus idaeus</i>	Cane spot	Major locus: <i>H</i> (cane pubescence) LG2	Glen Moy x Latham	SSRs	Graham et al. 2006.	

<i>Rubus idaeus</i>	Rust	QTL:LG2L Major locus: <i>H</i> LG2 ; QTLs :LG3L-44, LG5L-50	Glen Moy x Latham		Graham et al. 2006.
<i>Rubus idaeus</i>	Botrytis and spur blight	QTLs: 1 on LG1, 1 on LG2, 3 on LG3, 3 on LG5, 5 on LG6	Glen Moy x Latham		Graham et al. 2006.
<i>Rubus idaeus</i>	Cane spininess	QTLs : LG3L-58 & LG2L-59	Glen Moy x Latham		Graham et al. 2006.
<i>Rubus idaeus</i>	Aphid resistance	Major locus: <i>A</i> <sub>1</sub> , LG3-53.2	Malling Jewel x Malling Orion	SSR, AFLP	Sargent et al. 2007.
<i>Rubus idaeus</i>	Dwarfness	Major locus: <i>dw</i> ; LG6-36.5	Malling Jewel x Malling Orion	AFLPs	Sargent et al. 2007.
<i>Rubus idaeus</i>	RLSV and RVCV viral diseases	Major locus on LG 2 and 8	Glen Moy x Latham		Rusu et al. 2006.
<i>Rubus idaeus</i>	Cane spininess	QTL : Two regions with 16 markers on LG2, 46.5-56.1 cM, 48% ; and 108-120.9 cM, 50%	Glen Moy x Latham	AFLPs, SSRs	Graham et al. 2004.
<i>Rubus idaeus</i>	Root sucker density	QTL: LG8, 53%	Glen Moy x Latham	AFLPs and SSRs	Graham et al. 2004.
<i>Rubus idaeus</i>	Root sucker spread	QTLs: LG?, 79%, LG?, 33%	Glen Moy x Latham	AFLPs and SSRs	Graham et al. 2004.

## BLACKBERRY

None known

## ROSE

<i>Rosa x hybrida</i>	Black spot resistance	Major locus: Unlinked to <i>Rdr1</i> ; <i>Rdr2</i> (not yet assigned to linkage group)	'Chorale' x 'George Vancouver' (4x cultivated)	BSA in progress	Whitaker et al., 2008.
<i>Rosa x hybrida</i>	Black spot resistance	Major locus: <i>Rdr1</i> , LG1	<i>Rosa multiflora</i> (4x) x <i>R. x hybrida</i>	SSR	Von Malek et al., 2000.

<i>Rosa x hybrida</i>	Black spot resistance	Major locus: <i>Rdr2</i> , LG1	'Caramba' <i>Rosa multiflora</i>	SSR	Hattendorf et al., 2004.
<i>Rosa x hybrida</i>	Powdery mildew resistance	Major locus: <i>Rpp1</i> , LG3	<i>Rosa multiflora</i>	SCAR	Linde et al., 2004.
<i>Rosa x hybrida</i>	Powdery mildew resistance	28QTLs: >3.9; seven different linkage groups	<i>Rosa multiflora</i>	AFLP, CAPS, SCAR, RGA	Linde et al., 2006.
<i>Rosa x hybrida</i>	Powdery mildew resistance, flowering traits	8 QTLs	<i>Rosa wichuriana</i>	RAPD, SSR	Dugo et al., 2005.
<i>Rosa x hybrida</i>	Blooming date; Petal number	QTL <i>BD</i> , LG B4; Major locus: <i>NP</i> , LG A4, QTL <i>NP</i> , LG B4	<i>R. x hybrida</i> 'Zambra' (dihaploid) x <i>R. wichuriana</i>	SSR	Hibrand-Saint Oyant et al., 2008.
<i>Rosa x hybrida</i>	Recurrent bloom	Major locus <i>r4</i> , LG B4	<i>R. x hybrida</i> 'Zambra' (dihaploid) x <i>R. wichuriana</i>	AFLP	Crespel et al., 2002.
<i>Rosa x hybrida</i>	Flower double corollas	Major locus <i>d6</i> , LG A6	<i>R. x hybrida</i> 'Zambra' (dihaploid) x <i>R. wichuriana</i>	AFLP	Crespel et al., 2002.
<i>Rosa multiflora</i>	Double flowers	Major locus: <i>Blfo</i> , LG B4	F1 diploid mapping population	RAPD, AFLP	Debener and Matiesch, 1999.
<i>Rosa multiflora</i>	Flower color (pink)	Major locus: <i>Blfa</i> , LG B2	F1 diploid mapping population	RAPD, AFLP	Debener and Matiesch, 1999.

<i>Rosa x hybrida</i>	Stem thorn number	QTL <i>t4</i> , LG B4; QTL <i>tb4</i> , LG B4	<i>R. x hybrida</i> 'Zambra' (dihaploid) x <i>R. wichuriana</i>	AFLP	Crespel et al., 2002.
<i>Rosa x hybrida</i>	Petiole prickles	Major locus: <i>Pr</i> , LG 7	86-7 ( <i>R. wichuraiana</i> x <i>R. rugosa</i> ) x <i>R. x hybrida</i> 'Basye's Blueberry'	AFLP, SSR	Rajapakse et al., 2001.
<i>Rosa multiflora</i>	Plant vigor (10 traits)	42 QTLs: >2.5, 12-35% variation	F1 diploid mapping population	AFLP, SSR, PK, RGA, RFLP, SCAR	Yan et al., 2007.

<sup>a</sup> MAPS = marker-assisted parent selection ; MASS = marker-assisted seedling selection

## REFERENCES

- Abbott, A. G., Rajapakse, S., Sosinski, B., Lu, Z. X., Sossey-Alaoui, K., Gannavarapa, M., Reighard, G., Ballard, R. E., Baird, W. V., Scorza, R., and Callahan, A. 1998. Acta Hort. 465:41-49.
- Alston, F. H., and Briggs, J. B. 1968. Resistance to *Sappaphis devectora* (Wlk.) in apple. Euphytica 17:468-472.
- Alston, F. H., and Briggs, J. B. 1977. Resistance genes in apple and biotypes of *Dysaphis devectora*. Ann. Appl. Biol. 87:75-81.
- Arús, P., Ballester, J., Jáuregui, B., Joobeur, T., Truco, M.J., and de Vicente, M.C., 1999. The European *Prunus* mapping project : update on marker development in almond. Acta Hort. 484: 331-336
- Ballester, J., Boskovic, R., Battle, I., Arús, P., Vargas, F., and de Vicente, M.C., 1998. Location of self incompatibility gene in almond linkage map. Plant Breeding 117: 69-72.

- Ballester, J., Socias I, Company, Arús, P., and de Vicente, M.C., 2001. Genetic mapping of a major gene delaying blooming time in almond. *Plant Breeding* 120: 268-270.
- Banno, K., Ishikawa, H., Hamauzu, Y., and Tabira, H. 1999. Identification of a RAPD marker linked to the susceptible gene of black spot disease in Japanese pear. *J. Jap. Soc. Hort. Sci.* 68:476-481.
- Banno, K., Saito, S., Robbani, M., and Ishikawa, H. 2002. Introduction of new characteristics and genetic mapping using hybrids between Japanese pear cv. 'Osa Nijisseiki' and European pear F<sub>1</sub> cv. 'Oharabeni'. *Acta Hort.* 587:225-231.
- Battle, I., and Alston, F. H. 1996. Genes determining leucine aminopeptidase and mildew resistance from the ornamental apple, 'White Angel'. *Theor. Appl. Genet.* 93:179-182.
- Bliss, F.A., Arulsekar, S., Foolard, M.R., Beccera, V., Gillen, A.M., Warburton, M.L., Dandekar, A.M., Kocsin, G.M., and Mydin, K.K., 2002. An expanded genetic linkage map of *Prunus* based on an interspecific cross between almond and peach. *Genome* 45:520-529.
- Booi, S., van Dyk, M. M., du Preez, M. G., Rees, D. J. G., and Labuschagne, I. 2005. Molecular typing of red and green phenotypes of 'Bon Rouge' pear trees, with the use of microsatellites. *Acta Hort.* 671:293-297.
- Bénaouf, G., and Parisi, L. 2000. Genetics of host-pathogen relationships between *Venturia inaequalis* races 6 and 7 and *Malus* species. *Phytopathology* 90:236-242.
- Bus, V. G. M., Chagné, D., Bassett, H. C. M., Bowatte, D., Calenge, F., Celton, J.-M., Durel, C.-E., Malone, M. T., Patocchi, A., Ranatunga, A. C., Rikkerink, E. H. A., Tustin, D. S., Zhou, J., and Gardiner, S. E. 2008. Genome mapping of three major resistance genes to woolly apple aphid (*Eriosoma lanigerum* Hausm.). *Tree Genet. Genomes* 4:223-236.
- Bus, V. G. M., Laurens, F. N. D., van de Weg, W. E., Rusholme, R. L., Rikkerink, E. H., A., Gardiner, S. E., Bassett, H. C. M., Kodde, L. P., and Plummer, K. L. M. 2005a. The *VH8* locus of a new gene-for-gene interaction between *Venturia inaequalis* and the wild apple *Malus severii* is closely linked to the *Vh2* locus in *Malus pumila* R12740-7A. *New Phytologist* 166:1035-1049.

Bus, V., Ranatunga C., Gardener, S.E., Basset, H., Rikkerink, E. 2000 Marker assisted selection for pest and disease resistance in the New Zealand apple breeding programme. *Acta Hort* 538: 541-547

Bus, V. G. M., Rikkerink, E. H., van de Weg, W. E., Rusholme, R. L., Gardiner, S. E., Bassett, H. C. M., Kodde, L. P., Parisi, L., Laurens, F. N. D., Meulenbroek, E. J., and Plummer, K. M. 2005b. The *Vh2* and *Vh4* scab resistance genes in two differential hosts derived from Russian apple R12740-7A map to the same linkage group of apple. *Mol. Breed.* 15:103-116.

Calenge, F., Drouet, D., Denancé, C., Van de Weg, W. E, Brisset, M.-N., Paulin, J.-P., and Durel, C.-E. 2005. Identification of a major QTL together with several minor additive or epistatic QTLs for resistance to fire blight in apple in two related progenies. *Theor. Appl. Genet.* 111:128-135.

Calenge, F., and Durel, C.-E. 2006. Both stable and unstable QTLs for resistance to powdery mildew are detected in apple after four years of field assessments. *Mol. Breed.* 17:1-11. (doi: 10.1007/s11032-006-9004-7)

Calenge, F., Faure, A., Goerre, M., Gebhardt, C., Van de Weg, W. E., Parisi, L., and Durel, C.-E. 2004. Quantitative trait loci (QTL) analysis reveals both broad-spectrum and isolate-specific QTL for scab resistance in an apple progeny challenged with eight isolates of *Venturia inaequalis*. *Phytopathol.* 94:370-379.

Cevik, V., and King, G. J. 2000. Molecular genetic analysis of the *Sd1* aphid resistance locus in *Malus*. *Acta Hort.* 538:553-559.

Cevik, V., and King, G. J. 2002. High-resolution genetic analysis of the *Sd-1* aphid resistance locus in *Malus* spp. *Theor. Appl. Genet.* 105:346-354.

Chagne, D., Carlisle, C. M., Blond, C., Volz, R. K., Whitworth, C. I., Oraguzie, N. C., Crowhurst, R. N., Allan, A. C., Espley, R. V., Hellens, R. P, and Gardiner, S. E. 2007. Mapping a candidate gene (MdMYB10) for red flesh and foliage colour in apple. *BMC Genomics* 8:212.

Cheng FS, Weeden NF, Brown SK, Aldwinckle HS, Gardener SE, Bus VG (1998) Development of a DNA marker for *V<sub>m</sub>*, a gene conferring resistance to apple scab. *Genome* 41: 208-214

- Conner, P. J., Brown, S. K., and Weeden, N. F. 1998. Molecular-marker analysis of quantitative traits for growth and development in juvenile apple trees. *Theor. Appl. Genet.* 96:1027-1035.
- Costa, F., Stella, S., Van de Weg, W. E., Cuerra, W., Cecchinell, M., Dallavia, J., Koller, B., and Sansivini, S. 2005. Role of the genes *Md-ACO1* and *Md-ACS1* in ethylene production and shelf life of apple (*Malus domestica* Borkh.). *Euphytica* 141:181-190.
- Costa F, Stella S, Van de Weg WE, Dondini L, Pratesi D, Musacchi S, Sansavini S. Map position and functional allelic diversity of *Md-Exp7*, a new putative expansin gene associated with fruit softening in apple (*Malus x domestica* Borkh.) and pear (*Pyrus communis*). *Tree Genet. Genomes*. In Press
- Crespel, , L., Chirollet, M., Durel, C. E., Zhang, D., Meynet, J., and Gudin, S. 2002. Mapping of qualitative and quantitative phenotypic traits in *Rosa* using AFLP markers. *Theor. Appl. Genet.* 105:1207-1214.
- Davey et al 2006 .Genetics of Fruit Vitamin C. *Plant Physiol* 142 343-351
- Debener, T., and Mattiesch, L. 1999. Construction of a genetic linkage map for roses using RAPD and AFLP markers. *Theor. Appl. Genet.* 99:891-899.
- Denoyes-Rothan, B., Lerceteau-Köhler, E., Guérin, G., Bosseur, S., Bariac, J., Martin, E., and Roudeillac, P. 2004. QTL analysis for resistances to *Colletotrichum acutatum* and *Phytophthora cactorum* in octoploid strawberry (*Fragaria x ananassa*). *Acta Hort.* 663:147-151.
- Dondini, L., Pierantoni, L., Gaiotti, F., Chiodini, R., Tartarini, S., and Bazzi, C. 2004. Identifying QTLs for fire-blight resistance via a European pear (*Pyrus communis* L.) genetic linkage map. *Mol. Breed.* 14: 407-418.
- Durel, C. E., Parisi, L., Laurens, F., Van de Weg, W. E., Liebhard, R., and Jourjon, M. F. 2003. Genetic analysis of partial resistance to race 6 of *Venturia inaequalis* in apple. *Genome* 46:224-234.
- Durel, C. E., van der Weg, W. E., Venisse, J. S., and Parisi, L. 2000. Localization of a major gene for apple scab resistance on the European gene map of the Prima x Fiesta cross. *OILB/WPRS Bull.* 23:245-248.

Dugo, M. L., Satovic, Z., Millán, T., Cubero, J. I., Rubiales, D., Cabrera, A., and Torres, A. M. 2005. Genetic mapping of QTLs controlling horticultural traits in diploid roses. *Theor. Appl. Genet.* 111:511-520.

Dunneman, F., Bracker, G., Markussen, T., Roche, P. 1999 Identification of molecular markers for the major mildew resistance gene *Pl<sub>2</sub>* in apple. *Acta Hort* 484: 411-416.

Erdin, N., Tartarini, S., Broggin, G. A. L., Gennari, F., Sansavini, S., Gessler, C., and Patocchi, A. 2006. Mapping of the apple scab resistance gene *Vb*. *Genome* 49:1238-1245.

Etienne, C., Rothan, C., Moing, A., Plomion, C., Bodénè, C., Svanella-Dumas L., Cosson, P., Pronier, V., Monet, R., and Dirlwanger, E. 2002. Candidate genes and QTLs for sugar and organic acid contents in peach [*Prunus persica* (L.) Batsch]. *Theor. Appl. Genet.* 105:145-159.

Evans, K. M., and James, C. M. 2003. Identification of SCAR markers linked to *Pl-w* mildew resistance in apple. *Theor. Appl. Genet.* 106:1178-1183.

Foulongne, M., Pascal, T., Pfeiffer, F., and Kervella, J. 2003. QTLs for powdery mildew resistance in peach x *Prunus davidiana* crosses: consistency across generations and environments. *Mol. Breed.* 12:33-50.

Gao, Z. S., van de Weg, W. E., Shaart, J. G., Schouten, H. J., Tran, D. H., Kodde, L. P., van der Meer, I. M., van der Geest, A. H. m., Kodde, J., Breiteneder, H., Hoffman-Sommergruber, K., Bosch, D., and Gilissen, L. J. W. J. 2005. Genomic cloning and linkage mapping of the *Mal d 1* (*PR-10*) gene family in apple (*Malus domestica*). *Theor. Appl. Genet.* 111:171-183.

Gao, Z. S., van de Weg, W. E., Shaart, J. G., van Arkel, G., Breiteneder, H., Hoffman-Sommergruber, K., and Gilissen, L. J. W. J. 2005. Genomic characterization and linkage mapping of the apple allergen genes *Mal d 2* (thaumatin-like protein) and *Mal d 4* (profiling). *Theor. Appl. Genet.* 111:1087-1097.

Gao, Z. S., van de Weg, W. E., Shaart, J. G., van der Meer, I. M., Kodde, L., Laimer, M., Breiteneder, H., Hoffman-Sommergruber, K., and Gilissen, L. J. W. J. 2005. Linkage map positions and allelic diversity of two *Mal d 3* (non-specific lipid transfer protein) genes in the cultivated apple (*Malus domestica*). *Theor. Appl. Genet.* 110:479-491.



- Gardiner, S., Murdoch, J., Meech, S., Rusholme, R., Bassett, H., Cook, M., Bus, v., Rikkerink, E., Gleave, A., Crowhurst, R., Ross, G., and Warrington, I. 2003. Candidate resistance genes from an EST database prove a rich source of markers for major genes conferring resistance to important apple pests and diseases. *Acta Hort.* 622:141-151.
- Graham, J., Smith, K., MacKenzie, K., Jprgenon, L., Hackett, C., and Powell, W. 2004. the construction of a genetic linkage map of red raspberry (*Rubus idaeus* subsp. *idaeus*) based on AFLPs, genomic –SSR and EST-SSR markers. *Theor. Appl. Genet.* 109:740-749.
- Graham, J., Smith, K., Tierney, I., MacKenzie, K., and Hackett, C. A. 2006. Mapping gene *H* controlling cane pubescence in raspberry and its association with resistance to cane botrytis and spur blight, rust and cane spot. *Theor. Appl. Genet.* 112:818-831.
- Gygax, M., Gianfranceschi, L., Liebhard, R., Kellerhals, M., Gessler, C., and Patocchi, A. 2004. Molecular markers linked to the apple scab resistance gene *Vbj* derived from *Malus baccata jackii*. *Theor. Appl. Genet.* 109:1702-1709.
- Hattendorf, A., Linde, M., Mattiesch, L., Debener, T., and Kaufmann, H. 2004. Genetic analysis of rose resistance genes and their localization in the rose genome. *Acta Hort.* 651:123-130.
- Haymes, K. M., Van de Weg, W. E., Arens, P., Maas, J. L., Vosman, B., Den Nijs, A. P. M. 2000. Development of SCAR markers linked to a *Phytophthora fragariae* resistance gene and their assessment in European and North American strawberry genotypes. *J Am Soc Hortic Sci* 125:330–339
- Hemmat, M., Brown, S. K., Aldwinckle, H. S., Mehlenbacher, S. A., and Weeden, N. F. 2003. Identification and mapping of markers for resistance to apple scab from ‘Antonovka’ and ‘Hansen’s baccata #2. *Acta Hort.* 622:153-161.
- Hibrand-Saint Oyant, L., Crespel, L., Rajapakse, S., Zhang, L., and Foucher, F. 2008. Genetic linkage maps of rose constructed with new microsatellite markers and locating QTL controlling flowering traits. *Tree Genet. Genomes* 4:11-23.
- Ikeda, K., Watari, A., Ushijima, K., Yamane, H., Hauck, N. R., Iezzoni, A. F., and Tao, R. 2004. Molecular markers for the self-incompatible S<sub>4</sub> haplotype, a pollen-part mutant in sweet cherry (*Prunus avium* L.). *J. Amer. Soc. Hort. Sci.* 129:724-728.
- Iketani, H., Abe, K., Yamamoto, T., Kotobuki, K., Sato, Y., Saito, T., Terai, O., Matsuta, N., and Hayashi, T. 2001. Mapping of disease-related genes in Japanese pear using a molecular linkage map with RAPD markers. *Breed. Sci.* 51:179-184.

Inoue, E., Kasumi, M., Sakuma, f., Anzai, H., amino, K., and Hara, H. 2006. Identification of RAPD marker linked to fruit skin color in Japanese pear (*Pyrus pyrifolia* Nakai). *Sci. Hort.* 107:254-258.

Itai, A., Kotaki, T., Tanabe, K., Tamura, F., Kawaguchi, D., and Fukuda, M. 2003. rapid identification of 1-aminocyclopropane-i-carboxylate (ACC) sunthase genotypes in cultivars of Japanese pear (*Pyrus pyrifolia* Nakai) using CAPS markers. *Theor. Appl. Genet.* 106:1266-1272.

James, C. M., Clarke, J. B., and Evans, K. M. 2004. Identification of molecular markers linked to the mildew resistance gene *Pl-d* in apple. *Theor. Appl. Genet.* 110:175-181.

Jáuregui, B., 1998. PhD Thesis. University of Barcelona. Barcelona, Spain

Jáuregui, B., de Vincente, M.C., Messeguer, R., Felipe, A., Bonnet, A., Salesses, G., and Arús, P., 2001. A reciproca ; translocation between Garfi almond and Nemared peach. *Theor. Appl. Genet.* 102: 1169-1176.

Jia, Y. L., Wang, C. H., tian, Y. K., Dai, H. Y., and Wang, L. 2007. A SCAR marker linked to *PCDw* gene in pear. *Acta Hort. Sinica* 34:1531-1534.

Joobeur, T., Viruel, M.A., de Vincente, M.C., Jáuregui, B., Ballester, J., Dettori, M.J., Verdi, I., Truco, M.J., Messeguer, R., Battle, I., Quatar, R., Dirlwanger, E., and Arús, P., 1998. Construction of a saturated linkage map for *Prunus* using an almond x peach F<sub>2</sub> progeny. *Theor. Appl. Genet.* 97: 1034-1041.

Kellerhals, M., Dolega, E., Dilworth, E., Koller, B., and Gessler, C. 2000. Advances in marker-assisted apple breeding. *Acta Hort.* 538:535-540.

Kenis, K., and Keulemans, J. 2004. QTL analysis of growth characteristics in apple. *Acta Hort.* 663:369-374.

Kenis, K., and Keulemans, J. 2007. Study of tree architecture of apple (*Malus × domestica* Borkh.) by QTL analysis of growth traits. *Mol. Breed.* 19:193-208.

Khan, M. A., Durel, C. E., Duffy, B., Drouet, D., Kellerhals, M., Gessler, C., and Patocchi, A. 2007. Development of molecular markers linked to the 'Fiesta' linkage group 7 major QTL for fire blight resistance and their application for marker-assisted selection. *Genome* 50:568-577.

Khan, M. A., Duffy, B., Gessler, C., and Patocchi, A. 2006. QTL mapping of fire blight resistance in apple. *Mol. Breed.* 17:299-306.

Kim, D., Hwang, J. H., Shin, Y. U., Lee, H. J., Hong, S. S., and Kang, S. J. 2005. Development of molecular markers linked to several fruit traits in oriental pear. *Acta Hort.* 671:315-321.

King, G. J., Alston, F. H., Brown, L. M., Chevreau, E., Evans, K. M., Dunemann, F., Janse, J., Laurens, F., Lynn, J. R., Maliepaard, c., Manganaris, A. G., Roche, R., Schmidt, H., Tartarini, S., Verhaegh, J., and Vrielink, B. 1998. Multiple field and glasshouse assessments increase the reliability of linkage mapping of the *Vf* source of scab resistance in apple. *Theor. Appl. Genet.* 96:699-708.

King, G. J., Lynn, J. R., Dover, C. J., Evans, K. M., and Seymour, G. B. 2001. Resolution of quantitative trait loci for mechanical measures accounting for genetic variation in fruit texture of apple (*Malus pumila* Mill.). *Theor. Appl. Genet.* 102:1227-1235.

King, G. J., Maliepaard, C., Lynn, J. R., Alston, F. H., Durel, C. E., Evans, K. M., Griffon, B., Laurens, E., Manganaris, A. G., Schrevens, E., Tartarini, S., and Verhaegh, J. 2000. Quantitative genetic analysis and comparison of physical and sensory descriptors relating to fruit flesh firmness in apple (*Malus pumila* Mill.). *Theor. Appl. Genet.* 100:1074-1084.

Koller, B., Gianfranceschi, L., Seglias, N., McDermott, J., and Gessler, C. 1994. DNA markers linked to *Malus floribunda* 821 scab resistance. *Plant Mol. Biol.* 26:597-602.

Korban, S. S., and Chen, H. 1992. Apple. p. 203-227. In: F. A. Hammerschlag and R. E. Litz (eds.) . *Biotechnology of perennial fruit crops*. CAB International, Wallingford, UK.

Lawson, D. M., Hemmat, M., and Weeden, N. F. 1995. The use of molecular markers to analyze the inheritance of morphological and developmental traits in apple. *J. Amer. Soc. Hort. Sci.* 120:532-537.

Lerceteau-Köhler, E., Moing, A., Guérin, G., Renaud, C., Courlit, S., Camy, D., Praud, K., Parisy, V., Bellec, F., Maucourt, M., Rolin, D., Roudeillac, P., and Denoyes-Rothan, B. 2004. QTL analysis for fruit quality traits in octoploid strawberry (*Fragaria × ananassa*). *Acta. Hort.* 663: 331-336.

Lesemann, S., and Dunemann, F. 2006. Recent findings on the biodiversity of the apple powdery mildew pathogen. *Gesunde-Pflanzen* 58:117-123.

Liebhard, R., Kellerhals, M., Pfammatter, W., Jertmini, M., and Gessler, C. 2003a. Mapping quantitative physiological traits in apple (*Malus × domestica* Borkh.) *Plant Mol. Biol.* 52:511-526.

Liebhard, R., Koller, B., Patocchi, A., Kellerhals, M., Pfammatter, W., Jermini, M., and Gessler, C. 2003b. Mapping quantitative field resistance against apple scab in a 'Fiesta' × 'Discovery' progeny. *Phytopathol.* 93:493-501.

Linde, M., Hattendorf, A., Kaufmann, H., and Debener, T. 2006. Powdery mildew resistance in roses: QTL mapping in different environments using selective genotyping. *Theor. Appl. Genet.* 113:1081-1092.

Linde, M., Mattiesch, L., and Debener, T. 2004. *Rpp1*, a dominant gene providing race-specific resistance to rose powdery mildew (*Podosphaera pannosa*): molecular mapping, SCAR development and confirmation of disease resistance data. *Theor. Appl. Genet.* 109:1261-1266.

Liu, X., Reighard, G., Swire-Clark, G., Bridges, W., and Baird, V. 2008. Mapping the nuclear genomic region associated with the peach tree short life syndrome using microsatellite/SSR markers. *HortSci.* 43:1270.

Maliapaard, C., Alston, F. H., van Arkel, G., Brown, L. M., Chevreau, E., Dunemann, F., Evans, K. M., Gardiner, S., Guilford, P., van Heusden, A. W., Janse, J., Laurens, f., Lynn, J. R., Manganaris, a. G., den Nijs, A. P. M., Periam, N., Rikkerink, E., Roche, P., Ryder, C., Sansavini, S., Schmidt, H., Tartarini, S., Verhaegh, J. J., vrielink-vanginkel, M., and King, G. J. 1998. Aligning male and female linkage maps of apple (*Malus pumila* Mill) using multi-allelic markers. *Theor. Appl. Genet.* 97:60-73.

Maliapaard, C., Sillanpaa, M. J., van Ooijen, J. W., Jansen, R. C., Arjas, E. 2001. Bayesian versus frequentist analysis of multiple quantitative trait loci with an application to an outbred apple cross. *Theor. Appl. Genet.* 103:1243-1253

- Ogundiwin, E. A., Peace, C. P., Nicolet, C. M., Rashbrook, V. K., Gradziel, T. M., Bliss, F. A., Parfitt, D., and Crisosto, C. H. 2008. Leucoanthocyanidin dioxygenase gene (PpLDOX): a potential functional marker for cold storage browning in peach. *Tree Genet. Genomes* 4:543-554.
- Patocchi, A., Bigler, B., Koller, B., Kellerhals, M., and Gessler, C. *Vr<sub>2</sub>*: a new apple scab resistance gene. *Theor. Appl. Genet.* 109:1087-1092.
- Patocchi, A., Walser, M., Tartarini, S., Brogini, G. A. L., Gennari, F., Sansavini, S., and Gessler, C. 2005. Identification by genome scanning approach (GSA) of a microsatellite tightly associated with the apple scab resistance gene *Vm*. *Genome* 48:630-636.
- Peace, C. P., Crisosto, C. H., and Gradziel, T. M. 2005. Endopolycturonase: a candidate gene for *Freestone* and *Melting flesh* in peach. *Mol. Breed.* 16:21-31.
- Peil, A., Garcia-Libreros, T., Richter, K., Trognitz, F. C., Hanke, M.-V., and Flachowsky, H. 2007. Strong evidence for a fire blight resistance gene of *Malus robusta* located on linkage group 3. *Plant Breed.* 126:470-476.
- Pierantoni, L., Dondini, L., Cho, K. H., Shin, I. S., Gennari, F., Chiodini, R., tartarini, S. Kang, S. J., and Sansavini, S. 2007. Pear scab resistance QTLs via a European pear (*Pyrus communis*) linkage map. *Tree Genet. Genomes* 3:311-317.
- Rajapakse, S., Byrne, D. H., Zhang, L., Anderson, N., Arumuganathan, K., and Ballard, R. E. 2001. Two genetic linkage maps of tetraploid roses. *Theor. Appl. Genet.* 103:575-583.
- Rusholme-Pilcher, R. L., Celton, J.-M., Gardiner, S. E., and Tustin, D. S. 2008. Genetic markers linked to the dwarfing trait of apple rootstock 'Malling 9'. *J. Amer. Soc. Hort. Sci.* 2008 133: 100-106.
- Rusu, A. R., Pamfil, D., and Graham, J. 2006. Mapping resistance of red raspberry (*Rubus idaeus* subsp. *idaeus*) to viral diseases – Leaf spot (RLSV) and Vein chlorosis (RVCV) on the genetic linkage map. *Bul. Univ. Stiinte Agric. Med. Vet.-Cluj-Napoca, Ser. Zoo. Biotehn.* 62:318.
- Sánchez-Perez, R., Dicenta, F., and Martínez-Gómez, P. 2004. Identification of S-alleles in almond using multiplex PCR. *Euphytica* 138:263-269.

Sargent, D.J., Fernández-Fernández, F., Rys, A., Knight, V. H., Simpson, D. W., and Tobutt, K. R. 2007. Mapping of *A<sub>1</sub>* conferring resistance to the aphid *Amphorophora idaei* and *dw* (dwarfing habit) in red raspberry (*Rubus idaeus* L.) using AFLP and microsatellite markers. *BMC Plant Biol.* 7:15 (doi:10.1186/1471-12229-7-15)

Sargent, D. J., Davis, T. M., Tobutt, K. R., Wilkinson, M. J., Battey, N. H., Simpson, D. W. 2004. A genetic linkage map of microsatellite, gene specific and morphological markers in diploid *Fragaria*. *Theor. Appl. Genet.* 109:1385–1391.

Sicard, O., Marandel, G., Sorinao, J. M., Lalli, D. A., Lambert, P., Salava, J., Badenes, M. L. Abbott, A., and Decroocq. 2008. Flanking the major *Plum pox virus* resistance locus in apricot with co-dominant markers (SSRs) derived from candidate resistance genes. *Tree Genet. Genomes* 4:359-365.

Stankiewicz-Kosyl, M., Pitera, E., and Gawronski, S. W. 2005. Mapping QTL involved in powdery mildew resistance of the apple clone U 211. *Plant Breed.* 124:63-66.

Stoeckli, S., Mody, K., Gessler, C., Patocchi, A., Jermini, M., and Dorn, S. 2008. QTL analysis for aphid resistance and growth traits in apple. *Tree Genet. Genomes online* doi: 10.1007/s11295-008-0156-y.

Tartarini, S., Gennari, F., Pratesi, D., Palazzetti, C., Sansavini, S., Parisi, L., Fouillet, A., Fouillet, V., and Durel, C. E. 2004. Characterization and genetic mapping of a major scab resistance gene from the old Italian apple cultivar ‘Durello di Forli’. *Acta Hort.* 663:129-133.

Terakami, S., Adachi, Y., Iketani, H., Sato, Y., Sawamura, Y., Takada, N., MNishitani, C., and Yamamoto, T. 2007. Genetic mapping of genes for susceptibility to black spot disease in Japanese pears. *Genome* 50:735-741.

Terakami, S., Shoda, M., Adachi, Y., Gonai, T., Kasumi, M., Sawamura, Y., Iketani, H., Kotobuki, K., Patocchi, A., Gessler, C., Hayashi, T., and Yamamoto, T. 2006. Genetic mapping of the pear scab resistance gene *Vnk* of Japanese pear cultivar Kinchaku. *Theor. Appl. Genet.* 113:743-752.

Van de Weg, W. E. 1997. Resistance to *Phytophthora fragariae* var. *fragariae* in strawberry: The *Rpf2* gene. *Theor. Appl. Genet.* 94: 1092–1096.

Van de Weg, W.E., Haymes, K.M., Kodde, L.P., Maas J.L. (unpublished): A SCAR marker tightly to the *Rpf2* resistance gene conferring resistance to *Phytophthora fragariae* and its assessment in European and North American strawberry cultivars. Cooperative research between ARS-Beltsville and Wageningen-UR, The Netherlands. Manuscript to be submitted in 2009. Gene described in Van de Weg, 1997.

Verde, I., Quarta, R., Cedrola, C., and Dettori, M. T. 2002. QTL analysis of agronomic traits in a BC<sub>1</sub> peach population. *Acta Hort.* 592:291-297.

Vinatzer, B. A., Patocchi, A., Tartarini, S., Gianfranceschi, L., Sansavini, s., and Gessler, C. 2004. Isolation of two microsatellite markers from BAC clones of the *Vf* scab resistance region and molecular characterization of scab-resistant accession in *Malus* germplasm. *Plant Breed.* 123:321-326.

Viruel, M. A., Madur, D., Direlwanger, E., Pascal, T., and Kervella, J. 1998. Mapping quantitative trait loci controlling peach leaf curl resistance. *Acta Hort.* 465:79-87.

Von Malek, B., Weber, W. E., and Debener, T. 2000. Identification of molecular markers linked to *Rdr1*, a gene conferring resistance to blackspot in roses. *Theor. Appl. Genet.* 101:977-983.

Wang, D., Karle, R., and Iezzoni, A. F. 2000. QTL analysis of flower and fruit traits in sour cherry. *Theor. Appl. Genet.* 100:535-544.

Weebadde, C. K., Wang, D., Finn, C. F., Lewers, K. S., Luby, J. J., Bushakra, J., Sjulín, T. M., and Hancock, J. F. 2008. Using a linkage mapping approach to identify QTL for day-neutrality in the octoploid strawberry. *Plant Breed.* 7:94-101.

Weeden, N. F., Hemmat, M., Lawson, D. M., Lodhi, M., Bell, R. L., Manganaris, A. G., Reisch, B. I., Brown, S. K., and Ye, G. N. 1994. Development and application of molecular marker linkage maps in woody fruit crops. *Euphytica* 77:71-75.

Whitaker, V. M., and Hokanson, S. C. 2008. The inheritance of race-specific black spot resistance in tetraploid roses. *HortSci.* 43:1101.

Yamamoto, T., Kimura, T., Shoda, M., Imai, T., Saito, T., Sawamura, Y., Kotobuki, K., Hayashi, T., and Matsuta, N. 2002. genetic linkage maps constructed by using an interspecific cross between Japanese and European pears. *Theor. Appl. Genet.* 106:9-18.

Yamamoto, T., Kimura, T., Terakami, S., Nishitani, C., Sawamura, Y., Saito, T., Kotobuki, K., and Hayashi, T. 2007. Integrated reference genetic linkage maps of pear based on SSR and AFLP markers. *Breed. Sci.* 57:321-329.

Yan, Z., Visser, P. B., Hendriks, T., Prins, T. W., Stam, P., and Dolstra, O. 2007. QTL analysis of variation for vigour in rose. *Euphytica* 154:53-62.

Zini, E., Biasioli, F., Gasperi, F., Mott, d., Aprea, E., Märk, T. D., Patocchi, A., Gessler, C., and Komjanc, M. 2005. QTL mapping of volatile compounds in ripe apples detected by proton transfer reaction-mass spectrometry. *Euphytica* 145:269-279.