



Mapping and Characterizing of Simple and Complex Traits in Autotetraploid Roses (*Rosa hybrida* L.)

Yaniv Semel – EPPN summer school

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The Hebrew University of Jerusalem

Introduction

- Family: **Rosaceae** (*Prunus*, *Malus*, *Pyrus*....); Subfamily: **Rosoideae** (*Fragaria*, *Rubus*, ...); Genus: **Rosa** (over 100 species)
- Chromosome numbers in the genus **Rosa** are based on multiples of seven and range from $2n=2x=14$ to $2n=8x=56$
- Species: ***R. hybrida* L.** - The cultivated roses, perennial shrubs, highly heterozygosity, mostly autotetraploids ($2n=4x=28$), with a small genome estimated at about 550 Mb (0.57 pg/1C)
- Roses have been admired for their beauty and fragrance since the dawn of civilization and there are evidences that roses were cultivated 5000 years ago by the ancient civilization of China.
- The rose accounts for two-thirds of the international cut floral industry which exceeds 40 billion dollars per year.
- Surprisingly, **little is known about the rose genetics** because of **technical factors that make the rose difficult model for genetic analysis**: high heterozygosity, difficulties in sexual reproduction from pollination to seed germination and high ploidy levels.

Objectives

- Constructing genetic linkage maps of autotetraploid roses.
- Characterizing and mapping of simple and complex traits in autotetraploid roses.

Autotetraploid Linkage maps

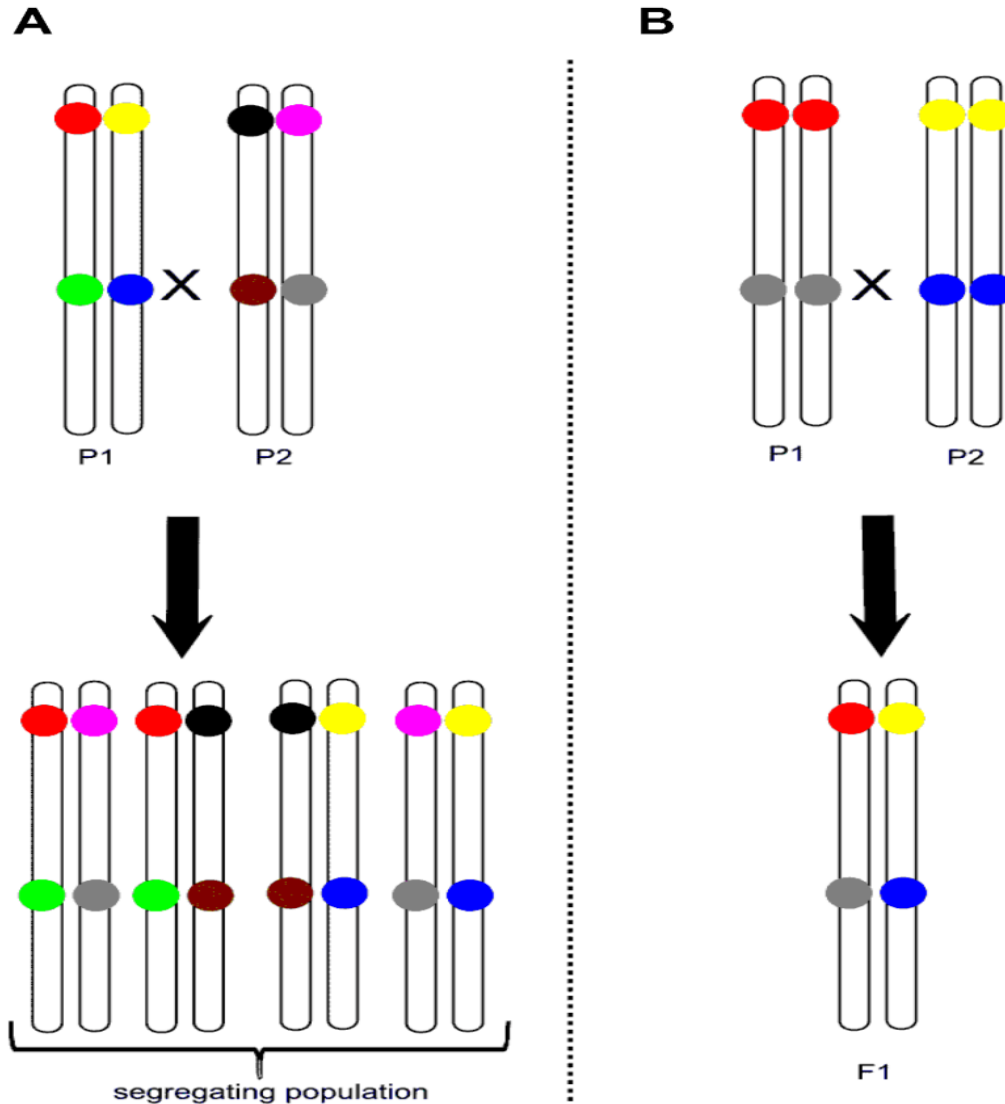


- **Golden Gate**[®] (GG) - Kordes
- Cutting variety
- Yellow- Carotenoids
- Sensitive to pests and disease
- Barely prickled
- Small-medium bud
- Lack a distinct odor
- Vase life – 12-14 days

- **Fragrant Cloud**[®] (FC)- Tantau
- Garden variety
- Red peach- Flavonoids
- Resistant to pests and disease
- Highly prickled
- Large bud
- Strong fruity scent
- Vase life - 5 days

Mapping population of 132 siblings- “GGFC”

Autotetraploid Mapping- Double pseudo testcross strategy



Gar et al. (2011)

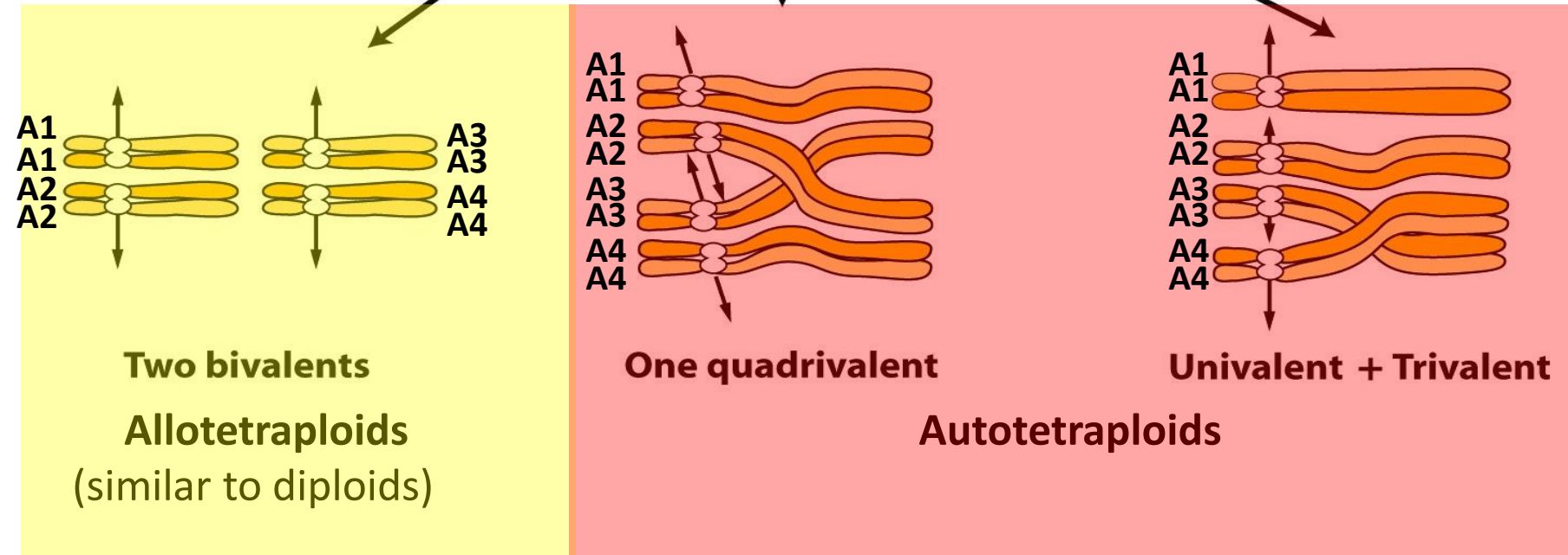
A close-up, top-down view of a dense arrangement of roses. The roses are in various stages of bloom, showing intricate petal patterns. The color palette is diverse, featuring bright yellows, warm oranges, soft pinks, deep purples, and hints of red. The lighting is even, highlighting the texture of the petals.

GGFC population

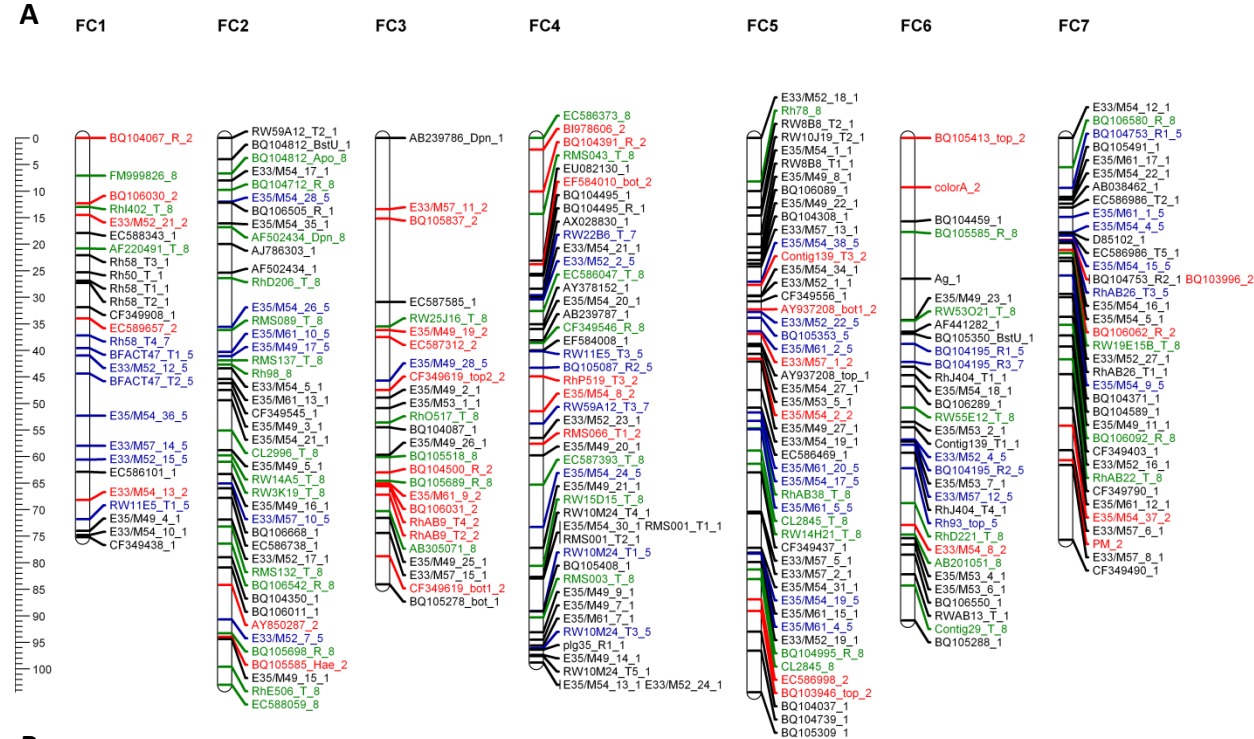
A1 A
A2 A
A3 a
A4 a

$$\textcircled{\times} \longrightarrow P(\text{aaaa}) = 1/36$$

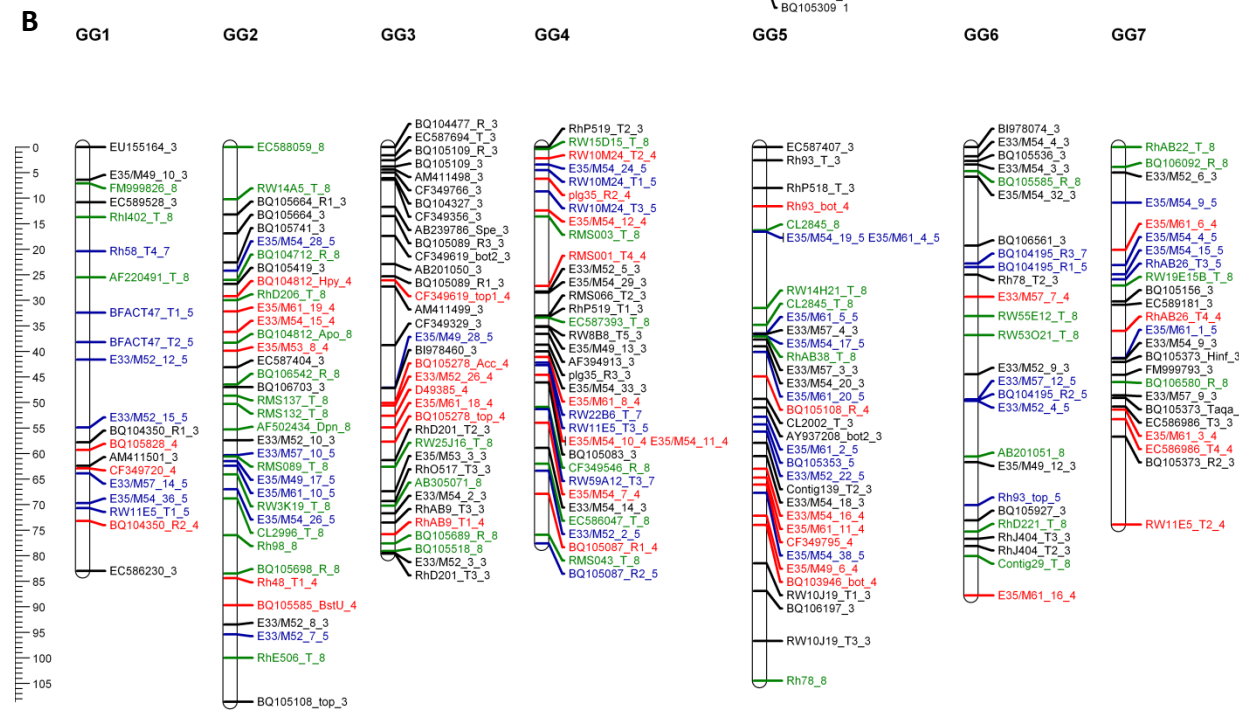
Pairing possibilities



- Simple genetic traits segregate up to 35:1 ratio compared to 3:1 in diploids models



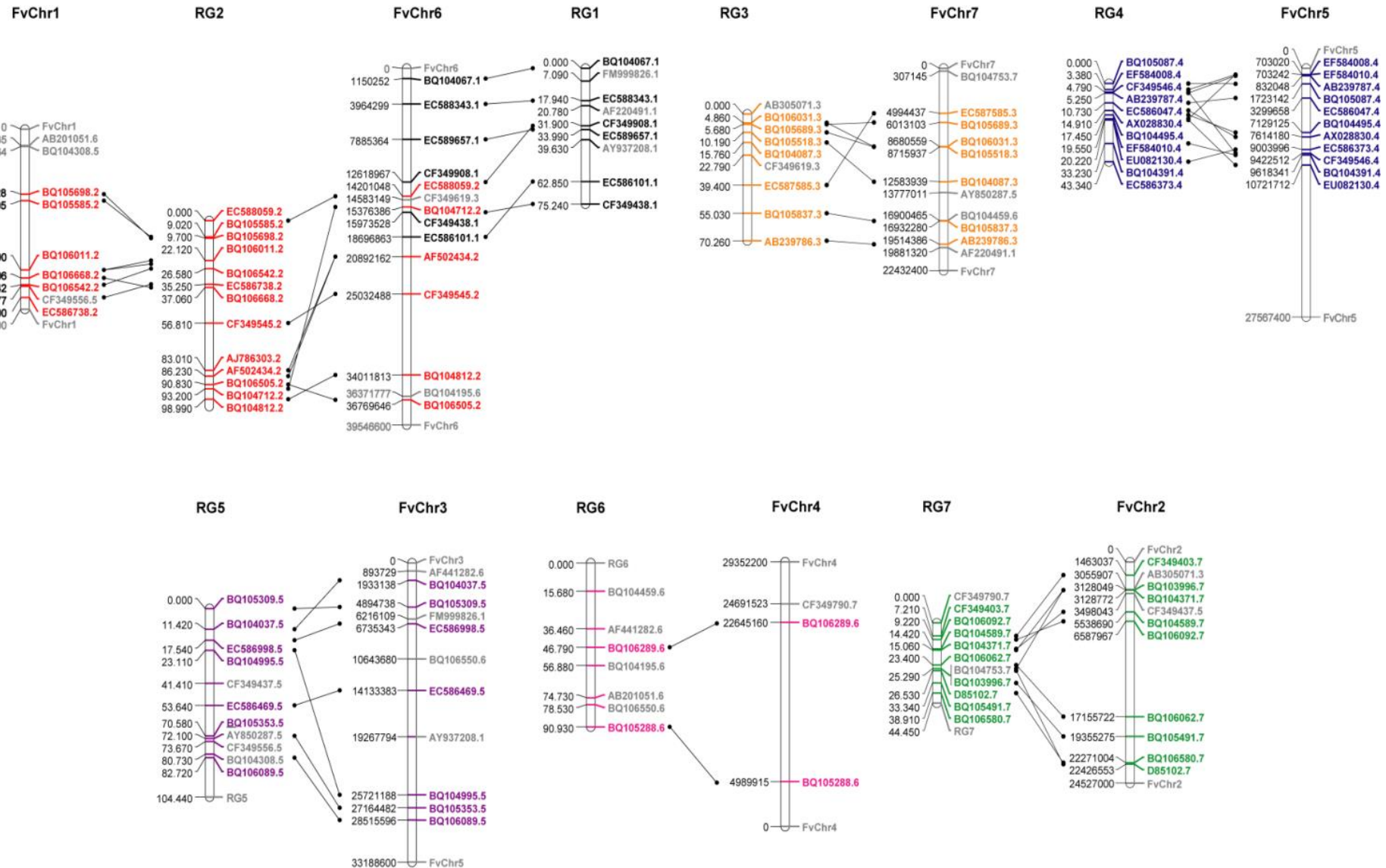
- FC- 259 markers covering 632 cM (A)
- GG- 210 markers covering 616 cM (B)
- 111 common markers.



- 128 sequence-based markers.

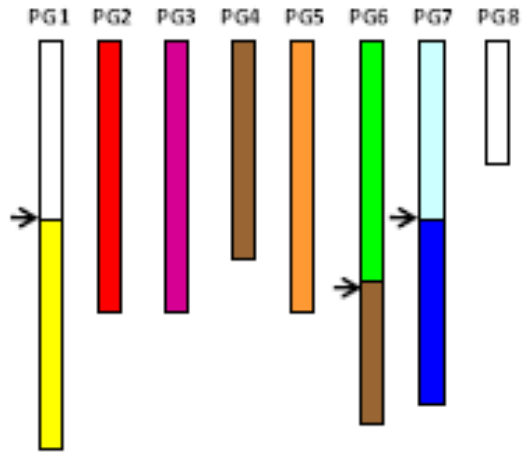
Gar et al. (2011)

Map validation- The synteny between *Rosa* and *Fragaria*

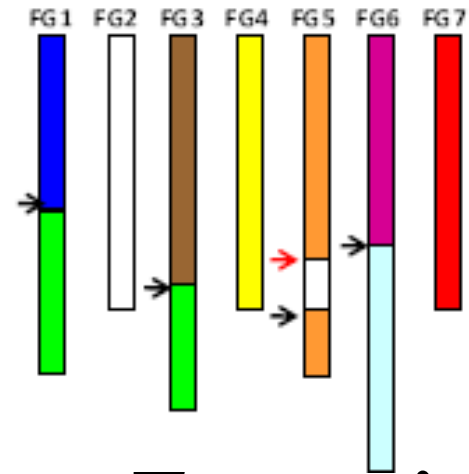


Gar et al. (2011)

Macro-synteny

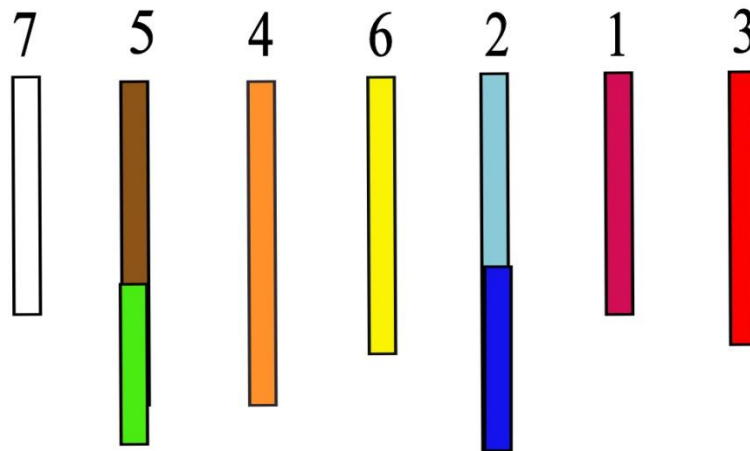


Prunus



Fragaria

Jung et al. (2012)



Rosa

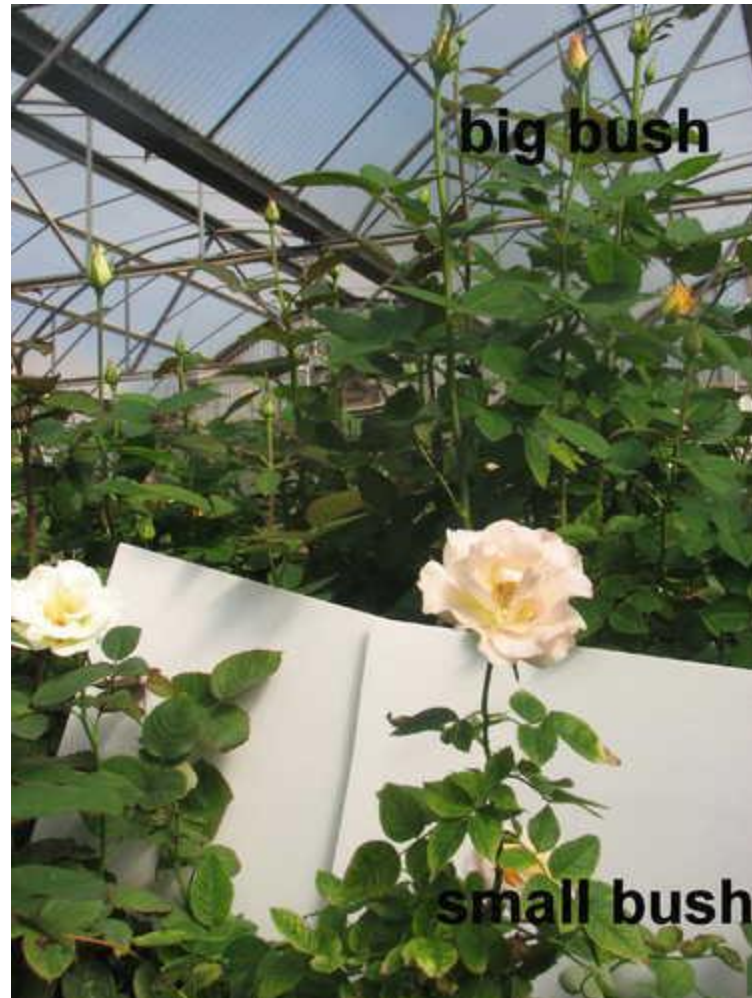
A close-up photograph of a yellow rose, showing the intricate layers of its petals. The petals are a vibrant yellow, with some areas appearing slightly lighter or more saturated than others, creating a sense of depth and texture. The word "Phenotyping" is overlaid in the center of the image in a bold, black, sans-serif font. The text is positioned horizontally and is clearly legible against the background of the rose's petals.

Phenotyping

Vase life experiments



Bush growth and development

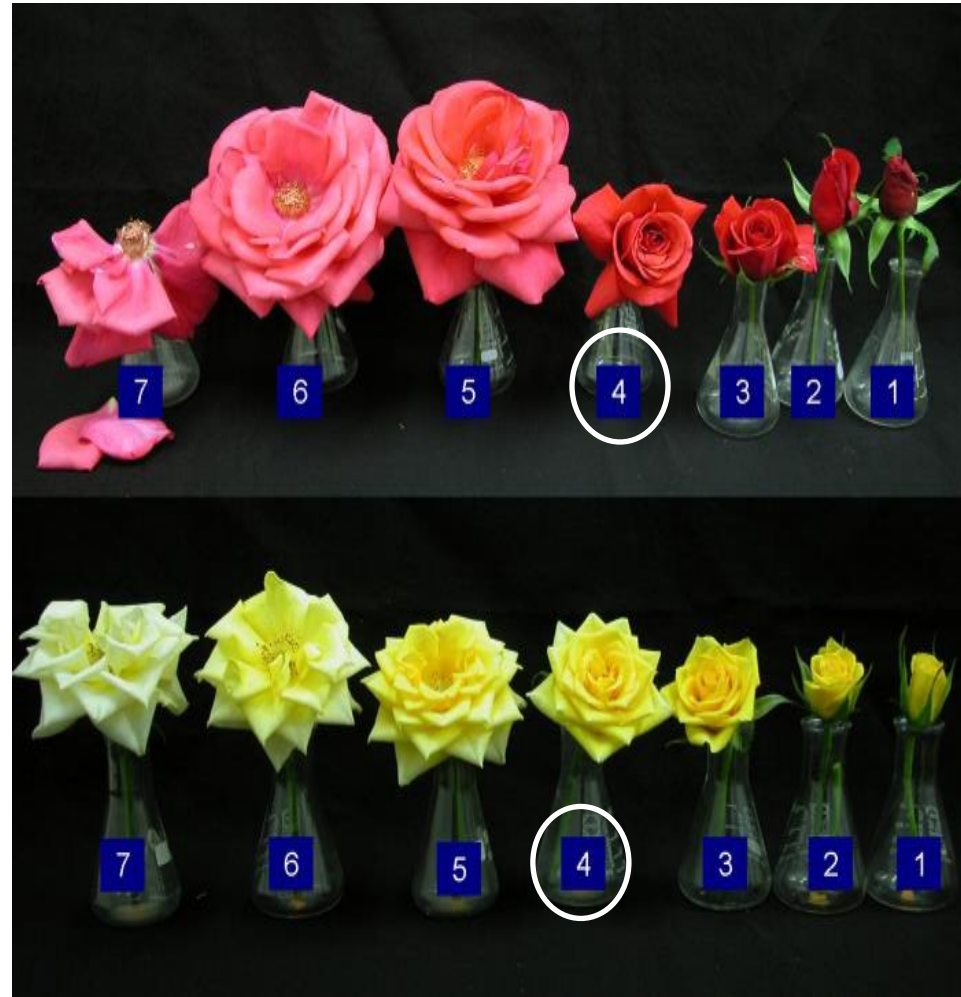


Number of prickles

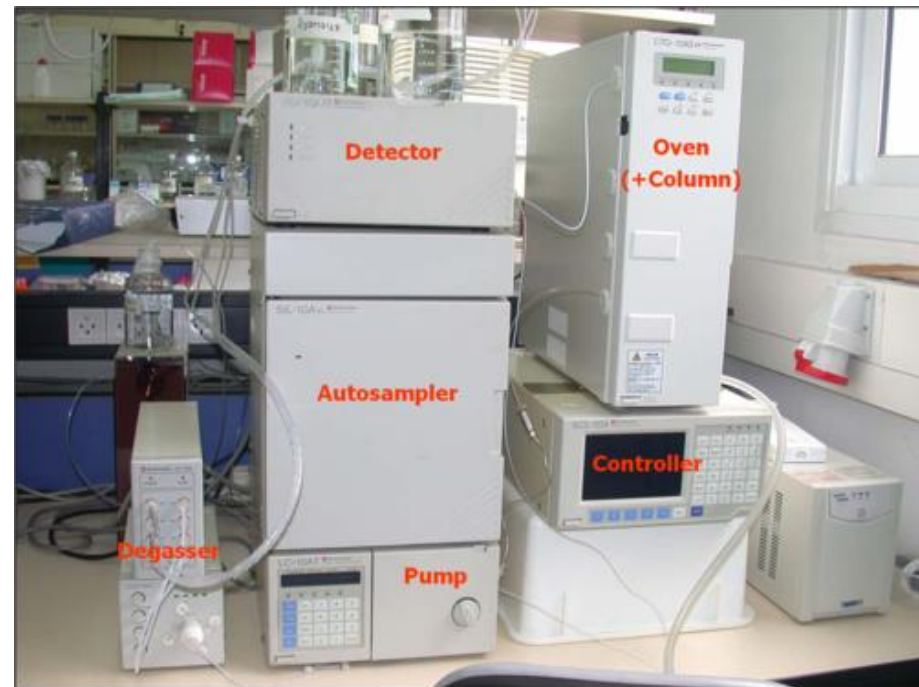
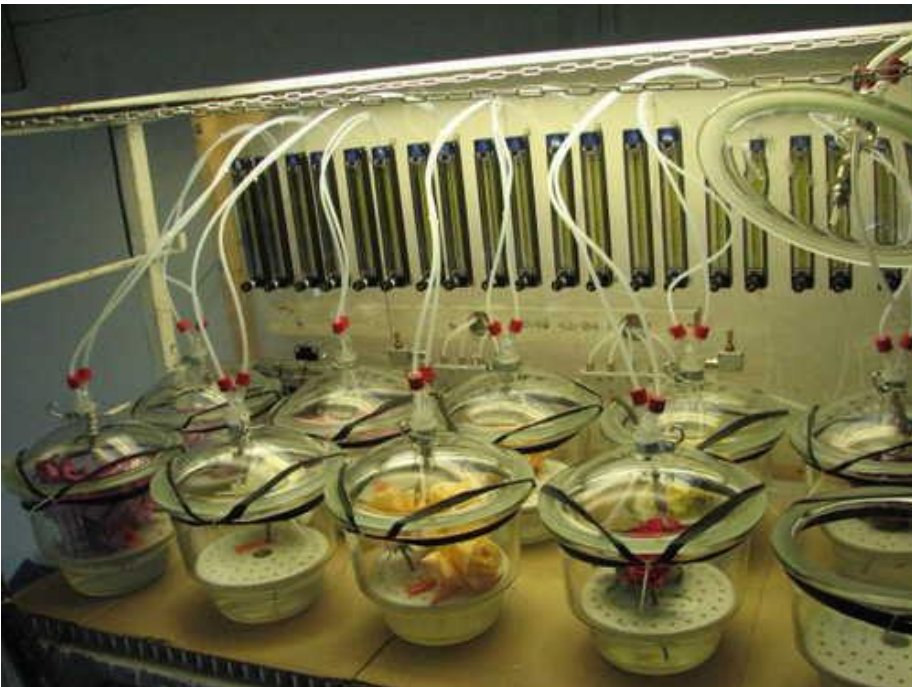


Metabolic profiling of rose scent

- Flowers at developmental stage 4 were collected.
- Some of which were analyzed for volatiles content using a GC-MS headspace.
- From the remained flowers, samples of **Petal tissues** (stage 4) were collected and sent to primary and secondary metabolic profiling.



Volatiles measured by GCMS (gas chromatograph mass spectrometry)



Secondary metabolites measured by LCMS (liquid chromatograph mass spectrometry)



Large Scale Phenotyping in Tetraploid Roses as a Tool for Locating and Mapping Traits at the Whole Plant Level

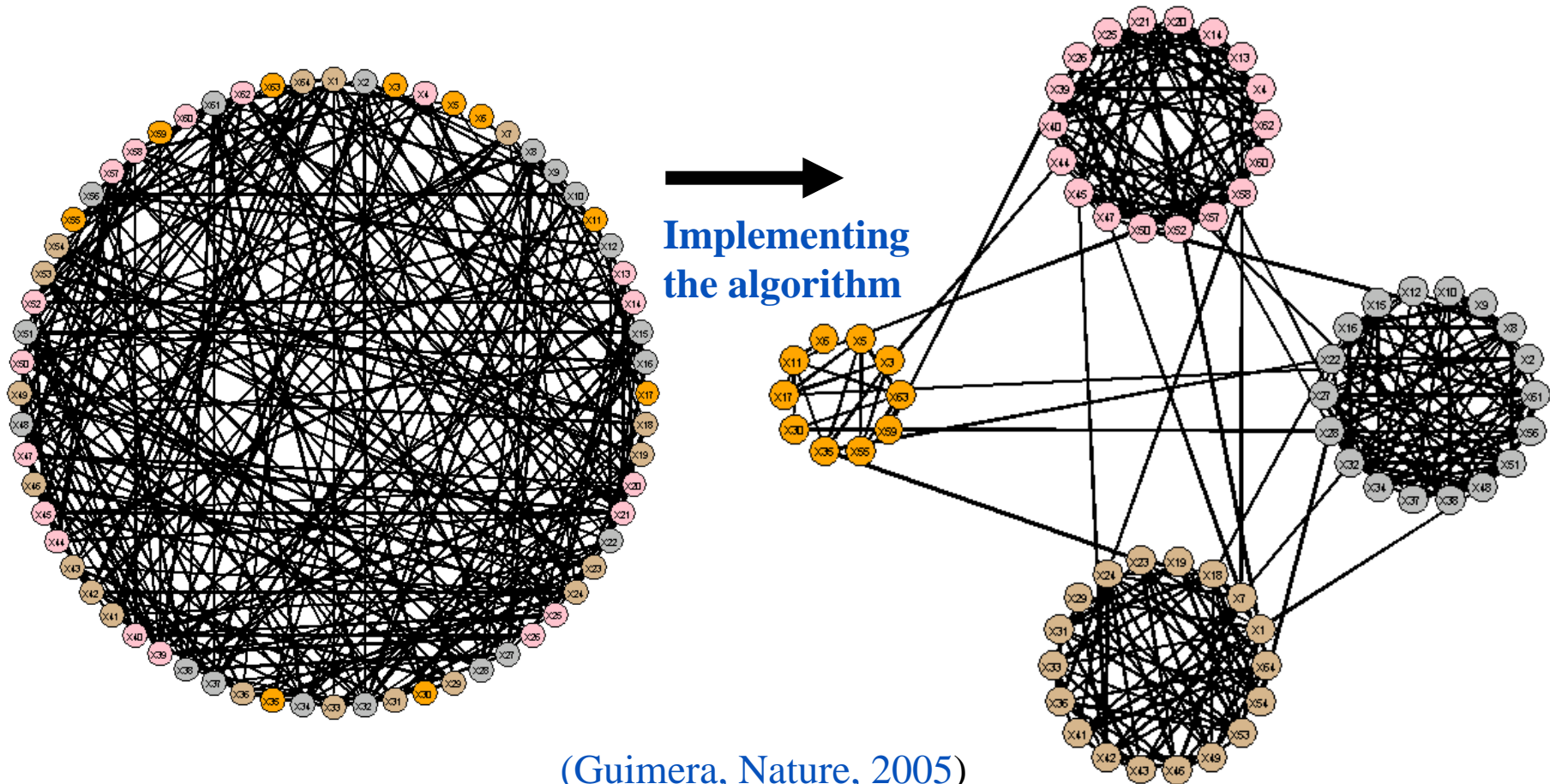
Character Groups	Number of characters	Collaboration	Measuring Tool
Flavonoids	8	Michal Oren-Shamir- Agricultural Research Organization	HPLC
Secondary Metabolites	116	Asaph Aharoni- Weizmann Institute	LCMS
Anatomy and Morphology	54		Manually
Fertility	4		Manually
Growth and Development	28		Manually
Quality	17		Manually
Yield	15		Manually
Primary Metabolites	85	Alisdair R Fernie- Max-Planck institute	GCMS
		Aaron Fait- Ben Gurion University	GCMS
Carotenoids	12	Joseph Hirschberg- Hebrew University	HPLC
Volatiles	102	Eyal Fridman- Hebrew University	GCMS –Headspace
Total	441		

Genotypes










Cartography of complex networks

The algorithm finds modules in networks. Each module represents a subset of nodes that are connected to each other more than to nodes in other modules.

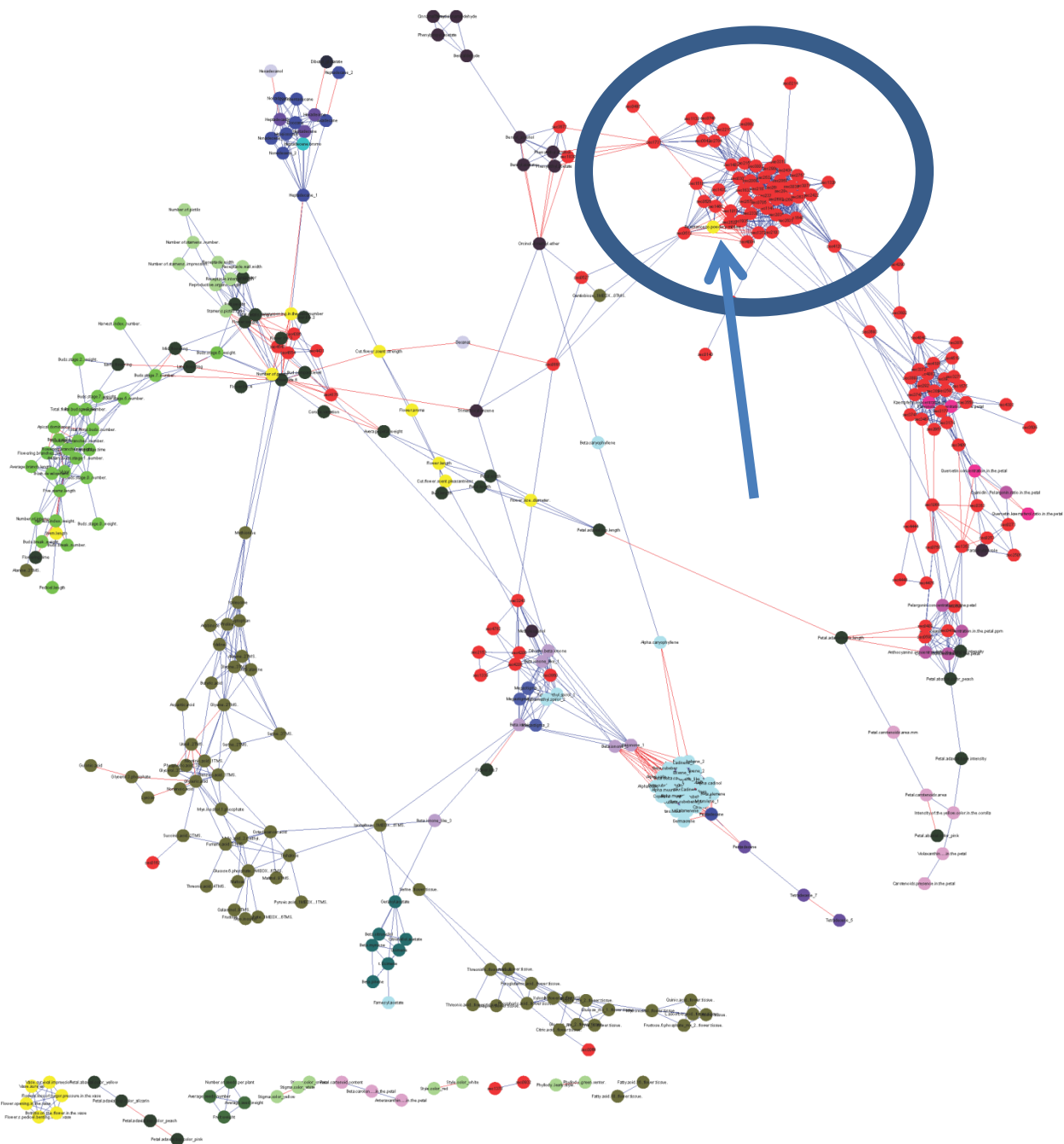
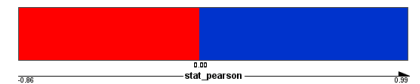


Node Label is displayed as ID

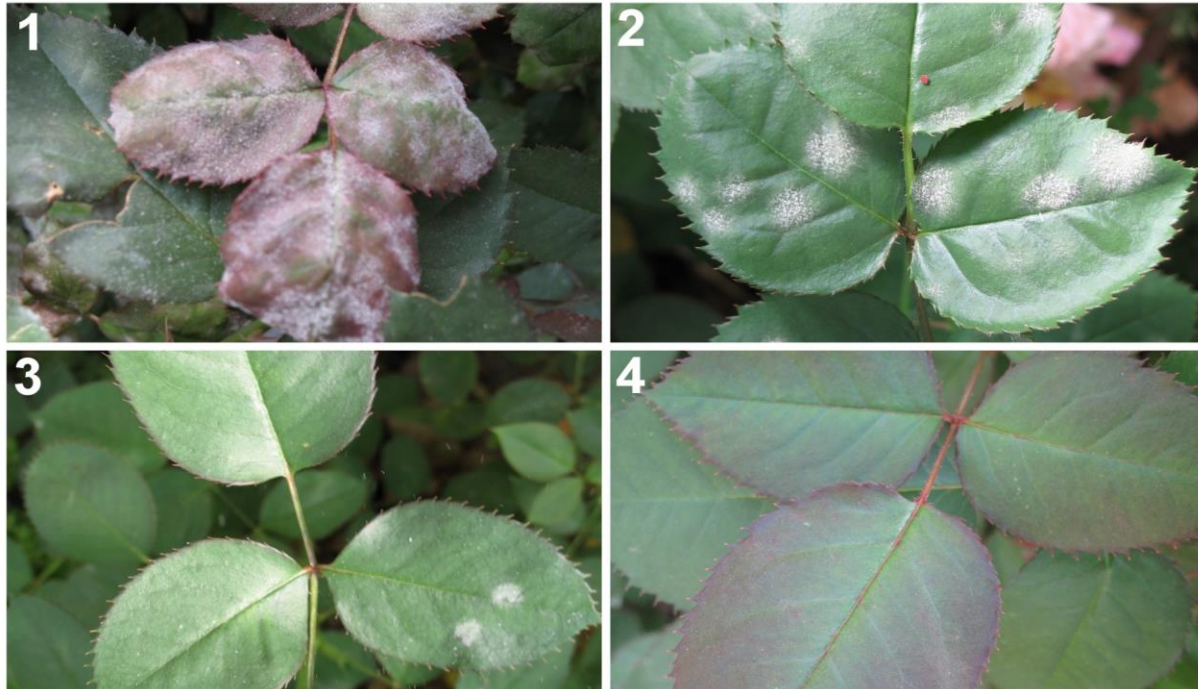
Node Color Mapping

Node Color	Column 2
	Biochemistry- Anthocyanins
	Biochemistry- Carotenoids
	Biochemistry- Flavonols
	Fertility
	Growth and development- Bush
	Growth and development- Flower
	Metabolites- Primary (Fat A.)
	Metabolites- Primary (Zamir D.)
	Metabolites- Secondary (Aharoni A.)
	Quality- Breeding
	Quality- Cutting product
	Quality- Resistances
	Quality- Scent
	Quality- Vase life
	Volatiles- Alkane
	Volatiles- Apocarotenoid
	Volatiles- Fatty acid derivative
	Volatiles- Halocarbon
	Volatiles- Monoterpene
	Volatiles- Norisoprene
	Volatiles- Olefin
	Volatiles- Organic acid
	Volatiles- Phenolic derivative
	Volatiles- Sesquiterpene
	Anatomy and morphology- Flower
	Anatomy and morphology- Leaf
	Anatomy and morphology- Prickles
	Anatomy and morphology- Reproductive organs
	Yield

Edge Color Mapping



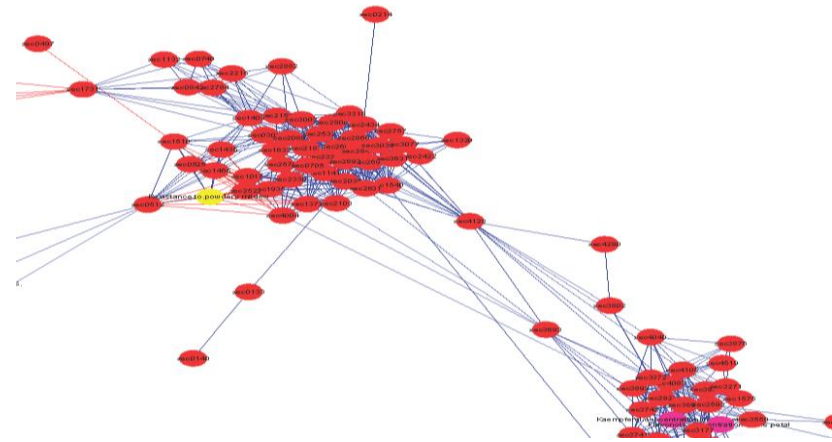
- **Powdery Mildew** disease in roses caused by the fungi *Sphaerotheca pannosa* (Wallr) Lev. var. *rosae* War. is one of the most common disease globally , damaging both greenhouse and open field roses.



Score	Disease Coverage (%)	Infected Tissues
1-	50-90	all plant tissues
2-	35-50	leaves, apices, sepals, receptacles, sometime stems pedicels and petals
3-	5-34	leaves, apexes, sometime sepals and receptacles
4-	0-4	only leaves

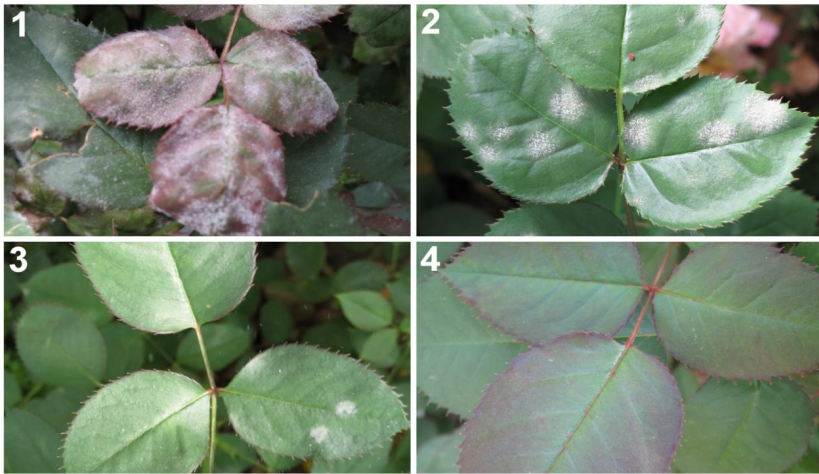
Correlation between some secondary metabolites and resistance to Powdery Mildew

- The LC-MS run yield 4807 fragments that were name secXXXX.
- By using statistical we were able to reduce it to 116 representative fragments.
- We detect seven fragments showing strong correlation with the level of the Powdery Mildew resistance.



Correlation between some secondary metabolites and resistance to Powdery Mildew

Powdery Mildew (leafs)



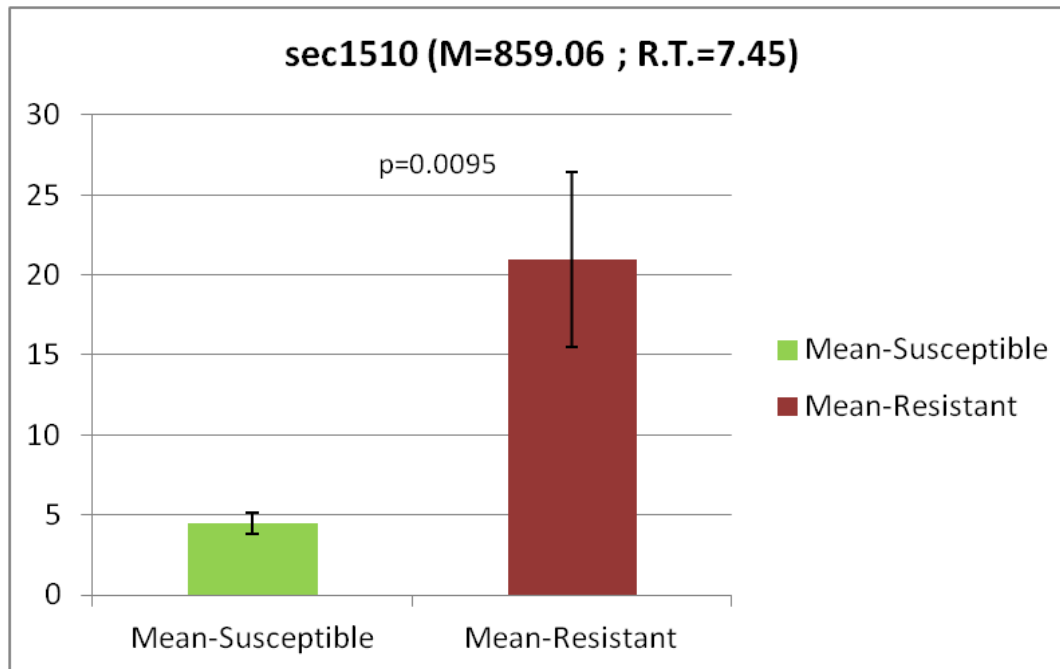
**Secondary metabolite
(flowers)**



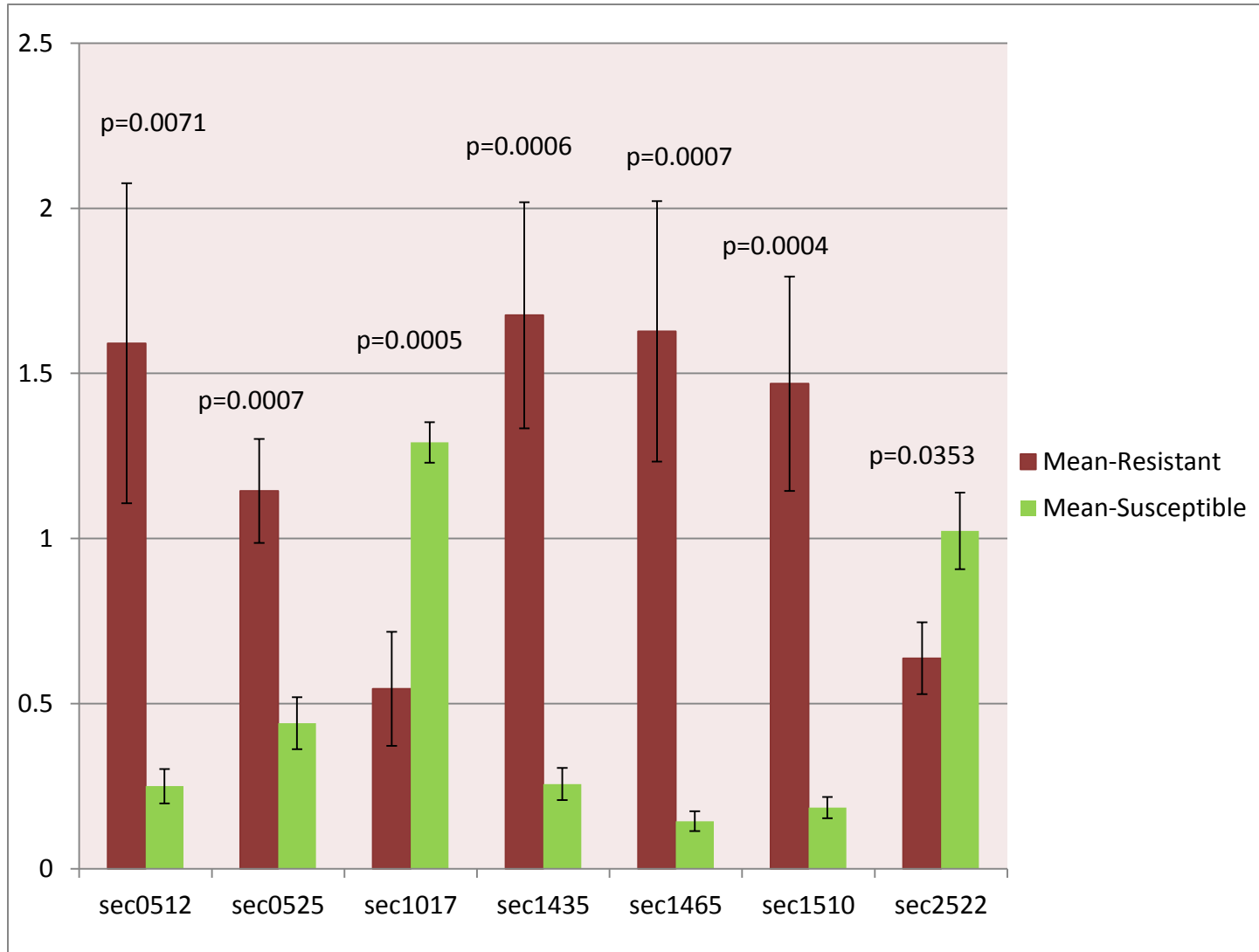
Second experiment

- To check the repeatability of the results in other rose varieties and to check the relationship with powdery mildew resistance.
- 15 resistant and 15 susceptible genotypes were selected.

Sec 1510



Petals LCMS Analysis- First run results for the selected varieties



Fragment 6

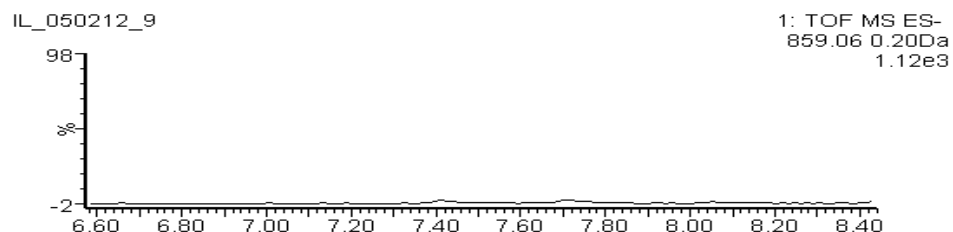
Petals LCMS Analysis

Name: **sec1510**

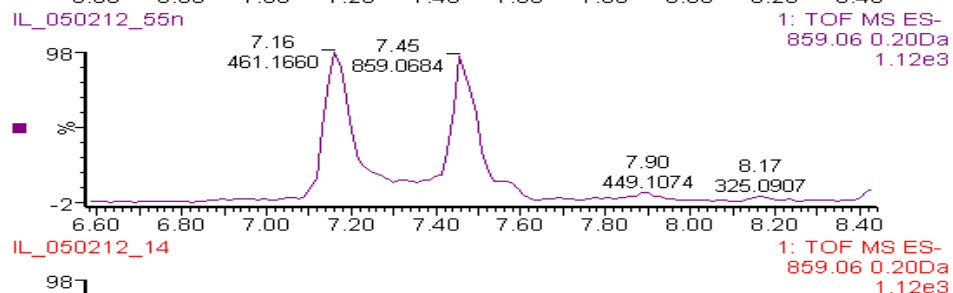
Comment: RT=7.16 & MZ=859.06

Formula: **C₇₅H₅₂O₄₈**

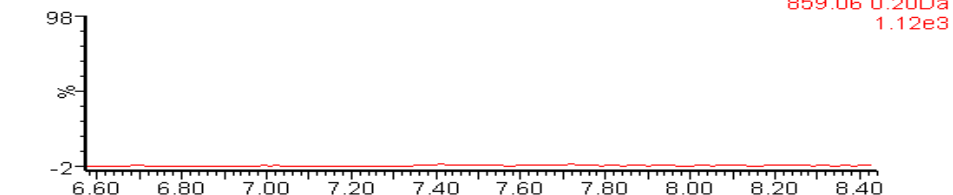
GF26-
susceptible



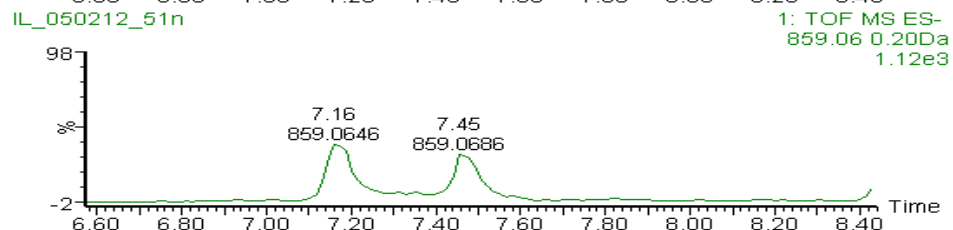
GF99-
resistant



GF97-
susceptible

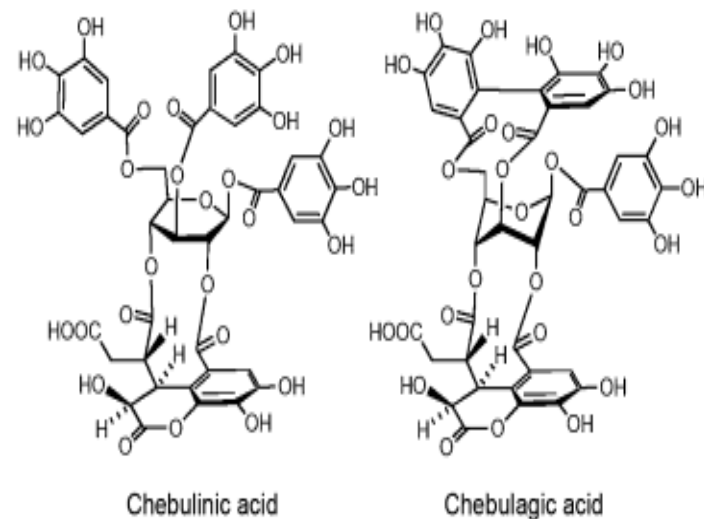


GF98-
resistant



- Following further analysis of the seven fragments according to their mass and R.T and the data of the related mass we determined that all fragments are hydrosable tannins.
- To be more specific **Ellagitannins.**

Chemistry and Biology of Ellagitannins



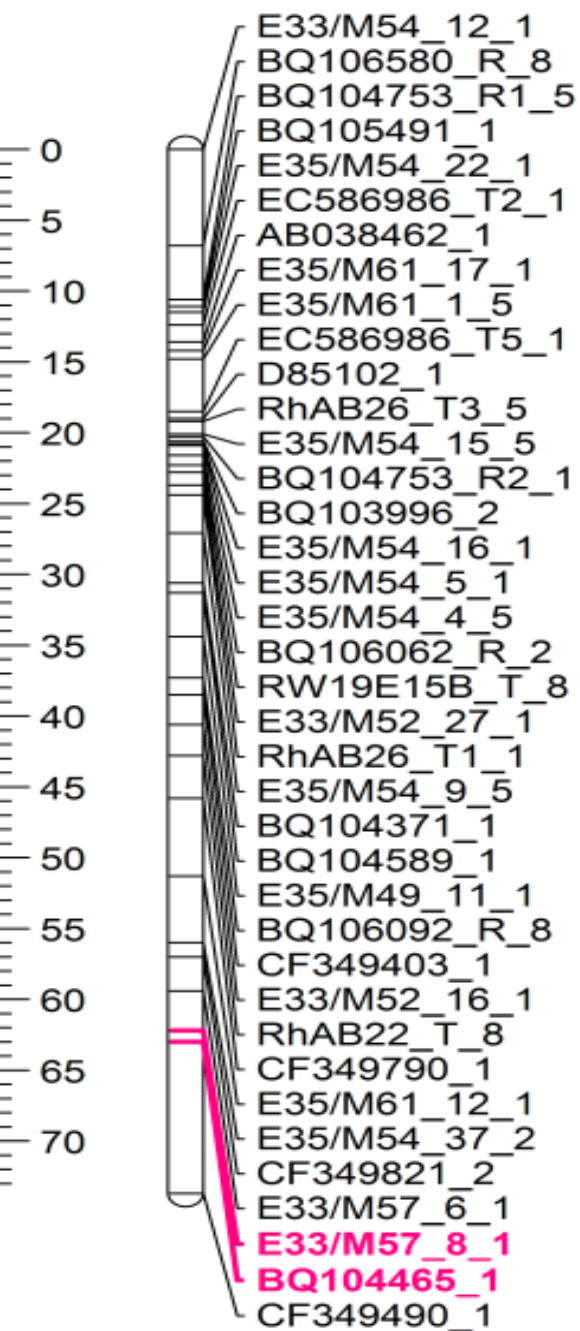
Ellagitannins in Rosaceae and Resistance to Powdery Mildew

- Ellagitannins may be important for the resistance to powdery mildew in strawberries as their accumulation was strong and rapid enough to inhibit the initial phases of infection.
- No firm conclusions can be drawn yet concerning the functions of ellagitannins in plant defense against pathogens.
- However, antimicrobial, antioxidant and protein binding capacity of ellagitannins suggest a protective role in plants.



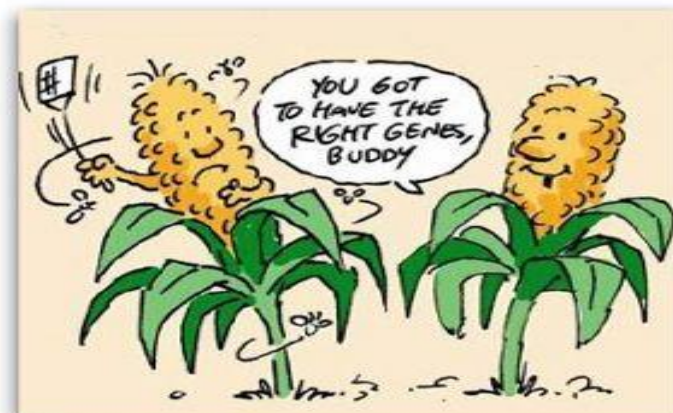
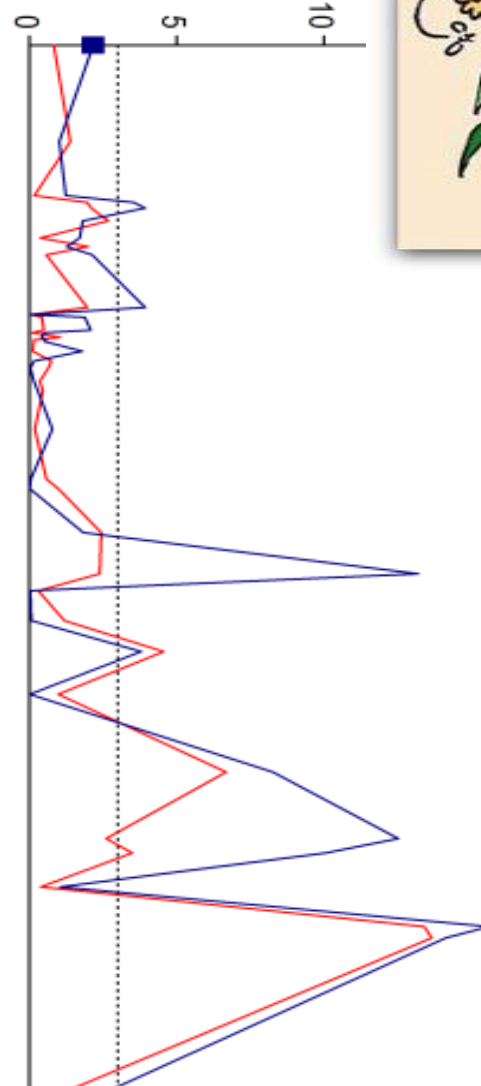
Hukkanen et al. (2007)

FC7_overall



PM

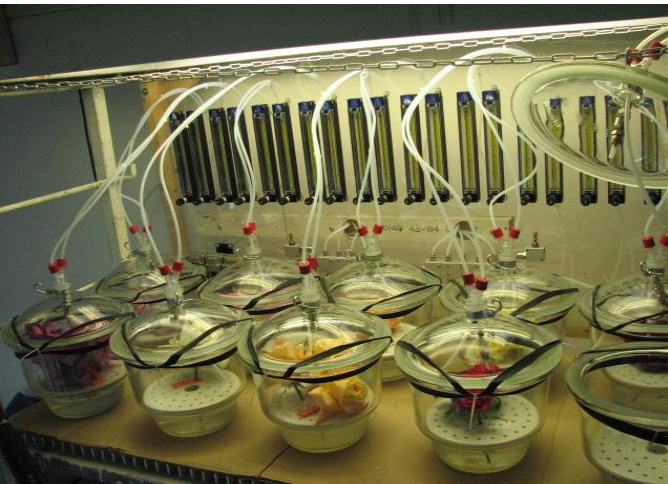
sec1510



PM
 sec1510

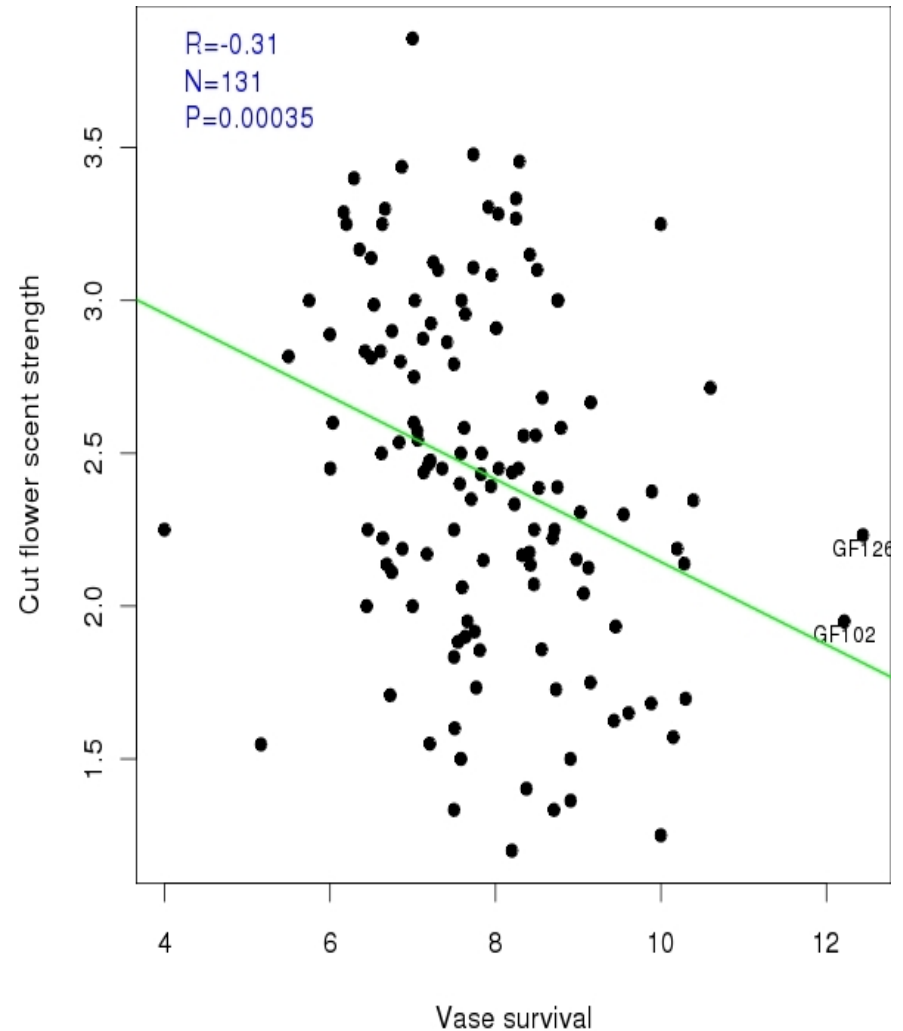
The recipe to rose scent

- Over a period of six years the roses segregating population was evaluated for smell strength by more than 50 students thus generating a reliable score for this trait.
- Using a GC-MS headspace technique we detect in the GGFC population 102 unique volatiles.

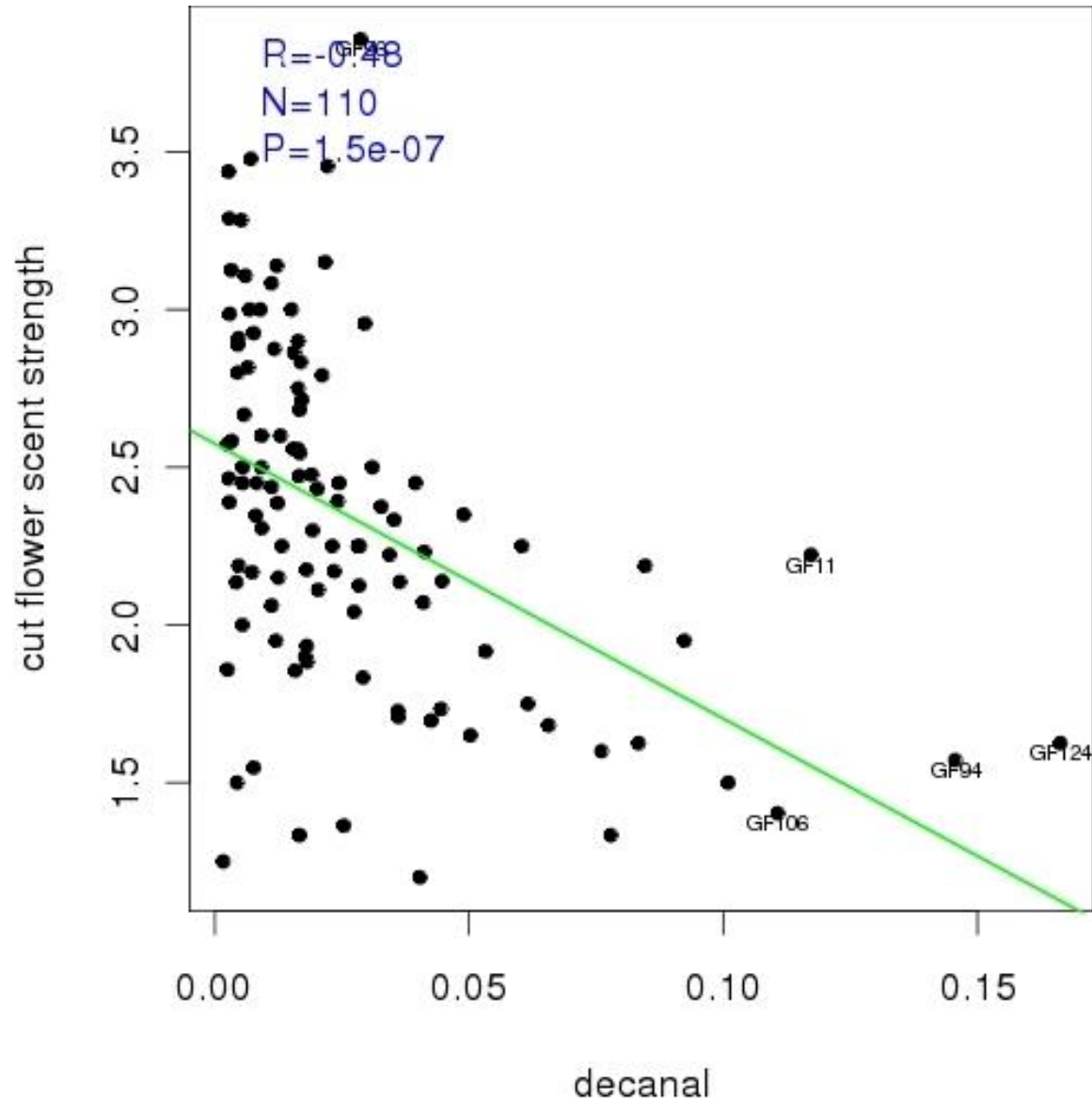


Reintroducing Fragrance to Cut Roses - Breeding Autotetraploids

- The intensive breeding efforts for cut roses with long vase life considerably narrowed the genetic basis of the crop and eliminated the scent of the flower.
- Consequently, most of the modern cut flower rose varieties today are odorless.

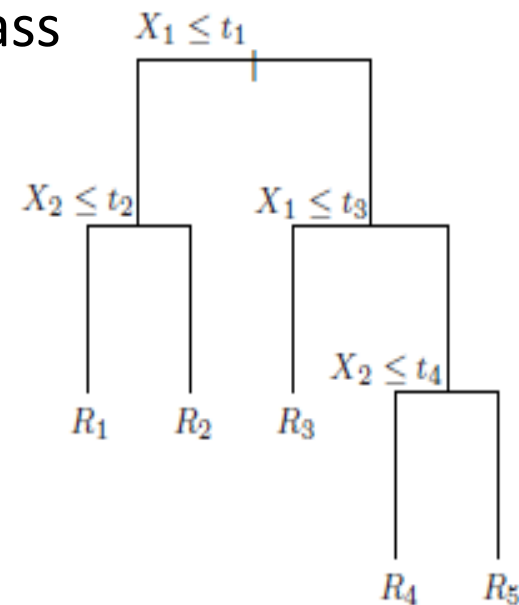
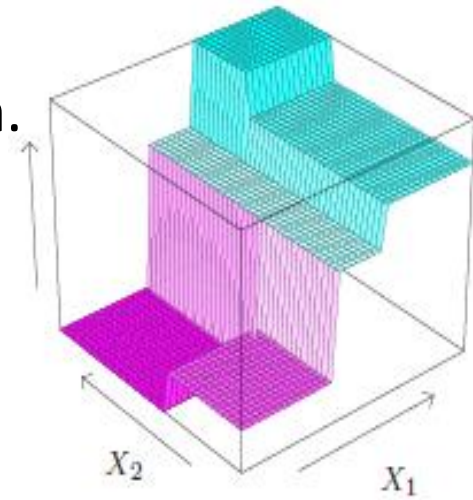


Correlation between a volatile and scent strength (example)

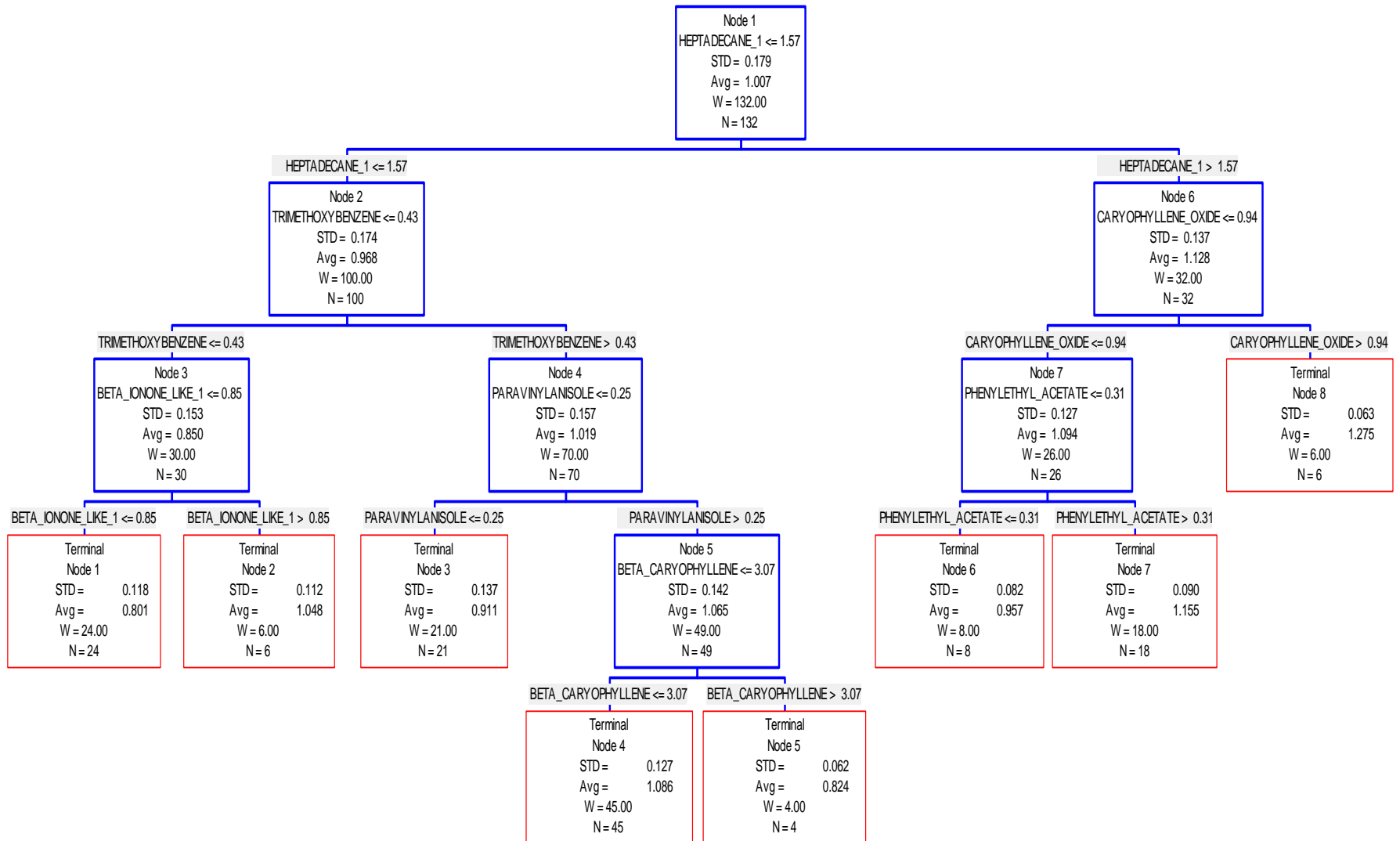


Classification and Regression Trees (CART)

- CART tree is generated by repeated dynamic partitioning of data set which automatically determines which variables and regions to focus on.
- **Seek the splitting variable j and split point s that minimize the inner variance of the sub-groups.**
- Path through the tree governed by the answers to questions or rules (Yes= left; No= right)
- At some point a given path ends in a terminal node, all records in the node are assigned to a single class
- “Regression trees have not been used much in genetics, but the natural way in which complex interactions can be expressed through regression trees does make them intriguing” (Broman & Sen 2009).

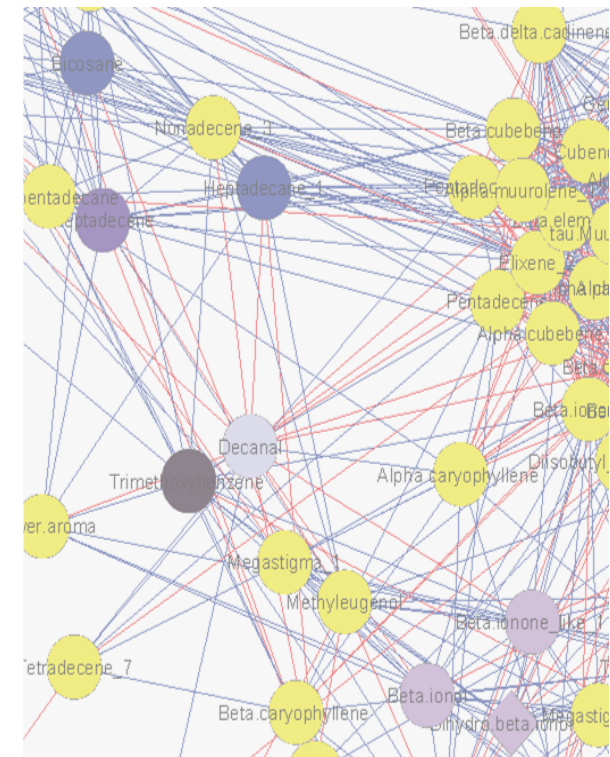
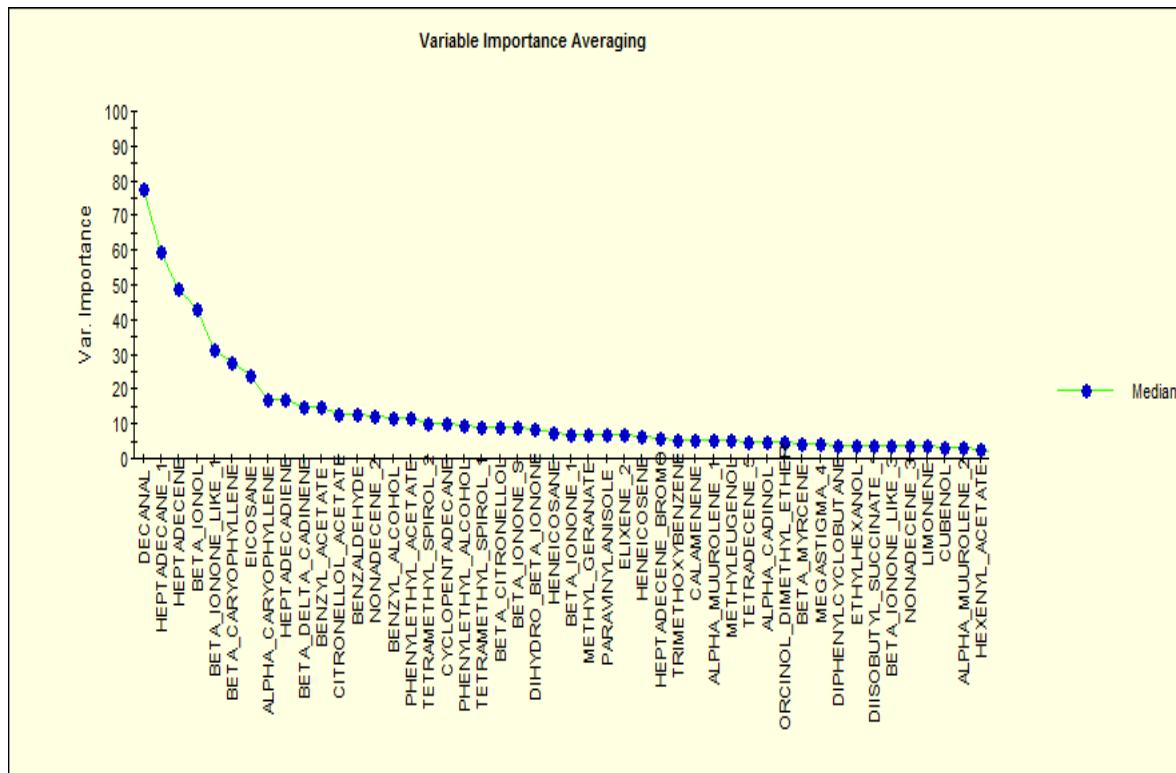


CART Analysis - Volatiles



CART Ensembles Analysis- Volatiles

- According to the CART ensembles analysis, the volatiles with the largest influence on the variance of fragrance in this population are: beta ionol, heptadecane, heptadecene and decanal.



Conclusion remarks

- Constructing autotetraploid maps.
- Comparative and evolutionary studies.
- PM resistance and Ellagitannins .
- Flower aroma and volatiles.
- CART as a tool for analyzing complex traits.
- “As we look ahead to developing new technologies, methodologies and training new minds, we must remember that no single technological advance or statistical method will unravel the genomic mystery. Instead, it will be the conglomeration of ideas, techniques and analyses that provide the end to this endeavor” Doerge (2002).



Rose database

Sign In

Select an organism from the right and then login

User Name:

Password:

[Forgot your password?](#)

[Sign Up](#)

[Log As Guest](#)

[Login](#)

Select an Organism



Tomato



Lisianthus

Rose



Rice



Arabidopsis



Barley



Maize



Melon



Mouse



Wheat

Tutorial

Demonstration videos

1. [Overview](#)
2. [Main system](#)
3. [Upload wizards](#)
4. [Breeding management](#)
5. [Phenotype analysis](#)
6. [Genetic analysis](#)

[Phenom-Networks company website](#)

Phenom-Networks

Genetic data analysis is a complex task. Phenom-Networks provides a platform on how to manage and analyze genetic data. The platform is designed to enable breeding data analysis and has developed a web-based platform into which scientists can import their raw data to enable analysis of simple and complex traits.

<http://phnserver.phenome-networks.com/>





Study Sets

Study Query

Description

Observations

Update

Images

Study Sets

Study Name	Variable Count
Breeding program	0
GG x FC population	0
Marker scores	0
Phenotypes	0
flower studies	0
GCMS studies	0
metabolites - Faculty	0
metabolites spring 06	0
metabolites fall 06	0
Secondary metabolites	0
volatiles - fall 2006	0
volatiles - winter 2006	0
HPLC studies	0
plant studies	0
quality studies	0
Drought resistance- summer 20	0
flower life - fall 2004	0
flower life - fall 2005	0
Powdery Mildew- fall 2009	0
Powdery Mildew- winter 2012	0
scent - winter 2005	0
vase - winter 2006	0
vase - winter 2007	0
yield studies	0

Add

Delete

Study Information

Name: volatiles - fall 2006

Start Date

Title: volatiles

End Date

Type:

GGFC experiments

Volatiles experiment

Type	Variable	Trait	Scale
Factor	genotype	Germplasm identification	variety name
Factor		Germplasm identification	GID
Factor		alpha_cadinol	arbitrary
Factor		alpha_calacorene	arbitrary
Factor		alpha_caryophyllene	arbitrary
Variate	alpha_cubebene_1	Alpha cubebene_1	arbitrary
Variate	alpha_cubebene_2	Alpha cubebene_2	arbitrary
Variate	alpha_Ionol_1	Alpha ionol	arbitrary
Variate	alpha_muurolene_1	Alpha muurolene_1	arbitrary
Variate	alpha_muurolene_2	Alpha muurolene_2	arbitrary
Variate	benzaldehyde	Benzaldehyde	arbitrary
Variate	benzyl_acetate_1	Benzyl acetate	arbitrary
Variate	benzyl_alcohol_1	Benzyl alcohol	arbitrary
Variate	beta_caryophyllene_1	Beta caryophyllene	arbitrary
Variate	beta_citronellol	Beta citronellol	arbitrary
Variate	beta_cubebene_1	Beta cubebene_1	arbitrary
Variate	beta_cubebene_2	Beta cubebene_2	arbitrary
Variate	beta_cubebene_3	Beta cubebene_3	arbitrary
Variate	beta_cubebene_4	Beta cubebene_4	arbitrary
Variate	beta_cubebene_5	Beta cubebene_5	arbitrary
Variate	beta_damascone_S	Beta damascone	arbitrary



Study Sets

Study Query

Description

Observations

Update

Images

Study: flower - summer 2005



Reload

Clear

Save



Filter



Columns

Split at:

2



Do split

Display



Tools

Observations

Rows per page

100



#	GID	col_low_l	col_low_in	col_up_l	col_up_int	pet_l	pet_wid	pet_wei	pet_wei_a	carp_l	carp_rec_l	flwr_rec_l	carp_wid	carp_wal	st_4num	col
1	222	5.9	2.3	27.8	3	33.6666667	29.3333333	3.6	0.09790909	1.26833333	0.80333333	9.3	0.82333333	0.20333333	4.7	
2	223	10.3	2	29.5	2	39.75	38.5	3.4	0.2	1.15	0.7475	9.5	0.8775	0.205	2.5	2
3	224	23.2	2	4.8	1.7	28	27.3333333	3.4	0.11064516	1.12666667	0.54	6.7	1.31833333	0.20333333	2.7	1.3
4	225	22.3	1.3	10	1	32.3333333	25.5833333	2.4	0.12327586	0.99833333	0.735	9.3	0.70166667	0.15416667	0.3	1
5	226	33.3	1	5.3	1	38.5	32.75	2.5	0.212	1.1	0.78666667	9.7	0.595	0.19083333	0.3	1.7
6	227	6.6	3	16.2	2	33.3333333	33.3333333	3.6	0.09790909	1.26833333	0.80333333	9.3	0.82333333	0.20333333	4.7	
7	228	4.3	2	27.3	3	39.75	38.5	3.4	0.2	1.15	0.7475	9.5	0.8775	0.205	2.5	2
8	230	35.7	1	4.5	1	28	27.3333333	3.4	0.11064516	1.12666667	0.54	6.7	1.31833333	0.20333333	2.7	1.3
9	231	0		33	1	32.3333333	25.5833333	2.4	0.12327586	0.99833333	0.735	9.3	0.70166667	0.15416667	0.3	1
10	232	11.7	1.3	18.2	2	38.5	32.75	2.5	0.212	1.1	0.78666667	9.7	0.595	0.19083333	0.3	1.7
11	233	6.5	1	25.8	2.3	33.6666667	30.5833333	4.5	0.07420769	1.17	0.605	7.7	1.05666667	0.22583333	3	2
12	234	7.9	2	25.8	1.3	35.25	36.125	4	0.14436364	0.85	0.42	6	0.8925	0.205	1	2
13	235	10.3	2	25	3	29.9583333	29.5416666	2.2	0.08666667	1.12416667	0.66	5.8	0.82916667	0.15583333	1.8	2
14	236	6.3	1	23.7	2	33.1	31.9	5.4	0.13038833	1.301	0.822	8.6	1.041	0.2125	6.6	1.8
15	237	7.3	2.6	25.8	2	37.25	32.125	4.2	0.15425926	1.125	0.6825	7.5	0.8525	0.20625	3	2
16	238	6.1	1.5	31.1	2	39.0833333	34.0833333	4.1	0.18560606	1.07333333	0.78666667	8	0.675	0.185	1.3	2
17	239	10.1	2	29	2	37.75	33.0833333	4.8	0.1452	1.175	0.58166667	5.3	1.15166667	0.215	1.3	2
18	240	6.4	2.3	31.3	1.7	35	34	2.6	0.14222222	1.07	0.395	5.7	0.6825	0.1925	3	2
19	241	6.5	1.5	4.2	1.5	34.0833333	30.8333333	3.3	0.16915254	1.03166667	0.53666667	5.3	0.96166667	0.22416667	1	2
20	242	7.2	2	26.9	3											

Raw data of an
experiment



Trait Query

Trait Studies

List of all traits

Save as Excel

Trait [Total Records 470]

#	Name	Abbr.	Description	Group	Ontology	Images?
1	Alanine (2TMS)		aliphatic amino acid	Metabolites- Primary (Fait A.)	TO:0002673	No
2	Alanine (3TMS)		aliphatic amino acid	Metabolites- Primary (Fait A.)	TO:0002673	No
3	Alpha cadinol		sesquiterpene alcohol	Volatiles- Sesquiterpene	TO:0000277	Yes
4	Alpha calacorene		sesquiterpene olefin	Volatiles- Sesquiterpene	TO:0000277	Yes
5	Alpha caryophyllene		sesquiterpene olefin	Volatiles- Sesquiterpene	TO:0000277	Yes
6	Alpha cubebene_1		sesquiterpene olefin	Volatiles- Sesquiterpene	TO:0000277	Yes
7	Alpha cubebene_2		sesquiterpene olefin	Volatiles- Sesquiterpene	TO:0000277	Yes
8	Alpha ionol		norisoprene alcohol	Volatiles- Norisoprene	TO:0000277	Yes
9	Alpha muurolene_1		sesquiterpene olefin	Volatiles- Sesquiterpene	TO:0000277	Yes
10	Alpha muurolene_2		sesquiterpene olefin	Volatiles- Sesquiterpene	TO:0000277	Yes
11	Amino acids in the petal		flower's primary metabolites content	Metabolites- Primary (Fait A.)	TO:0002673	No
12	Anteraxanthin (%) in the petal		petal's carotenoids content (% from the total amount of caroteno	Biochemistry- Carotenoids	TO:0000496	No
13	Anther color		main color of an anther	anatomy and morphology- Reproduc	TO:0000187	Yes
14	Anthocyanins concentration in the pe		petal's anthocyanins content	Biochemistry- Anthocyanins	TO:0000071	Yes
15	Apical dominance		shoot apical dominance	Growth and development- Bush	TO:0000660	Yes
16	Arabinose		cell wall component	Metabolites- Primary (Fait A.)	TO:0000291	No
17	Ascorbic acid		organic acid	Metabolites- Primary (Fait A.)	TO:0000281	No
18	Asparagine		neutral amino acid	Metabolites- Primary (Fait A.)	TO:0002673	No
19	Asparagine (flower tissue)		neutral amino acid	Metabolites- Primary (Zamir D.)	TO:0002673	No
20	Aspartic acid		acidic amino acid	Metabolites- Primary (Fait A.)	TO:0002673	No
21	Aspartic acid (flower tissue)		acidic amino acid	Metabolites- Primary (Zamir D.)	TO:0002673	No
22	Average branch length		average branches length per bush	Growth and development- Bush	TO:0000576	Yes
23	Average petal weight		total petals weight divided by total number of petals	anatomy and morphology- Flower	TO:0000275	No
24	Average seed weight		the average weight of the seed	Fertility	TO:0000181	Yes
25	Average seeds number		total number of seeds per plant divided by total number of fruit	Fertility	TO:0000445	Yes
26	Axillary buds stage 1 (number)		number of axillary buds in floral developmental stages 0-1	Growth and development- Bush	TO:0000646	Yes
27	Axillary buds stage 6 (number)		number of axillary buds in floral developmental stages 6-7	Growth and development- Bush	TO:0000646	Yes

Ontology column

New Trait



Trait Query

Trait Studies

Trait Details

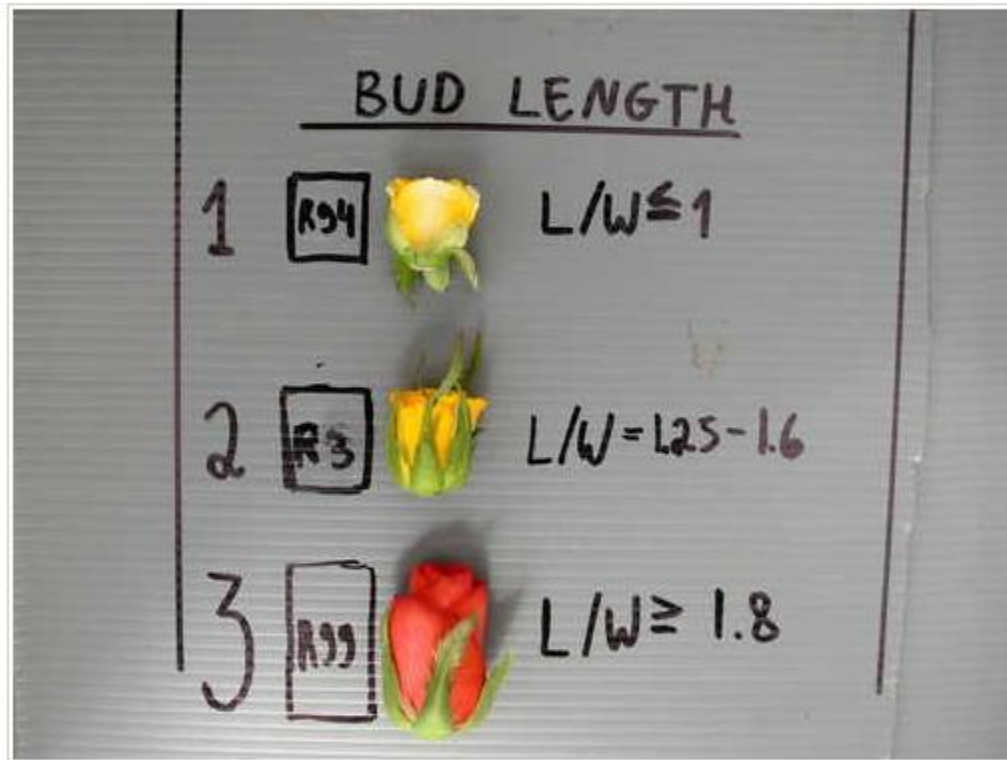
Trait Images

< Previous Next

Trait: Bud length



Description + Images
for each trait



Bud length 1

[Download Original](#)

null

Fields list Lines Traits Groups Nursery Field map Field plots **Observations**



Field: fall 2012 (BR12)

Refresh Clear Save Filter Split at: 2 Do split Display Category Edit Tools Reports

Selections and Observations

Rows per page 30


Columns

#	GID	Line name	Source	Plot name	Fruit extracted	Seed extracted	Plant	Observation date	Selected	Prickly pedicel	location	Cut flower scent pleasant	Flower color-major	Flower color-minor
1	222	GF1	GG / FC	1	No	No		20121105	No	3			pink	3
2	270	GF2	GG / FC	2	No	No		20121105	No	3				
3	281	GF3	GG / FC	3	No	No		20121105	No	2	2		dark pink	2
4	292	GF4	GG / FC	4	No	No		20121105	No	3	1		cream	1
														
5	222	GF1	GG / FC	1	No	No		20121105	No	3	1		pink	1
														
6	303	GF5	GG / FC	5	No	No		20121106	No	3	1		light pink	peach
7	314	GF6	GG / FC		No	No			No	3	1		pink	
8	325	GF7	GG / FC		No	No			No	3	3		peach	pink
9	336	GF8	GG / FC		No	No			No	3	1		pink	
10	347	GF9	GG / FC		No	No			No	3	1		pink	

Phenotypes + Images
for each genotype

Analysis page: analyze the entire database using many statistical and graphical methods

1) Select experiments



1) Select experiments

Study	Title						Date	End
+	Breeding program	-					6	0
-	GG x FC population	-					5	0
+	Marker scores	-					6	0
-	Phenotypes	-		0	0		20110327	0
+	flower studies	-		0	0		20091207	0
-	GCMS studies	-		0	0		20091119	0
	metabolites - Faculty	primary metabolites	-	E	2	28	0	0
	metabolites spring 06	metabolites - tissue flower	-	E	5	51	20060301	2006
	metabolites fall 06	metabolites - flower tissue	-	E	5	51	0	0
	Secondary metabolite	LCMS analysis of secondar	-	E	4	116	0	0
	volatiles - fall 2006	volatiles	-	E	2	102	0	0
	volatiles - winter 2006	volatiles	-	E	2	99	0	0

Search for study



Search



Clear

Name

Type

Experiment

Filter

Cancel

Alpha cubeben arbitrary Volatiles- Sesquite

Alpha ionol arbitrary Volatiles- Norisopropyl

Alpha muroler arbitrary Volatiles- Sesquite



Form

Result 3



Studies filter Basic filter Advanced filter Germplasm filter Normalize Traits display Search trait History

- Univariate
 - Distribution
 - Bars
 - Fit Y by X
 - Distribution II
- Multivariate
 - Means heatmap
 - Bars
 - HCA cluster
 - Cloud 3d
 - PCA cluster
 - Multiple ttests
 - Decision tree
- Nominals
- Correlations
 - Heatmap
 - Traits pairwise
 - Genetic background
- Network
 - Static image
 - Dynamic network
- Correlate Y by X
 - XY plot
 - Table
 - XY plot II
- Study comparison
- Pedigree
- QTL
- Query
- Marker

2) Select statistical method

Variables				
	Type	Trait	Scale	Trait Group
+	Factor	Chromatogram	Not Specified	design factor
+	Factor	Derivative nam	Not Specified	design factor
+	Factor	Full name	Not Specified	design factor
+	Factor	Full name	Not Specified	design factor
+	Factor	Full name	Not Specified	design factor
+	Factor	Germplasm set	Not Specified	design factor
+	Factor	Observation d	Not Specified	design factor
+	Factor	Observation n	Not Specified	design factor
+	Factor	Parent 1	Not Specified	design factor
+	Factor	Parent 2	Not Specified	design factor
+	Factor	Physical locati	Not Specified	design factor
+	Factor	Plot	Not Specified	design factor
+	Factor	Plot in a row	Not Specified	design factor
+	Factor	Sample ID	Not Specified	design factor
+	Factor	Source	Not Specified	design factor
+	Variate	Alanine (2TMS) Aaron GCMS prot	Metabolites- Prima	
+	Variate	Alanine (3TMS) Aaron GCMS prot	Metabolites- Prima	
+	Variate	Alpha cadinol	arbitrary	Volatiles- Sesquite
+	Variate	Alpha calacore	arbitrary	Volatiles- Sesquite
+	Variate	Alpha caryoph	arbitrary	Volatiles- Sesquite
+	Variate	Alpha cubeben	arbitrary	Volatiles- Sesquite
+	Variate	Alpha cubeben	arbitrary	Volatiles- Sesquite
+	Variate	Alpha ionol	arbitrary	Volatiles- Norisopri
+	Variate	Alpha muuroler	arbitrary	Volatiles- Sesquite

Y, Variables >>

X, Variables >>

Variate:

Variate:

Display image map

☒ Display image map?

Heatmap dendrogram

Cluster by Rows and cols

Correlation type

Correlation type: Pearson

☐ Logarithmic data☒ Remove outliers that are 3 standard than the mean

Figure size



Form

Result 3



Studies filter Basic filter Advanced filter Germplasm filter Normalize Traits display Search trait History

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Variables

	Type	Trait	Scale	Trait Group
+	Factor	Chromatogram	Not Specified	design factor
+	Factor	Derivative nam	Not Specified	design factor
+	Factor	Field row	Not Specified	design factor
+	Factor	GCMS run	Name	design factor
+	Factor	Generation	Not Specified	design factor
+	Factor	Germplasm ide	GID	germplasm identifi
+	Factor	Germplasm ide	variety name	germplasm identifi
+	Factor	Germplasm set	Not Specified	design factor
+	Factor	Observation d	Not Specified	design factor
+	Factor	Observation n	Not Specified	design factor
+	Factor	Parent 1	Not S	
+	Factor	Parent 2	Not S	
+	Factor	Physical locati	Not S	
+	Factor	Plot	Not S	
+	Factor	Plot in a row	Not Specified	design factor
+	Factor	Sample ID	Not Specified	design factor
+	Factor	Source	Not Specified	design factor
+	Variate	Alanine (2TMS)	Aaron GCMS protc	Metabolites- Prima
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+	Variate	Alpha cubeben	arbitrary	Volatiles- Sesquite
+	Variate	Alpha cubeben	arbitrary	Volatiles- Sesquite
+	Variate	Alpha ionol	arbitrary	Volatiles- Norisopri
+	Variate	Alpha muuroler	arbitrary	Volatiles- Sesquite

Y, Variables >>

X, Variables >>

3) Classify variables
(traits) to boxes

Variate:

Alanine (2TMS)

Alanine (3TMS)

Alpha cadinol

Variate:

Apical dominance

Arabinose

Ascorbic acid

Display image map

☒ Display image map?

Heatmap dendrogram

Cluster by Rows and cols

Correlation type

Correlation type: Pearson

☐ Logarithmic data☒ Remove outliers that are 3 standard
than the mean

Figure size



Form

Result 3



Save



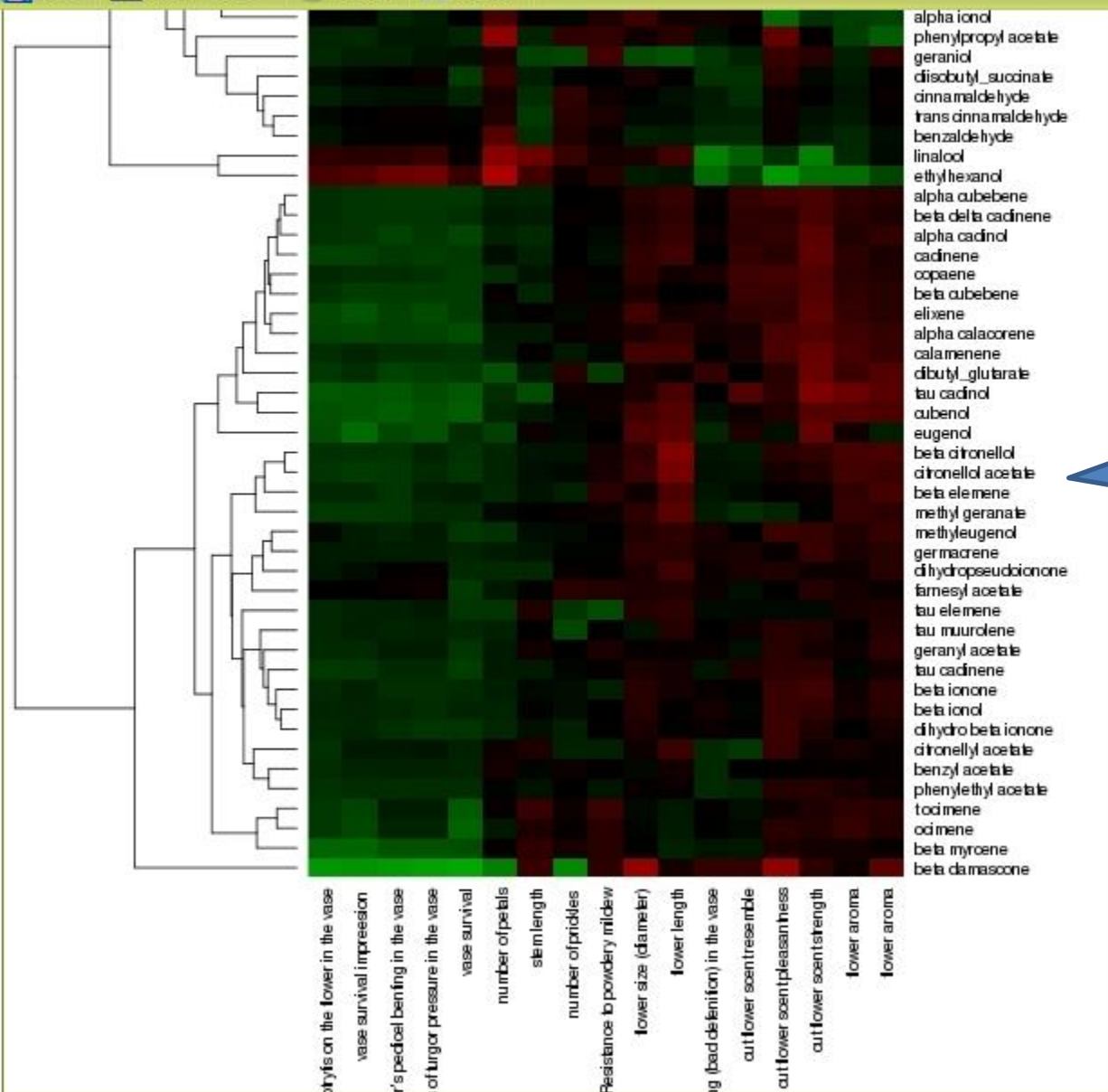
Query table ▾



History



Details



Output example

Acknowledgments

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- **Yinon and Ester**
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- Rinat Ovadia

Weizmann Institute

- **Asaph Aharoni**
- Ilana Rogachev
- Sagit Meir



Golden Gate (GG)
no scent
vase life-12 days



Fragrant Cloud (FC)
strong scent
vase life- 5 days



07-1130
strong scent
vase life- 15 days

07-655
strong scent
vase life- 19 days



07-296
strong scent
vase life- 13 days



07-2087
light scent
vase life- 18 days



07-1336
strong scent
vase life-14 days



07-1449
strong scent
vase life-12 days



07-1773
strong scent
vase life- 12 days



07-133
light scent
vase life- 14 days



07-1465
light scent
vase life- 11 days



07-2506
strong scent
vase life- 18 days



07-2811
light scent
vase life- 22 days

rosita vendela
no scent
vase life- 11 days

*"Don't hurry.
Don't worry.
You're only here
for a short visit.
So don't forget to
stop and smell
the roses."*

(Walter Hagen)

The Israeli president Mr. Shimon Peres
smelling our roses (June, 2011)

