



The First Version of the Whole-Genome Sequencing of the
Muscadine Grape
(*Muscadinia rotundifolia* cv. Noble)

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Our Muscadine Breeding Program

- ❖ Currently muscadine grapes is commercially grown in 12 states in the US.
- ❖ There is a demand for new muscadine cultivars with enhanced fruit/vinification qualities.
- ❖ Over the years, conventional breeding strategy has resulted in several muscadine cultivars suitable for table and wine markets with improved quality traits.
- ❖ However, the procedure is lengthy, laborious and expensive, which does not match with the rapid, ever-increasing industry needs.
- ❖ Our goal is capitalizing on the latest achievements of genomic technologies to develop breeding platform that meets the growing industry demands.

Grapevine Cultivars & Germplasm at CVSFR



Background	Total No. of Vines	No. of Individual Genotypes	No. of Cultivars	Adult ^a	Juvenile ^b
<i>Muscadinia</i> hybrids	3925	2537	51	664	1877
<i>Vitis</i> hybrids	3380	2432	172	589	1843
<i>Muscadinia</i> x <i>Vitis</i> (MV)	106	106	-	103	3

a) 3-year old and /or elder; Reproductive Phase.

b) 2-year old and or younger; Growth Phase.

Grape

***Muscadinia* Genus (3)**

M. rotundifolia

M. munsoniana

M. popenoei

40 chromosomes ($n=20$)



Vineyard-BMP-Supreme (*muscadinia*)-fresh fr



Vineyard-BMP-Darlene (*muscadinia*)-fresh fru

***Euvitis* Genus (14)**

V. vinifera

V. labrusca

V. riparia

V. aestivalis

V. rupestris

38 chromosomes ($n=19$)



d-BMP-Stover (*Vitis* hybrid) white wine



Vineyard-BMP-Cynthiana/Norton (*Vitis* hybrid)-rec

Morphological Differences:

Leaves, flower type, tendrils, cluster size, berry size, and age of cutting for propagation.

***Muscadinia* sp. hold particular agronomical and reproductive traits:**

- ❖ Resistance to most of diseases that limit the worldwide production of *V. vinifera*.
- ❖ Adapted to poor soil.
- ❖ Grow well under warm and humid conditions.
- ❖ Exotic fruit quality traits (berry size, unique flavor spectrum, therapeutic phytochemicals content).
- ❖ Unique vinification qualities.

Muscadines have obviously different challenges from those of traditional '*vinifera*' grapes

- ❖ Cold hardiness.
- ❖ Extended ripening season.
- ❖ Perfect flower with large berry size.
- ❖ Fruit quality traits (diversity in berry color & shape, seedlessness, large cluster, thinner skin, dry stem scar, uniform ripening, extended shelf-life).



Male



Female



Perfect



Wet stem scar

Whole – Genome Sequencing of Muscadine Grape Noble cv.



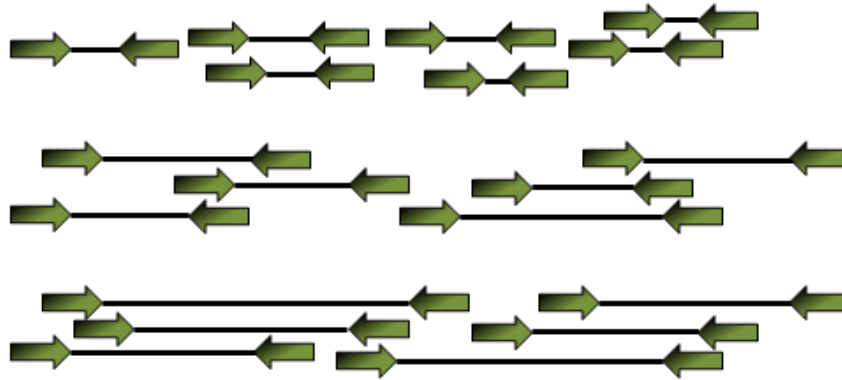
Main red wine muscadine cultivar released by Nesbitt, 1973.

Thomas (♀) x Tarheel (♂)

- ❖ High vigor growth.
- ❖ Disease resistant.
- ❖ Self-fertile (perfect flower).
- ❖ Midseason.
- ❖ Uniform ripening.
- ❖ Black small berries (3.8 – 4.3 g).
- ❖ High yield (9.6 – 10.7 tone/acre).
- ❖ Stable purple pigments.
- ❖ Suitable for wine and juice production.

Muscadine Genome Sequencing & Assembly

Sequencing

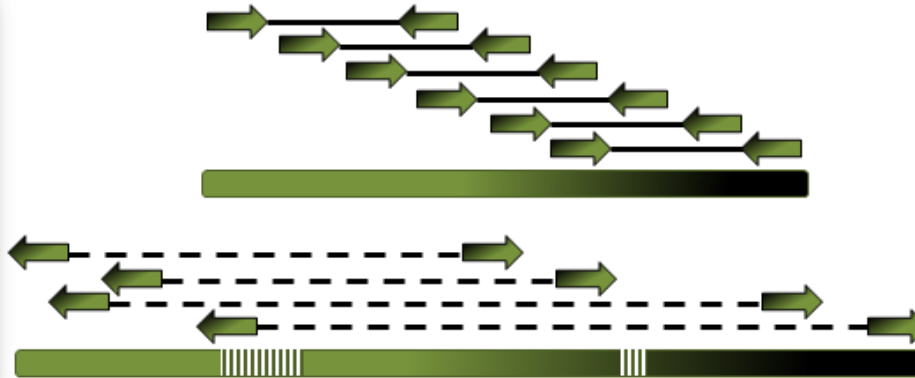


Illumina HiSeq 2500
(80 M reads)

Chicago
(181 M reads)

Hi-C
(223 M reads)

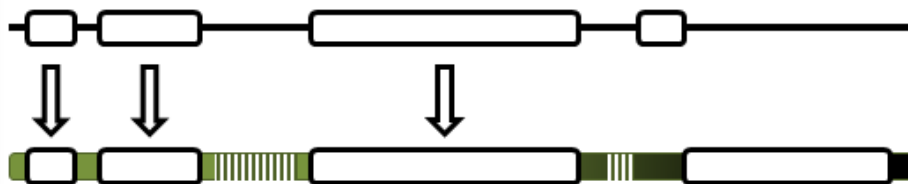
Genome HIRise Assembly



Contigs

Scaffold

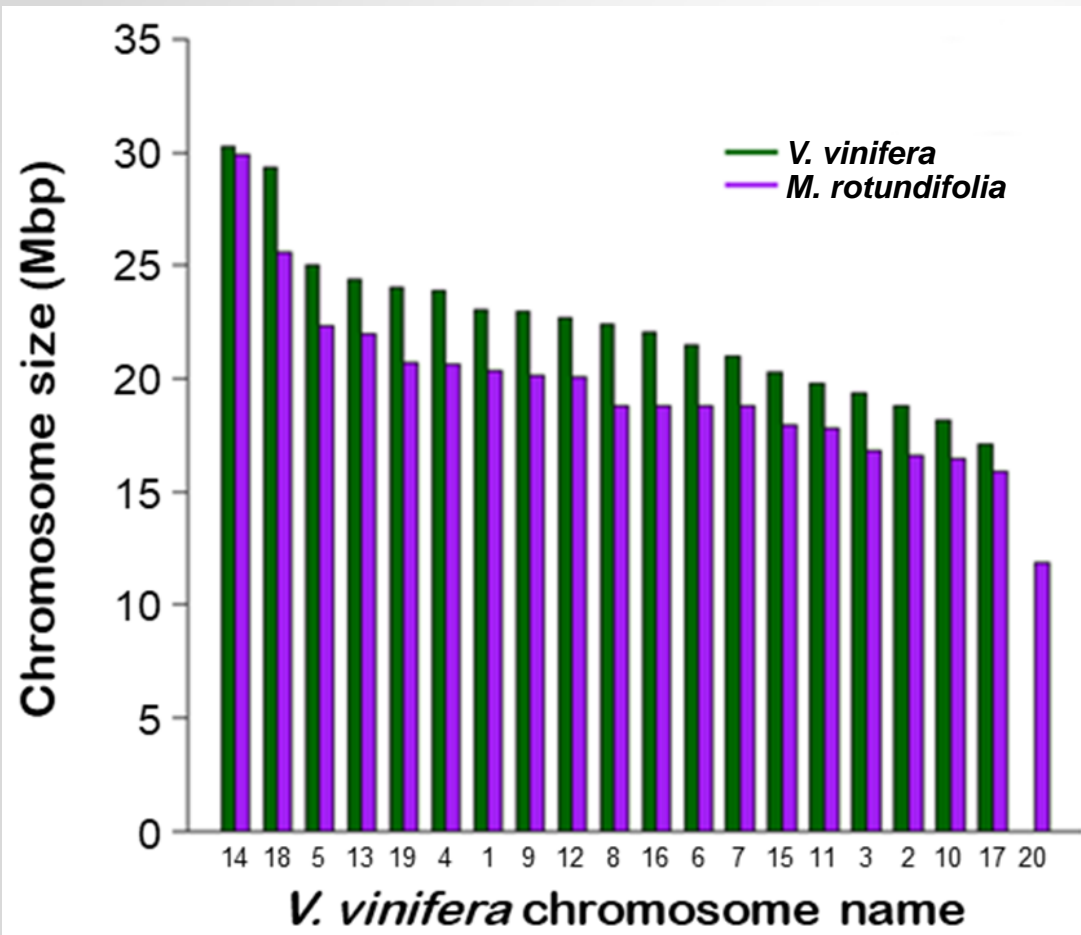
Annotation



Related Genome

Annotated Draft Genome

Muscadine Genome Assembly Statistics



Estimated genome size	414 Mb (82%)
Assembly Size	400.26 Mb (96.6%)
Contig N50	107 kb
Contig L50	1,011 contigs
Scaffold N50	20.045 Mb
Scaffold L50	9
Scaffold N90	16.486 Mb
Scaffold L90	18
Number of scaffolds	1,629
Heterozygosity	1.47%
Unplaced sequence	2.47%
Contig gaps	8,168

Benchmarking Universal Single-Copy Ortholog (BUSCO)

Assessment of muscadine genome assembly and gene set using BUSCOs indicates highly-complete representation of protein-coding genes.

Statistics searching Embryophyta BUSCOs:

1382 Complete BUSCOs (96%).

❖ 1339 Complete and single-copy BUSCOs (93%).

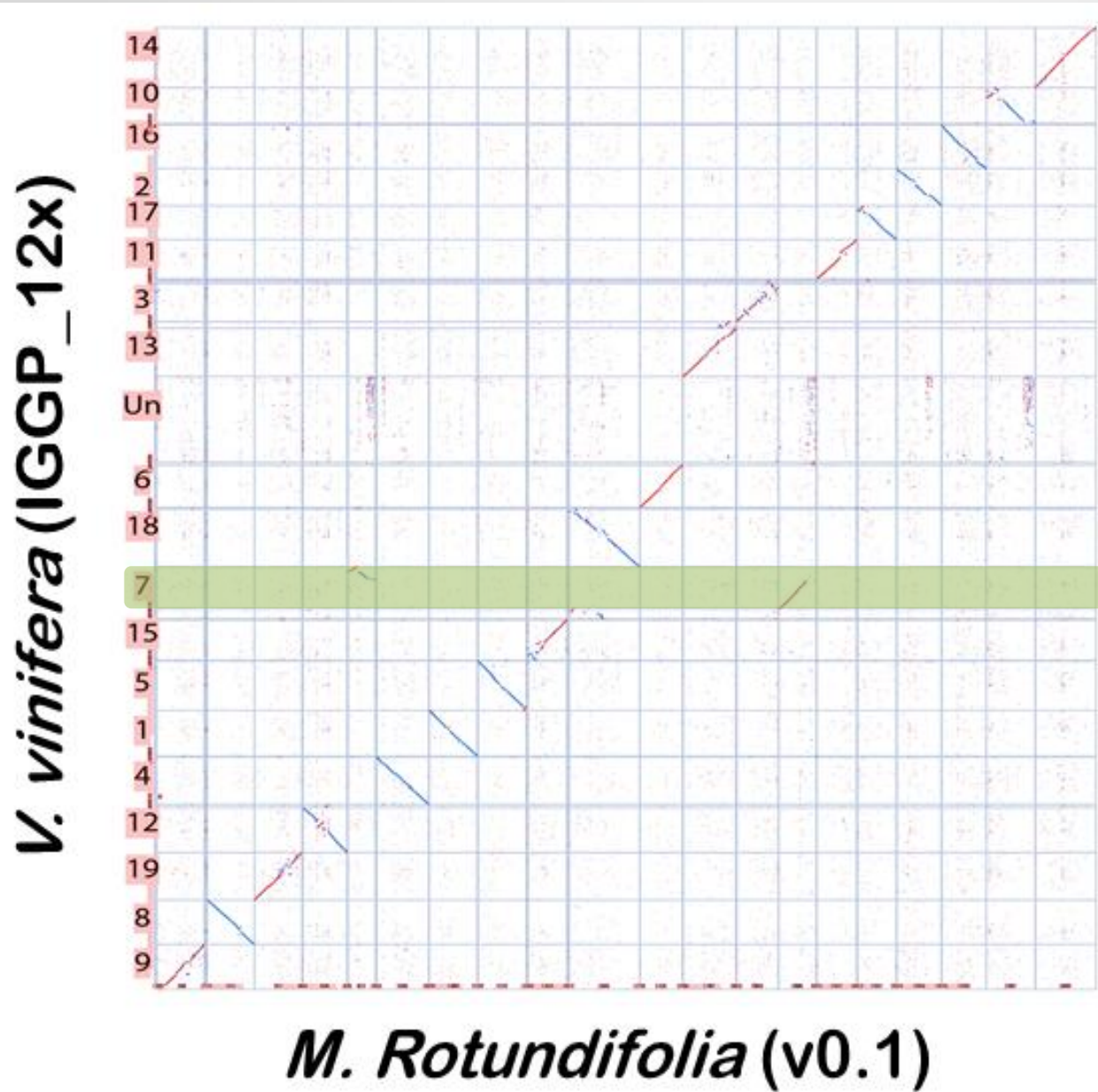
❖ 43 Complete and duplicated BUSCOs (3%).

17 Fragmented BUSCOs (1.2%).

41 Missing BUSCOs (2.8%).

1440 Total BUSCO groups searched.

M. rotundifolia and *V. vinifera* display high colinearity



The extra *Muscadinia* chromosome is composed of two large segments homologous to *Vitis* chromosome 7

Characterization of muscadine population

A population of 400 muscadine genotypes (50 standard cultivars and 350 lines) has been carefully selected to ensure diversity and subjected to:

1. Phenomic Characterization

- ❖ Fertility traits (bud fertility, bud fertility coefficient, position of first fruiting bud);
- ❖ Cluster physical traits (intensity, size, weight, number and weight of berries/cluster);
- ❖ Berry physical traits (scar pattern, color, size, weight, number and weight of seeds/berry, firmness);
- ❖ 50 berries traits (berries weight, pomace weight, juice volume);
- ❖ Yield traits (No. of clusters/vine, yield/vine); and
- ❖ Berry quality traits (TSS, acidity, pH).

Characterization of muscadine population

2. Phenology traits

- ❖ Bud break.
- ❖ Green shoot visible.
- ❖ Visible inflorescences.
- ❖ Partial bloom.
- ❖ Full bloom.
- ❖ Fruit-set.
- ❖ Cluster closure.
- ❖ Veraison.
- ❖ Post-Veraison.
- ❖ Ripening.

3. Biotic stress traits

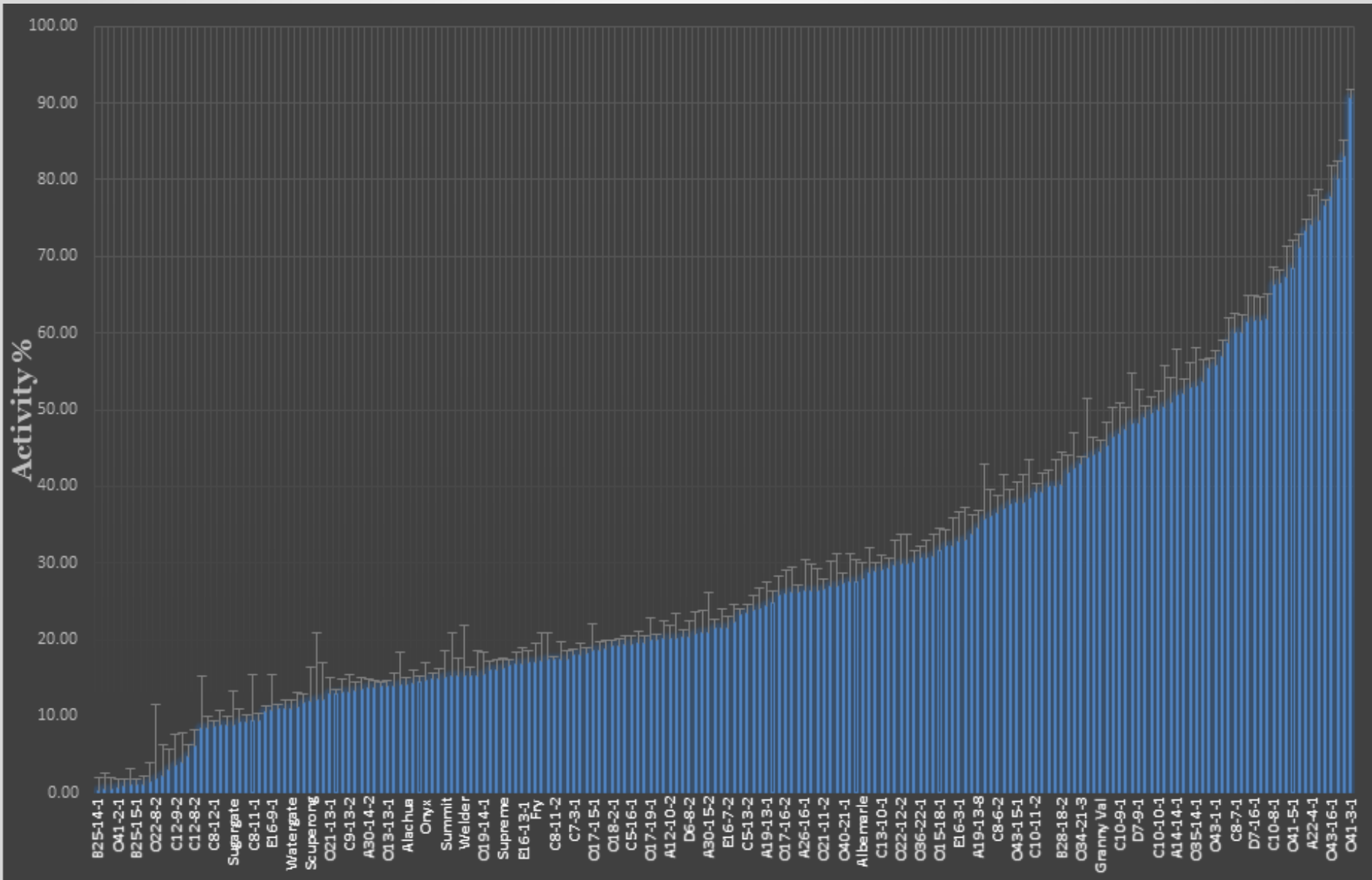
- ❖ Leaf roll virus (several strains); and
- ❖ Fungal diseases (powdery and downy mildew).

Characterization of muscadine population

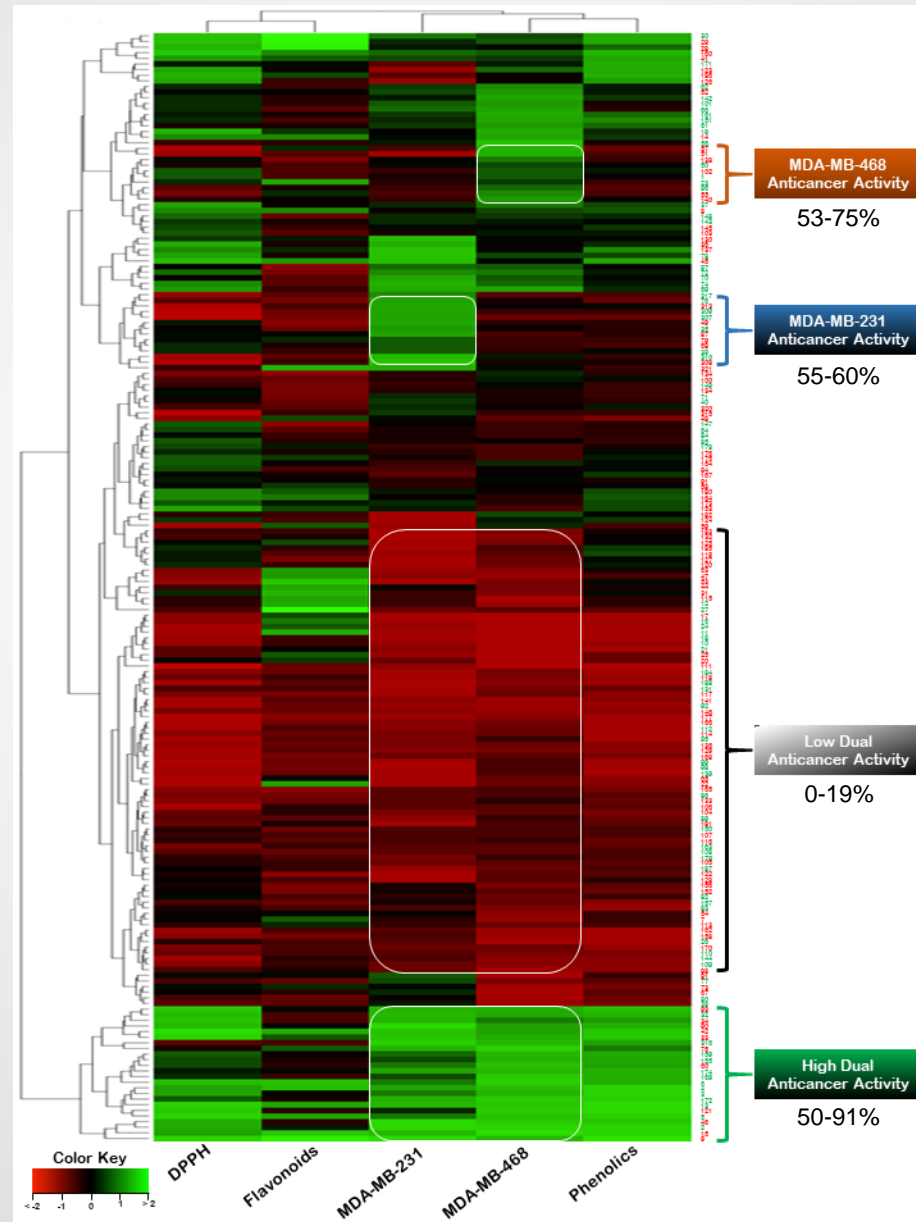
4. Metabolomic and sensorial traits

- ❖ Aroma volatile esters production.
- ❖ Total phenolic.
- ❖ Total flavonoid.
- ❖ Antioxidant activity.
- ❖ Antimicrobial activity (*E. coli*).
- ❖ Anticancer activity (Breast cancer cell lines MDA-MB-231 and MDA-MB-468).

Anticancer activity of muscadine extracts using African American breast cancer cell lines



Classification of muscadine genotypes based on their anticancer activity



Undergoing

- ❖ Generating Pacbio Iso-seq and Illumina RNA-seq data on multiple tissues to generate accurate and comprehensive gene predictions.
- ❖ Performing ATAC-seq (Assay for Transposase-Accessible Chromatin using sequencing) to determine the chromatin accessibility of muscadine genome.
- ❖ Analyzing of *M. rotundifolia* genes' PAV (presence-absence variation) and their predicted functions compared to *V. vinifera*.
- ❖ Complete QC analysis and submission to Phytozome.



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Team Members



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