### A Genetic Study of Grapevine Rooting from Dormant Harwood Cuttings

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# **Rooting practice**

INCREASING VINES IN THE VINEYARD



- Mark vines with good vigor/yield before harvest
- Take cutting during the dormant season (December or January)
- Store in a cold place at temps slightly above freezing ( $\sim 4^{\circ}C$ )
- Early spring, after soil warms, cuttings can be placed in nursery



Missouri State

UNIVERSITY

### **Problem** NOT ALL VINES ARE EASY TO ROOT





#### NO TIME TO PROPAGATE GREEN CUTTINGS



https://rickshory.wordpress.com/2018/01/05/growing-grapes-part-1-propagation/

### Problem

#### **ADVENTITIOUS ROOTING**

- Complex developmental process that requires plasticity to adjust to stressful conditions
- Respond to certain stimuli (moisture, minerals, light & lack thereof)
- Requires at least three processes:
  - De-differentiation (committed cells grow in reverse, unlearn, etc.)
  - Cell division (during a time they normally were not)
  - AR = root primordia initiation, development and outgrowth



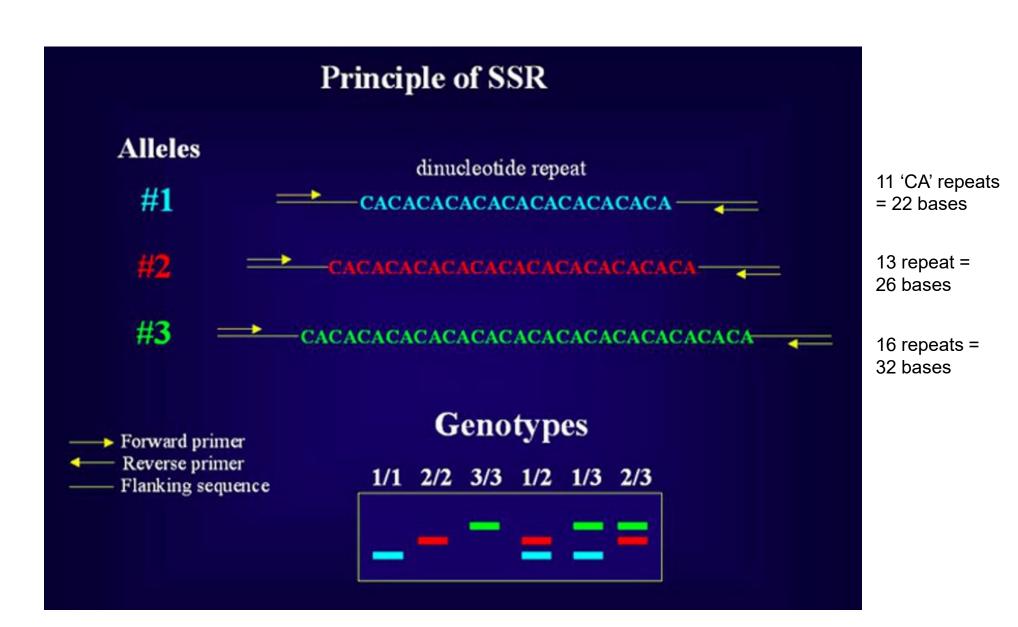
### What we do

#### **PRECISION BREEDING (MOLECULAR)**



- Cross parents of interest
- Genotype parents and progeny using DNA markers
- Identify traits of interest
- Associate DNA markers with traits of interest for future markerassisted selection







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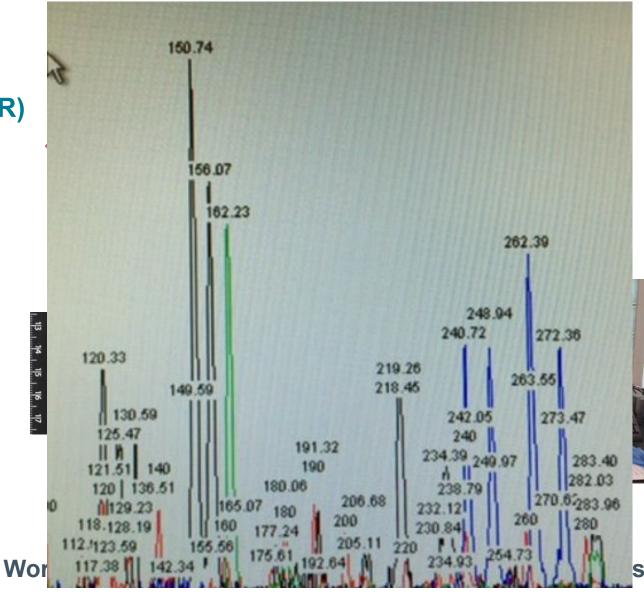
### What we do

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#### **PRECISION BREEDING (MOLECULAR)**







Length of fragment

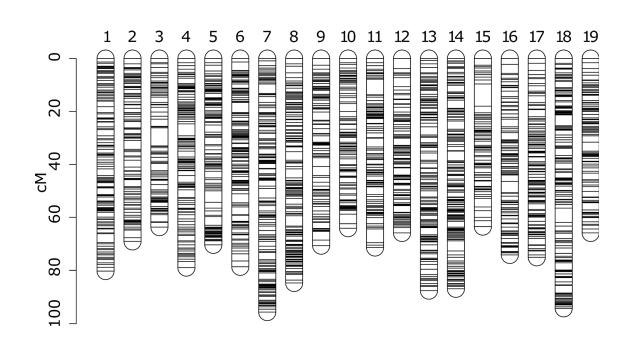
HISTORY

- 2005 Vitis aestivalis 'Norton' x V. vinifera 'Cabernet Sauvignon'
  = 74 interspecific hybrids
- •2011 hybrid progeny were expanded to 310 genotypes
- The mapping population for this study was 189



#### **CONSENSUS LINKAGE MAP**

- A map was constructed using 406 SSR markers and 1,659 single nucleotide polymorphic (SNP) markers – generated by GBS
- Map size 1,678.6 cM



Chromosome

Hammers et al. 2017. Molecular Breeding 37:64 Sapkota et al. 2019. Theo. and App. Genetics 132:137 Negaus et al. 2021. Fruit Research 1:8



VITIS AESTIVALIS – 'NORTON'

- Missouri's state wine grape
- Resistant to fungal diseases
- Cold hardy
- Difficult-to-root



- Not resistant
- Doesn't grow well in Missouri
- Easy-to-root

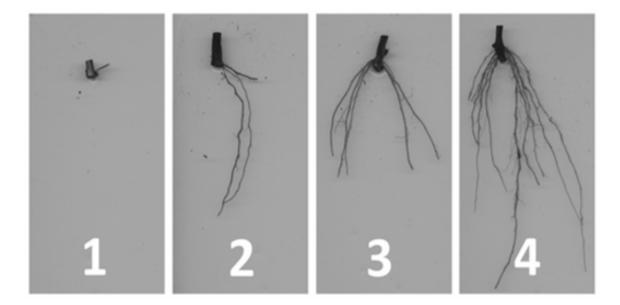




#### **PROGENY PHENOTYPES**





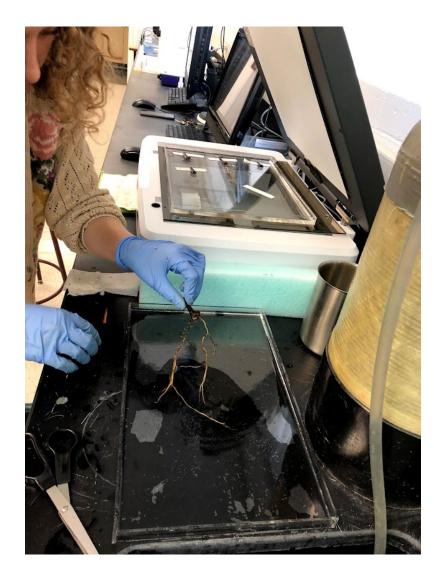


Representative phenotypes segregating for rooting ability

**Phenotyping** MEASURING ROOTING ABILITY

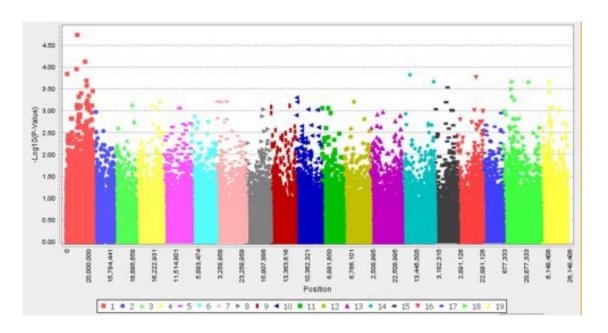
- Average root number
- Longest root length
- Average total root length
- Lateral : Primary root ratio
- Percent rooting
- •Rank 1-4





## Results

#### **FIRST YEAR**



Manhattan plot using the mixed linear model in TASSEL5.0

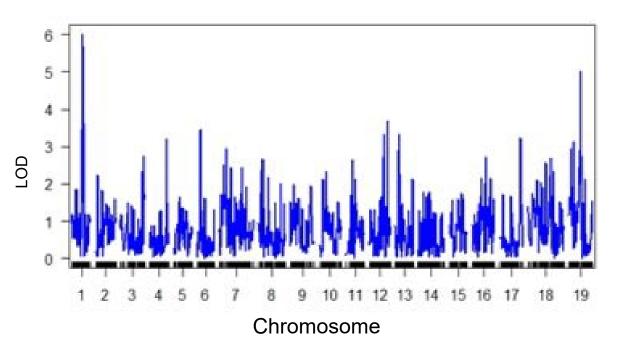
Analysis shows linkage group 1 to have the best SNP-trait association (4.7). Each colored bar represents a linkage group. Each point represents a marker on the respective linkage group



### Results

**FIRST YEAR** 

 Analysis of average root number (ARN) per genotype reveals that linkage group 1 (LOD 6.13) may contain a QTL



R/QTL analysis using SSR & SNP markers



### Conclusion

PRELIMINARY

- •A potential QTL for root number has been identified on Linkage Group 1
- •Flanking SSR markers for possible QTL are ctg8034 and VMCN2g7
- •An additional SNP marker, between SSRs, is S1\_18502204



### Future

2022-23

- Increase density of map using new genotyping platform (rhAmpSeq)
- Add another year of data, totaling 3 years data
- Analyze the other rooting phenotypes
- Discover quantitative trait loci (QTL) for dormant rooting ability



### **Thank You**

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