

Metabolomics approaches for tracking biotic and abiotic stress performance in tree improvement programs

Anna Conrad¹, Jared Westbrook², Tatyana Zhebentyayeva³, Luis Rodriguez-Saona⁴, Pierluigi Bonello⁴, Joseph James⁵, Steven Jeffers³, Paul Sisco⁶, Frederick Hebard⁶, Laura Georgi², Margaret Staton⁷, Jean-Marc Audergon⁸, Veronique Decroocq⁹, Zongrang Liu¹⁰, Christopher Dardick¹⁰, C. Dana Nelson^{1,11}, and Albert Abbott¹

¹University of Kentucky, Forest Health Research and Education Center, ²The American Chestnut Foundation, ³Clemson University, ⁴The Ohio State University, ⁵Chestnut Return Farm, ⁶Retired-The American Chestnut Foundation, ⁷University of Tennessee, ⁸Institut National de la Recherche Agronomique, Avignon, ⁹Institut National de la Recherche Agronomique, Bordeaux, ¹⁰USDA-ARS Appalachian Fruit Research Station, ¹¹USDA-FS Southern Research Station



Outline

- An introduction to metabolite-based screening
- Evaluating chemical fingerprinting as a tool to screen chestnut for disease resistance
- Targeted metabolomics to track developmental progression in peach and apricot
- Conclusions



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Plant secondary metabolites (PSMs)

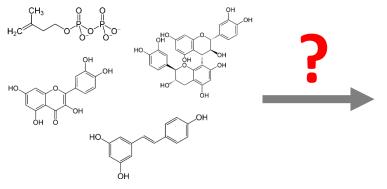
- PSMs in particular are known for their association with tree stress response
- Types and amounts are tree-species dependent
- Temporally and spatially variable





Chemical biomarkers

- Certain chemicals may be associated with a trait of interest
 - e.g. disease resistance/susceptibility
- Goal is to identify chemicals that may serve as biomarkers
- In some cases, statistical models can be developed to predict the trait based solely on a tree's chemical composition



ages from: http://www.wikipedia.or



https://www



Metabolite-based screening approaches

- Identify and quantify specific metabolites or measure general profiles
- Different tools for different objectives/questions
- Examples of different analytical approaches include:
 - FT-IR spectroscopy
 - HPLC-MS





Overall objectives

- Evaluate general metabolomic profiles for the identification of chemical fingerprints linked to pathogen resistance in American x Chinese chestnut hybrids
- Evaluate specific intermediates within the phenylpropanoid pathway as biomarkers for developmental progression linked to annual climatic cycling in fruit trees



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What is chemical fingerprinting?

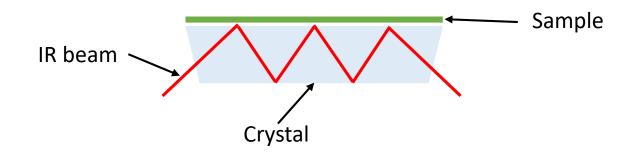
- **Chemical fingerprinting** is a comprehensive analysis of all the chemicals present within a given sample
- Individual chemicals are not separated, identified, or measured
- Provides a "snap-shot" of the chemical composition of a given tissue at a given time





Fourier transform infrared spectroscopy

- Fourier transform infrared (FT-IR) spectroscopy is one method of chemical fingerprinting
- FT-IR spectroscopy measures how a sample absorbs light over a wide spectral window
- Differences in chemical composition/concentration will affect the FT-IR spectrum





FT-IR spectrometers

Benchtop spectrometer

"Portable" spectrometer





Handheld spectrometer





Chestnut chemical fingerprinting

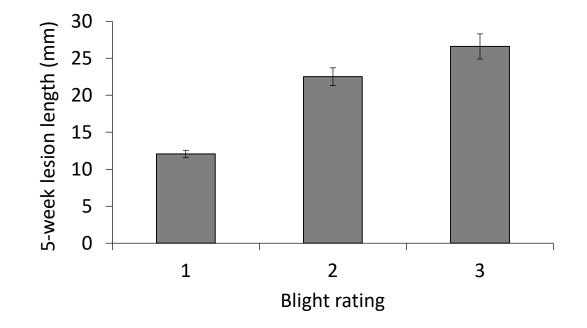
Evaluate if chemical fingerprinting can be used to screen hybrid chestnut for disease resistance prior to infection with chestnut blight or Phytophthora root rot



Photo credit: Linda Haugen, USDA Forest Service, Bugwood.org



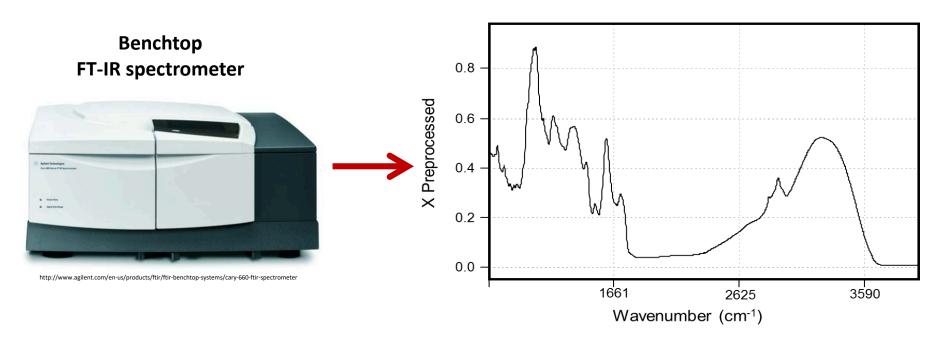
Phenotypic variation is necessary to build predictive models for disease resistance



Average (± standard error) lesion length of BC_3F_3 hybrids derived from Clapper for each blight rating group (N = 41). Phenotypic data and tissue samples provided by J. Westbrook (TACF).



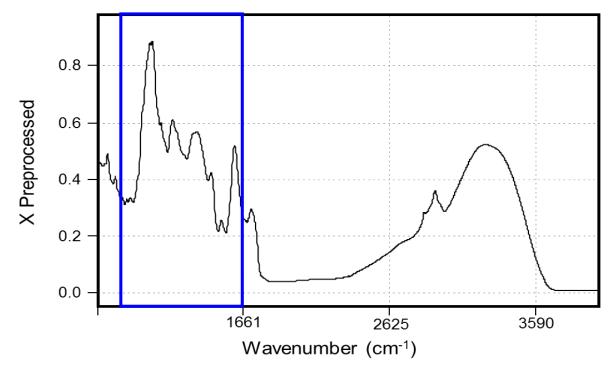
Chestnut chemical fingerprints



A representative hybrid chestnut spectrum collected from the mid-infrared region.



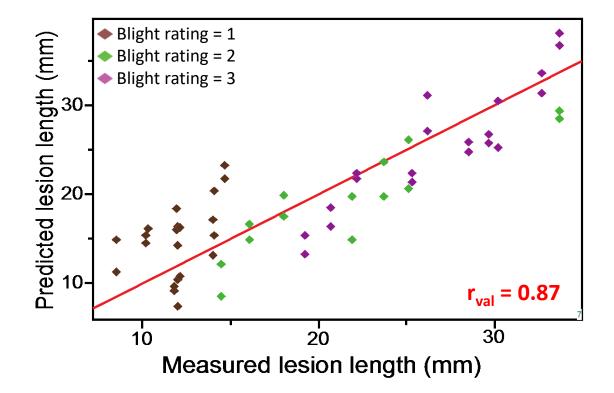
Focus on a specific spectral region



For blight analysis, focused on spectral range: 901 – 1622 cm⁻¹.



Susceptibility of 'Clapper' hybrids to blight can be estimated using spectral data



Correlation plot from 7-factor partial least squares regression analysis of Clapper data set showing the relationship between measured and predicted lesion lengths, based on transformed spectral data. N = 55, with 2 technical replicates per biological replicate and outliers trimmed based on preliminary analysis.



Application of chemical fingerprinting

- Chemical fingerprinting has the potential to be a useful tool for screening hybrid chestnut for disease resistance
- Additional evaluations are planned for summer 2017
- Potential for more highthroughput analysis in the field/forest



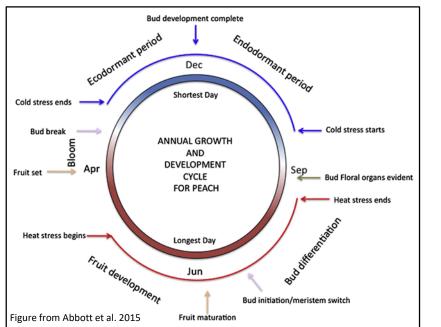


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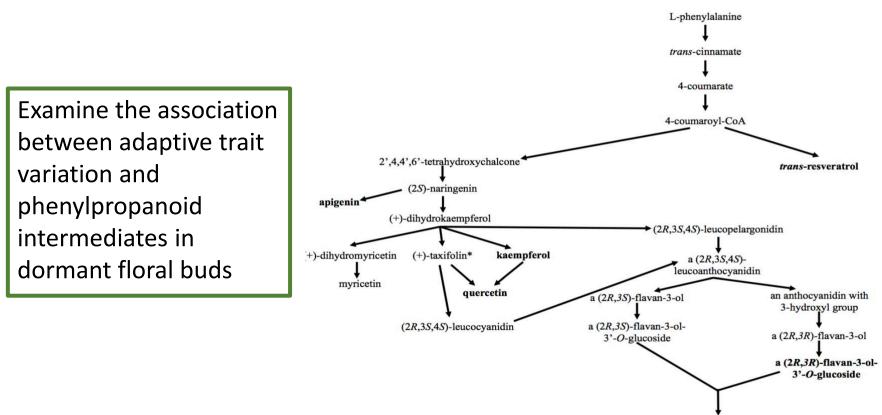
Developmental progression and adaptive traits



- Phenology is the study of cyclic events and the influence of climatic and seasonal variability on these events
- Dormancy, bloom date, and chilling requirement are examples of adaptive phenological traits
- Trait variability may be of interest to breeding programs



Identifying chemical markers for developmental progression



a proanthocyanidin

Simplified phenylpropanoid biosynthetic pathway in peach.



Adaptive traits

Species	Phenotype	Variety/Individual
Peach	Low-chill	A209
	Low-chill	A340
	High-chill	A318
	High-chill	A323
Apricot	Early flowering	A2312
	Early flowering	A2137
	Early flowering	A1956
	Late flowering	A660
	Late flowering	A1267

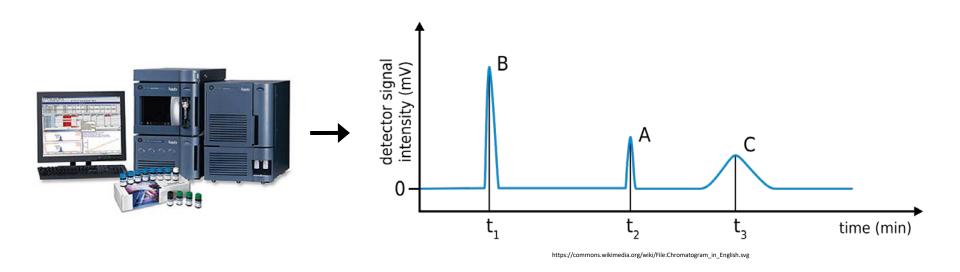




Photo credit: (TOP) Carroll E. Younce, USDA Agricultural Research Service, Bugwood.org; (BOTTOM) Howard F. Schwartz, Colorado State University, Bugwood.org



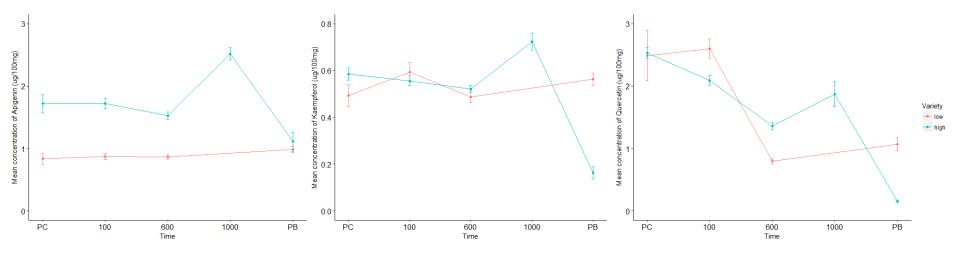
Targeted metabolomics analysis



Flavonoids: apigenin, kaempferol, quercetin **Flavan-3-ols**: epicatechin-3'-*O*-glucoside; Procyanidin B1, B2, B3 **Stilbene**: resveratrol



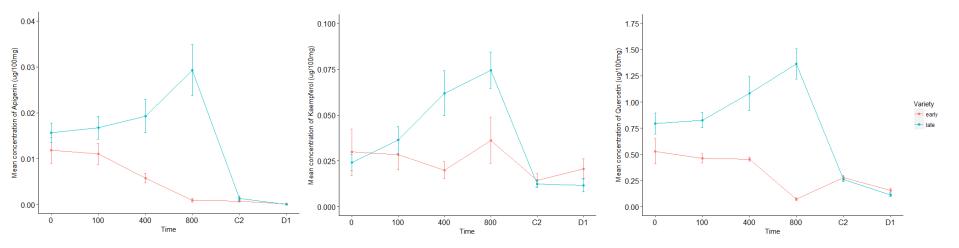
Flavonoid concentrations change during dormancy and are impacted by tree genotype in peach



Concentrations of flavonoid aglycones, apigenin, kaempferol, and quercetin, were significantly affected by the interaction of time (at 100, 600, and PB) and chilling requirement (i.e. genotype). Repeated measures ANOVA (P < 0.05).



A similar pattern is observed in apricot



Concentrations of flavonoid aglycones, apigenin, kaempferol, and quercetin, were significantly affected by the interaction of time and chilling requirement (i.e. genotype). Repeated measures ANOVA (*P* < 0.05).

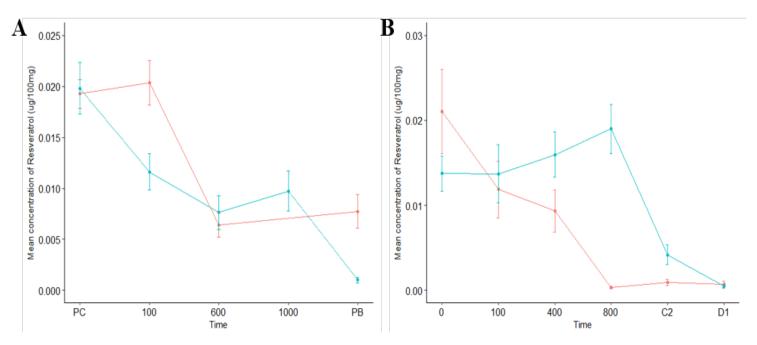


Phenylpropanoids as biomarkers for developmental progression

- Concentrations of phenylpropanoid intermediates change as dormancy progresses
- Pronounced differences in concentration between endodormant and post-dormant buds
- Timing of changes is impacted by phenotype/genotype



Changing in response to environmental cues?



Resveratrol in peach (**A**) and apricot (**B**) buds during dormancy in trees that varied in chill requirement (CR) and bloom date (BD). **Red**: low. **Blue**: high.

Concentrations of resveratrol are known to be impacted by changes in water availability



Application of chemical biomarkers

- Alternative method for determining developmental stage
- Predict low versus high chill genotypes/phenotypes



Photo credit: University of Georgia Plant Pathology , University of Georgia, Bugwood.org



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Conclusions

- Different metabolomics approaches can be used to generate chemical phenotypes (chemotypes) of trees
- Chemotypes may be associated with trait of interest
 - Disease resistance/susceptibility (e.g. chestnut)
 - Developmental progression (e.g. peach and apricot)
- Application for tracking biotic and abiotic stress performance in tree improvement programs

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