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Enzymatic Testing in a Wine Laboratory

What to look out for and how to do it well

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Webinar Formalities

- This Webinar is being recorded and will be published on the Enartis website
- Please refrain from using the chat box during the presentation, there will be 15 minutes for questions at the end of the presentation
- If you are having technical difficulties please use chat box 2, Whitney will be there to help
- Please complete the survey at the end of the webinar



Malic Acid
Glucose and Fructose
Acetic Acid

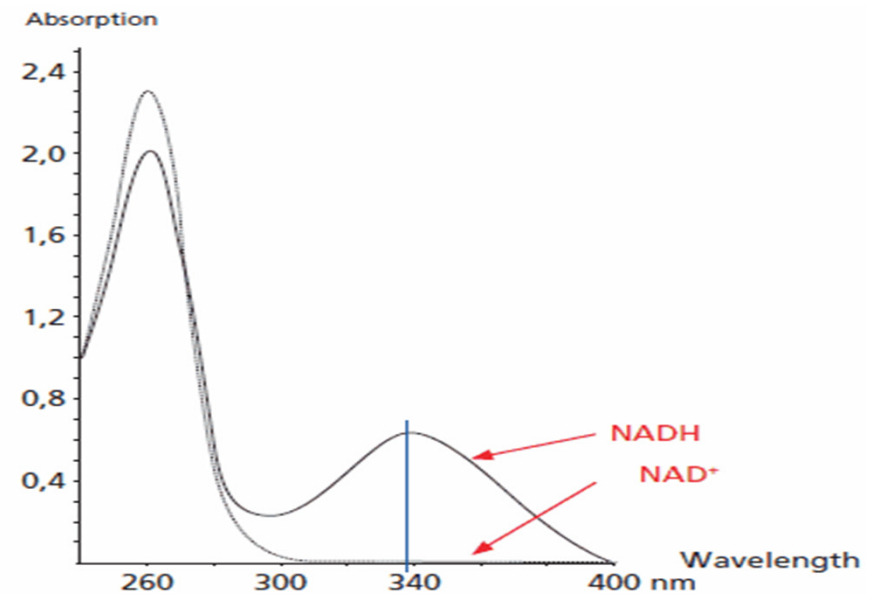


Reduction of NAD⁺
to NADH

Ammonia
Citric Acid



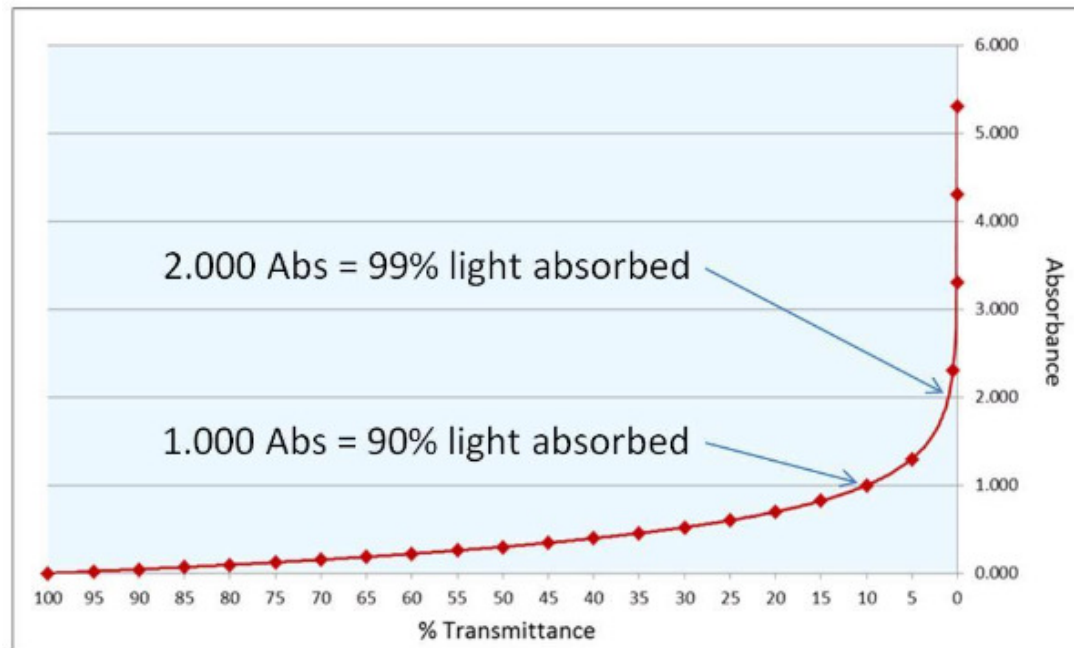
Oxidation of NADH to
NAD⁺





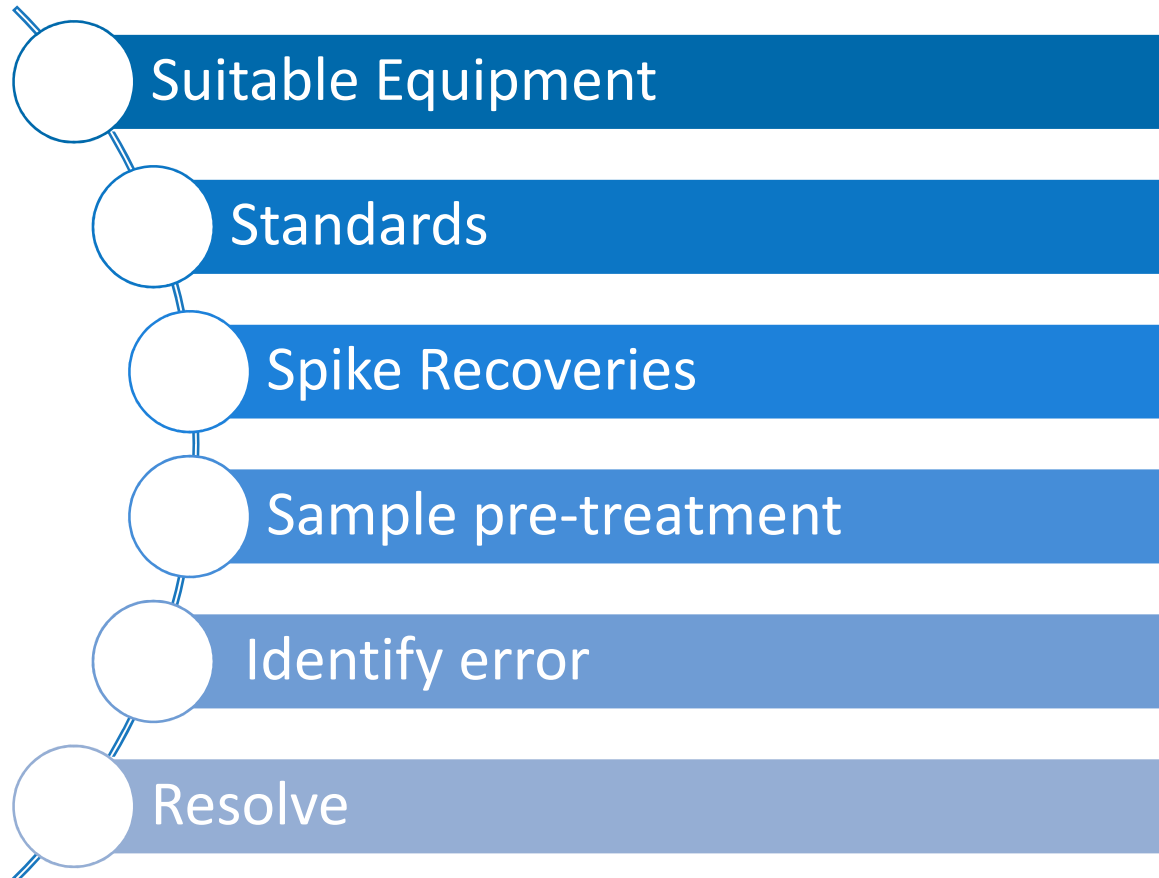
A closer look at absorbance...

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Spectrophotometers measure the light that is **not** absorbed by the sample

<1.5 ABS



What are you trying to achieve?



Is your equipment suitable?

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Spectrophotometer

- Research thoroughly
- Calibrate as specified by manufacturer
- Follow a maintenance schedule
- Service as recommended





Is your equipment suitable?

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Pipettes

- Clean
- Calibrated and checked
- Handled appropriately
- Tips fit well and are clean





Is your equipment suitable?

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Enzymes/Kits

- Stored cold
- In date
- No mixing of batches
- Lids shut when not in use
- Suitable for desired outcome





Is your equipment suitable?

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Water source

- Free of mould and metal contamination

Cuvettes

- Free of contamination
- Right path length
- No scratches/marks



Confidence in results

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Model Wine Solutions

- Model standard of the analyte
- Clear matrix
- Can include small amounts of ethanol
- Commercially available
- Easy to make your own water/analyte mix to desired concentration
- Used in spike recoveries





Secondary Wine Standards

- Ensures no matrix interference occurring
- Checks necessary pre-treatments are correct
- Choose a wine/s that represents a similar matrix and at a level of expected concentration of analyte
- Plot on a control chart to identify outliers and bias easily



Spike Recovery

Test: Malic acid determined by discrete analyser, calibration range 0 – 3g/L

Procedure: Analyse sample (**Y**)

Add 1mL of the top calibration standard (3g/L) to 4mL of sample

Analyse this spiked sample (**Z**)

Calculation: % Recovery = $\frac{z}{(0.8 \times y) + (0.2 \times 3)} \times 100$

Interpretation: For most tests, a recovery of between 95-105% is acceptable

✓ **Measures trueness & identifies possible bias**



Pre-Dilution

- Review recommended dilution rates in instructions
- Dilute so sample concentration is under the upper limit of the kit/method
- **Don't over dilute**
- Use the lowest possible dilution to reduce error
- Check accuracy of dilutions using standards



Degassing

Bubbles can cause..

- Reduced assay transmittance
- Sensing errors in DA's
- Random error in pipettes and DA probes occurring during aspiration



Decolourisation

- Coloured wines can reduce assay transmittance
- PVPP
- *May* get away with just pre-dilution (if needed as well)
- Avoid using alternative methods without trialling



Filtering

- Highly turbid wines reduce assay transmittance
- Solids block tubing and probes in DA'S
- Filter through 0.45um



$$\text{Concentration} = \frac{(\text{Final volume} \times \text{MW})}{(\epsilon \times \text{light path} \times \text{sample volume} \times 1000)} \times \Delta A$$

✓ **Always perform a blank and a duplication!**





Watch out for **low recoveries**

- Kit expired or stored incorrectly
- Reagents too cold
- Working outside kit testing range
- Incorrect sample prep
- Short incubation time
- Incorrect reagent prep
- High blank recovery




Watch out for **high recoveries**

- Contamination (eg. Cuvettes, pipette tips)
- Incubation time incorrect
- Incorrect reagent prep
- Poor photometric accuracy
- Too much sample added



Watch out for **fluctuating absorbance**

- 
- Incorrect sample pre-treatment
 - Instrument faults
 - Cuvette lined up incorrectly
 - Unstable power supply



Watch out for **random error**



- Contaminated cuvettes
- Inaccurate pipetting
- Method discrepancies
- Cuvette facing wrong way
- Human error



Watch out for **bias**

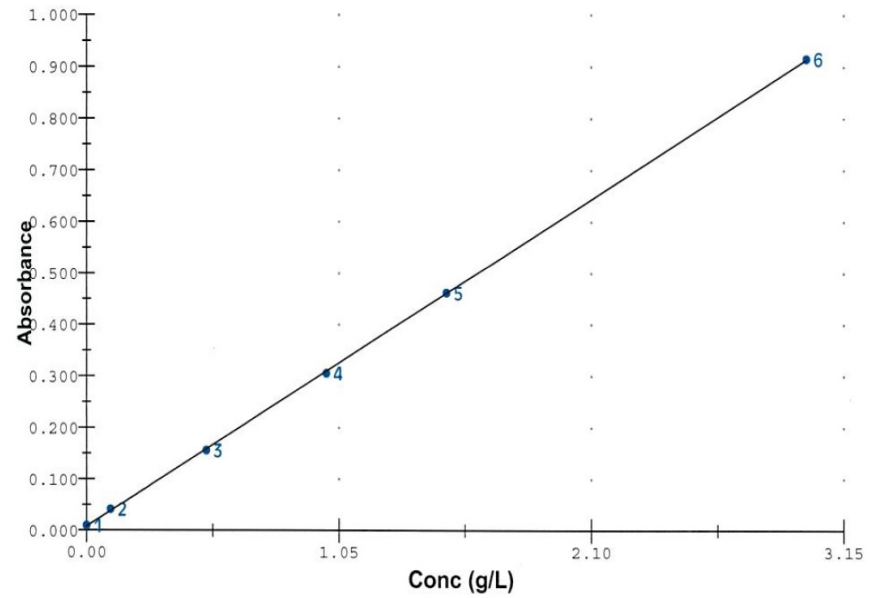
- Poor kit storage
- Gradual contamination over time
- Pipettes dispensing high or low
- Pre-dilutions are inaccurate



The Automated Way



Assay Curve
Abs = 0.0075664 + 0.30239 * Conc
Correlation (r squared) = 0.99993





How to test well

- ✓ Trial method first
- ✓ Responses that range from 0 to 1.
- ✓ Keep enzyme VS sample ratios the same
- ✓ Calibrate before every run
- ✓ Analyse your data
- ✓ Clean routinely

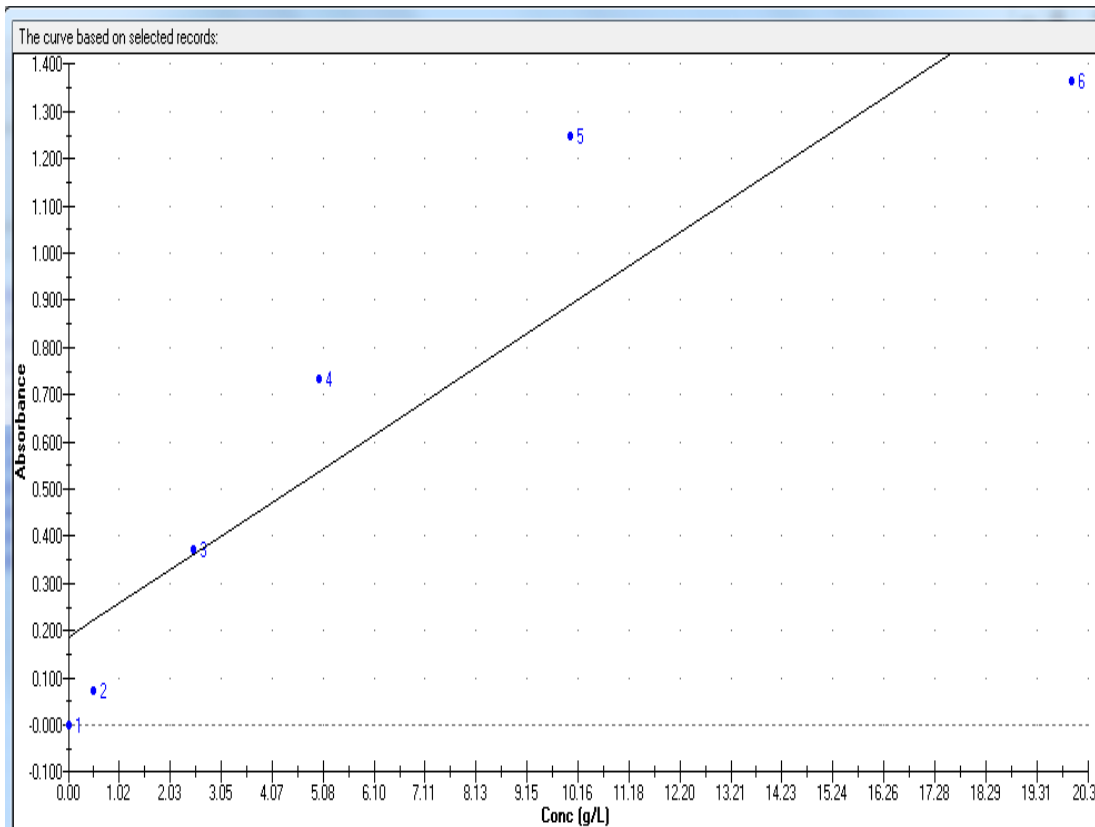


Watch out for...

- ✗ Mixing old reagents with new
- ✗ Evaporation of reagents
- ✗ Contamination
- ✗ Poor calibrations



Watch out for...

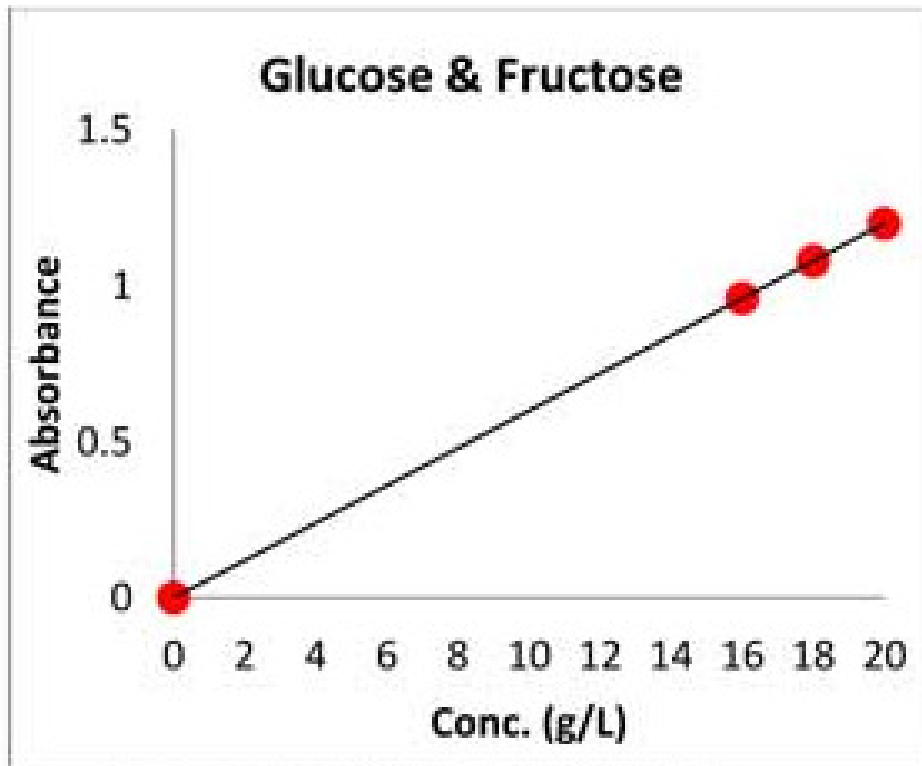


Name	Absorbance	Concentration	Interpretation	Note	Referent
22/05/2018 2:55:40 PM					
<input checked="" type="checkbox"/> Vint Comb STD 1	0.000 Abs => 0.000 Abs	0.00 g/L => 0.00 g/L			0.00
<input checked="" type="checkbox"/> Vint Comb STD 1	-0.003 Abs => -0.003 Abs	0.00 g/L => 0.00 g/L			0.00
<input checked="" type="checkbox"/> Vint Comb STD 2	0.073 Abs => 0.073 Abs	0.00 g/L => 0.00 g/L			0.50
<input checked="" type="checkbox"/> Vint Comb STD 2	0.076 Abs => 0.076 Abs	0.00 g/L => 0.00 g/L			0.50
<input checked="" type="checkbox"/> Vint Comb STD 3	0.381 Abs => 0.381 Abs	2.76 g/L => 2.76 g/L			2.50
<input checked="" type="checkbox"/> Vint Comb STD 3	0.360 Abs => 0.360 Abs	2.46 g/L => 2.46 g/L			2.50
<input checked="" type="checkbox"/> Vint Comb STD 4	0.758 Abs => 0.758 Abs	8.13 g/L => 8.13 g/L			5.00
<input checked="" type="checkbox"/> Vint Comb STD 4	0.709 Abs => 0.709 Abs	7.44 g/L => 7.44 g/L			5.00
<input checked="" type="checkbox"/> Vint Comb STD 5	1.250 Abs => 1.250 Abs	15.15 g/L => 15.15 g/L			10.00
<input checked="" type="checkbox"/> Vint Comb STD 5	1.245 Abs => 1.245 Abs	15.08 g/L => 15.08 g/L			10.00
<input checked="" type="checkbox"/> Vint Comb STD 6	1.363 Abs => 1.363 Abs	16.76 g/L => 16.76 g/L			20.00
<input checked="" type="checkbox"/> Vint Comb STD 6	1.368 Abs => 1.368 Abs	16.83 g/L => 16.83 g/L			20.00

✓ See solutions for low recoveries



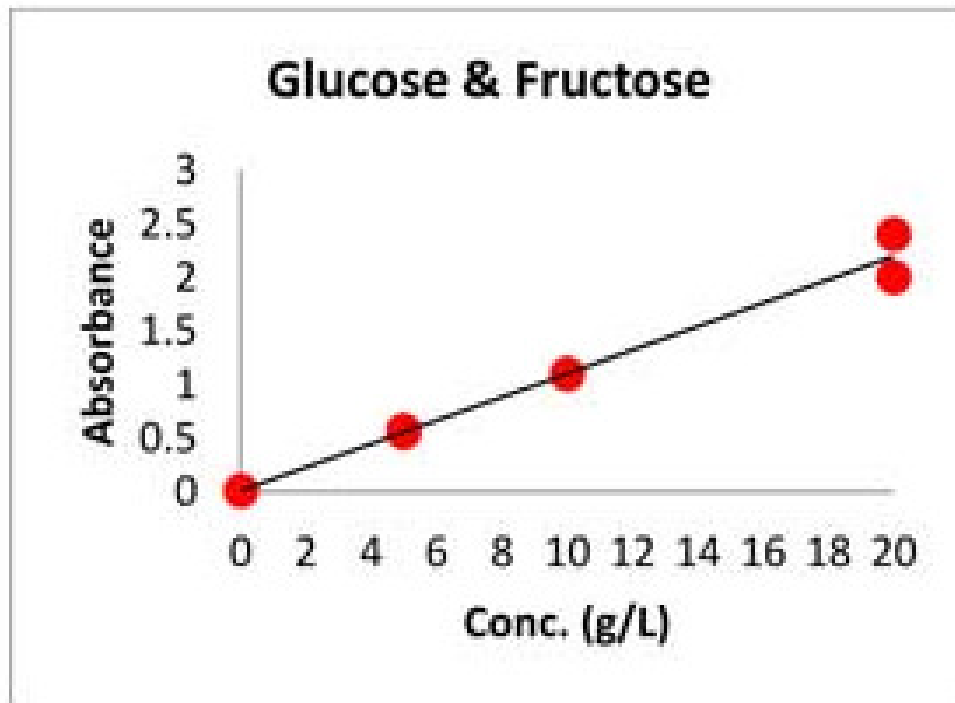
Watch out for...



✓ Trial any new methods



Watch out for...



✓ Remember,
<1.5 ABS



What should you do if you get a poor calibration?

- Re calibrate!
- Omit large single outliers
- Ensure reagents are loaded into the right position
- Check you're using the right method
- Investigate enzymes/kit



The right results the first time!







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