

# Where did my analyte go?

Coping with poor solubility and non-specific binding

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**175**  
years

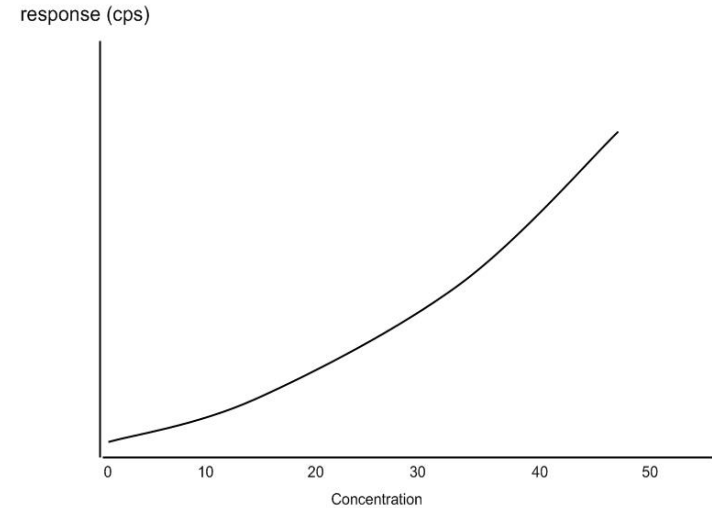
# Poor solubility and non-specific binding

- **Often overlooked issues**
- **Prominent in low-protein matrices, *i.e.* urine, CSF**
- **Can cause:**
  - > Non-linear standard lines
  - > Inaccuracy of QC's
  - > Assay variability
  - > Loss of sensitivity
- **Difficult to identify**
- **Difficult to resolve**



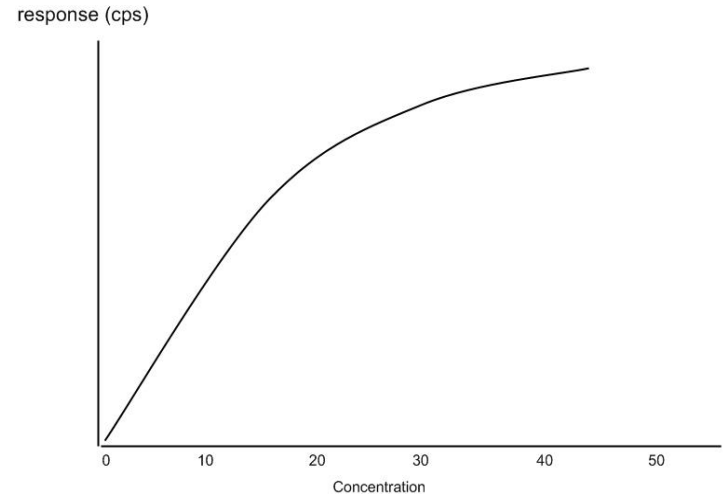
# Non-specific binding

- Adsorption of analyte onto the surface area of the container
- Higher impact at lower analyte concentrations
- Non-linear calibration line
- Adsorption depends on:
  - > Analyte properties
  - > Temperature
  - > Matrix / solvent
  - > Container



# Poor solubility

- **Poor dissolution of analyte in solvent**
- **Higher impact at higher analyte concentrations**
- **Non-linear calibration line**
- **Analyte variability from a single well**
- **Solubility depends on:**
  - > Analyte properties
  - > Temperature
  - > Matrix / solvent



# **Non-specific binding**

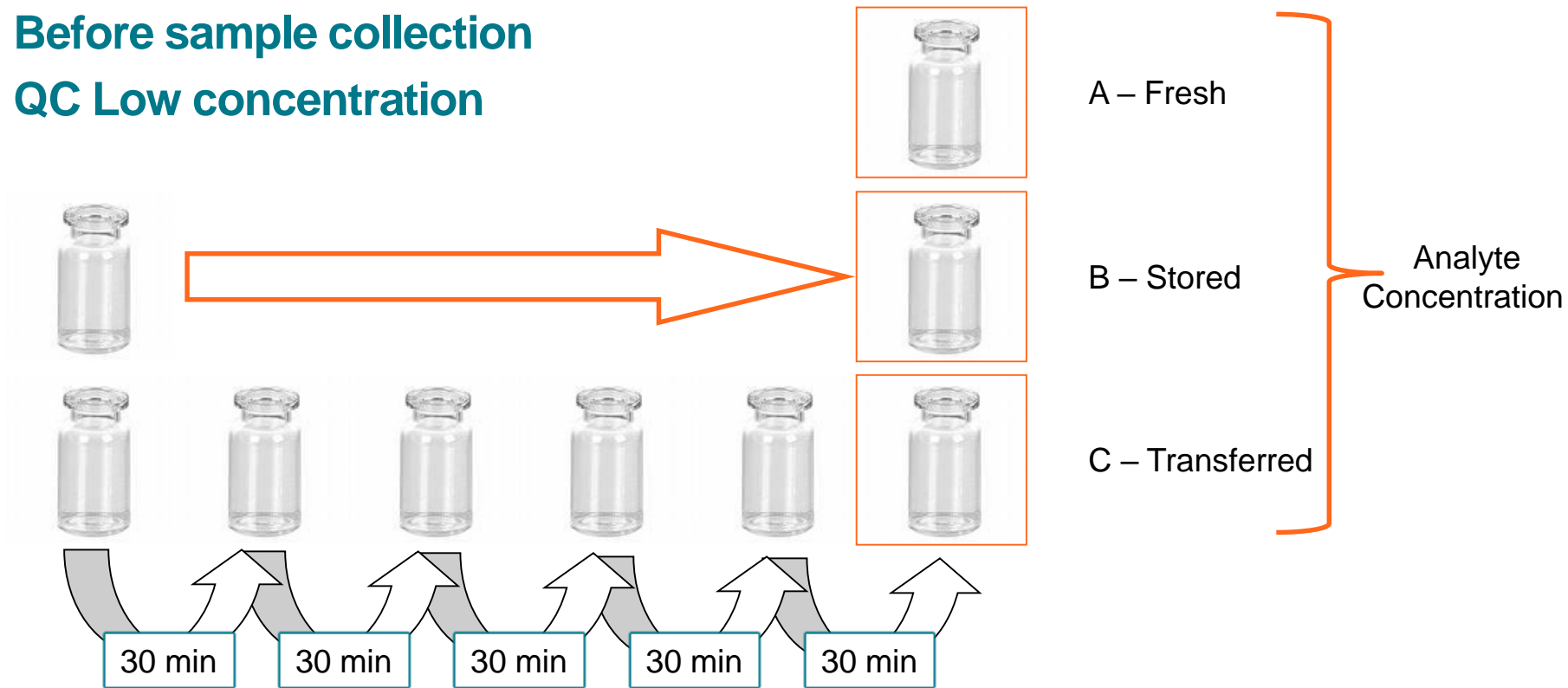
# Non-specific binding

- **Analyte adsorbs onto container surface**
  - > Electrostatic interactions
  - > Non-specific interactions
- **Binding may be affected by:**
  - > Matrix (ionic strength, pH, lipids and protein content)
  - > Container material
  - > Container brand or quality
  - > Freeze/thaw or long term storage



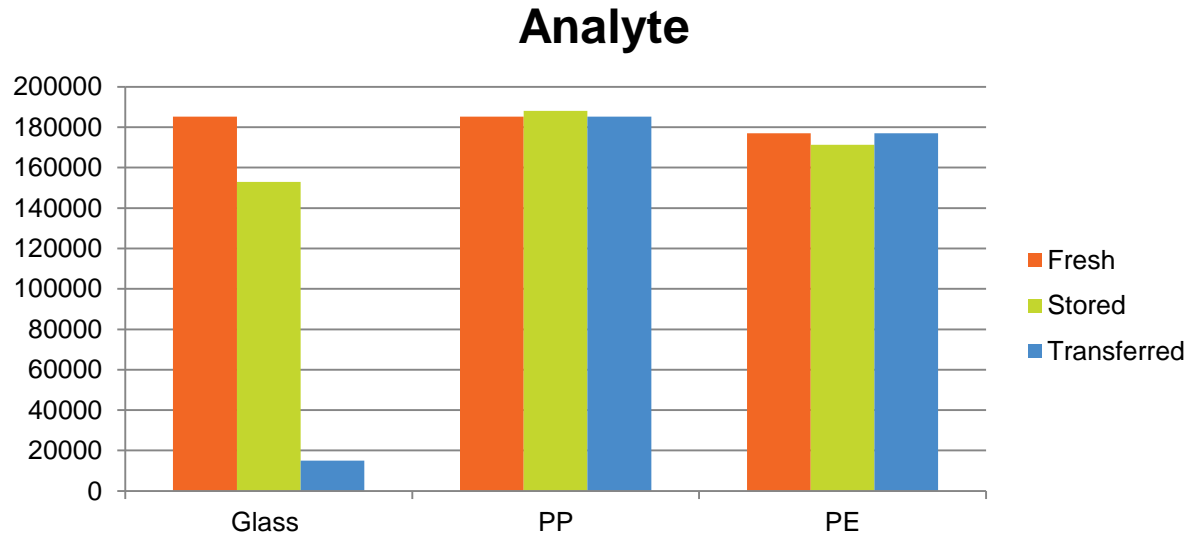
# Non-specific binding - assessment

- During method development
- Before sample collection
- QC Low concentration



# Non-specific binding - assessment

- Glass, polypropylene, polyethylene
- Water, dark urine, light urine





# Non-specific binding - solutions

- **Change of container**

- > Simplest solution
- > Same material different container
- > Eppendorf LoBind<sup>®</sup> - modified PP
- > Wheaton<sup>®</sup> AntiBIND<sup>™</sup> - modified PP surface



- **Addition of organic**

- > MeOH, MeCN, DMSO, DMF or IPA
- > Decrease urine polarity and therefore increase analyte solubility
- > Solvent may attach to binding sites?
- > % solvent needs to be optimised



# Non-specific binding - solutions

- **Addition of acid / buffers**

- > Decreases electrostatic charge from surface?

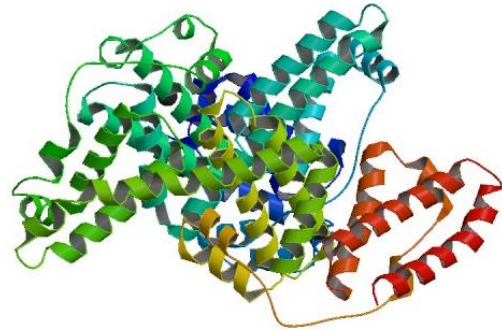
- > pH control

- Urine pH varies between 4.5 and 8.
    - Change of pH can affect **ionisation state** of analyte and therefore adsorption/solubility



# Non-specific binding - protein

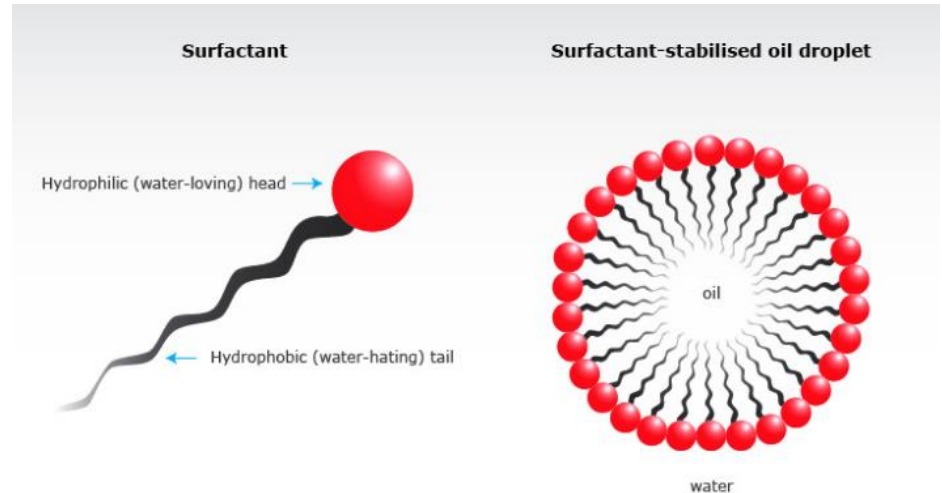
- **Addition of protein**
  - **Bovine Serum Albumin (BSA)**
    - > Serum albumin protein derived from cows
    - > Mainly used with large molecules
    - > Expensive
    - > Can interfere with the extraction
    - > Can contain biomarkers!!



# Non-specific binding - solutions

- **Addition of surfactants**

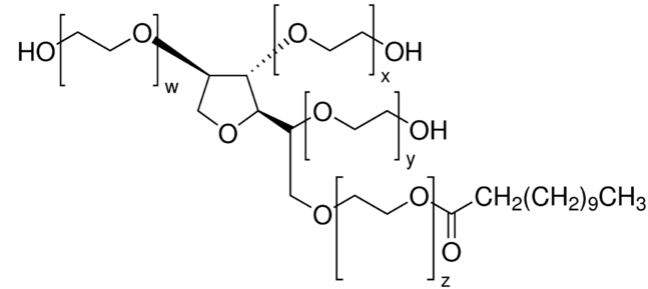
- > Detergents
- > Organic compounds that contain hydrophobic (*tails*) and hydrophilic groups (*heads*)
- > Coat surface?
- > Increase analyte solubility
- > May reduce analyte sensitivity
- > May be difficult to clean from MS



# Non-specific binding - surfactants

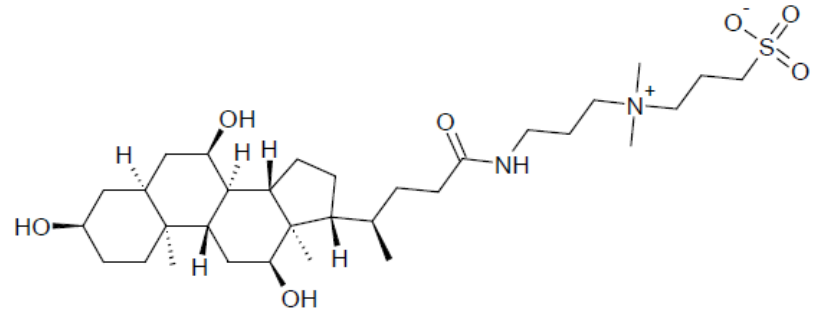
- **Tween 20 & Tween 80**

- > Most commonly used in our labs
- > Small concentration required
- > Cheap and non-toxic



- **CHAPS**

- > Zwitterionic surfactant
- > Used to solubilise biological macromolecules
- > Expensive



# Non-specific binding – solutions

- Anti-adsorbing agents (AAA) - pre or post sample collection?
- **Post-collection**
  - > Ideal scenario
  - > Reversible adsorption



Urine volume calculated from weight

# Non-specific binding - post-collection

- **Advantages**

- > Same % of AAA added to each sample
- > Standards and QC's treated the same way
- > No need to correct sample concentrations
- > Urine collection bottles don't need to be pre-filled with chemicals

- **Disadvantages**

- > Primary containers may need to be shipped to the laboratories
- > Additional sample analysis time with sample pre-treatment



# Non-specific binding – pre-collection

- Pre-collection

> When non-specific binding is irreversible



Urine volume calculated from density



# Non-specific binding – pre-collection

- **Advantages**

- > Easier sample pre-treatment

- **Disadvantages**

- > Not all AAA suitable ( $\text{H}_3\text{PO}_4$ , FA, etc)

- > Individual sample concentrations need to be corrected

- > A range of % AAA needs to be validated



# Solubility

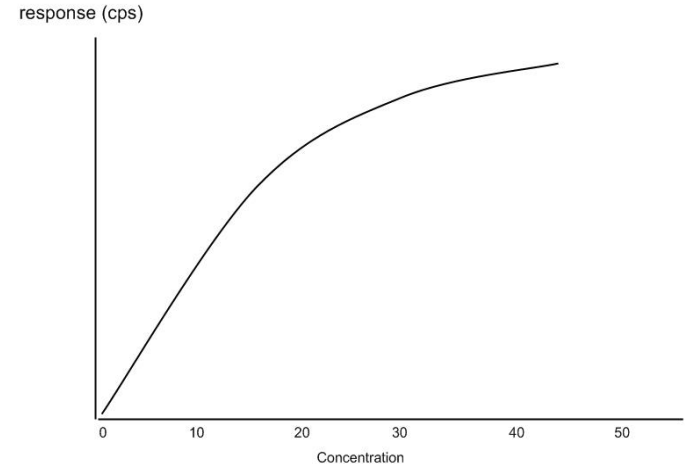
# Solubility

- **A compound is soluble when it forms a homogeneous solution with a solvent**
- **Solubility depends on:**
  - > Analyte properties
    - Molecules with a similar polarity to water will dissolve well in water
  - > Temperature
    - Solids become more soluble as temperature increases
  - > Matrix
    - pH and protein content



# Poor solubility - assessment

- **Poor solubility of an analyte results in a solution that is not homogeneous**
- **Solution **may** look homogeneous**
- **More difficult assessment than non-specific binding**
  - > Non-linear calibration lines
  - > Poor CV's from repeat injections
- **Poor solubility may appear after storage**
  - > Lower temperatures upon storage
  - > Slow analyte precipitation



# Poor solubility - solutions

- **Similar to non-specific binding**
- **Addition of organic**
  - > MeOH, MeCN, DMSO, DMF or IPA
  - > Decrease urine polarity and therefore increase analyte solubility
  - > % solvent needs to be optimised
- **Addition of acid / buffers**
  - > pH control
    - Urine pH varies between 4.5 and 8.
    - pH of urine can affect **ionisation state** of analyte and therefore adsorption/solubility
- **Addition of surfactants**



# Poor solubility - solutions

- **Sonication**

- > Sound energy agitates particles using ultrasonic frequencies
- > Can be used to speed dissolution, by breaking intermolecular interactions **if the solute is soluble**
  
- > However:
  - **Supersaturated** solutions may be formed – precipitation/ crystallisation may occur upon storage
  - Prolonged sonication increases sample temperature – analyte stability?
  - Inconsistent results
  
- > SONICATION SHOULD BE AVOIDED



# Conclusions

- **Solubility and non-specific binding are difficult to identify**
- **Prominent in low protein matrices**
- **Always investigated during method development and prior to sample collection**
- **Interconnected issues – similar solutions**



