

SAMPL8 Preview GCC 2020 and SAMPL Satellite Workshop

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Overview - Goals

SAMPL8 Project



- Aim #1: Develop automated approach for determining ionization constants (pK_a) on multiple small molecules
- Aim #2: Develop automated high throughput method for measuring distribution coefficients (logD)
- Aim #3: Measure logD for 20-30 compounds for a variety of solvent pairs

Why?

- Understanding the protonation state of a small molecule helps to build computational models for predicting how macromolecules will behave (https://doi.org/10.1007/s1082 2-018-0168-0)
- Distribution coefficient values provide for computational models that predict protein-ligand binding affinities or hydration/solvation free energies (http://dx.doi.org/10.1101/757393)

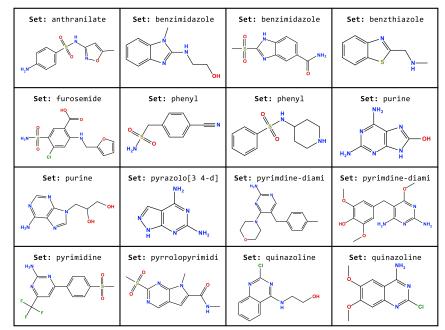
88 Compounds Selected for Evaluation



Molecular weight ranges from 135 to 476 Dalton

Several Key Characteristics

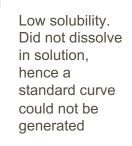
- Two long range polar groups (separated by > 3 bonds)
- Molecules chosen for having a wide variety of scaffolds
- 100 mg of available powder needed for full testing

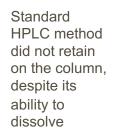


Example representative structures

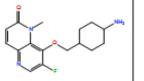
Experimental pK_a Challenges

- Not all of the selected compounds were successful in pKa determination...
- The team started with 88 small molecules, but succeeded in measuring 24... Why?
 - Some of the compounds had degraded over time, while in storage
 - Some of the compounds provided had insufficient amounts available for testing
 - Several of the compounds were semisolids and could no be dispensed for testing









Only able to

solubility which

cannot be used

of compound

to determine pK_a

determine

intrinsic

Why Automate?





Current Manual Process

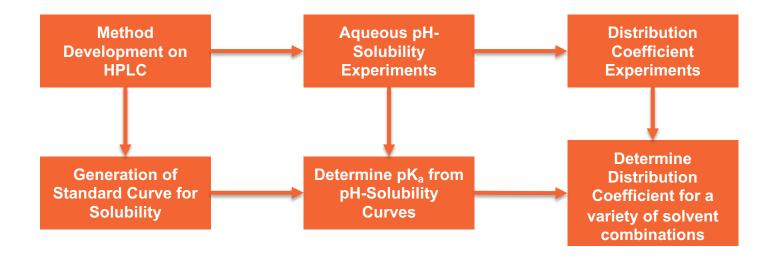
- Less data generated
- Only one or two reactions at one time
- Time consuming
- Data inaccuracy / human error
- Resource and material heavy



- Greater data generation
- Higher throughput
- Greater accuracy / precision
- Reduce resource required
- Combine multiple analytical techniques

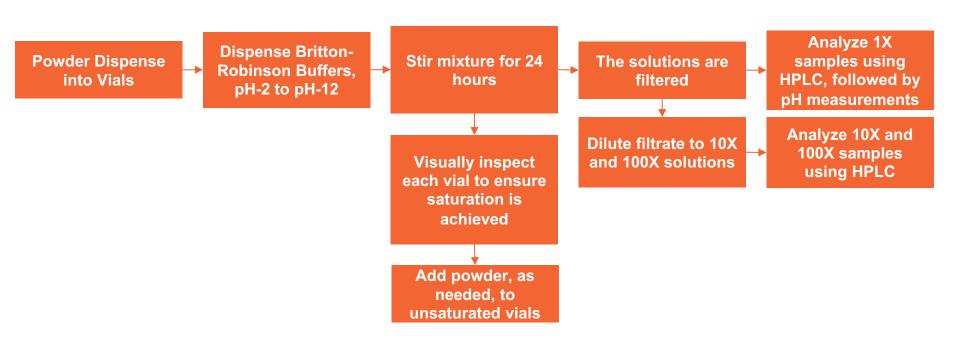
General Overview of the High Throughput Process





pKa Determination: Generation of pH Solubility Curve



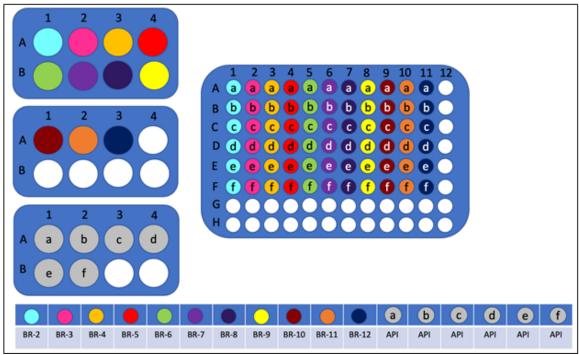


pKa Determination: Generation of pH Solubility Curve



Using Freeslate (Unchained Labs) CM3, samples are prepared using several source and destination vials in a mapping scheme illustrated here



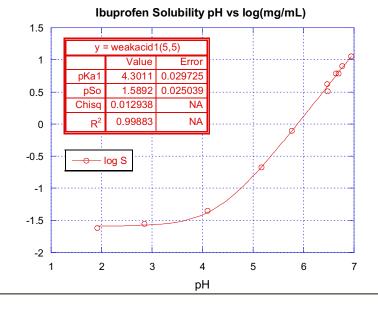


Curve Fitting to Determine pK_a



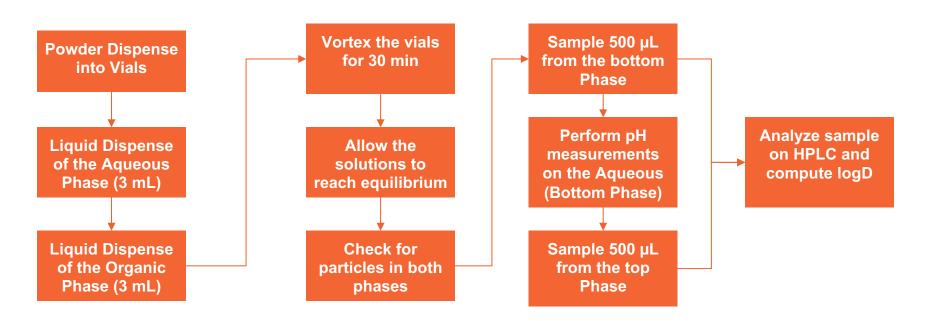
Henderson-Hasselbalch Equations $pH = pK_a + log_{10} \frac{[A^-]}{[HA]}$ $pH = pK_a + log_{10} \frac{S - S_0}{S_0}$ Weak mono-acids $pH = pK_a + log_{10} \frac{S_0}{S - S_0}$ Weak mono-bases

Where S_0 is the intrinsic solubility (equilibrium solubility) and the S is the total solubility



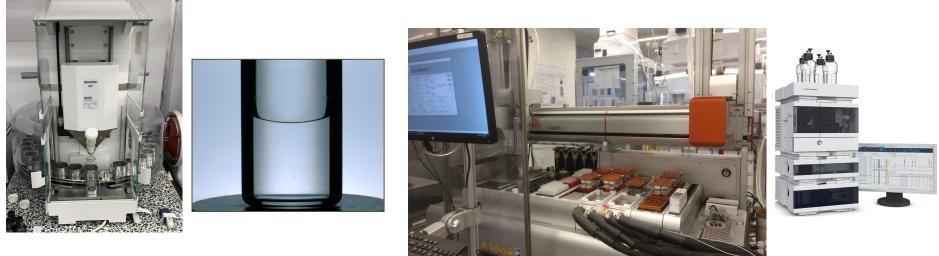
log S

High Throughout Approach for Distribution Coefficient Measurements





High Throughout Method for Distribution Coefficient -Automated Platforms

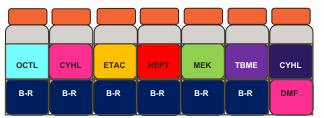


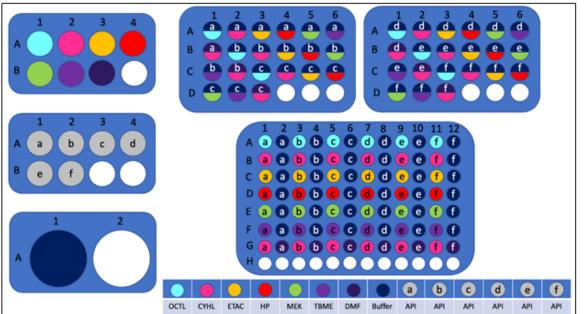
Experimental Sample Preparation



High Throughout Method for Distribution Coefficient Measurement

 Using Freeslate (Unchained Labs) CM3, samples are prepared using several source and destination vials in a mapping
scheme illustrated here







Workflow in Action



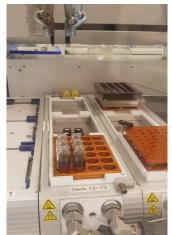
Density vials weighed



Samples transferred to pre-tared vials



Images taken of reaction vials



Density sample taken from reaction vials





- Created a high-throughput approach to HPLC method development
- Developed an automated pH-solubility workflow for pKa determination
- Developed a fully automated distribution coefficient workflow
- Successfully sampled from organic and aqueous phases
- Increased amount of data generated in a short time

Acknowledgements



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- David Mobley Professor of Pharmaceutical Sciences
- Teresa Danielle Bergazin PhD Candidate



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