



Skyline Tutorial Webinar #22

Welcome to the next in the series of tutorial webinars designed to help you get the most out of Skyline mass spectrometry software.

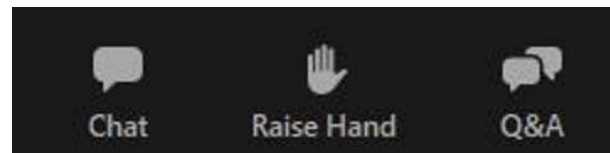
We'll begin shortly ...

- **Webinar Audio:** Make sure you have audio during the webinar.

Click **Audio Settings** in the lower left-hand corner of the screen :

- Computer audio (preferred) – through your computer's speakers or a headset
- Phone call (US and International numbers available) By phone, use conference ID: **977 3811 3390**

****Note: we currently have music playing which you should be able to hear.****



- **Submit Skyline Questions:** the Q&A form in Zoom to submit Skyline questions to the presenters
- **Have logistic issues?** Use the **Chat** applet in Zoom for report webinar software (Zoom) issues or other problems.



Tutorial Webinar #22

Using DIA Data To Create SRM Methods

With

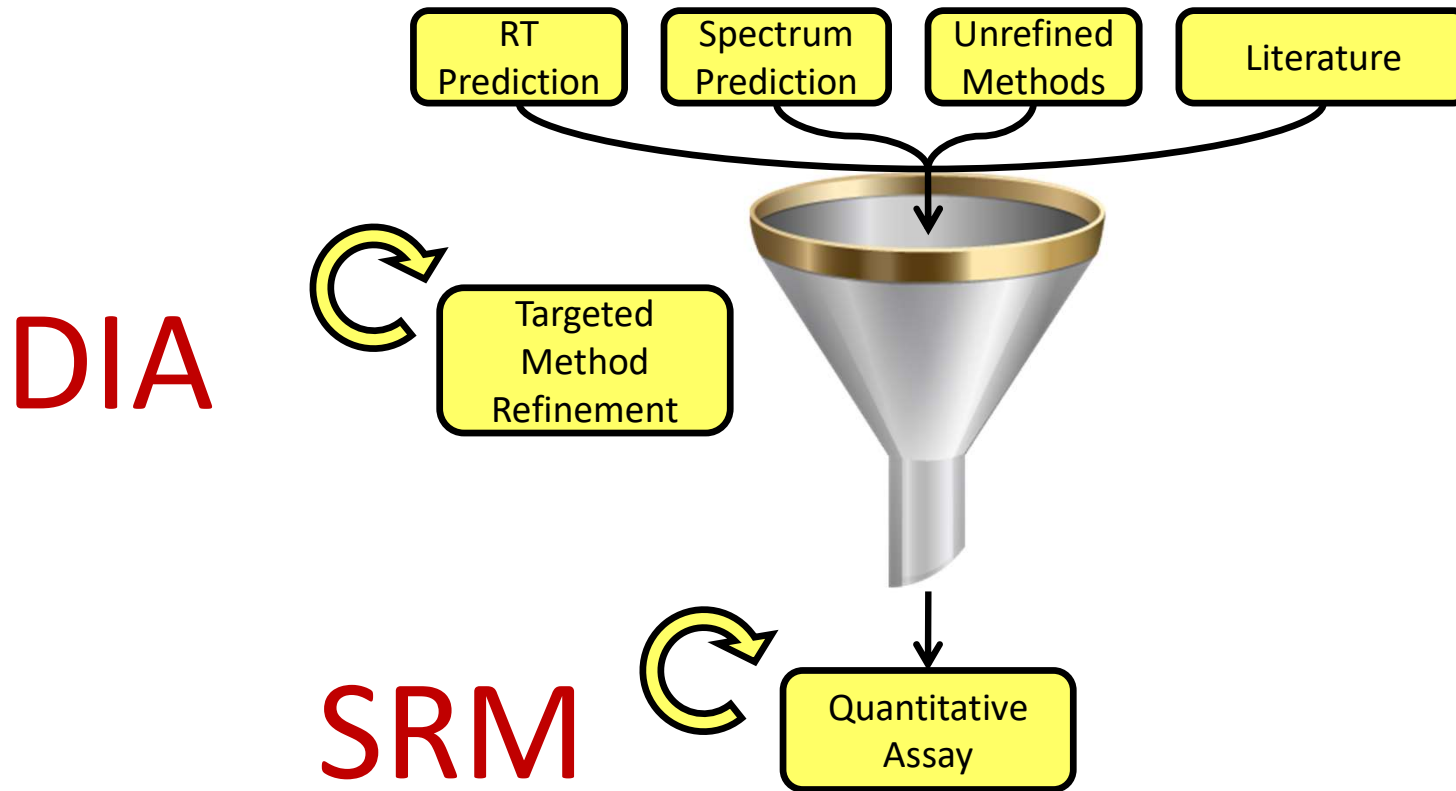
Brendan MacLean (Principal Developer, Skyline)
Deanna Plubell (Targeted proteomics researcher)
Mark Belanger (Project Manager, Skyline)



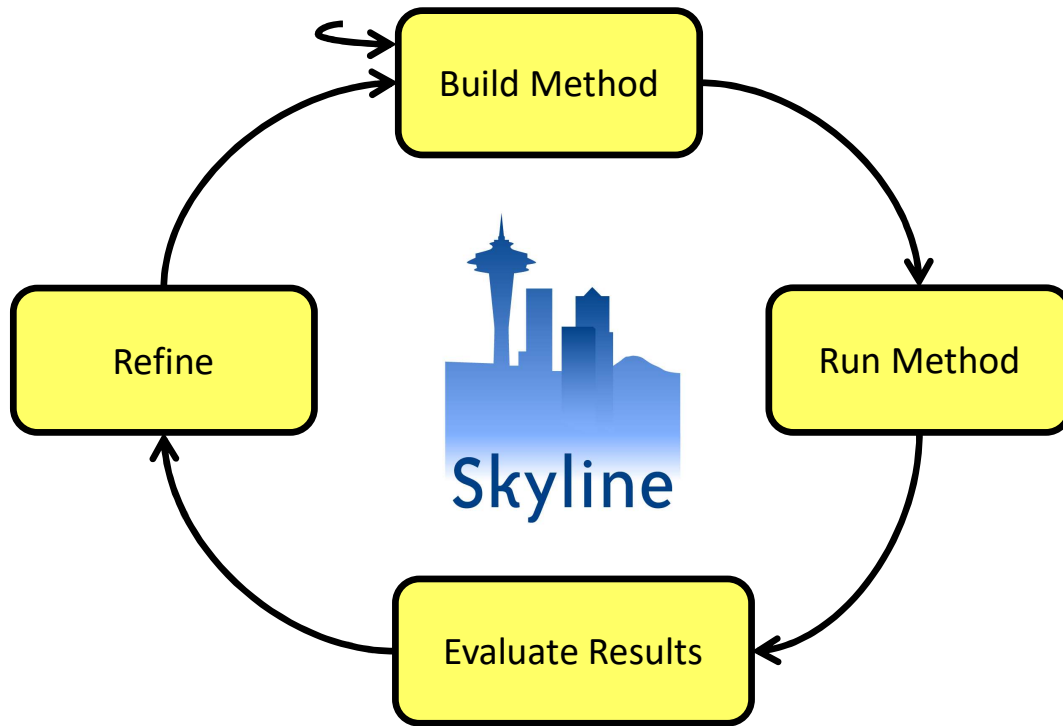
Agenda

- Welcome 395+ registered, from the Skyline team!
- Using DIA Data To Create SRM Methods
 - Introduction with **Brendan MacLean**
 - Overview and Tutorial with **Deanna Plubell**

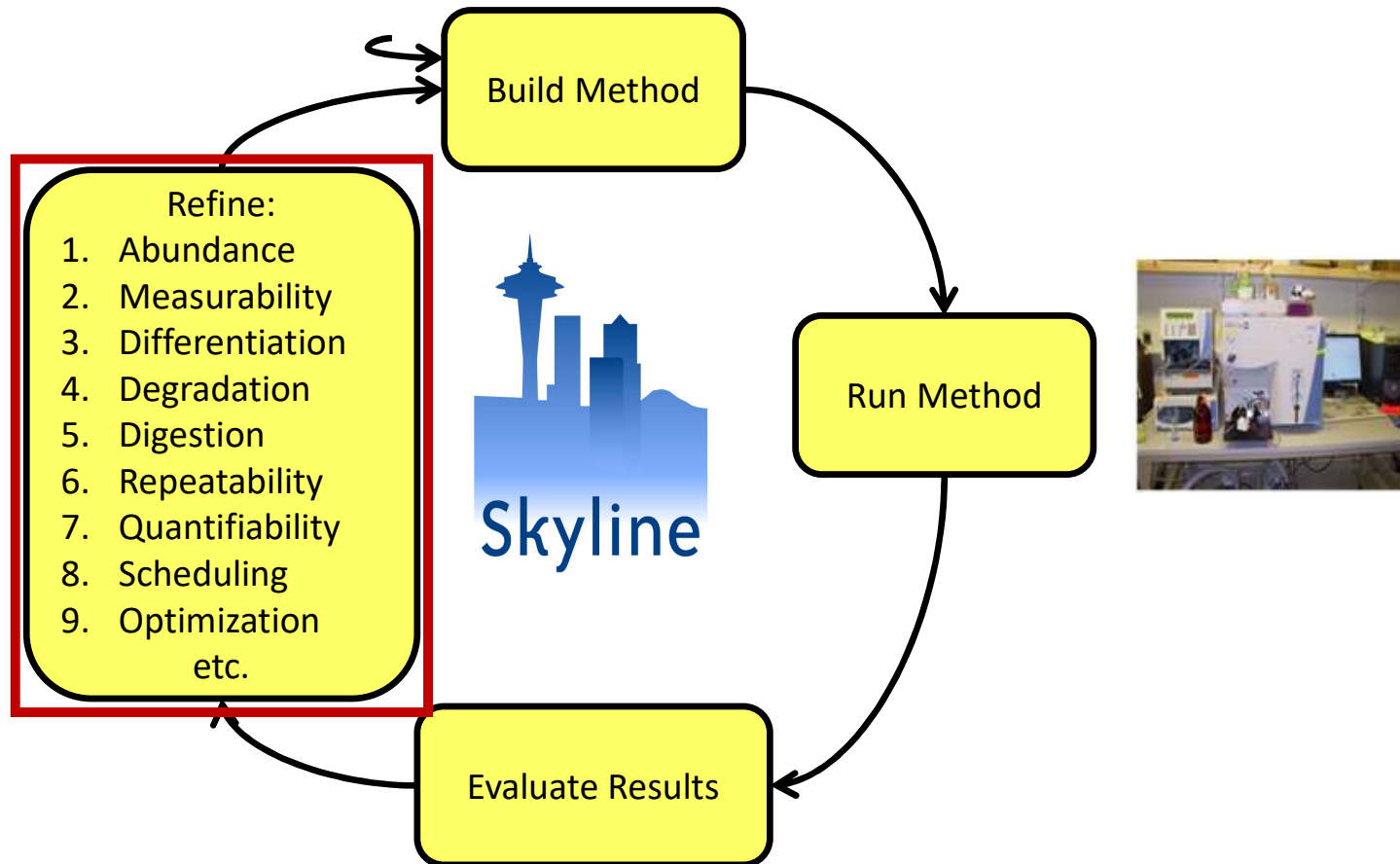
Supporting a Broader Hypothesis



Targeted Method Refinement

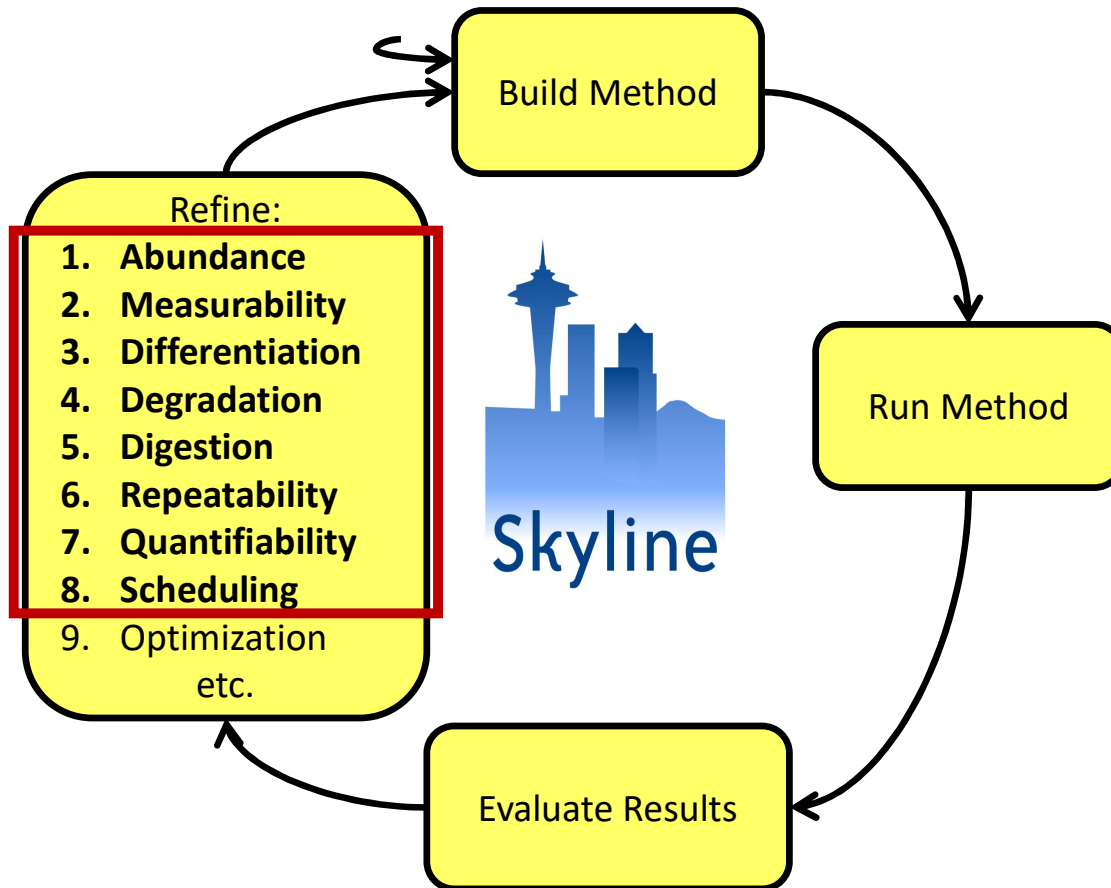


Targeted Method Refinement



Targeted Method Refinement

DIA

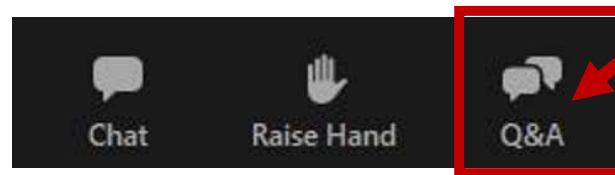


Deanna Plubell

- Overview
- Tutorial

Questions?

- Ask any questions at the Q&A form in Zoom:



- Take the post-webinar survey:

<https://skyline.ms/survey4webinar.url>



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This ends this Skyline Tutorial Webinar.

Please give us feedback on the webinar at the following survey:

<https://skyline.ms/survey4webinar.url>

A recording of today's meeting will be available shortly at the Skyline website.

We look forward to seeing you at a future Skyline Tutorial Webinar.