

Question	Answers (Time location in video - min:sec)
Could you please explain (again) what "peak area" is measuring?	Live Answered (Ashwood - 21:33)
Can you please post a link to the PDF instructions for developing small molecule analysis methods that Virag is referring to?	We have used this one for the between person comparison: https://skyline.ms/_webdav/home/software/Skyline/@files/tutorials/SmallMolecule-20_1.pdf But there are two more small molecule examples on the website. See: https://skyline.ms/tutorial.url
We love LipidCreator!! Thank you all so much for your hardwork developing this tool!	Live answered (Ahrends - 7:29)
Very cool! Have you looked at differences in results using LipidCreator for mining untargeted data for specific molecules vs. other lipidomics annotation tools?	Live answered (Ahrends - 7:40)
Very good and clear presentation! How would you estimate the "real" heavy leucine precursor pool enrichment in living organisms?	Live answered (Basisty - 20:20)
Is the tool applicable for a bolus or continuous infusion labeling typically used in people?	Live answered (Basisty - 23:17)
Do you have a pipeline to use heavy-leucine in the control group?	Live answered (Basisty - 25:25)
Thank you, what human peptides do you use for normalization, as human proteome can also be affected by the viral infection? Did you observe higher reproducibility of your data after normalization compared to unnormalized data?	Live answered (Kennedy - 19:26)
<p>Question: DNA viruses vs RNA viruses</p> <p>1)Background</p> <p>a)Herpes Simplex -- is a double-stranded DNA virus</p> <p>b)The Coronavirus -- is a single-stranded RNA virus</p> <p>Single-stranded RNA viruses mutate much, much faster than DNA viruses</p> <p>c)Are your techniques applicable to single stranded viruses like the Corona virus ?</p> <p>d)Are there notable differences in working with single-stranded RNA viruses vs double-stranded viruses</p>	Live answered (Kennedy - 22:00)
what was your LC/MS system?	
Would you agree that sensitivity by qPCR is still higher compared to PRM for single protein targets?	Live answered (Kennedy - 23:45)