Oncohost Tube Bridging Experiment

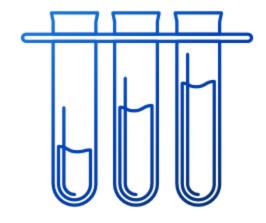
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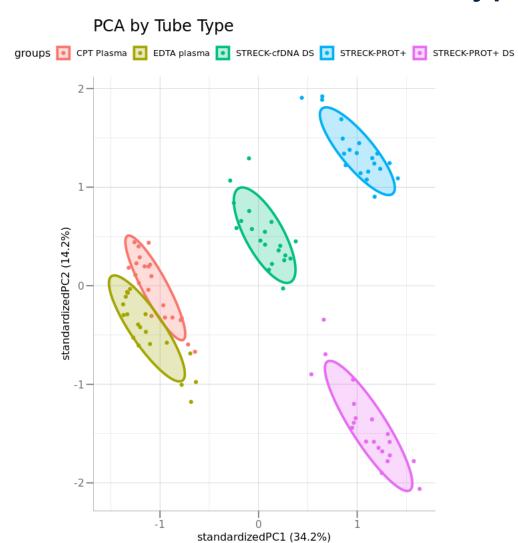


Overview of Experiment

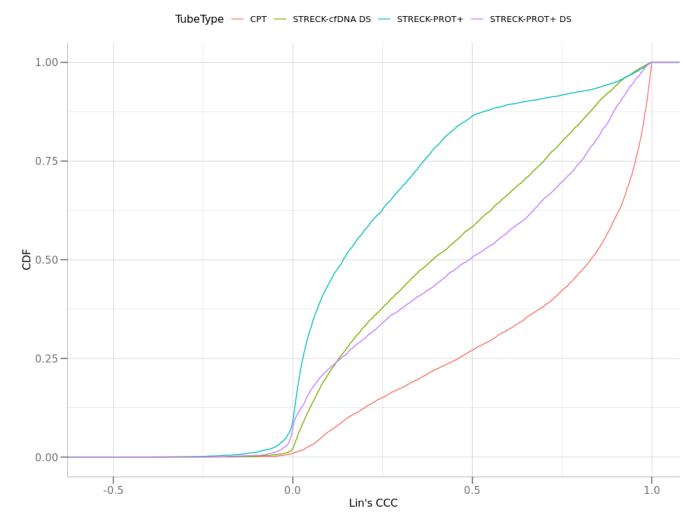
- Different plasma tubes have shown differing proteomic signatures
- The standard for the SomaScan assay is EDTA tubes
- Bridging methods give the possibility of making alternate tube types comparable to EDTA plasma results
- Oncohost ran several different tube types with the goal of bridging the data to the SomaScan EDTA standard



Variance Between Tube Types



Distribution of Concordance to EDTA Plasma



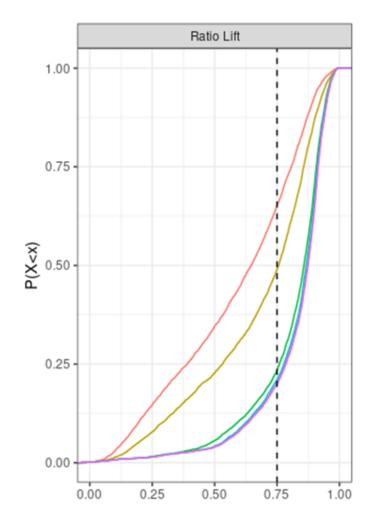
The Bridging Method

- The SomaScan team have previously tested several bridging methods for bridging cross-assay version
- For our purposes, we found a ratio lift method to perform best
- Take a ratio of the reference type divided by the type to be bridged

$$R_i = \frac{\widehat{\mu}_{7k}}{\widehat{\mu}_{11k}},$$

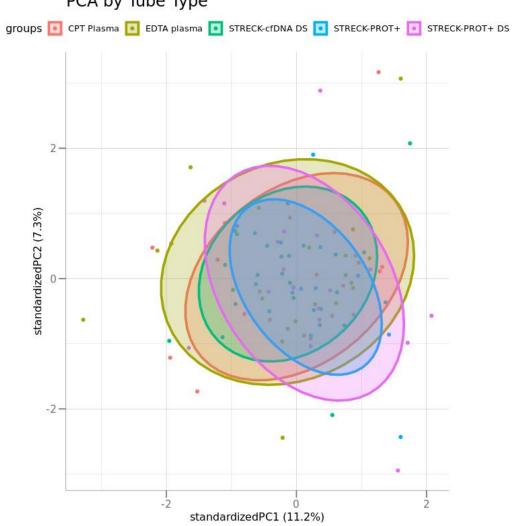
- Resulting scalar can be multiplied by the values to be bridged to lift the data
- Assessed by Lin's CCC

$$CCC = \frac{2\rho\widehat{\sigma}_x\widehat{\sigma}_y}{\left(\widehat{\mu}_x - \widehat{\mu}_y\right)^2 + \widehat{\sigma}_x^2 + \widehat{\sigma}_y^2},$$

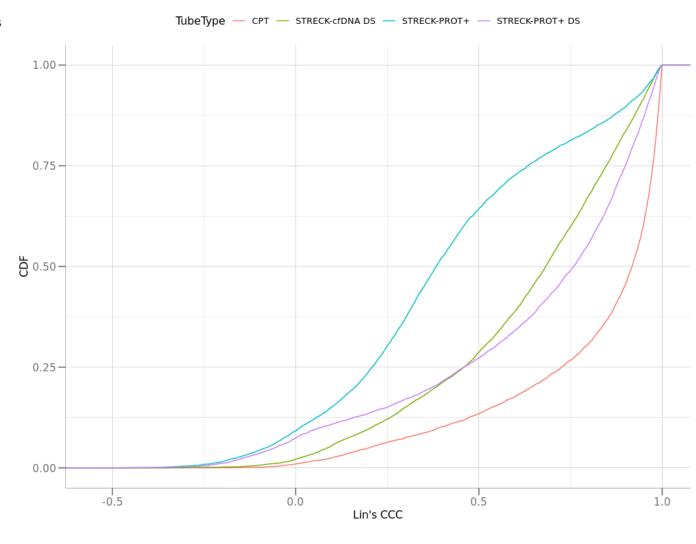


Post-Bridging

PCA by Tube Type



Distribution of Concordance to EDTA Plasma



Preliminary Results

Median CCC	СРТ		STRECK Protein+	STRECK Protein+ DS
Pre-Bridge	0.82	0.39	0.14	0.49
Post-Bridge	0.91	0.68	0.38	0.76

- CPT tubes bridge very well
- In an Estimated Marginal Means (EMM) model, only two analytes are significant for CPT post-bridge
- STRECK Protein+ bridges the worst by our methods
- In an EMM for STRECK Protein+, 80 analytes are significant post-bridge

Next Steps

- We can incorporate the new data to see what results are
- We can also share the scalars, CCC's, and statistical results
- Could also use CV (with a method like Bland-Altman) as a different way of representing reproducibility





Questions?