

Harmonization of Individual Samples

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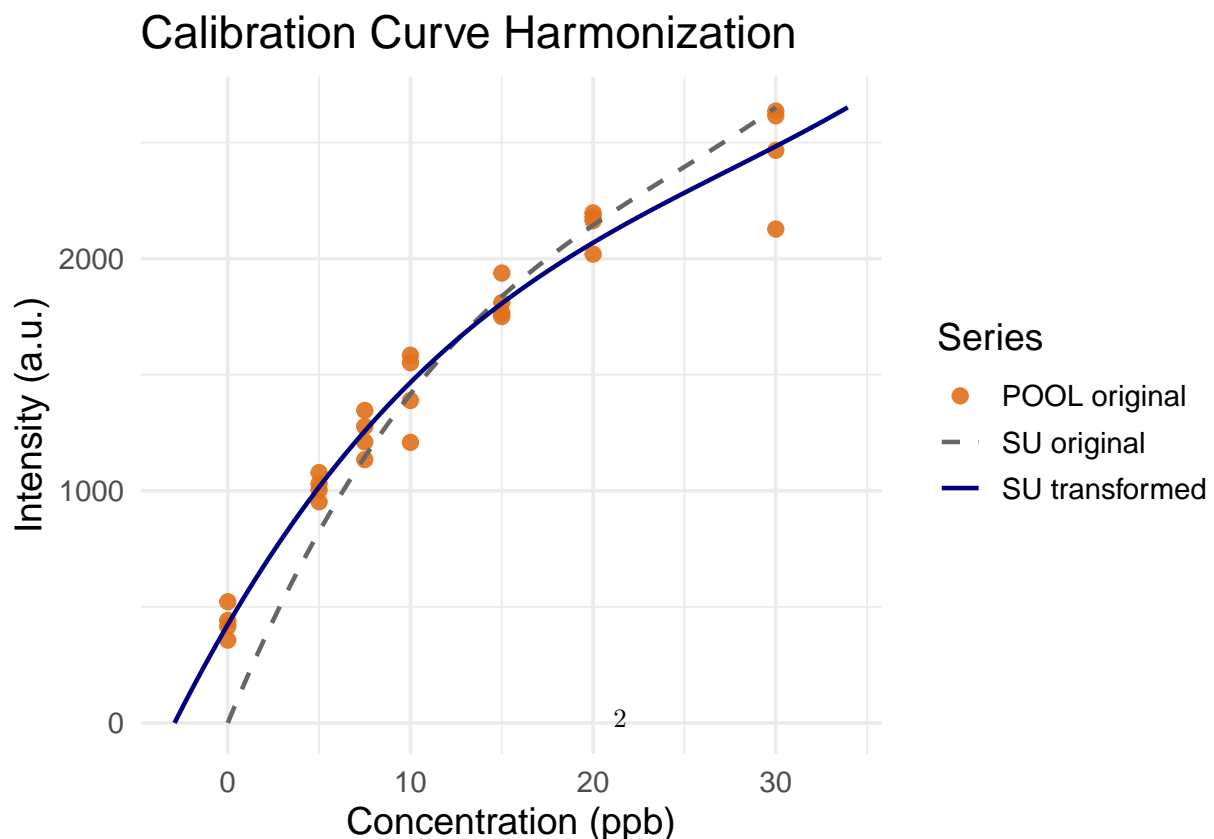
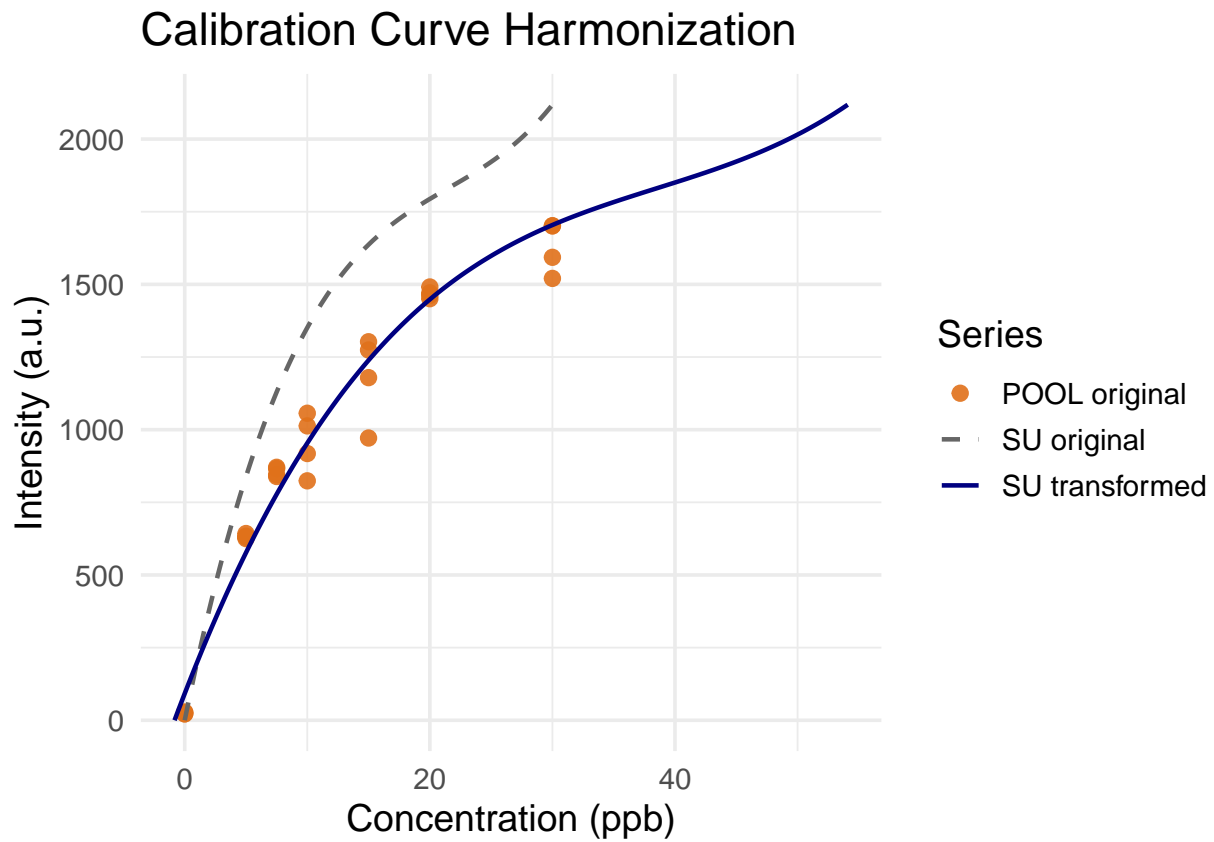
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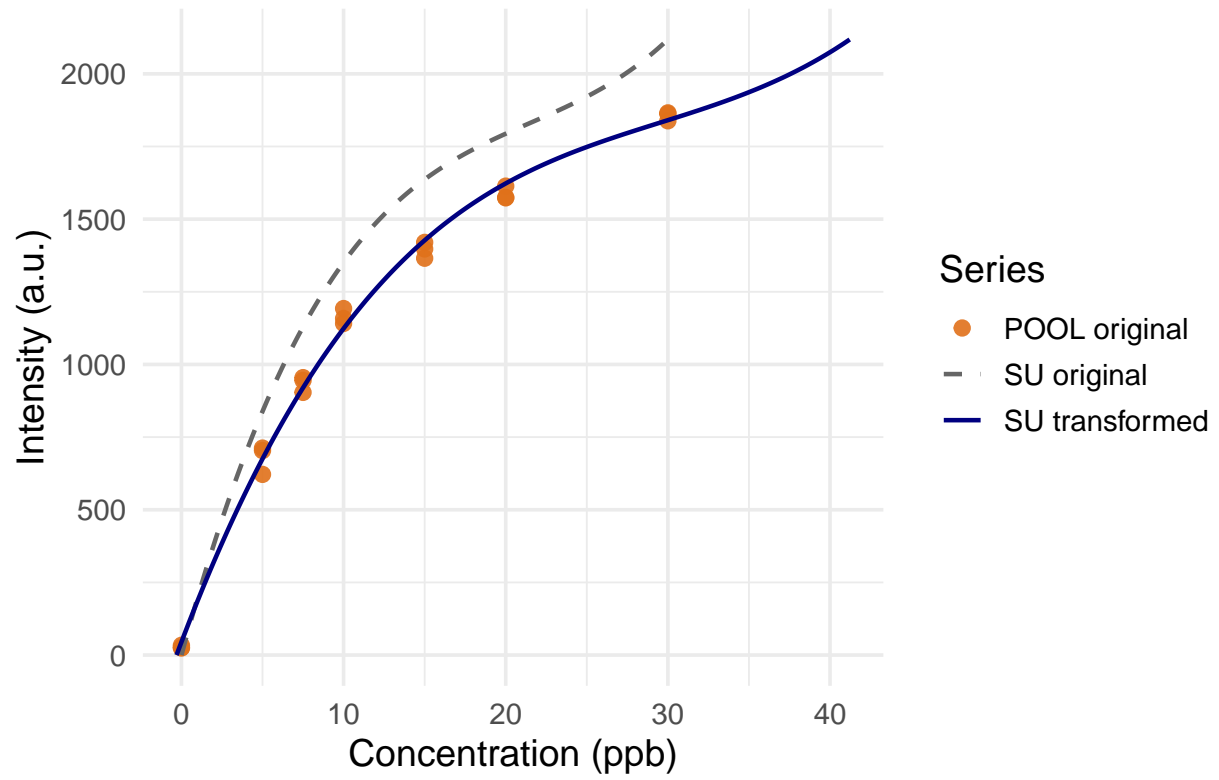
Load Data

2. Harmonization of all samples

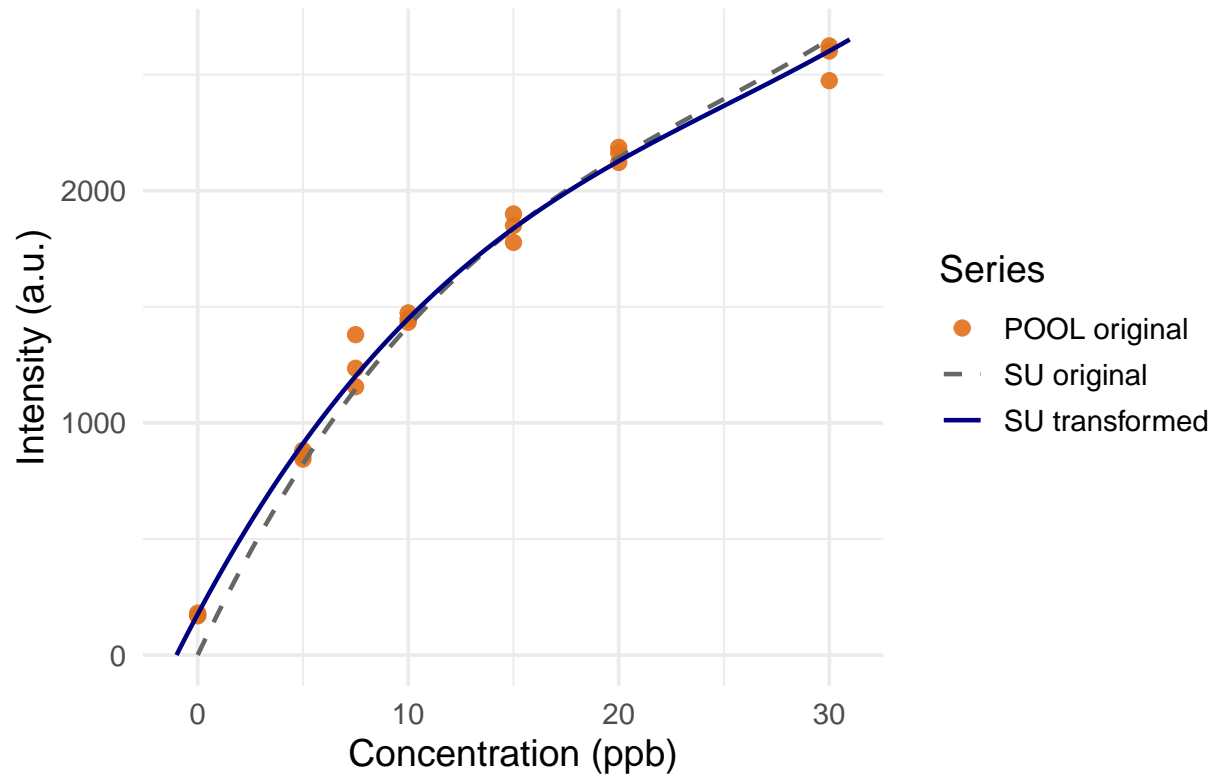
3. Plot for each sample and analyte

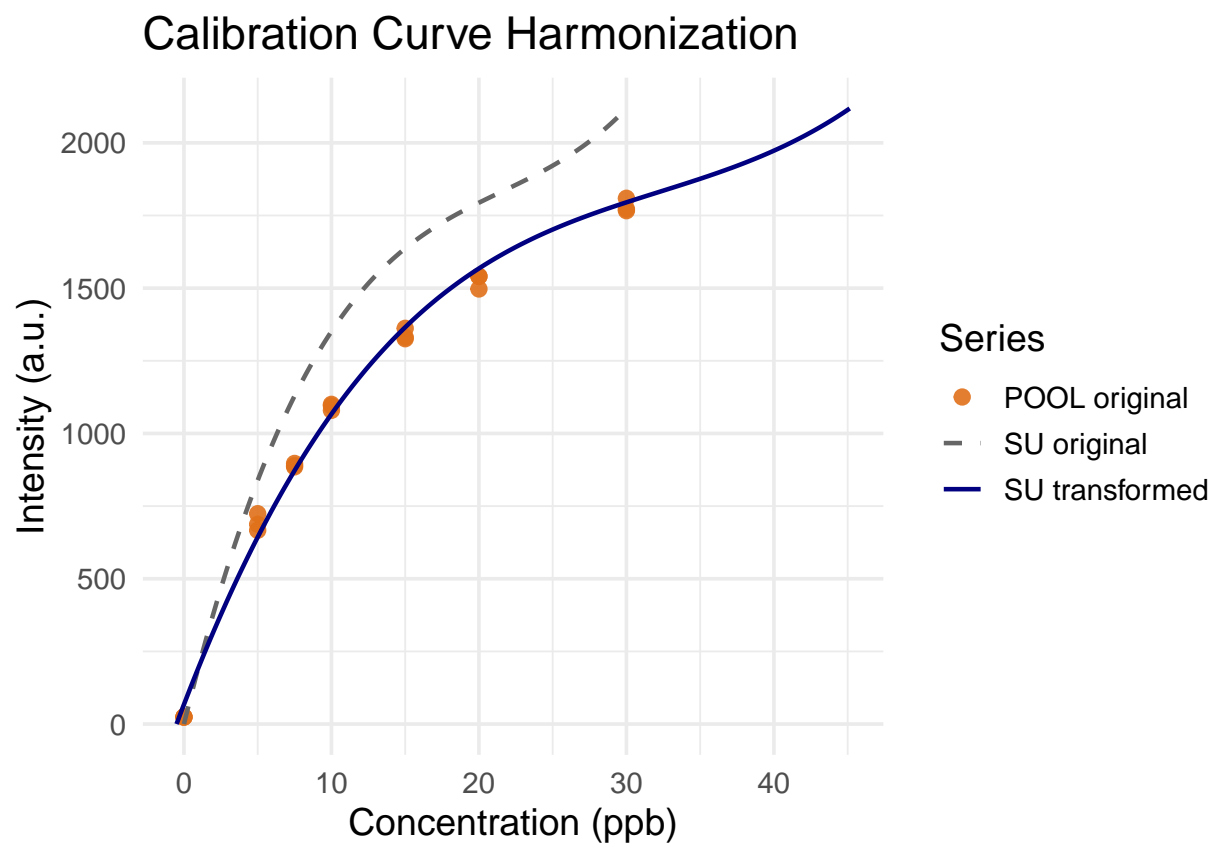


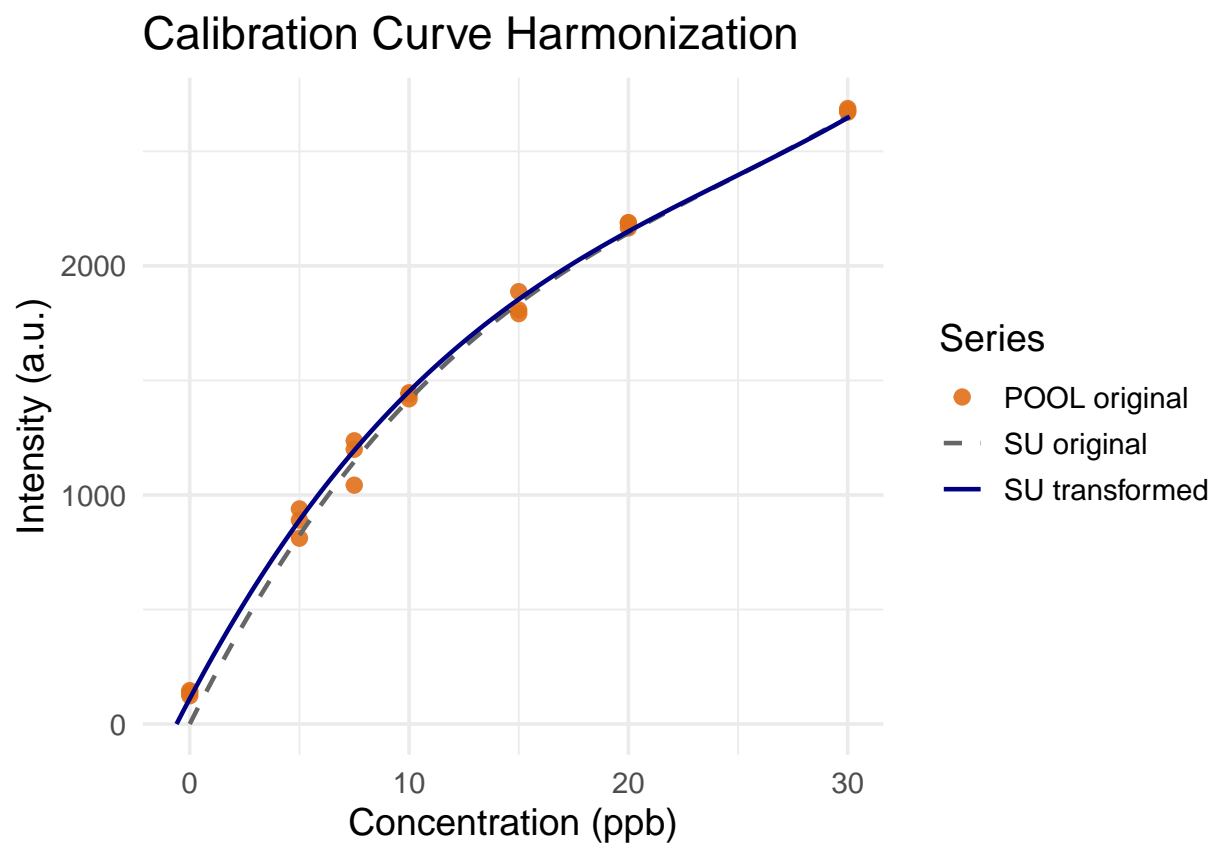
Calibration Curve Harmonization



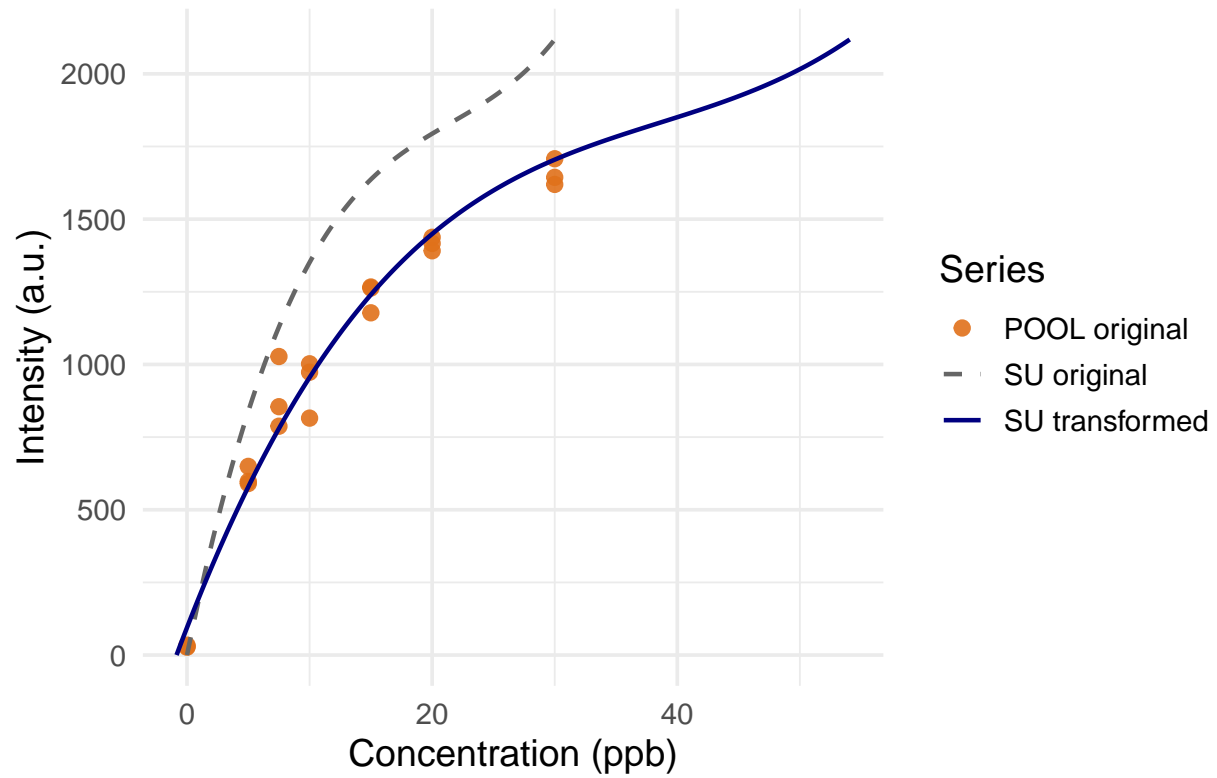
Calibration Curve Harmonization



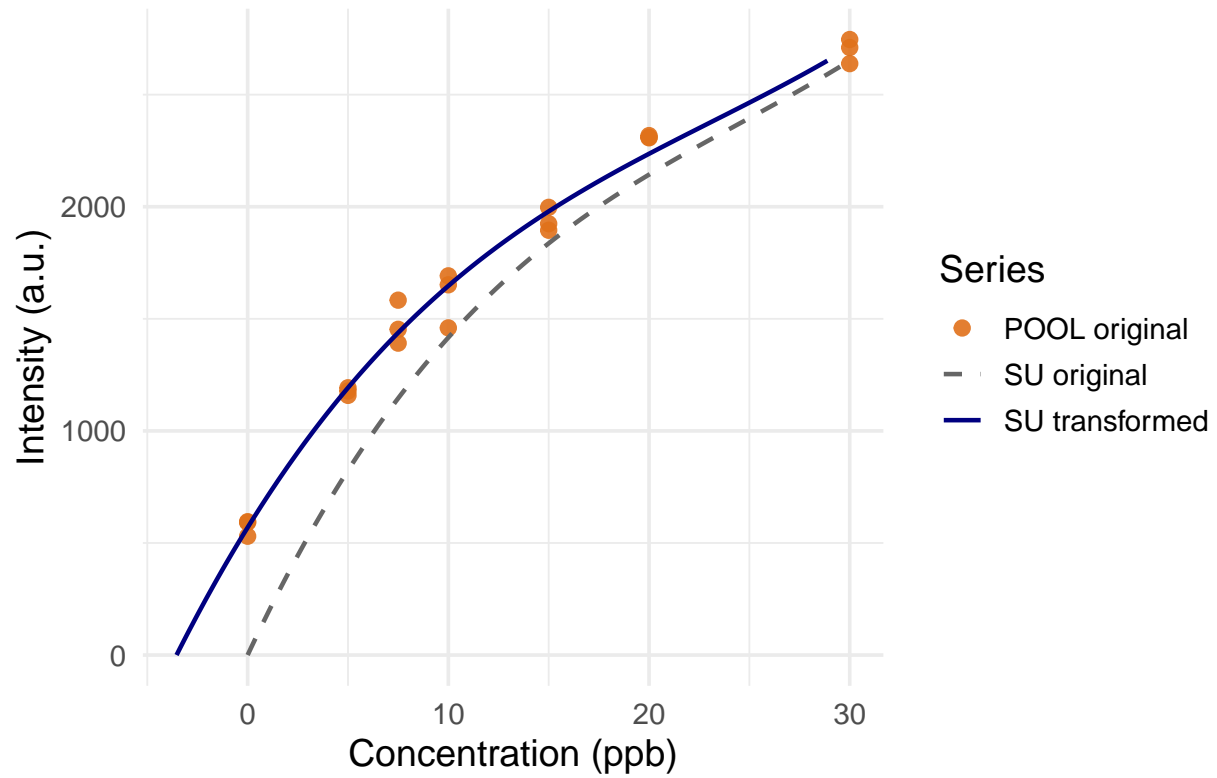


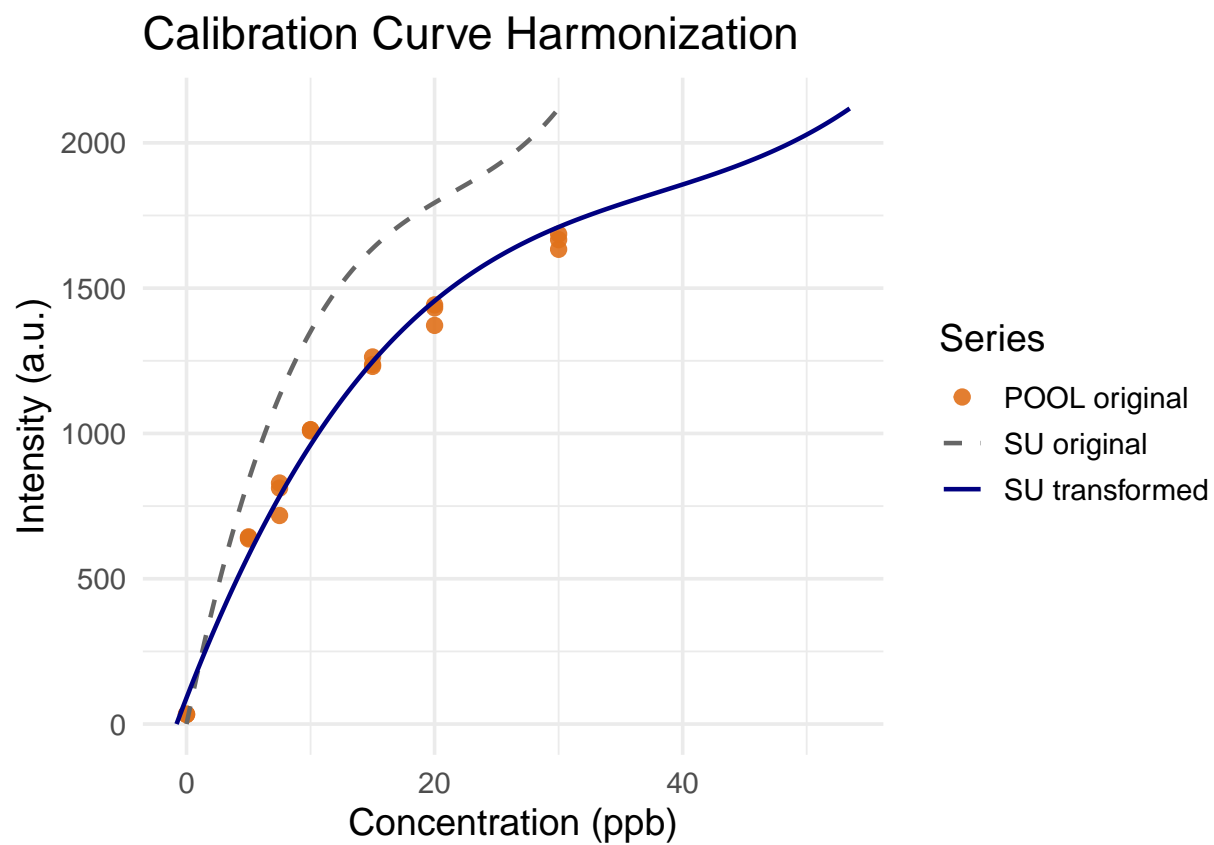


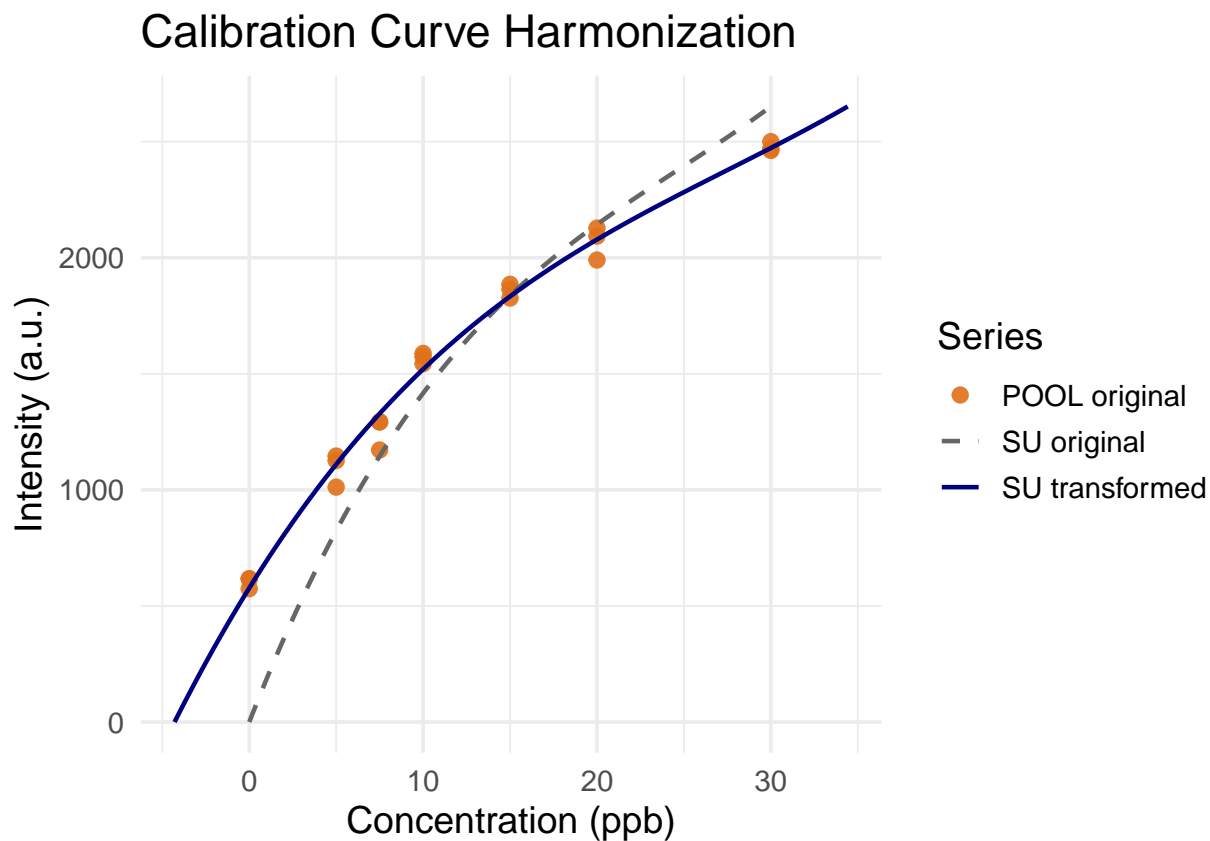
Calibration Curve Harmonization



Calibration Curve Harmonization







4. Summary table of all scale/shift/errors

Table 1: Summary of harmonization parameters

Sample	Analyte	Scale	Shift	Scale_Error	Shift_Error
pool	anisole	0.5462	0.8291	0.1185	1.9442
pool	heptanone	0.8141	2.9146	0.1557	2.2427
s1	anisole	0.7226	0.3015	0.0561	0.5790
s1	heptanone	0.9382	1.0050	0.0769	0.8080
s2	anisole	0.6573	0.5025	0.0564	0.6807
s2	heptanone	0.9774	0.6030	0.0651	0.6354
s3	anisole	0.5462	0.8543	0.1043	1.7238
s3	heptanone	0.9251	3.5427	0.0949	1.2776
s4	anisole	0.5528	0.8040	0.0672	1.0894
s4	heptanone	0.7749	4.2965	0.0778	1.3011

5. Final plot: scaling factors across samples

Scaling Factors for All Samples

Dashed line = Pool scale

