Supporting Information

Quantitative accuracy and precision in multiplexed single-cell proteomics.

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Figure S1: Re-analysis of public single cell data, (a) Identified proteins, peptide groups, PSMs, number of MS/MS scans, ID-rates and (b) delta between expected and acquired carrier to single cell ratio across all MS/MS scans or PSMs of published SCoPE-MS (light brown) and SCoPE2 (dark brown). Median and median absolute deviation (mad) is shown.



Figure S2: Ratio compression for SCoPE, TMT10-plex and TMTpro samples at various carrier spikes. Log₁₀ median RI intensity of all MS/MS scans and binned ratios between 'single cells' and the carrier is displayed. Identified and unidentified MS/MS scans are indicated in grey or pink, respectively. The expected bin (carrier +/- 50 %) is highlighted in black, and the percent of MS/MS scans within those is indicated.



Figure S3: Measurement stability and RI variability at various carrier ratios. RI intensity distributions based on all MS/MS scans for **(a-b)** SCoPE, **(c-d)** TMT10-plex and **(e-f)** TMTpro experiments at indicated carrier spikes.



Figure S4: Measurement stability and variance of public SCoPE-MS and SCoPE2 data. (a-b) RI intensity distributions based on all MS/MS scans and (c-d) percent CV across 'single cell' channels and log₁₀ mean RI S/N for publicly available SCoPE data. The horizontal solid line and dashed lines indicate median S/N across all MS/MS scans or post-S/N filtering, respectively. The vertical blue line specifies the S/N filter cut-off. Colors reflect the number of missing 'single cell' RIs per MS/MS scan. (e) Percent missing quantitative data in publicly available SCoPE data per PSM.



Figure S5: Measurement accuracy of TMT_{zero} *experiments.* (a-c) RI intensity distribution across all MS/MS scans in TMT_{zero} experiments. Colors indicate different TMT channels. (d-f) Median percent CV and log_{10} mean RI S/N at indicated TMT_{zero} carrier spikes. Horizontal solid and dashed lines indicate median S/N across all MS/MS scans or post-S/N filtering, respectively. The vertical blue line indicates the S/N filter cut-off. Colors indicate the number of missing 'single cell' RIs per MS/MS scan.



Figure S6: Ratio compression in DIA-TMT experiments. Log₁₀ median RI intensity across all MS/MS scans is displayed with binned 'single cell' to carrier channels ratios. The expected ratio (carrier +/- 50%) is highlighted in black, and percent of MS/MS scans within this bin is indicated.