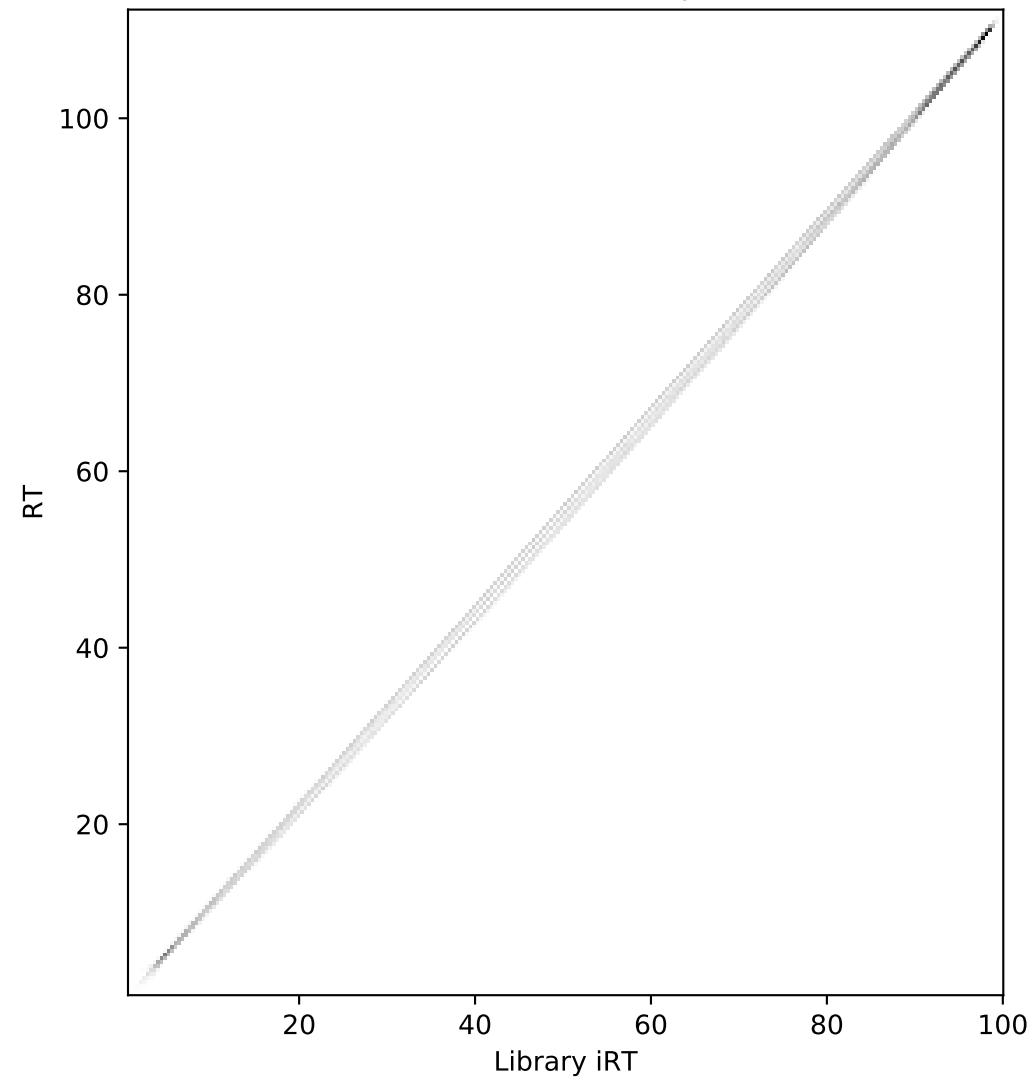
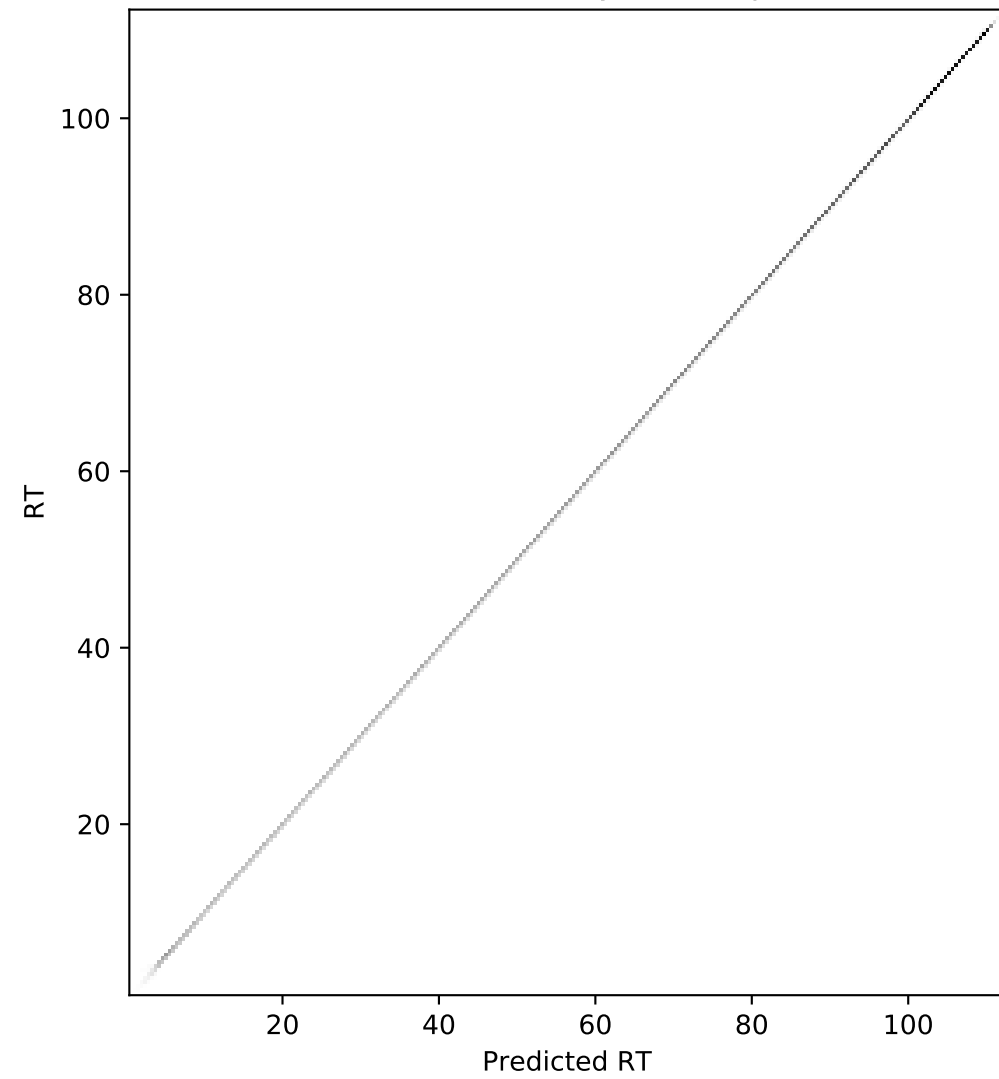


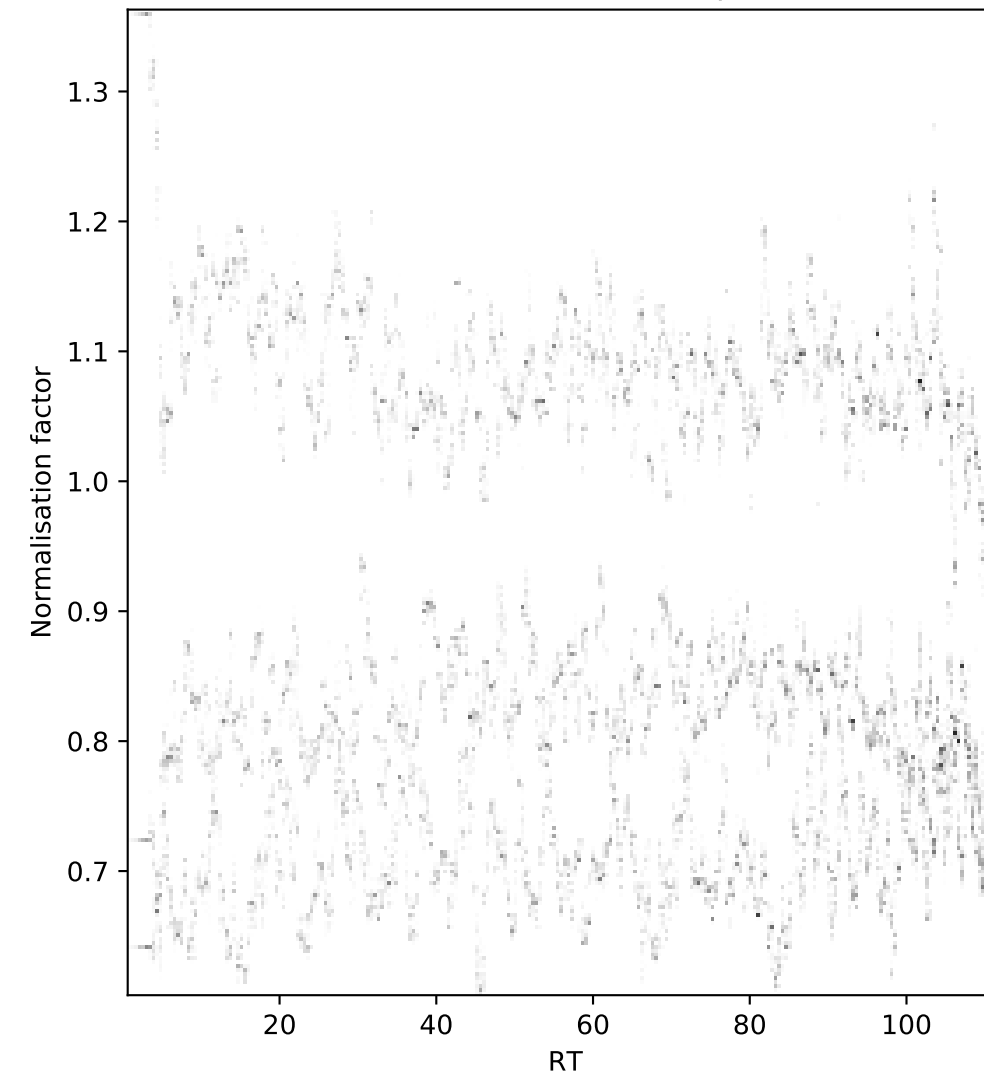
Retention times heatmap, all runs

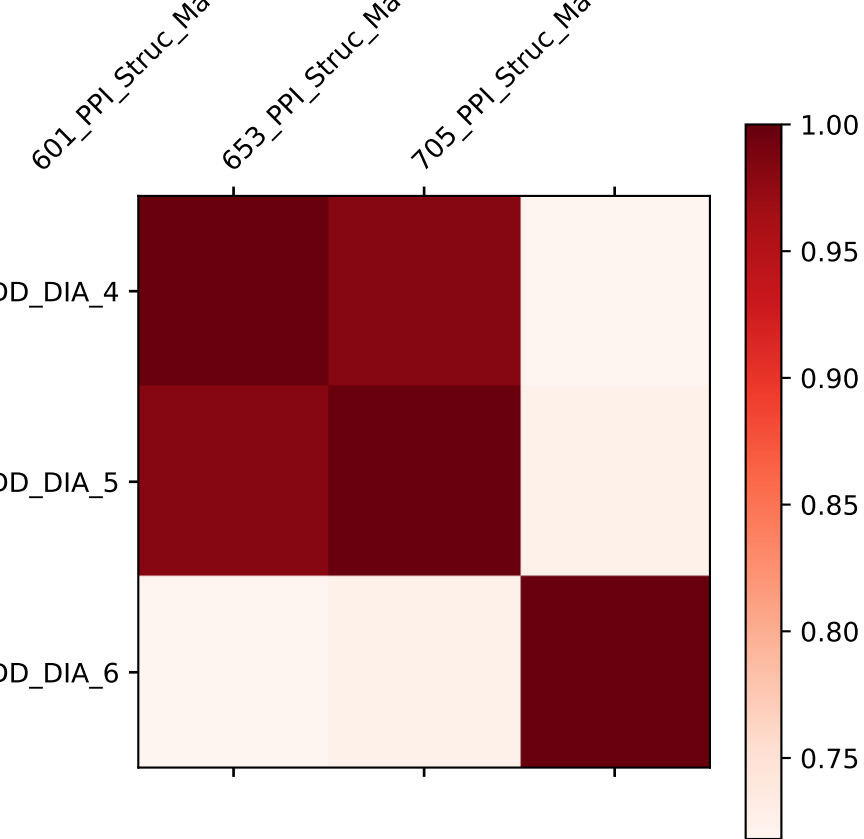


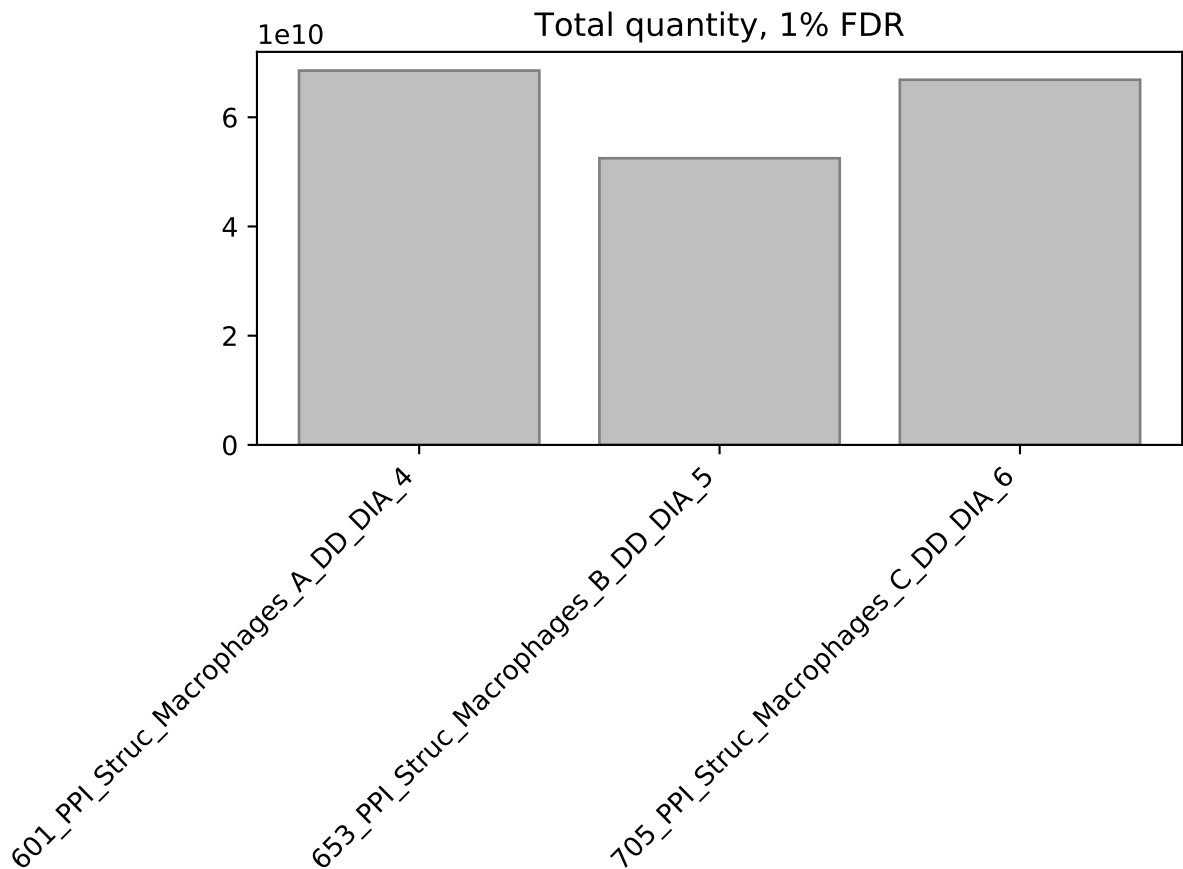
Retention time accuracy heatmap, all runs

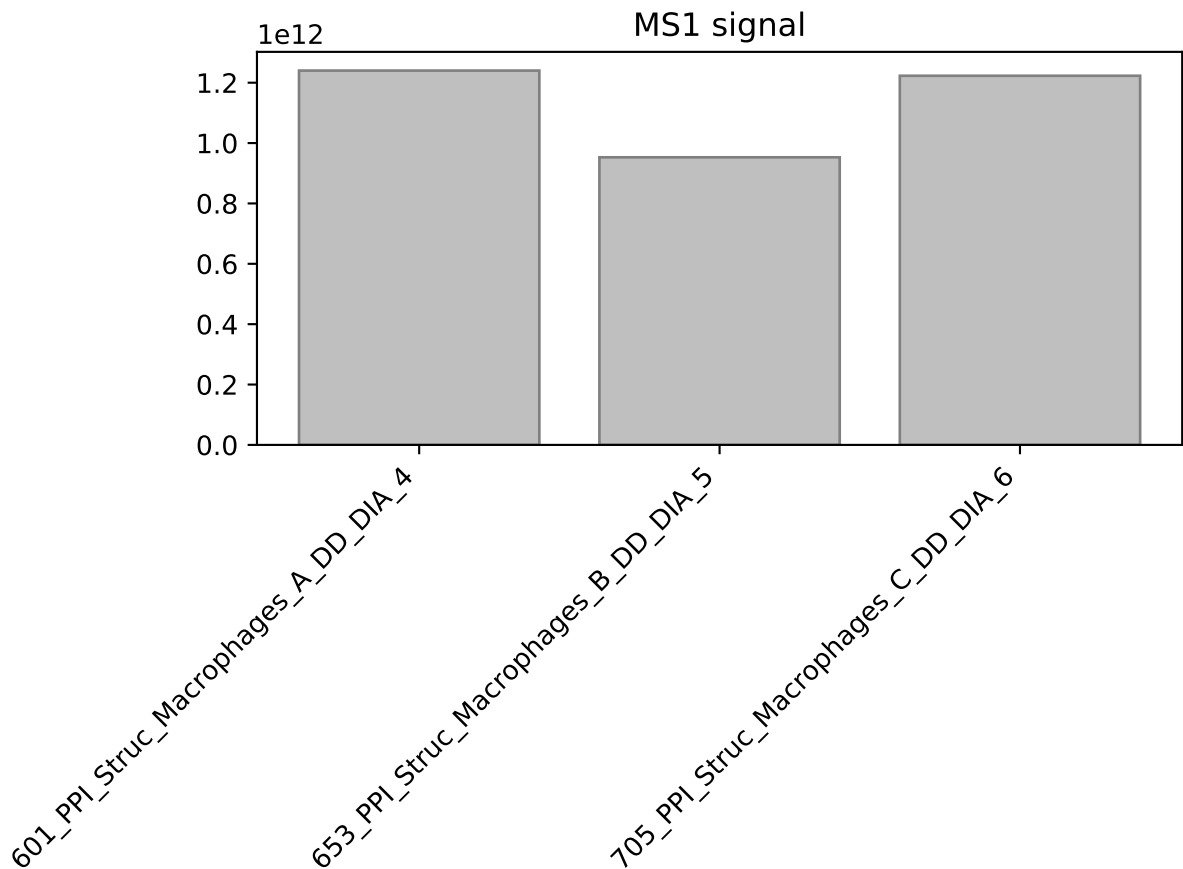


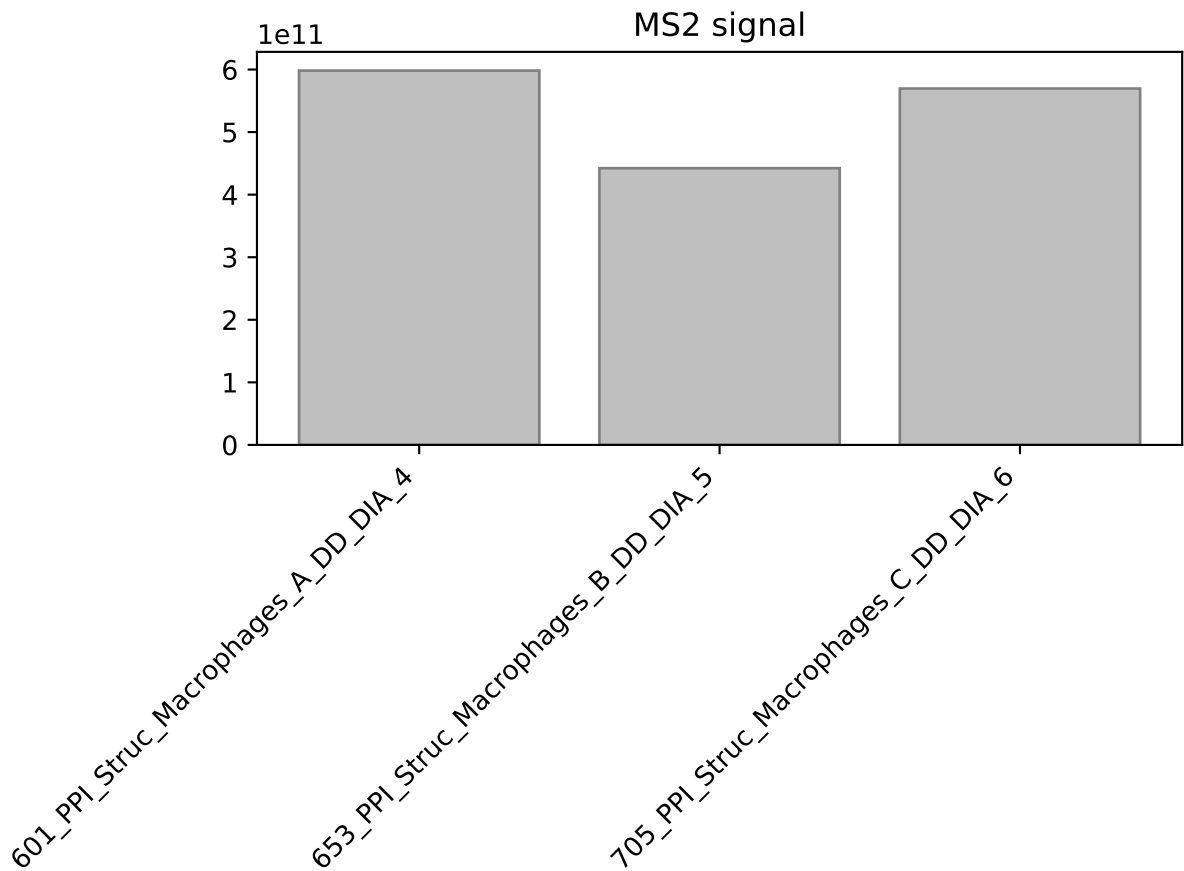
Normalisation factor heatmap, all runs



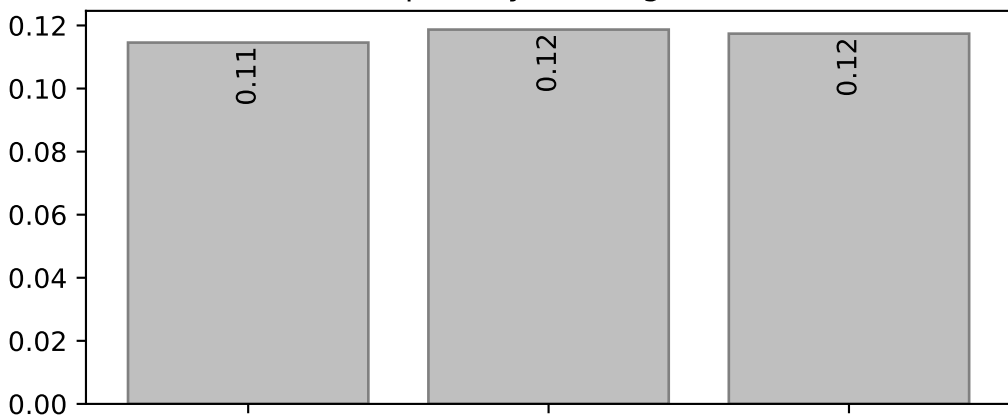








Total quantity/MS2 signal ratio

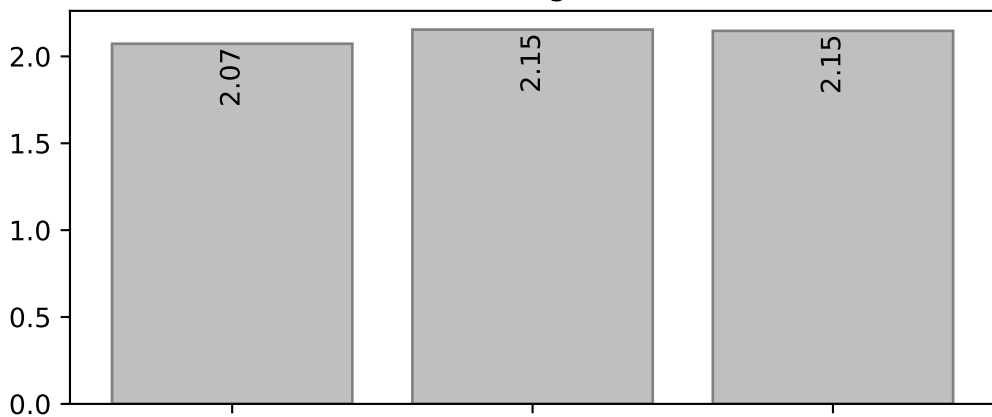


601\_PPI\_Struc\_Macrophages\_A\_DD\_DIA\_4

653\_PPI\_Struc\_Macrophages\_B\_DD\_DIA\_5

705\_PPI\_Struc\_Macrophages\_C\_DD\_DIA\_6

MS1/MS2 signal ratio

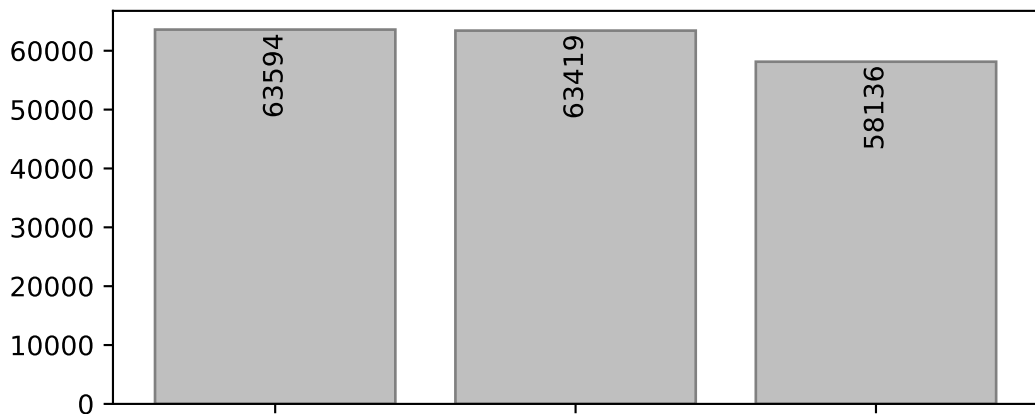


601\_PPI\_Struc\_Macrophages\_A\_DD\_DIA\_4

653\_PPI\_Struc\_Macrophages\_B\_DD\_DIA\_5

705\_PPI\_Struc\_Macrophages\_C\_DD\_DIA\_6

Precursors, 1% FDR

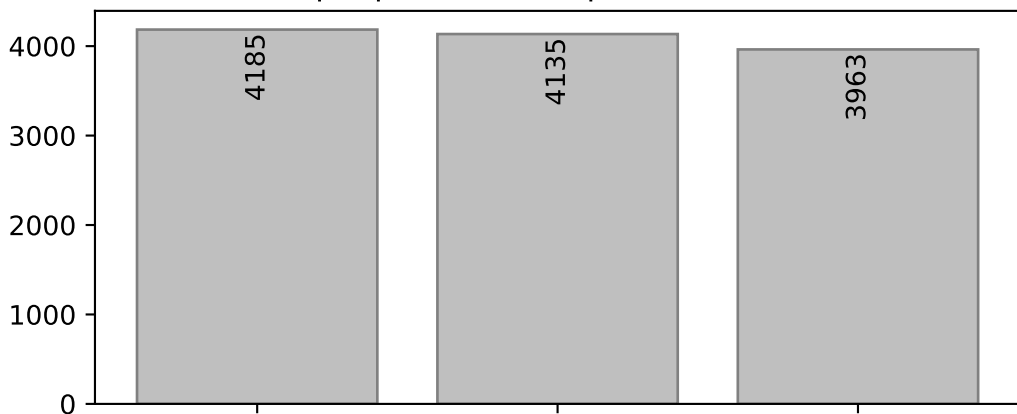


601\_PPI\_Struc\_Macrophages\_A\_DD\_DIA\_4

653\_PPI\_Struc\_Macrophages\_B\_DD\_DIA\_5

705\_PPI\_Struc\_Macrophages\_C\_DD\_DIA\_6

Unique proteins, 1% protein-level FDR

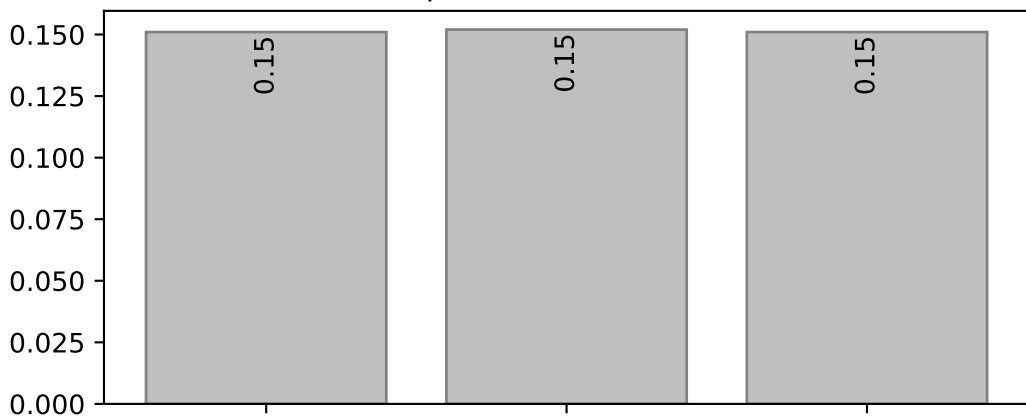


601\_PPI\_Struc\_Macrophages\_A\_DD\_DIA\_4

653\_PPI\_Struc\_Macrophages\_B\_DD\_DIA\_5

705\_PPI\_Struc\_Macrophages\_C\_DD\_DIA\_6

Mean peak FWHM, in minutes

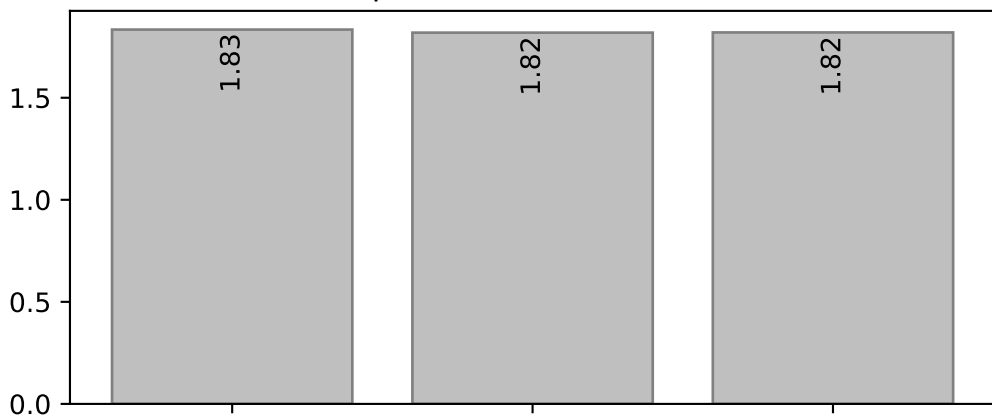


601\_PPI\_Struc\_Macrophages\_A\_DD\_DIA\_4

653\_PPI\_Struc\_Macrophages\_B\_DD\_DIA\_5

705\_PPI\_Struc\_Macrophages\_C\_DD\_DIA\_6

Mean peak FWHM, in MS2 scans

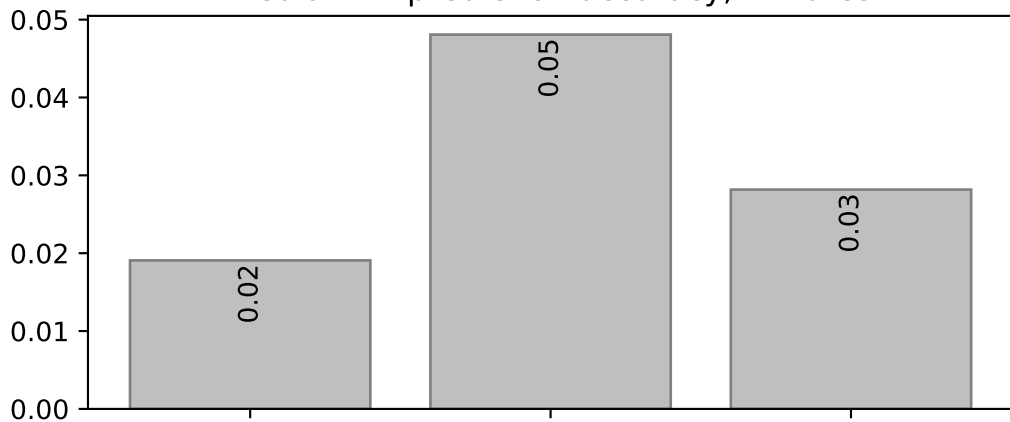


601\_PPI\_Struc\_Macrophages\_A\_DD\_DIA\_4

653\_PPI\_Struc\_Macrophages\_B\_DD\_DIA\_5

705\_PPI\_Struc\_Macrophages\_C\_DD\_DIA\_6

Median RT prediction accuracy, minutes

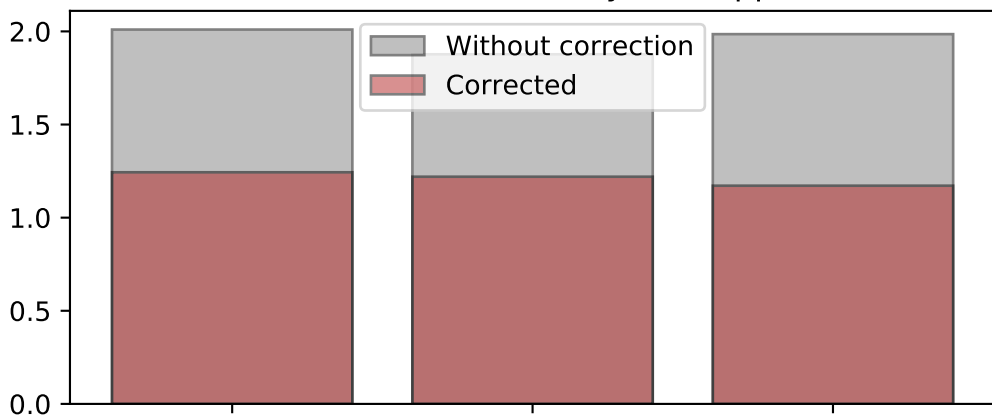


601\_PPI\_Struc\_Macrophages\_A\_DD\_DIA\_4

653\_PPI\_Struc\_Macrophages\_B\_DD\_DIA\_5

705\_PPI\_Struc\_Macrophages\_C\_DD\_DIA\_6

Median mass accuracy, MS2, ppm

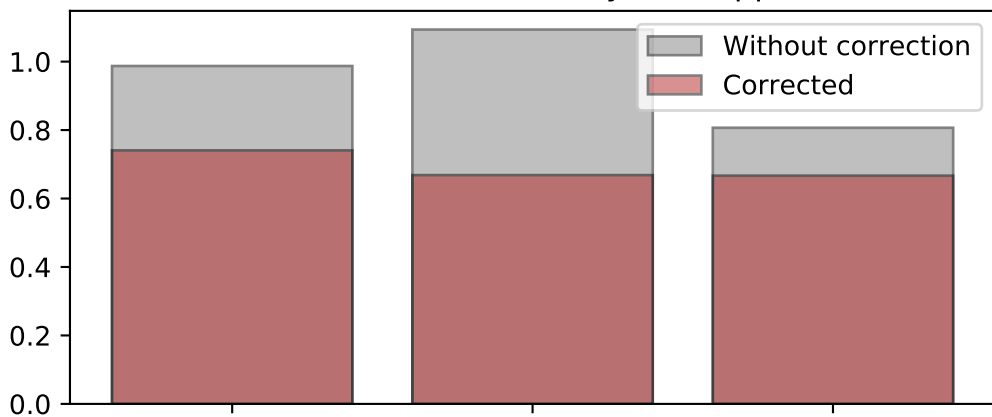


601\_PPI\_Struc\_Macrophages\_A\_DD\_DIA\_4

653\_PPI\_Struc\_Macrophages\_B\_DD\_DIA\_5

705\_PPI\_Struc\_Macrophages\_C\_DD\_DIA\_6

Median mass accuracy, MS1, ppm

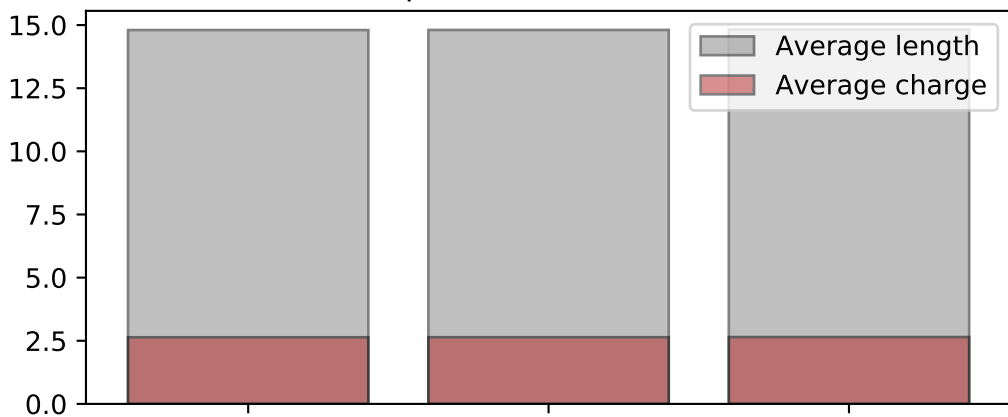


601\_PPI\_Struc\_Macrophages\_A\_DD\_DIA\_4

653\_PPI\_Struc\_Macrophages\_B\_DD\_DIA\_5

705\_PPI\_Struc\_Macrophages\_C\_DD\_DIA\_6

Peptide characteristics

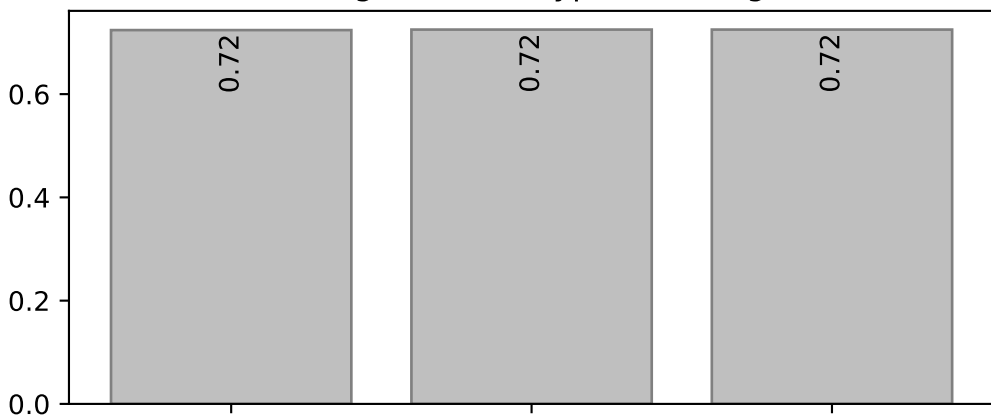


601\_PPI\_Struc\_Macrophages\_A\_DD\_DIA\_4

653\_PPI\_Struc\_Macrophages\_B\_DD\_DIA\_5

705\_PPI\_Struc\_Macrophages\_C\_DD\_DIA\_6

Average missed tryptic cleavages



601\_PPI\_Struc\_Macrophages\_A\_DD\_DIA\_4

653\_PPI\_Struc\_Macrophages\_B\_DD\_DIA\_5

705\_PPI\_Struc\_Macrophages\_C\_DD\_DIA\_6