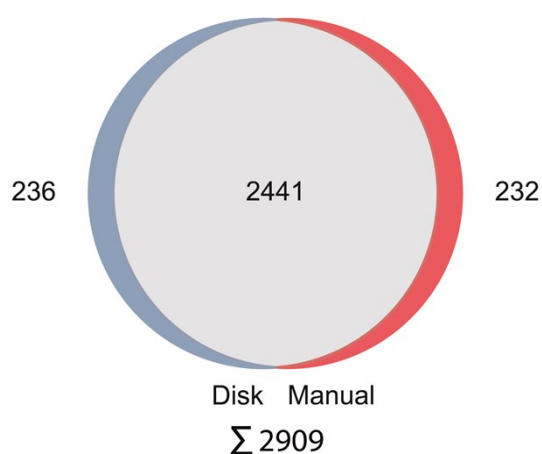


Supplementary Information

1 Protein level results of the explorative LC-MS/MS experiment

In total, 2909 proteins were identified in the two sets, with 84% common identifications. As can be seen in Supplementary Figure 1, identifications in the individual sets are comparable.



Supplementary Figure 1 Venn diagram illustrating the identifications on protein level for both desalting approaches.