

Supplementary Figure 3

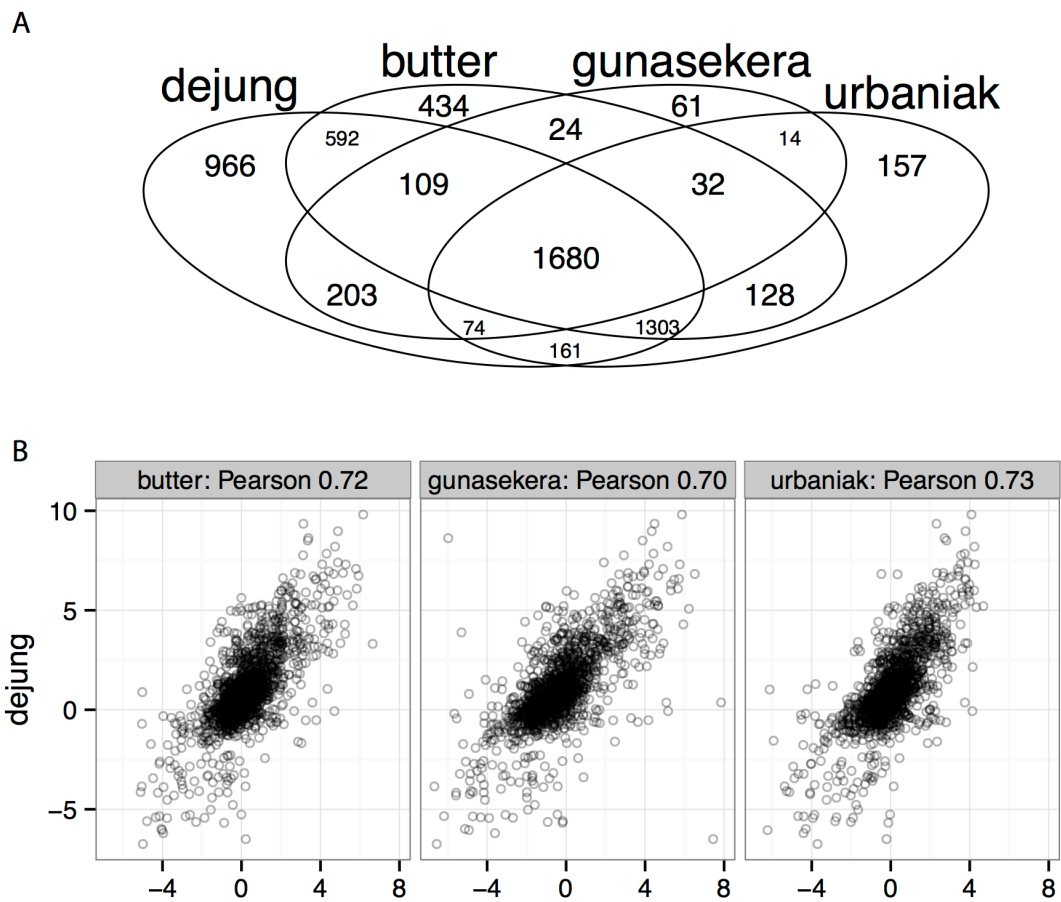


Fig S3: Comparison with available SILAC proteomes of PF and LS states. **(A)** Venn diagram with the number of individual protein IDs comparing to previously published datasets. **(B)** Scatterplot of LS/PF enrichment of 1680 protein groups overlapping with all previous studies. Each plot compares to the corresponding study. For each case, we obtain a pearson coefficient of $r \sim 0.7$.