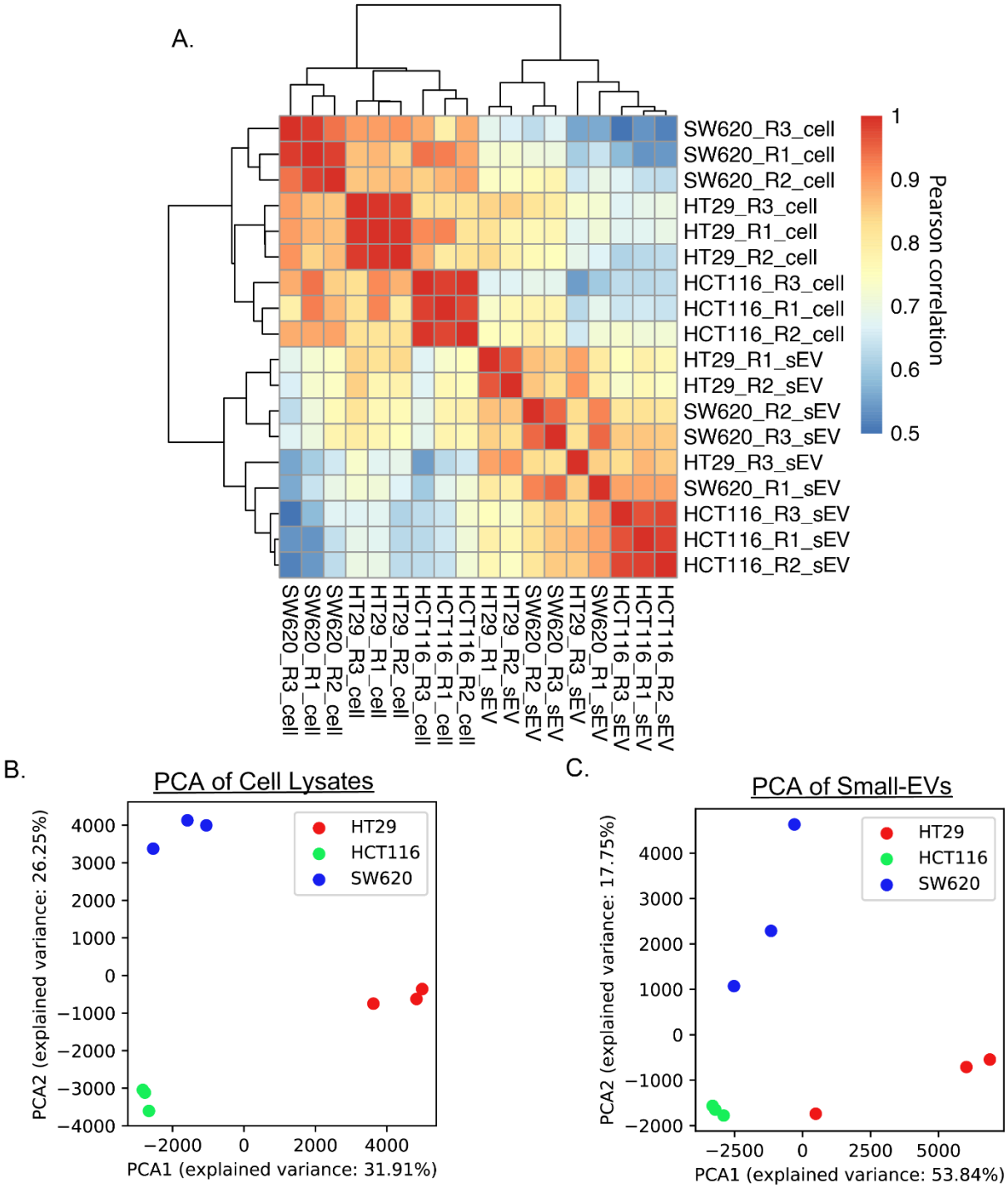

Figure S2: Replicates from shotgun proteomics cluster together with Pearson correlation and Principal Component Analysis (PCA): (A) Heatmap showing pearson correlation for all sample comparisons using the logarithm of normalized abundance values. (B) and (C) PCA for whole cell lysates (B) and small-EVs (C) displaying proteomic profile for the three replicas in each sample type.