

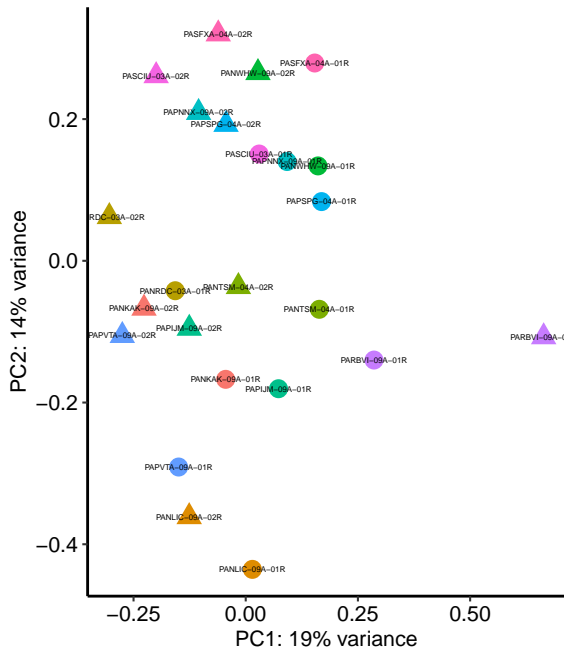
PCA plot showing the first principal component (PC1) accounting for 19% of the variance. The x-axis is labeled 'PC1: 19% variance' and ranges from -0.2 to 0.4. The y-axis is unlabeled. Data points are labeled with protein names and their corresponding domain types (e.g., -02R, -01R, -03A, -01R).

Protein Name	Domain Type	PC1 (approx.)	PC2 (approx.)
PANRDC-03A-02R	-02R	0.05	0.55
PAPVTA-09A-02R	-02R	-0.15	0.50
PANKAK-09A-02R	-02R	-0.10	0.45
PASCIU-03A-02R	-02R	0.05	0.45
PANRDC-03A-01R	-01R	0.25	0.45
PAPVTA-09A-01R	-01R	0.05	0.40
PAPNNX-09A-02R	-02R	0.10	0.40
PANRDC-03A-02R	-02R	-0.15	0.35
PANRDC-03A-01R	-01R	-0.15	0.30
PANKAK-09A-01R	-01R	0.05	0.35
PANWHH-03A-02R	-02R	0.20	0.30
PASCIU-03A-01R	-01R	0.35	0.30
PAPNNX-09A-01R	-01R	0.30	0.25
PANWHH-03A-01R	-01R	0.40	0.20
PANLIC-09A-01R	-01R	-0.05	0.25
PAPLUN-09A-01R	-01R	-0.05	0.20
PANRDC-03A-01R	-01R	-0.15	0.15
PASFXA-04A-01R	-01R	-0.15	0.10
PANRDC-03A-01R	-01R	-0.10	0.10
PANRDC-03A-01R	-01R	0.20	0.10
PANRDC-03A-01R	-01R	-0.15	0.05
PANRDC-03A-01R	-01R	-0.15	0.00

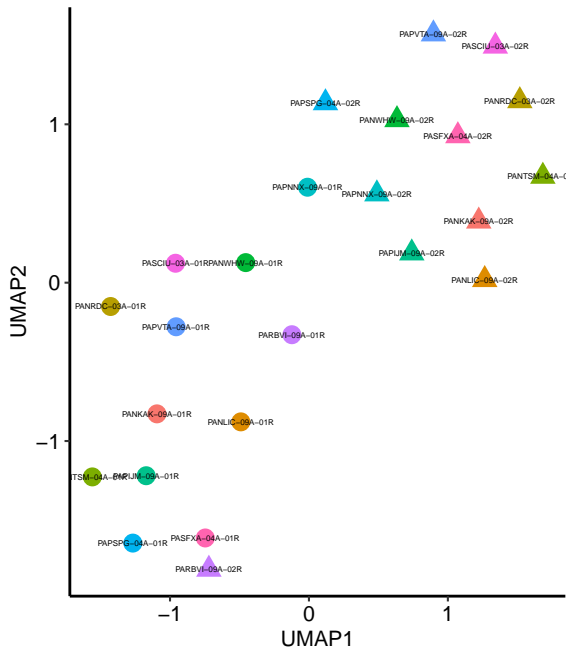
●

▲

PCA: RUVg output ($k = 2$)

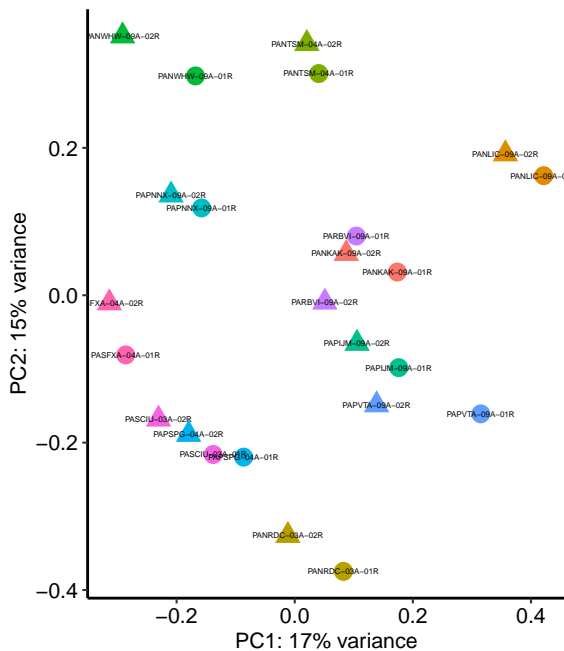


UMAP: RUVg output ($k = 2$)

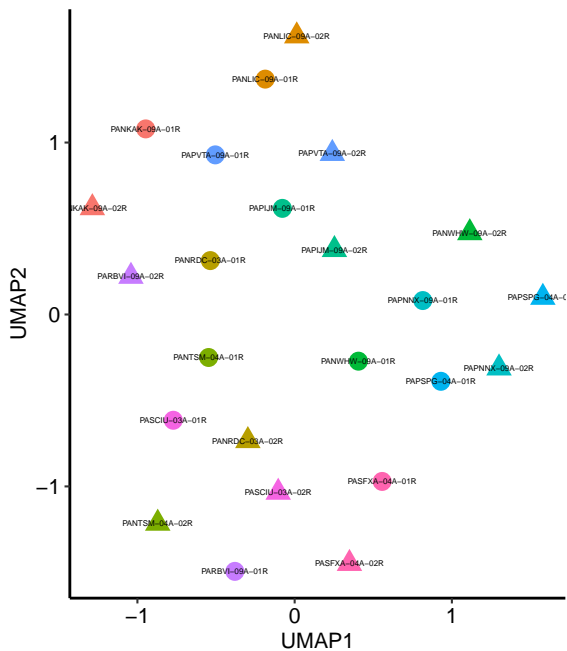


rna_library ● poly-A stranded ▲ stranded

PCA: RUVg output ($k = 3$)

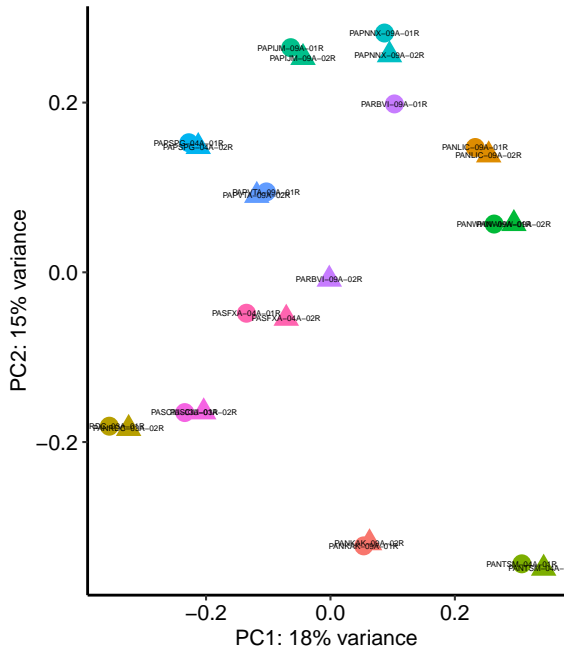


UMAP: RUVg output ($k = 3$)

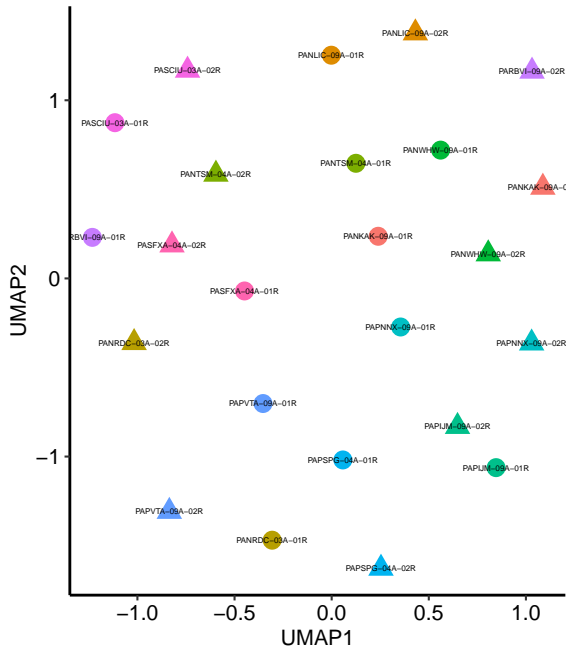


rna_library ● poly-A stranded ▲ stranded

PCA: RUVg output (k = 4)

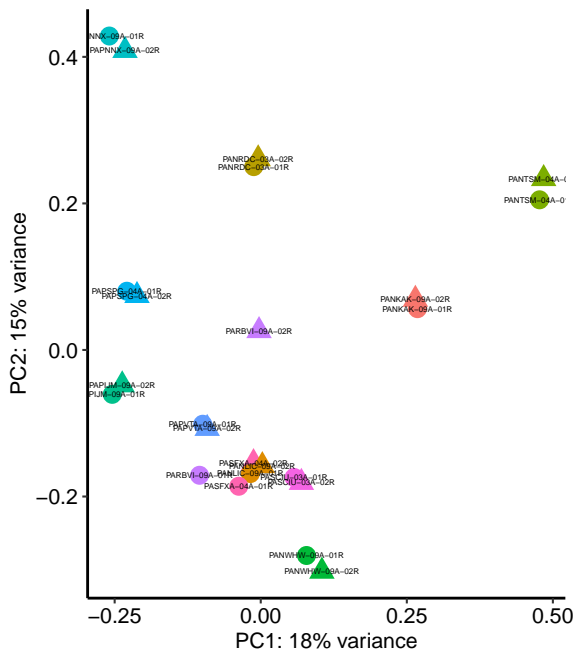


UMAP: RUVg output (k = 4)

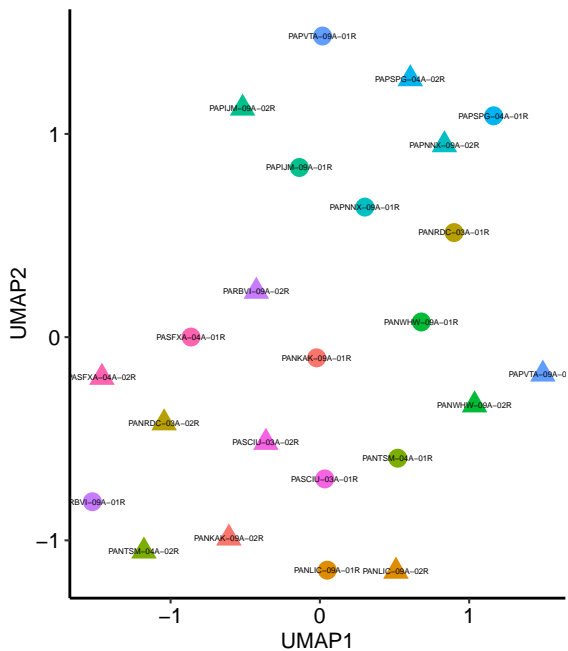


rna_library ● poly-A stranded ▲ stranded

PCA: RUVg output ($k = 5$)




UMAP: RUVg output ($k = 5$)



rna_library

- poly-A stranded

 stranded