Protein Modification All SiRCle Clusters PLOD1 P4HA1 VAE Integrated Rank (Mean) P4HA2 -0.25 0.00 0.25 PLOD3 0.50 Condition BAP1 PLOD2 PBRM1 P4HB SLC25A1 0.0 0.5 1.0 -0.5VAE integrated value