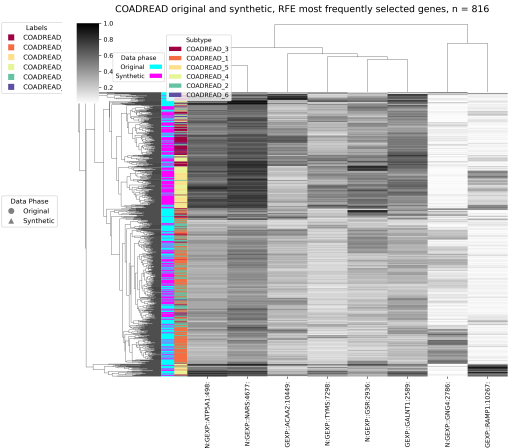
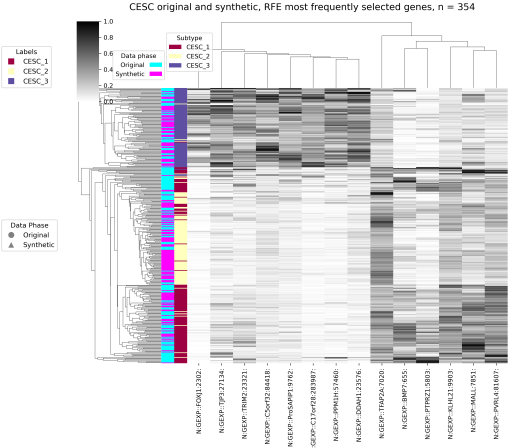
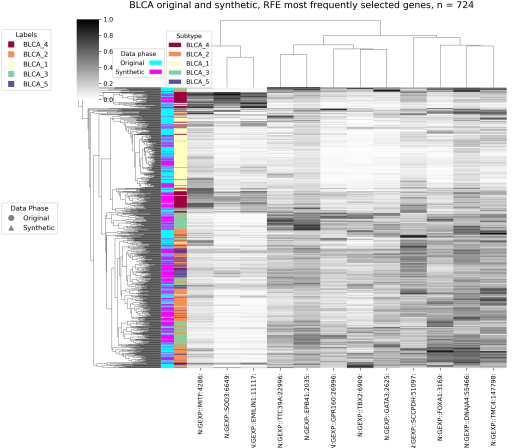
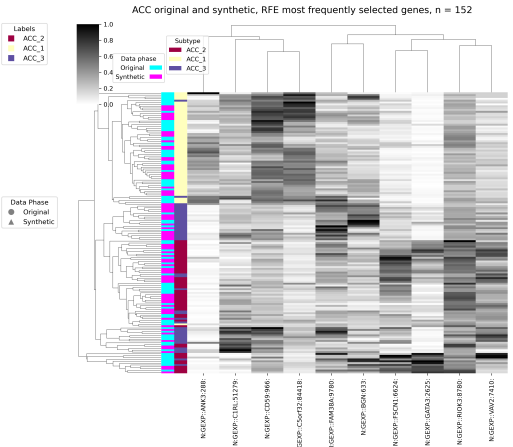
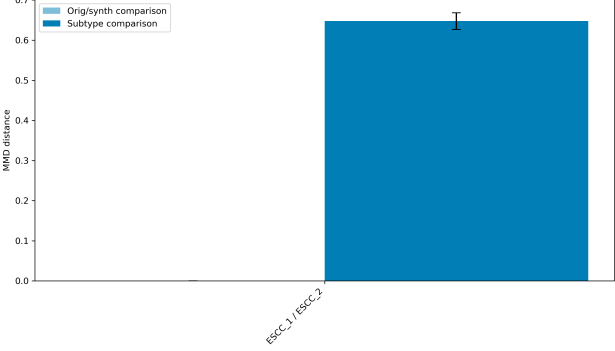
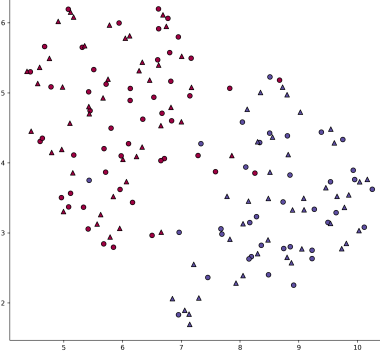




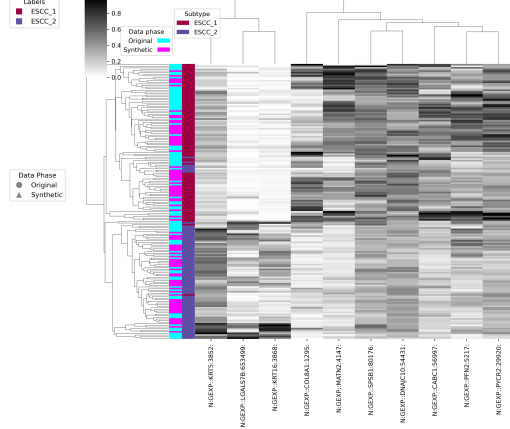
ESCC maximum mean discrepancy comparison



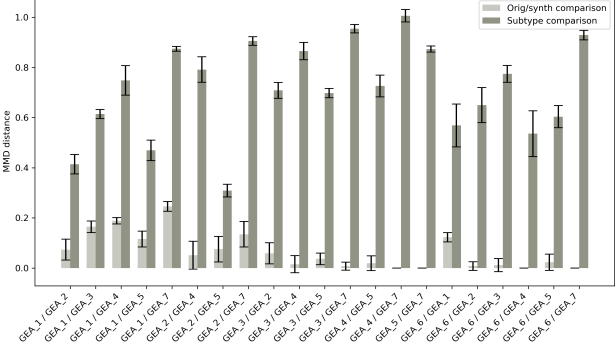
Esophageal carcinoma - synthetic and original samples cluster by subtype



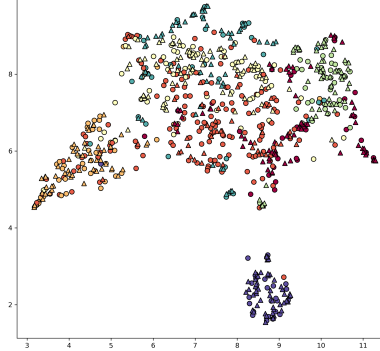
ESCC original and synthetic, RFE most frequently selected genes, n = 172



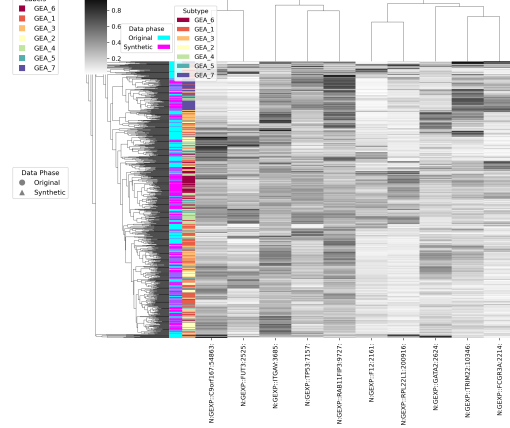
GEA maximum mean discrepancy comparison



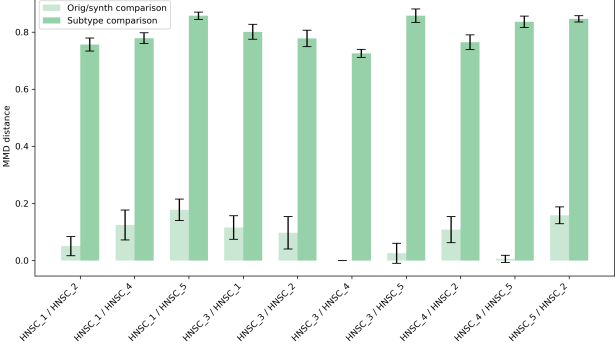
Gastroesophageal adenocarcinoma - synthetic and original samples cluster by subtype



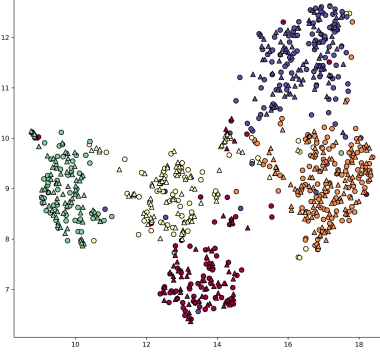
GEA original and synthetic, RFE most frequently selected genes, n = 812



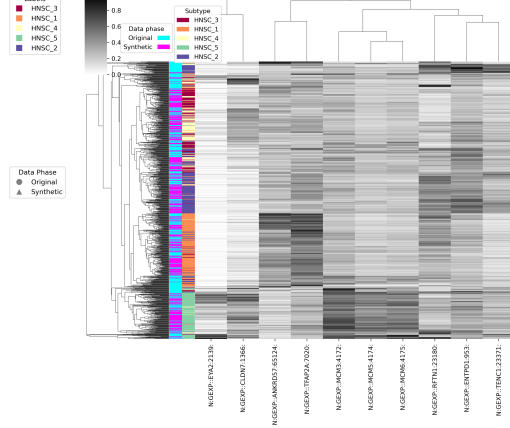
HNSC maximum mean discrepancy comparison



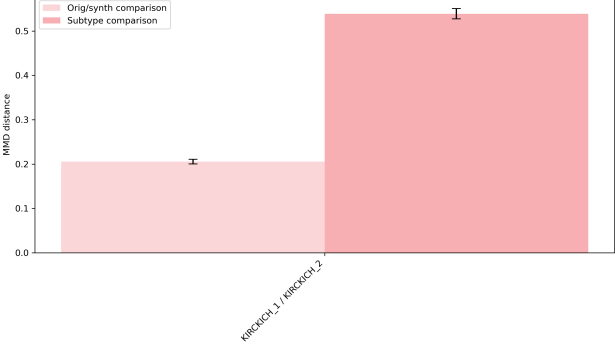
Head & Neck squamous cell - synthetic and original samples cluster by subtype



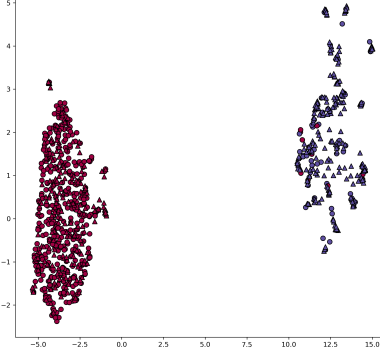
HNSC original and synthetic, RFE most frequently selected genes, n = 870



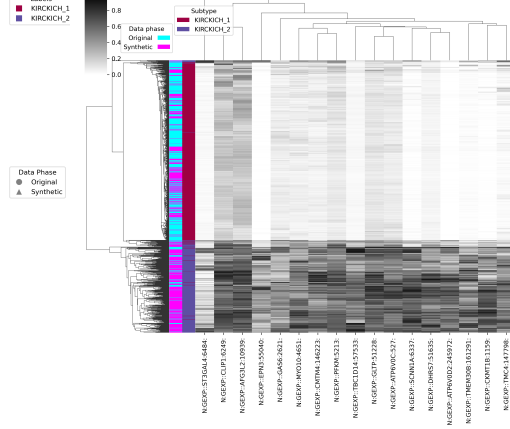
KIRC1CH maximum mean discrepancy comparison



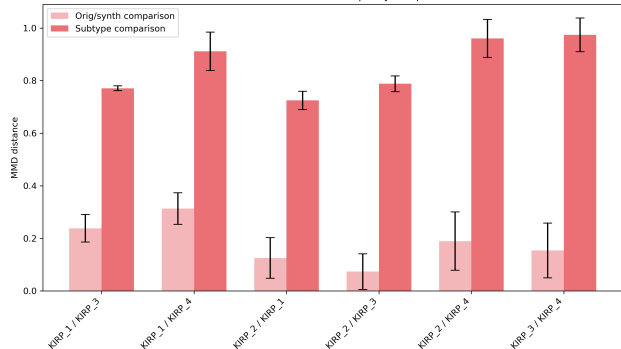
Kidney Chromophobe & RCC - synthetic and original samples cluster by subtype



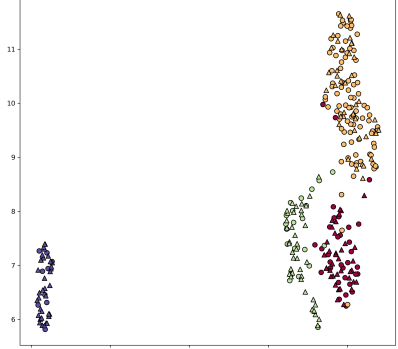
KIRC1CH original and synthetic, RFE most frequently selected genes, n = 952



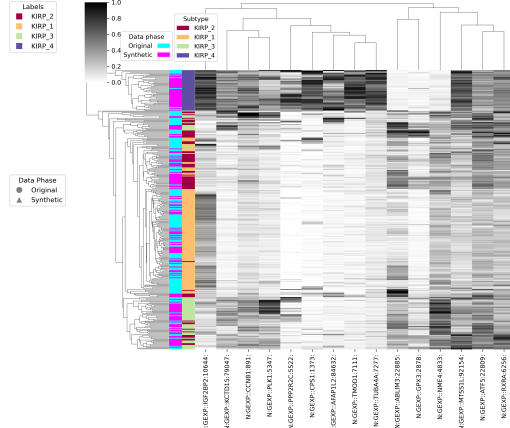
KIRP maximum mean discrepancy comparison



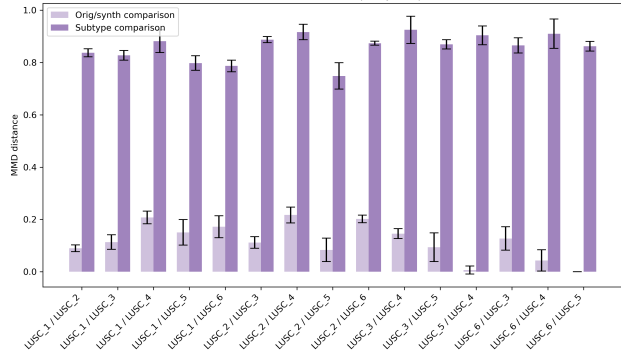
Kidney renal papillary cell - synthetic and original samples cluster by subtype



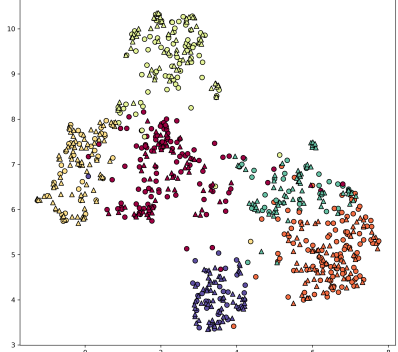
KIRP original and synthetic, RFE most frequently selected genes, n = 310



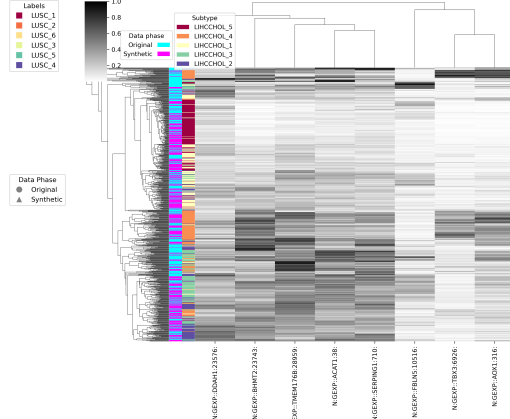
LUSC maximum mean discrepancy comparison



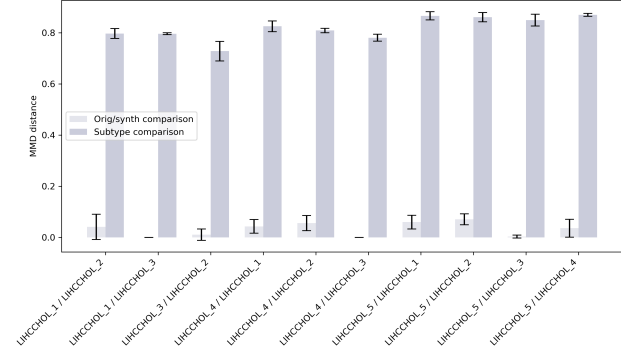
Lung squamous cell carcinoma - synthetic and original samples cluster by subtype



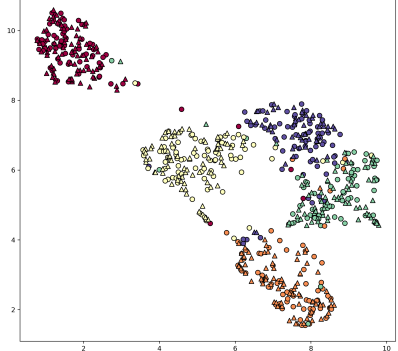
LIHCCHOL original and synthetic, RFE most frequently selected genes, n = 720



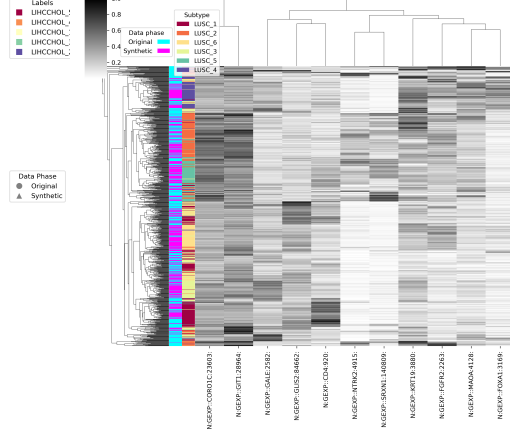
LIHCCHOL maximum mean discrepancy comparison



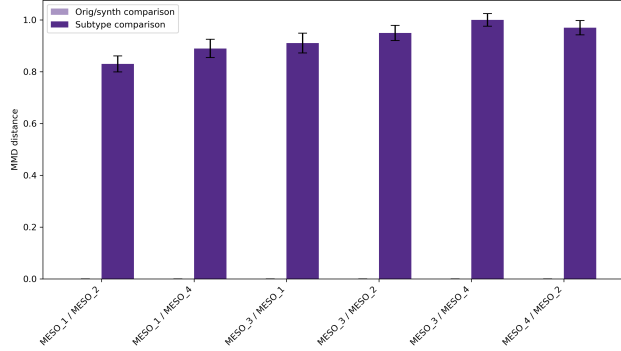
Liver hepatocellular & cholangio - synthetic and original samples cluster by subtype



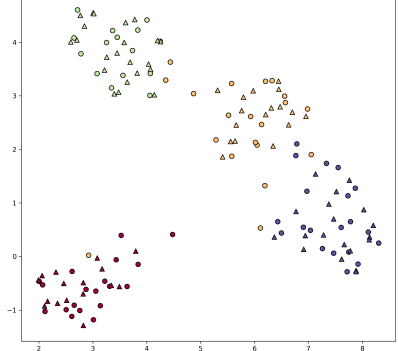
LUSC original and synthetic, RFE most frequently selected genes, n = 834



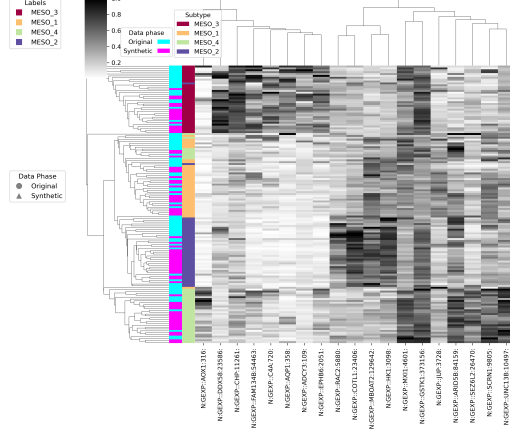
MESO maximum mean discrepancy comparison



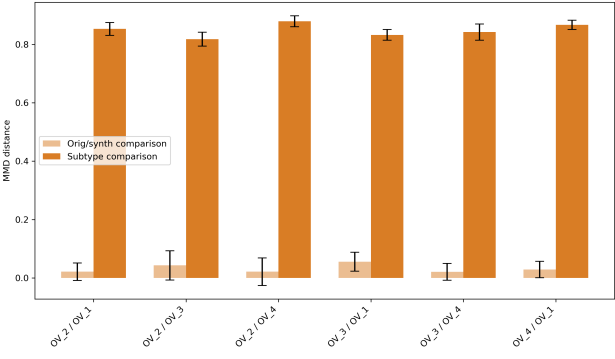
Mesothelioma - synthetic and original samples cluster by subtype



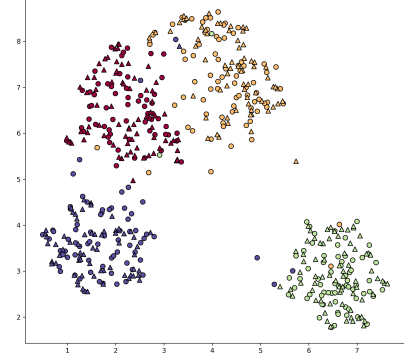
MESO original and synthetic, RFE most frequently selected genes, n = 148



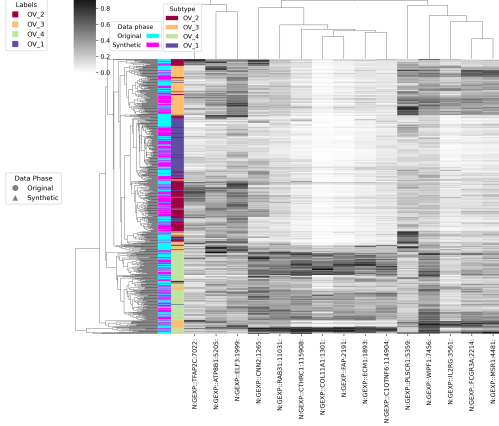
OV maximum mean discrepancy comparison



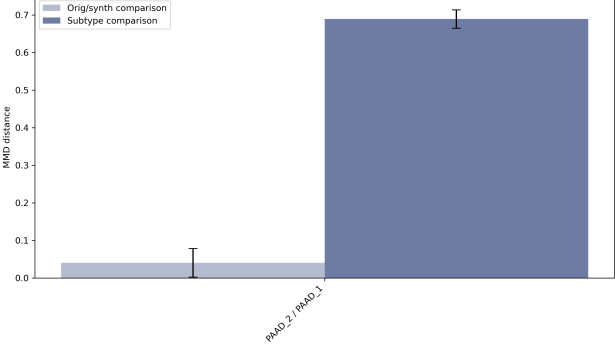
Ovarian serous cystadenocarcinoma - synthetic and original samples cluster by subtype



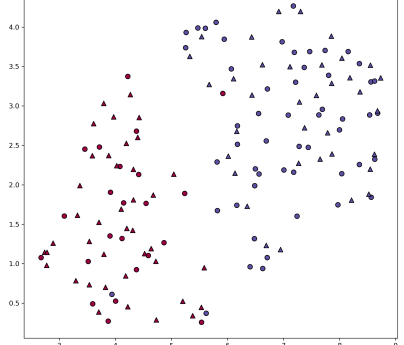
OV original and synthetic, RFE most frequently selected genes, n = 556



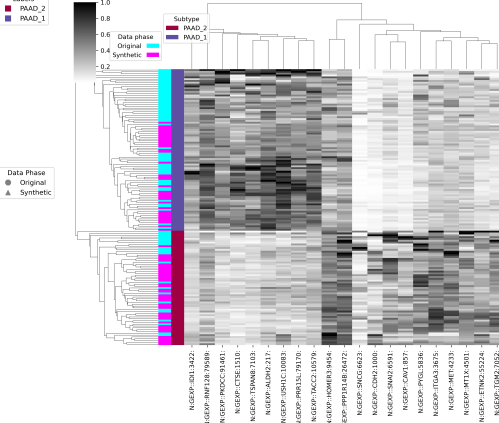
PAAD maximum mean discrepancy comparison



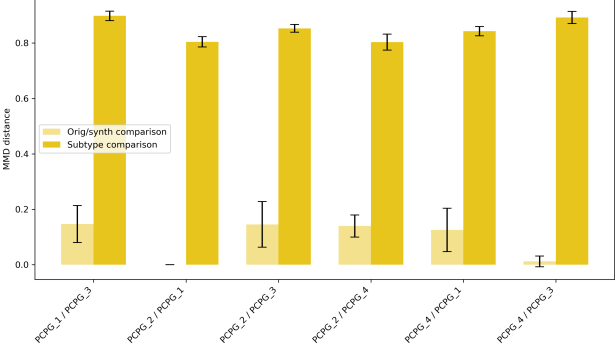
Pancreatic adenocarcinoma - synthetic and original samples cluster by subtype



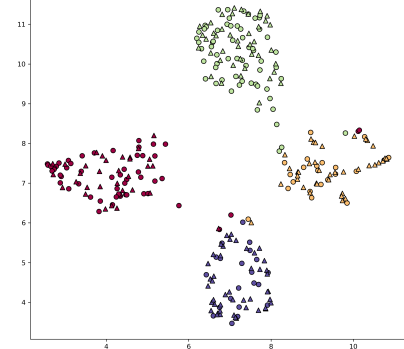
PAAD original and synthetic, RFE most frequently selected genes, n = 152



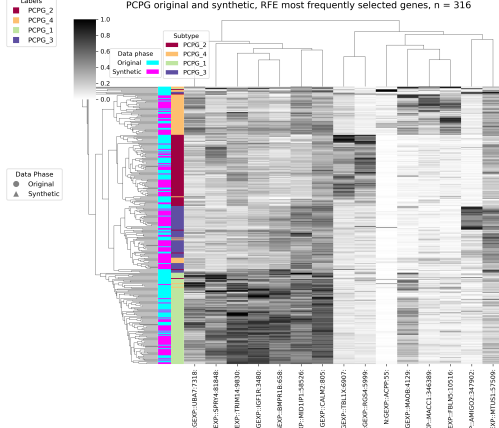
PCPG maximum mean discrepancy comparison



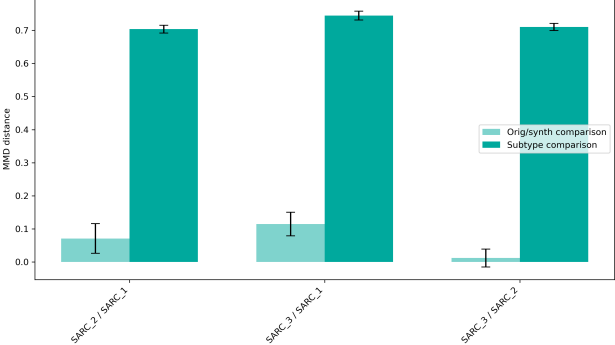
Pheochromocytoma & Paraganglioma - synthetic and original samples cluster by subtype



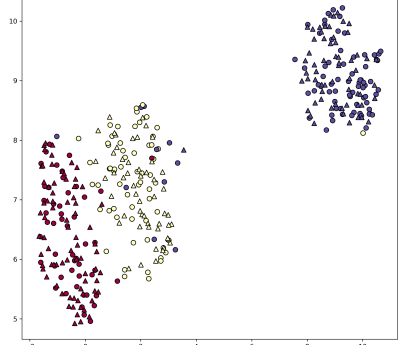
PCPG original and synthetic, RFE most frequently selected genes, n = 316



SARC maximum mean discrepancy comparison



Sarcoma - synthetic and original samples cluster by subtype



SARC original and synthetic, RFE most frequently selected genes, n = 382

