Pan-cancer genomic and proteomic analyses

Monday, January 13, 2020
3:30 PM - 4:20 PM
UTHealth School of Public Health, Room E305
RAS Building, 1200 Pressler St., Houston

Presenter:
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Pan-cancer molecular analyses seek to identify commonalities, differences, and emergent themes across tumor lineages. This talk will present findings regarding the molecular landscape of human cancers across many different tissues of origin, using data from The Cancer Genome Atlas (TCGA) consortium, the Clinical Proteomic Tumor Analysis Consortium (CPTAC), and the Pan-cancer Analysis of Whole Genomes (PCAWG) consortium.