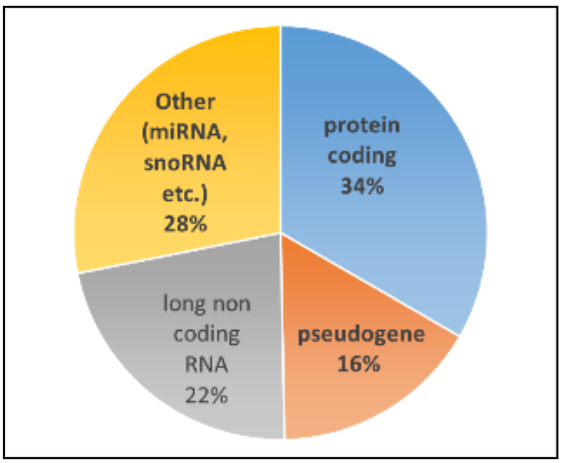
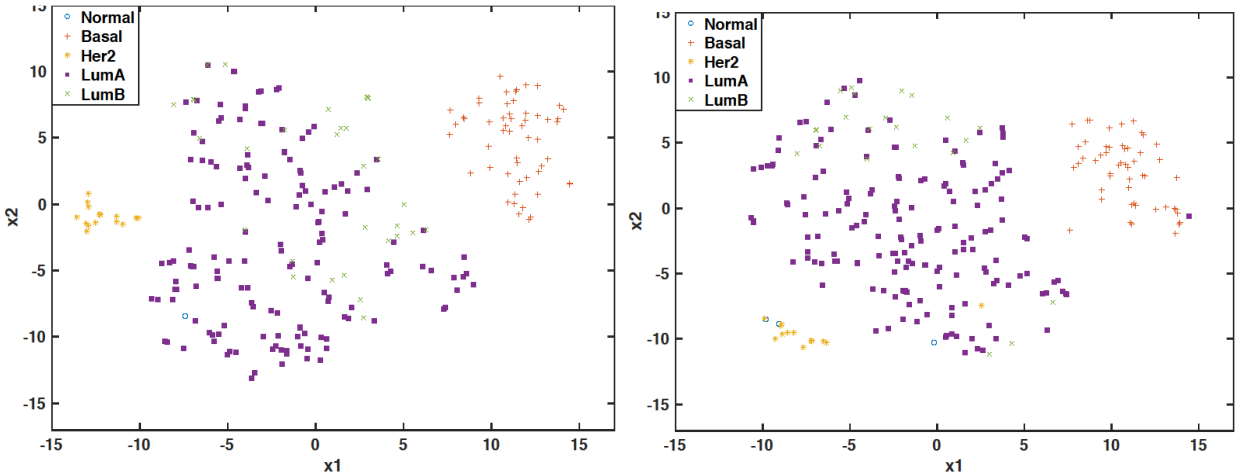


Supp. Fig 1. Transcripts Distribution in GENCODE (v23)

Supp. Fig 2. Relationship of breast cancer intrinsic subtypes and IHC biomarker status



(a) (b)

Supp. Fig 3. Visualization of breast cancer subtypes using selected 29 non-coding (a) and 36 all gene features for 253 TCGA RNAseq testing set.