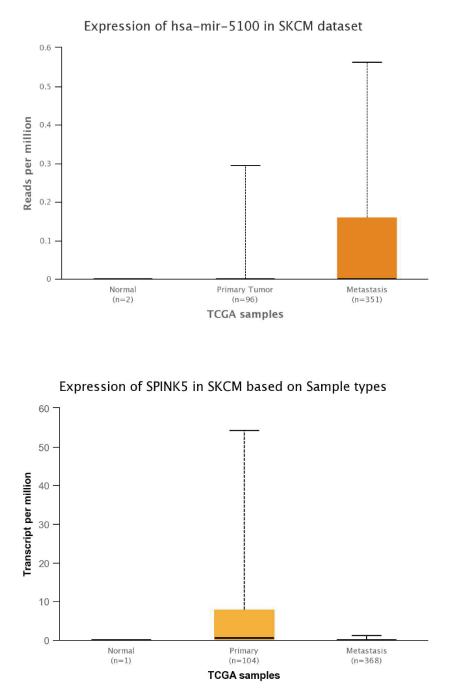
Supplementary data 2



TCGA data showed the expression patterns of miR-5100 and SPINK5 in normal tissue, primary melanoma and metastic melanoma.