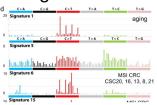
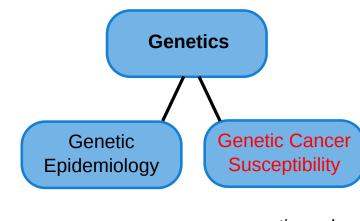
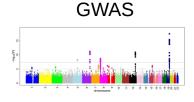


environmental exposures and mutational signatures







somatic and germline variations

