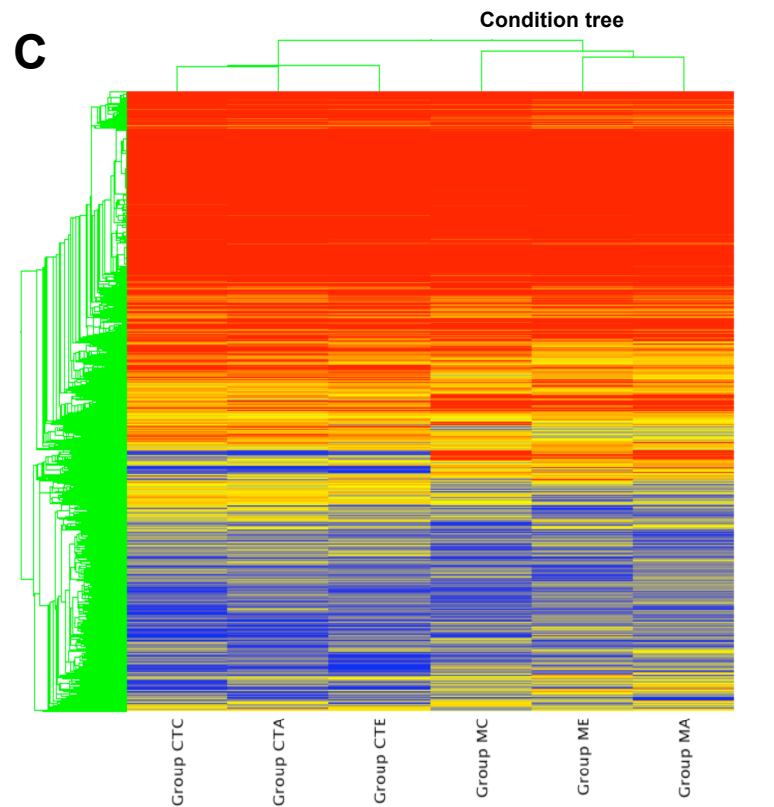


10 μg of fragmented cRNA was hybridized to each array



Data was analyzed using GeneSpring Software (Silicon Genetics).
Genes present on at least 1 array (26,029) are displayed as normalized to the median intensity of the array.
Red=High expression, Yellow=Medium expression, Blue=Low expression