

FIGURE 1A

324 miRNAs profiled in 289 Caucasian samples

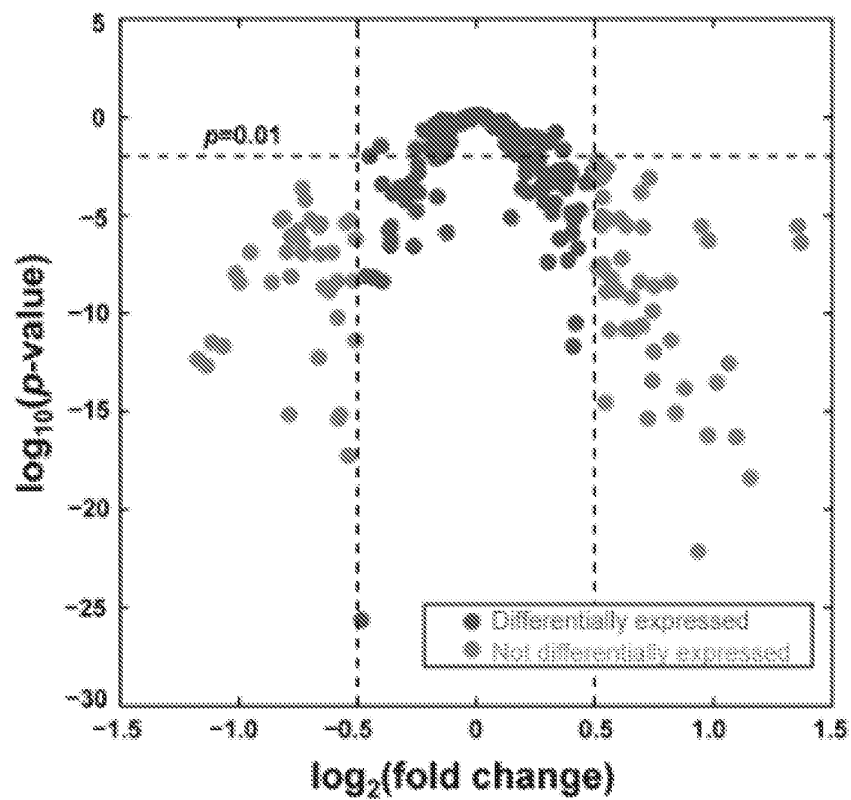


FIGURE 1B

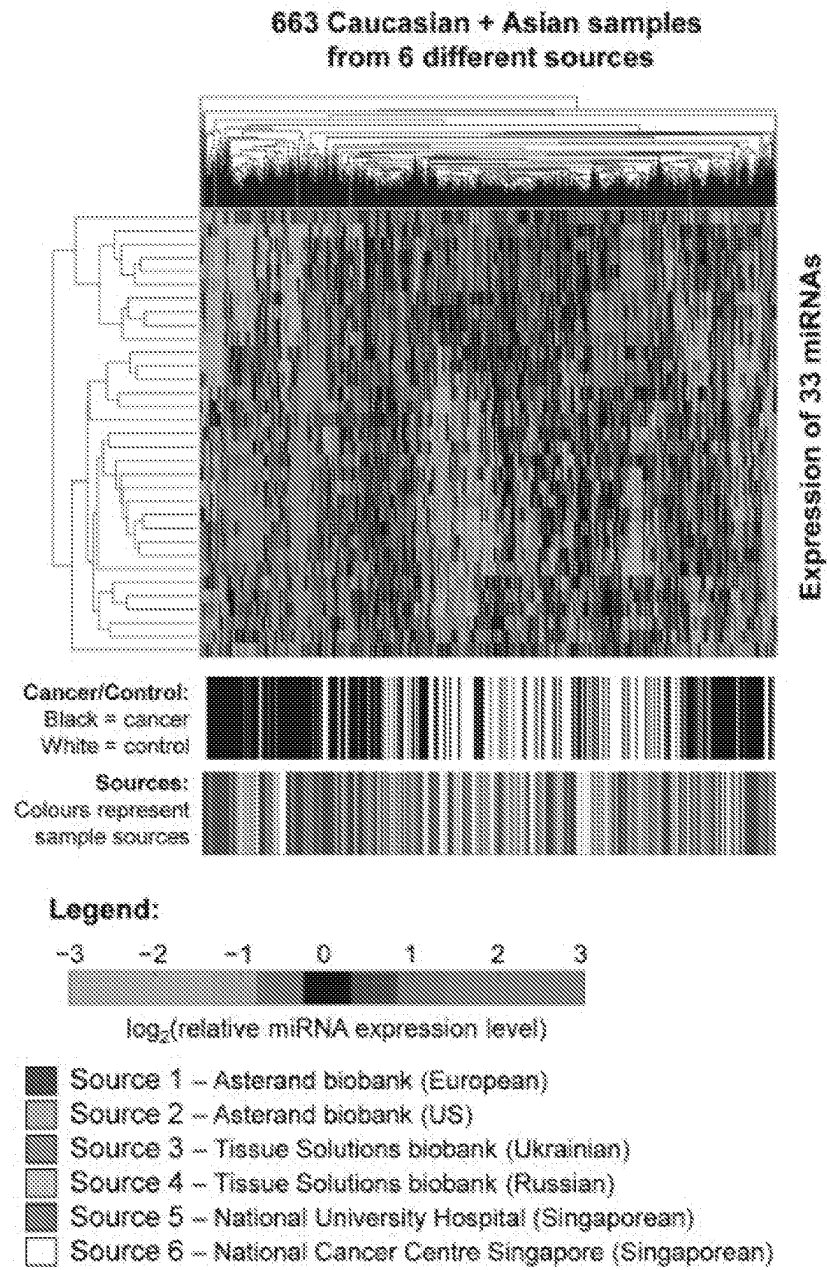


FIGURE 1C

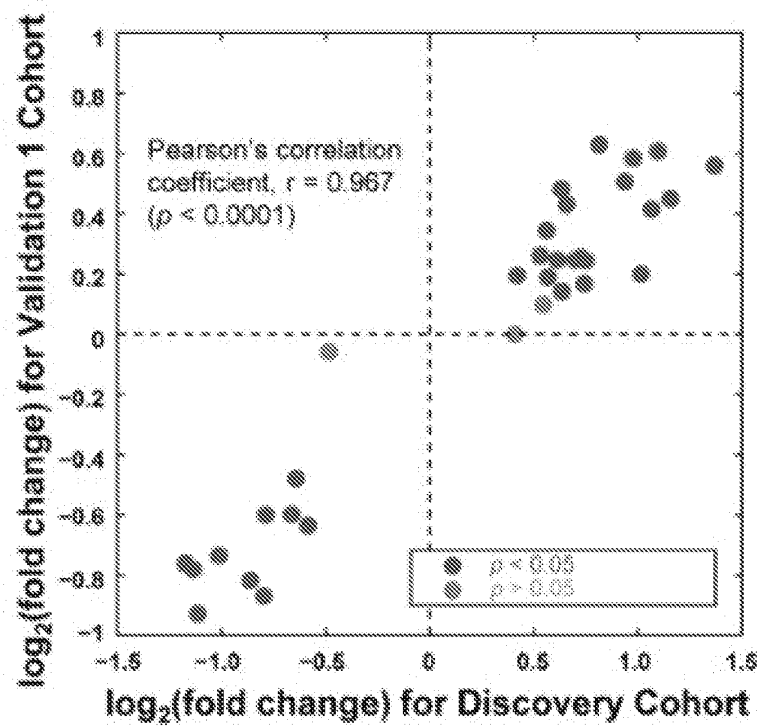


FIGURE 2A

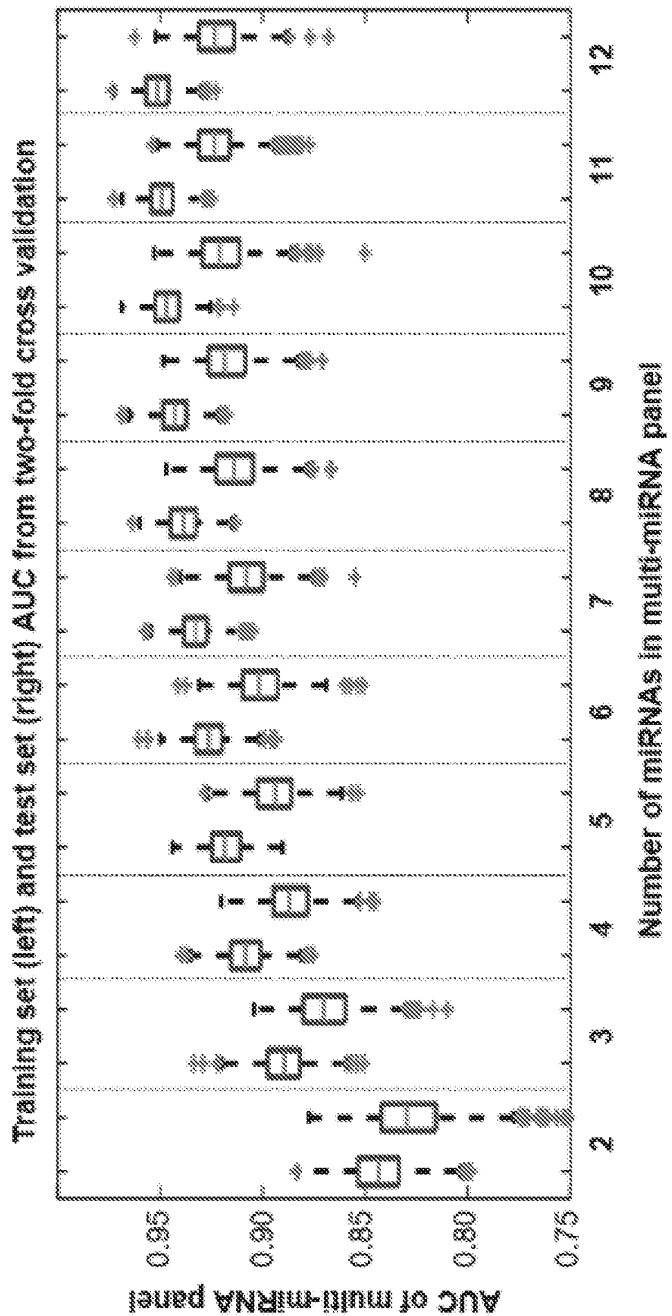


FIGURE 2B

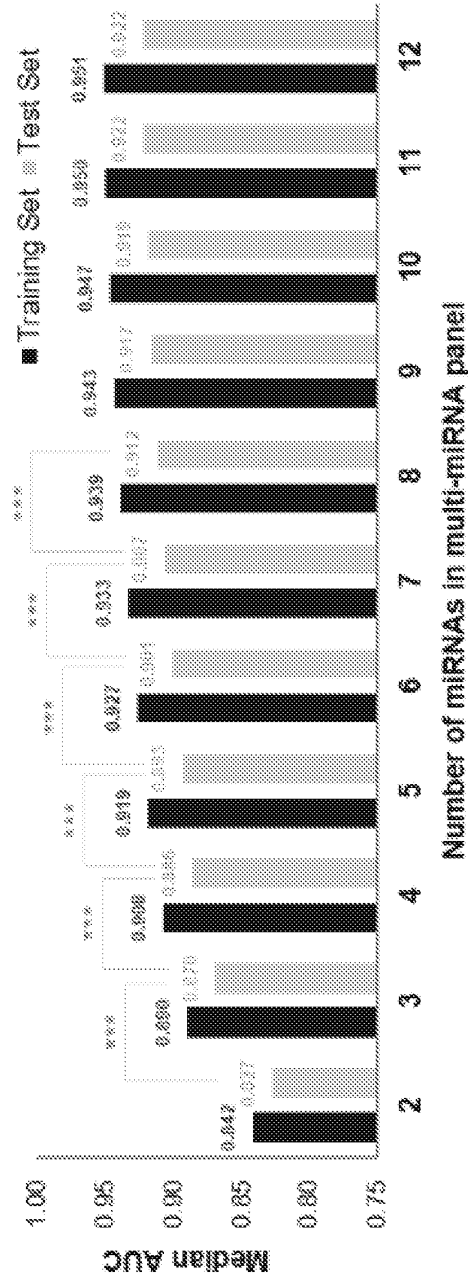


FIGURE 2C

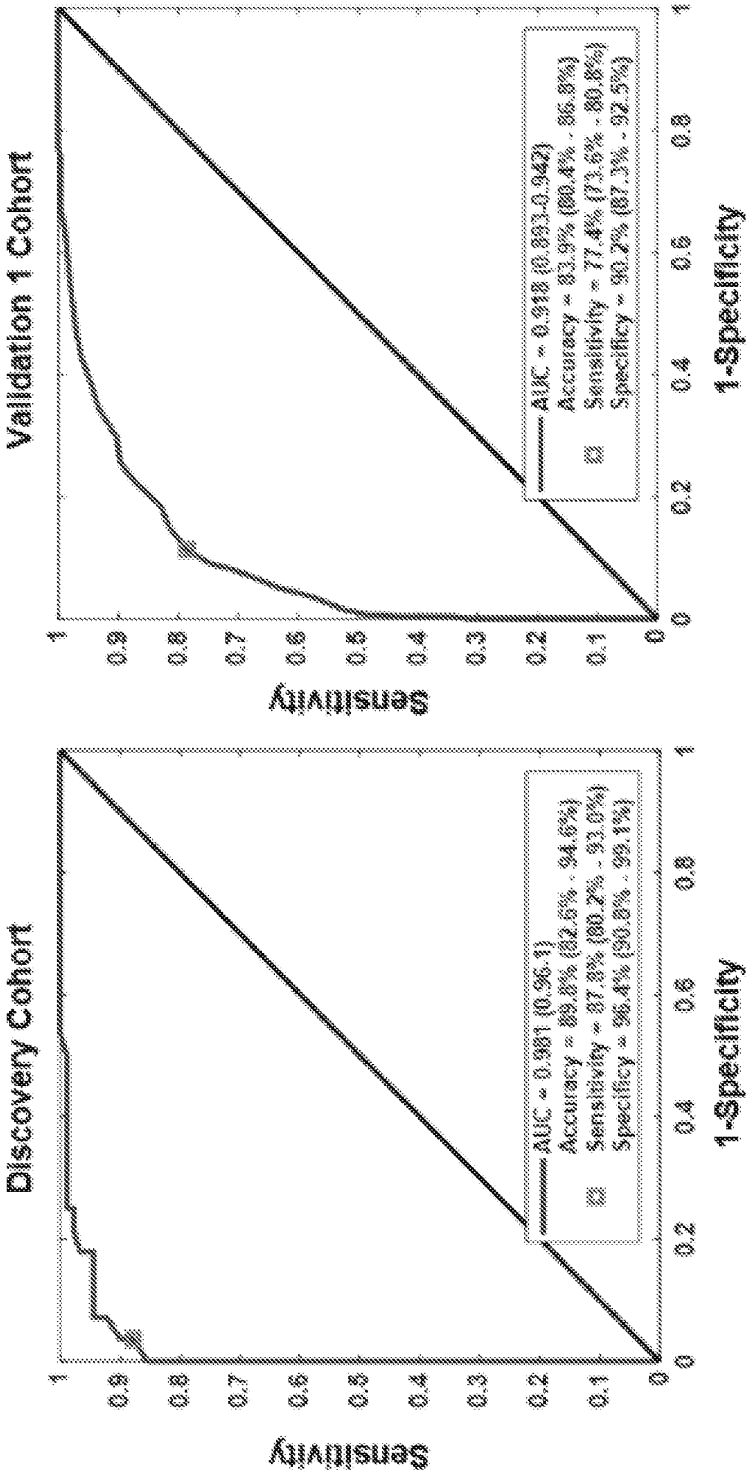


FIGURE 3A

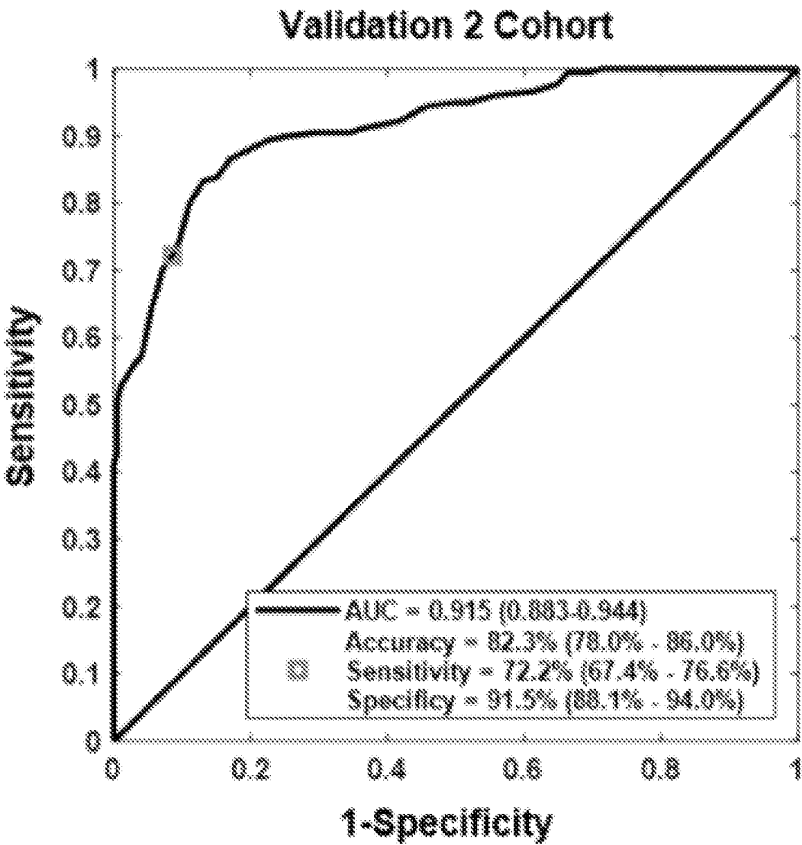


FIGURE 3B

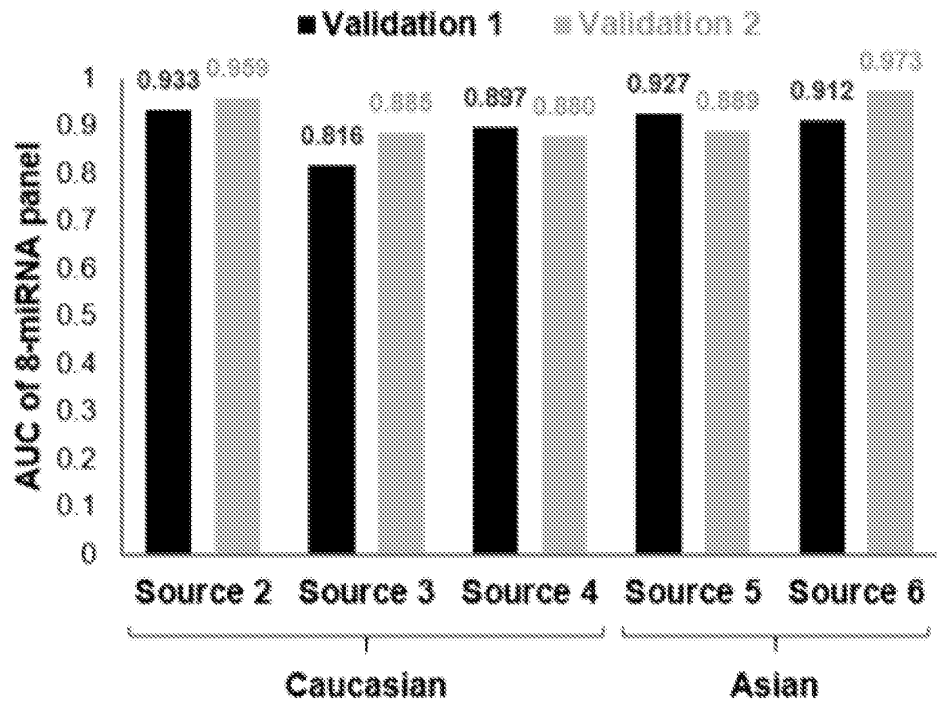


FIGURE 3C

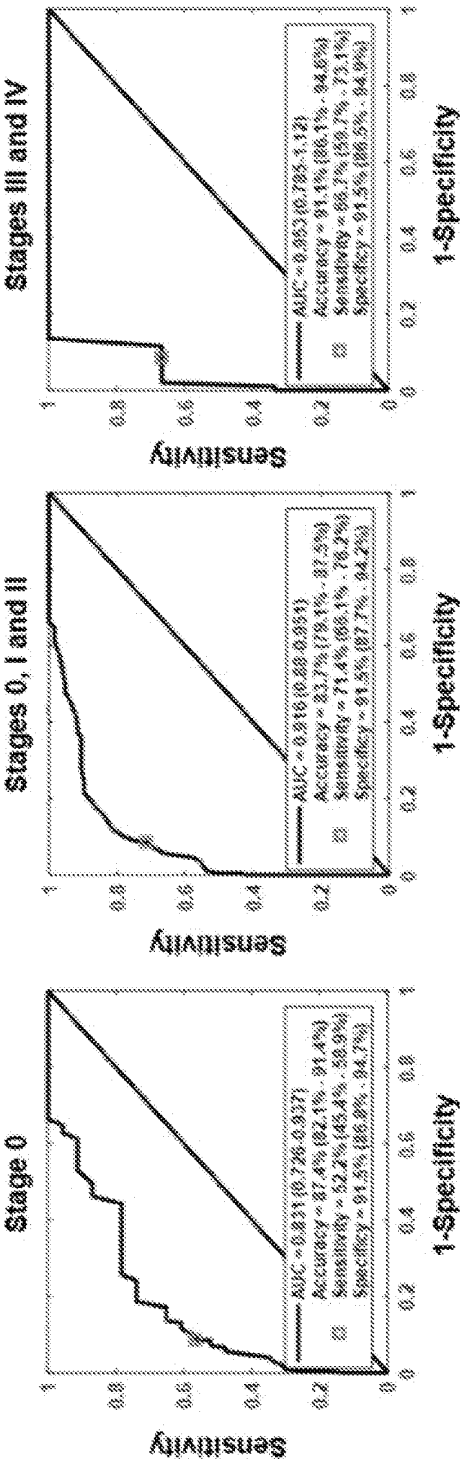


FIGURE 4A

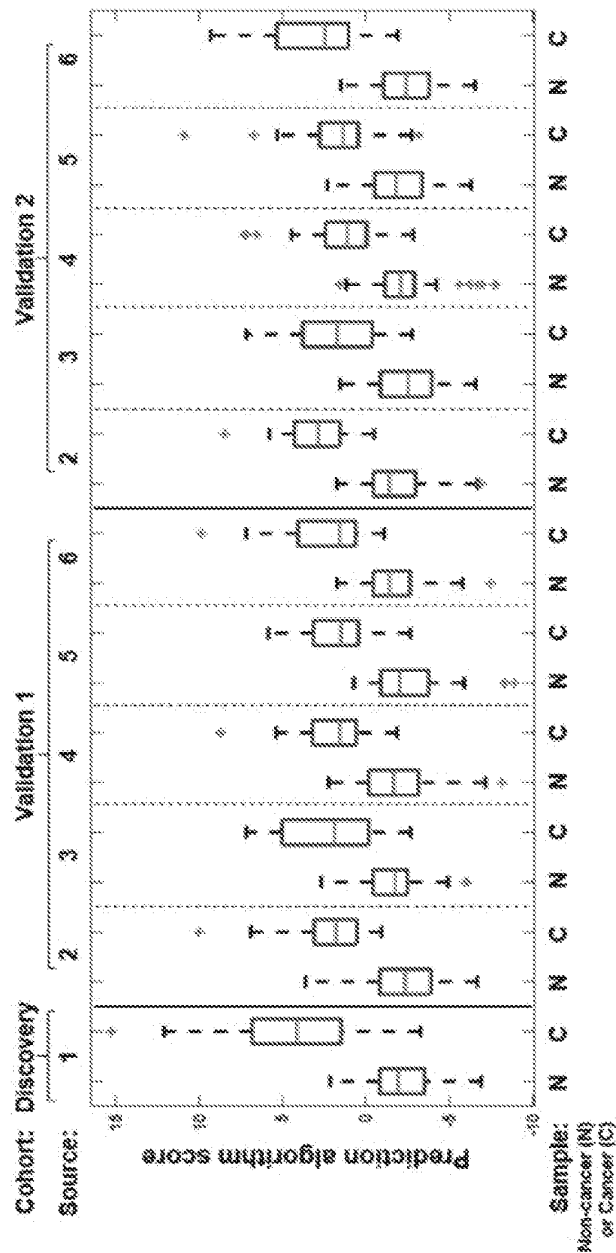


FIGURE 4B

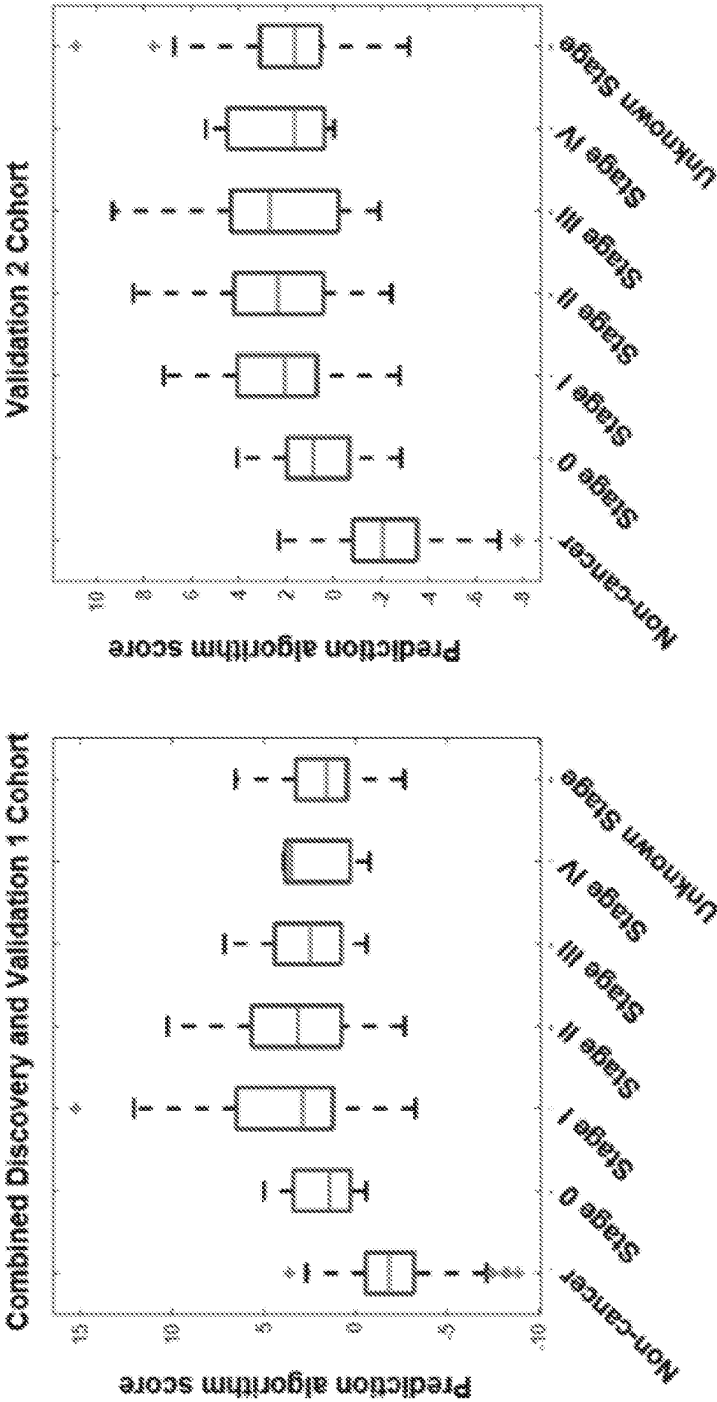


FIGURE 5

Before normalization (some of the samples have overall higher or lower expression)

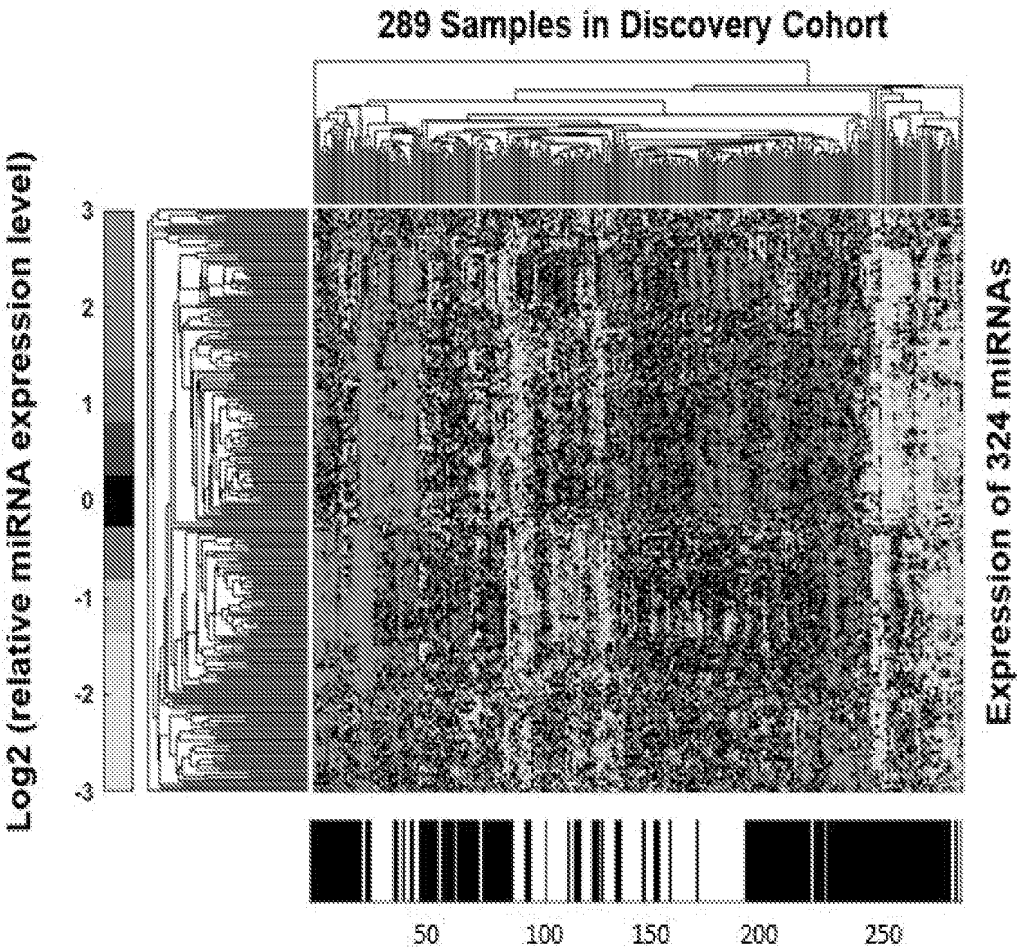


FIGURE 5 continued

After normalization with
geNORM and NormFinder.

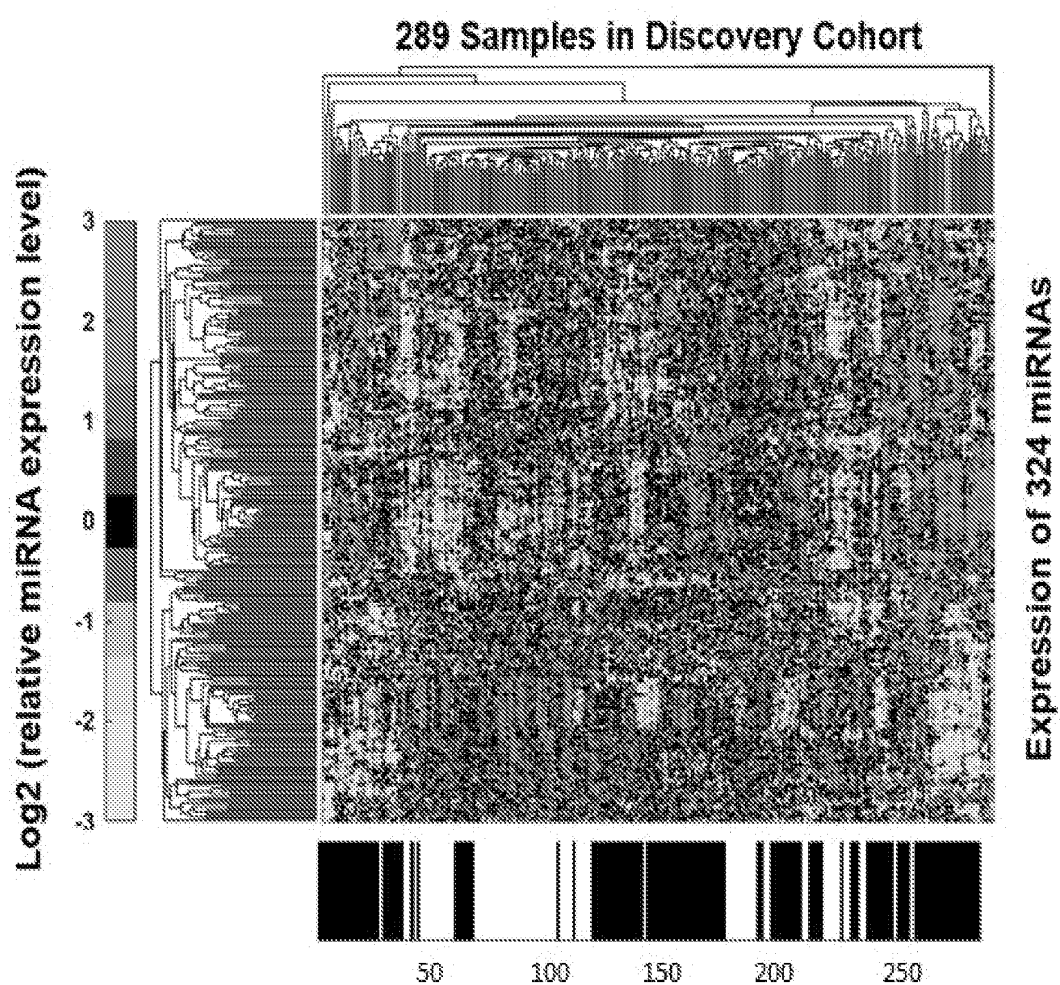


FIGURE 6

