A
p = 0.021
The expression of hsa-miR-29b-1-5p Log 2 (RPM+1)
<=60
>60
Age
B
p = 0.039
The expression of hsa-miR-29b-1-5p Log 2 (RPM+1)
Favorable
Poor
Cytogenetic risk
C
p = 0.002
The expression of hsa-miR-29b-1-5p Log 2 (RPM+1)
Negative
Positive
FLT3 mutation
D
p < 0.001
The expression of hsa-miR-29b-1-5p Log 2 (RPM+1)
Negative
Positive
NPM1 mutation
E
p = 0.022
The expression of hsa-miR-217 Log 2 (RPM+1)
Female
Male
Gender
F
p = 0.04
The expression of hsa-miR-217 Log 2 (RPM+1)
Negative
Positive
NPM1 mutation
G
p < 0.001
The expression of hsa-miR-128-1-5p Log 2 (RPM+1)
Negative
Positive
NPM1 mutation
H
p = 0.008
The expression of MAPK3 Log 2 (TPM+1)
<=60
>60
Age
I
p = 0.023
The expression of CD44 Log 2 (TPM+1)
Negative
Positive
FLT3 mutation