

Epimastigote

Trypomastigote

Amastigote

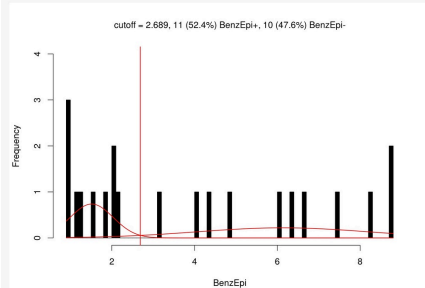
A

Analysis results

Cutoff Finder analysis of "TCruzIECOFF" using R version 2.15.0 (2012-03-30).

21 patient data sets loaded.
21 data sets with biomarker measurements.
0 data sets with outcome information.
0 data sets with survival information.

Optimal cutoff value using method "distribution": 2.689

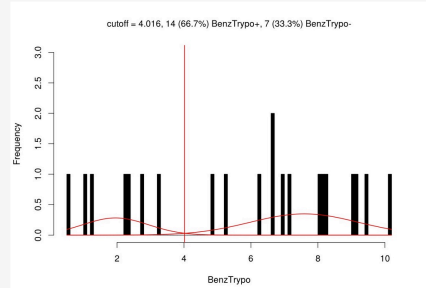


Analysis results

Cutoff Finder analysis of "TCruzIECOFF" using R version 2.15.0 (2012-03-30).

21 patient data sets loaded.
21 data sets with biomarker measurements.
0 data sets with outcome information.
0 data sets with survival information.

Optimal cutoff value using method "distribution": 4.016

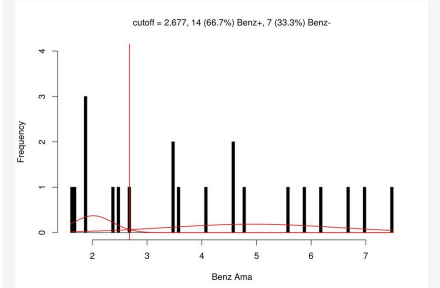


Analysis results

Cutoff Finder analysis of "TCruzIECOFF" using R version 2.15.0 (2012-03-30).

21 patient data sets loaded.
21 data sets with biomarker measurements.
0 data sets with outcome information.
0 data sets with survival information.

Optimal cutoff value using method "distribution": 2.677



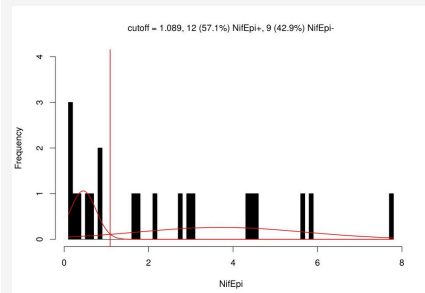
B

Analysis results

Cutoff Finder analysis of "TCruzIECOFF" using R version 2.15.0 (2012-03-30).

21 patient data sets loaded.
21 data sets with biomarker measurements.
0 data sets with outcome information.
0 data sets with survival information.

Optimal cutoff value using method "distribution": 1.089

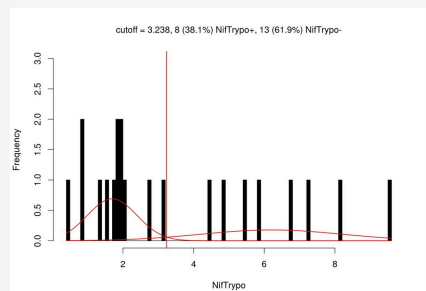


Analysis results

Cutoff Finder analysis of "TCruzIECOFF" using R version 2.15.0 (2012-03-30).

21 patient data sets loaded.
21 data sets with biomarker measurements.
0 data sets with outcome information.
0 data sets with survival information.

Optimal cutoff value using method "distribution": 3.238



Analysis results

Cutoff Finder analysis of "TCruzIECOFF" using R version 2.15.0 (2012-03-30).

21 patient data sets loaded.
21 data sets with biomarker measurements.
0 data sets with outcome information.
0 data sets with survival information.

Optimal cutoff value using method "distribution": 2.123

