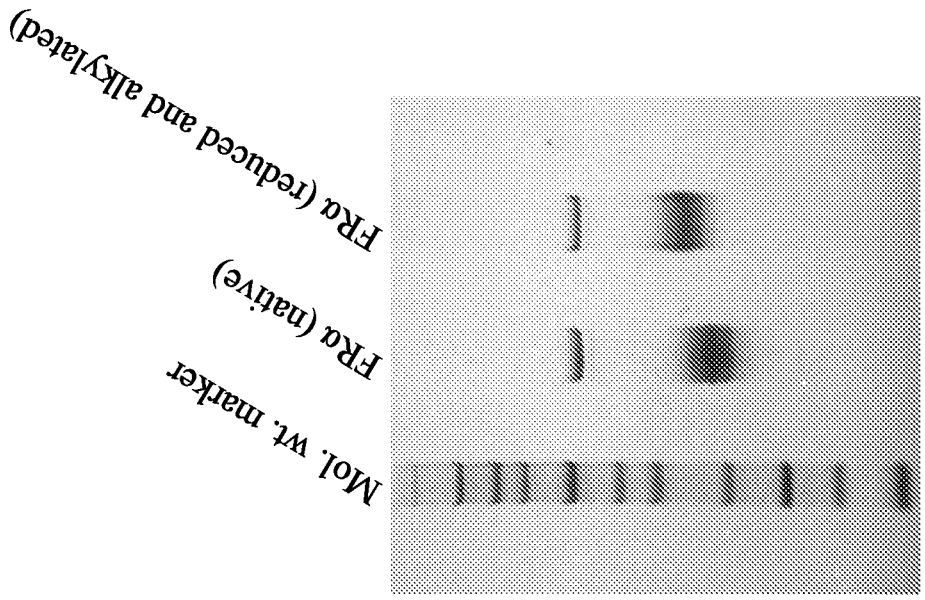
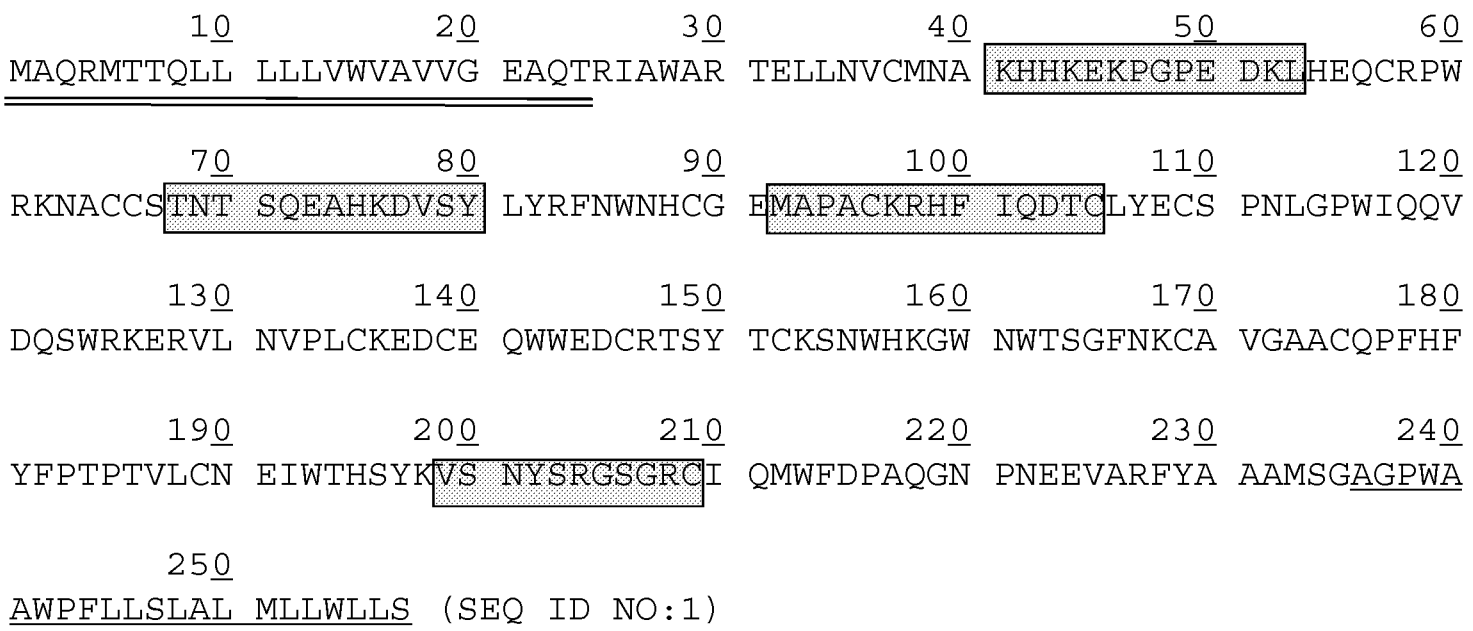


Figure 1



Legend

Residues 1-24 (double underscore) = Signal Sequence (not present in rh-FRα)
Residues 41-53 (shaded) = MAb 9F3 (PTA-11887) epitope
Residues 68-80 and 92-105 (shaded) = MAb 24F12 (PTA-11886) epitope
Residues 199-209 (shaded) = MAb 26B3 (PTA-11885) epitope
Residues 236-257 (underscore) = GPI-anchor Sequence (not present in rh-FRα)

Figure 2

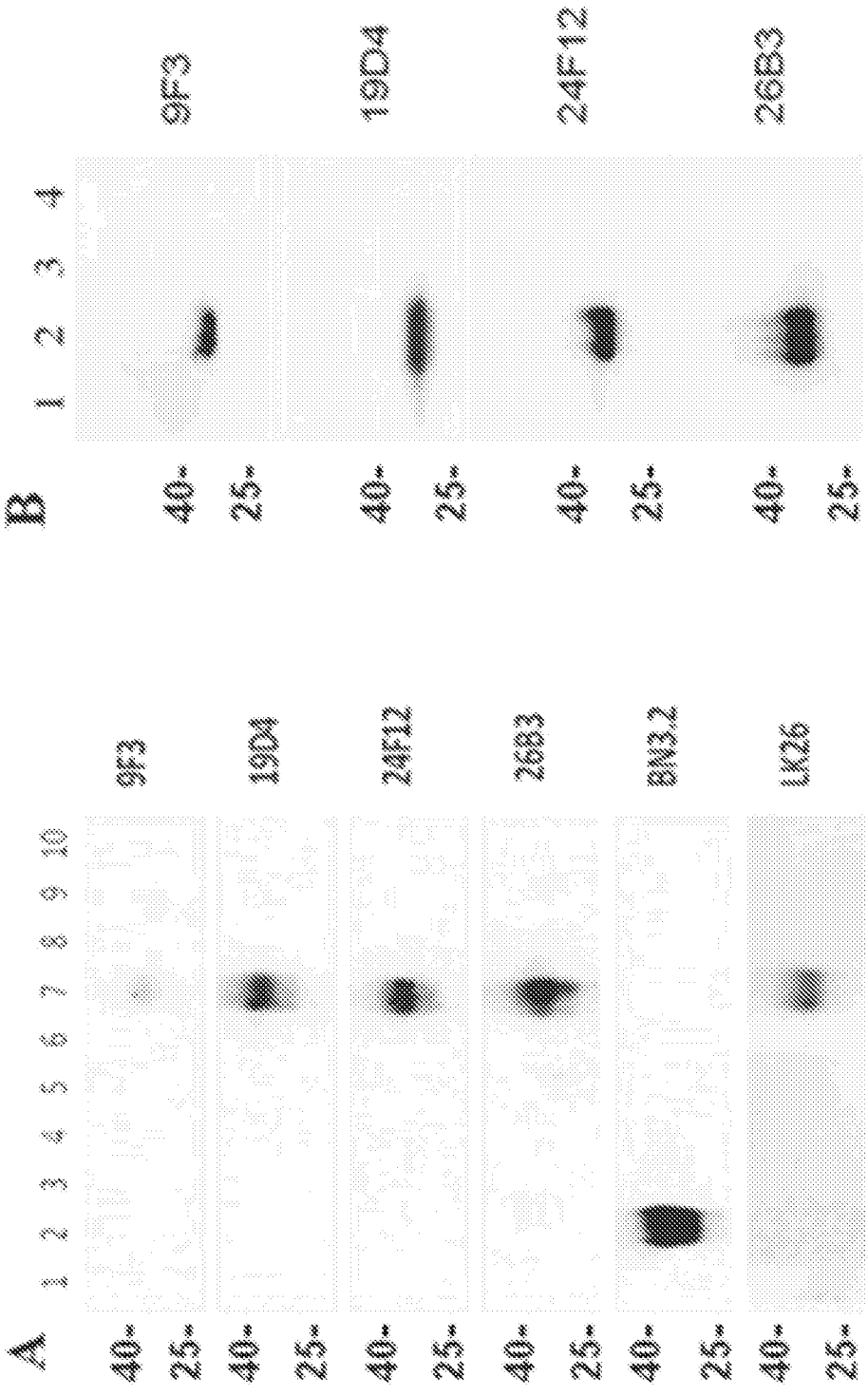

A
1 2 3 4 5 6 7 8 9 10
9F3
19D4
24F12
26B3
BN3.2
LK26
40-
25-
B
1 2 3 4
9F3
19D4
24F12
26B3
40-
25-

Figure 3

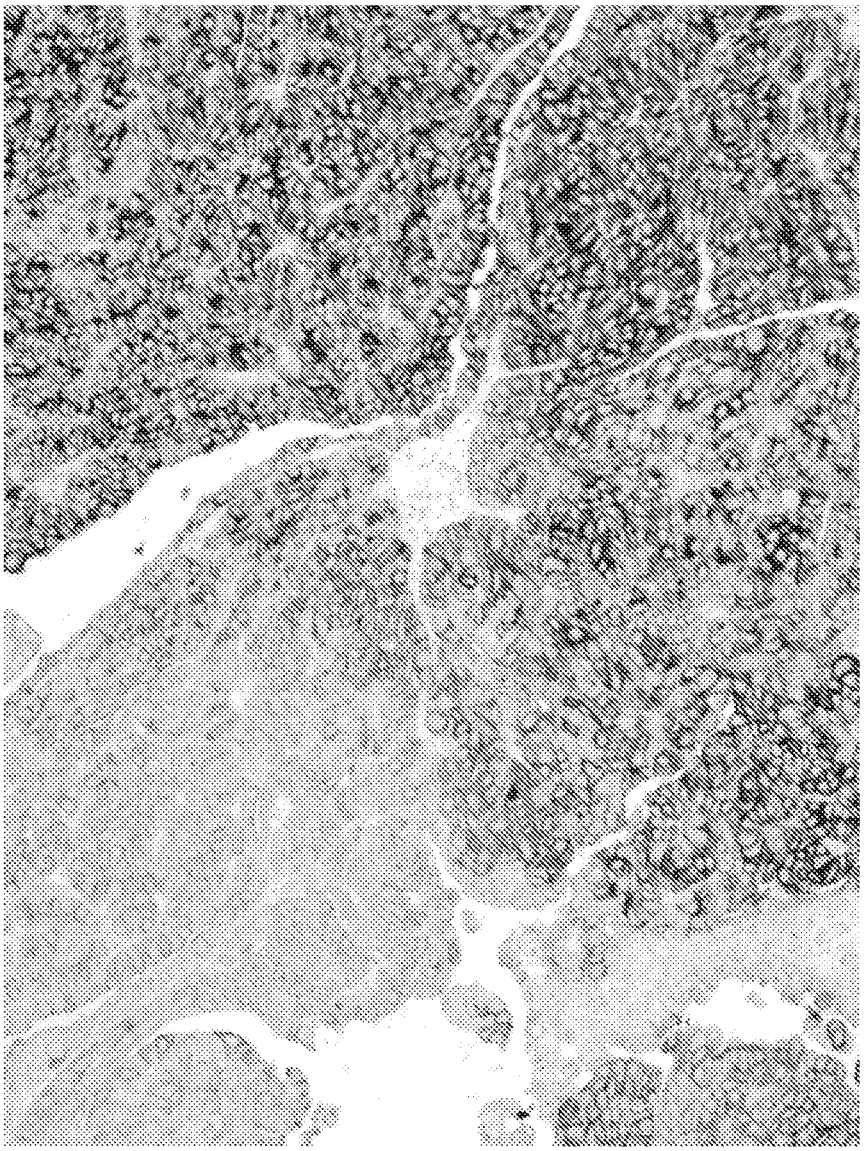
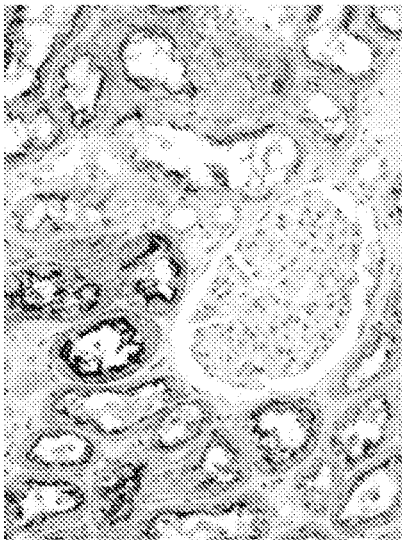
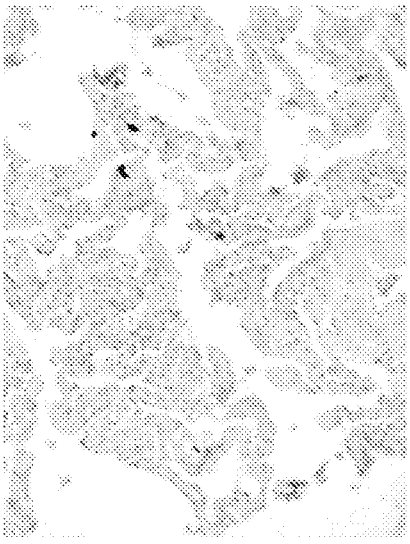


Figure 4



B.



A.

Figure 5

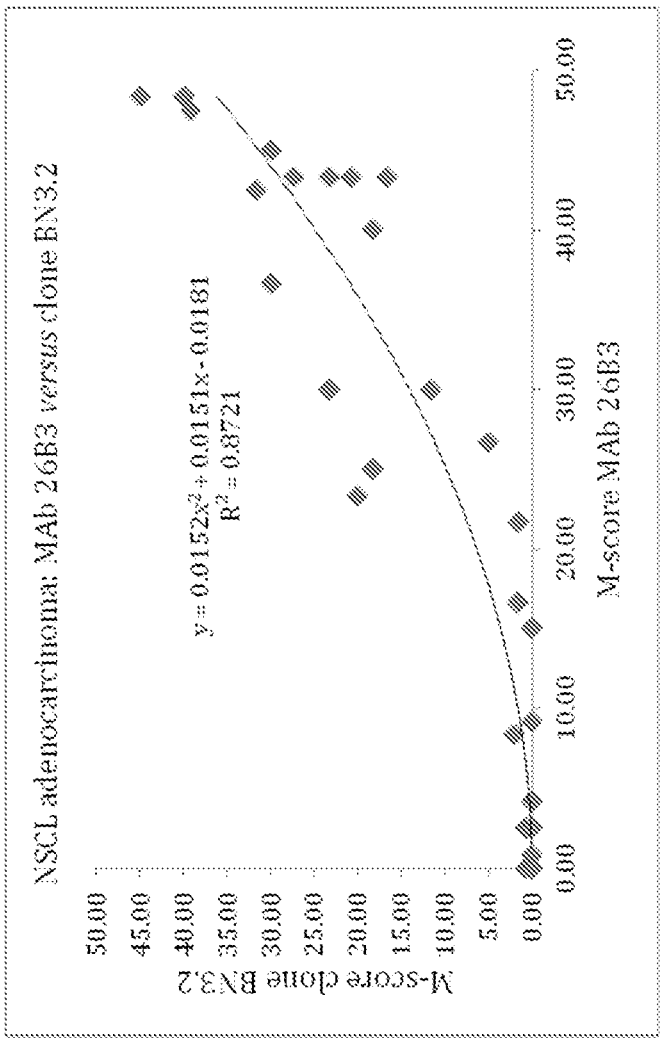

NSCL adenocarcinoma: MAb 26B3 *versus* clone BN3.2
$y = 0.0152x^2 + 0.0151x - 0.0181$
$R^2 = 0.8721$
M-score clone BN3.2
M-score MAb 26B3
0.00
5.00
10.00
15.00
20.00
25.00
30.00
35.00
40.00
45.00
50.00
0.00
10.00
20.00
30.00
40.00
50.00

Figure 6

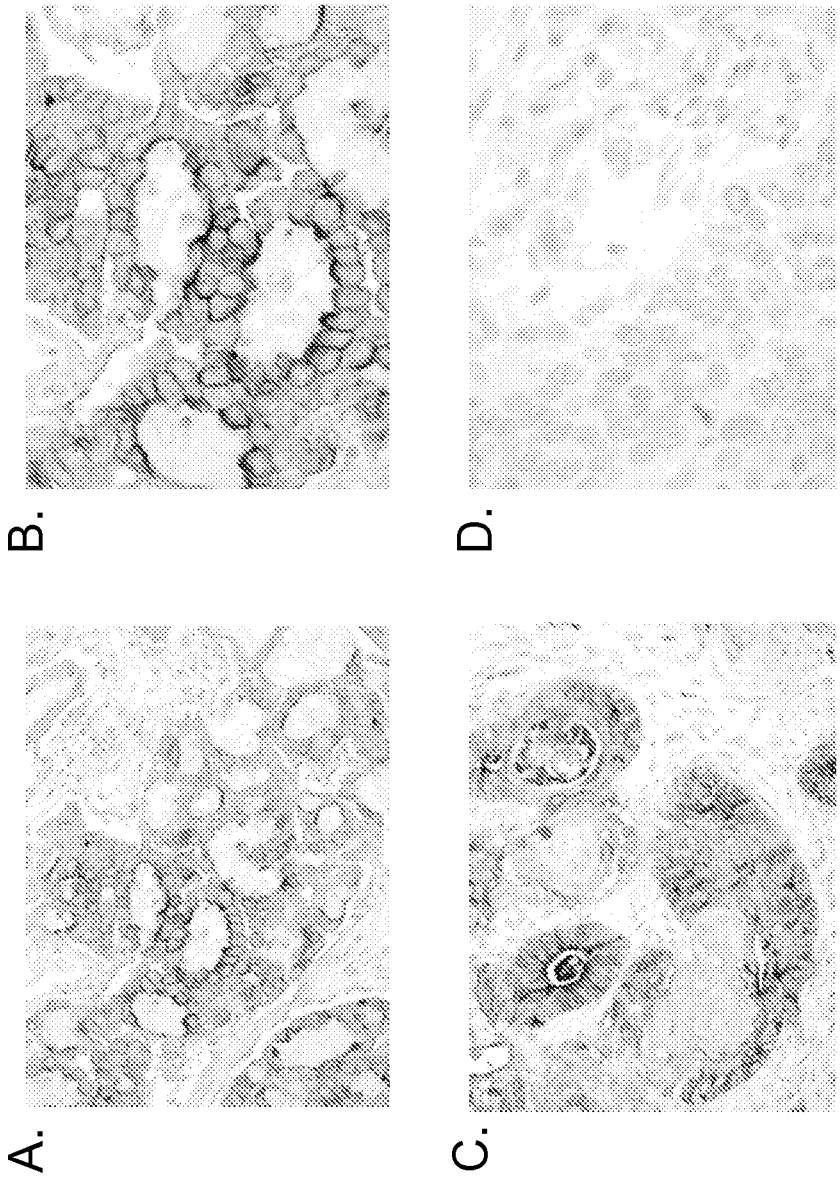


Figure 7

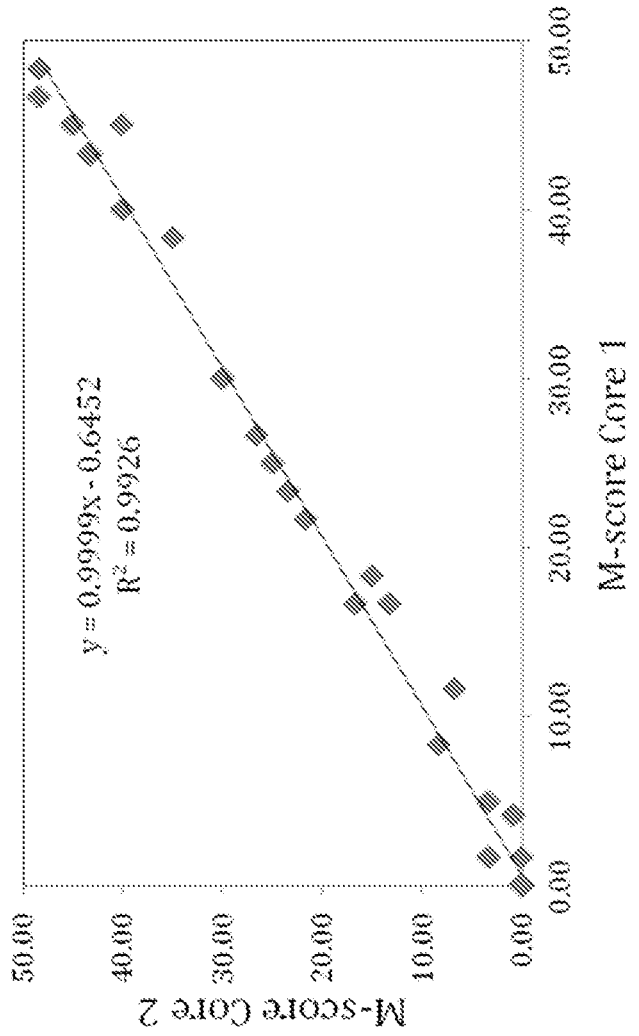

y = 0.9999x - 0.6452
R² = 0.9926
M-score Core 1
M-score Core 2
0.00
10.00
20.00
30.00
40.00
50.00

Figure 8

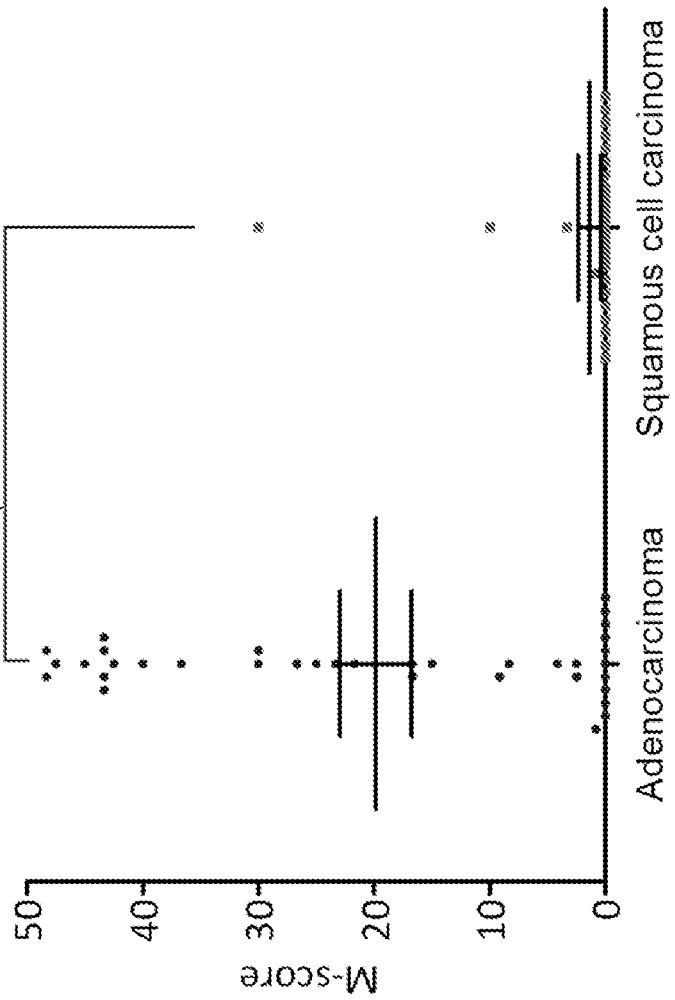

p < 0.0001
M-score
50
40
30
20
10
0
Adenocarcinoma
Squamous cell carcinoma
Lung Histology

Figure 9

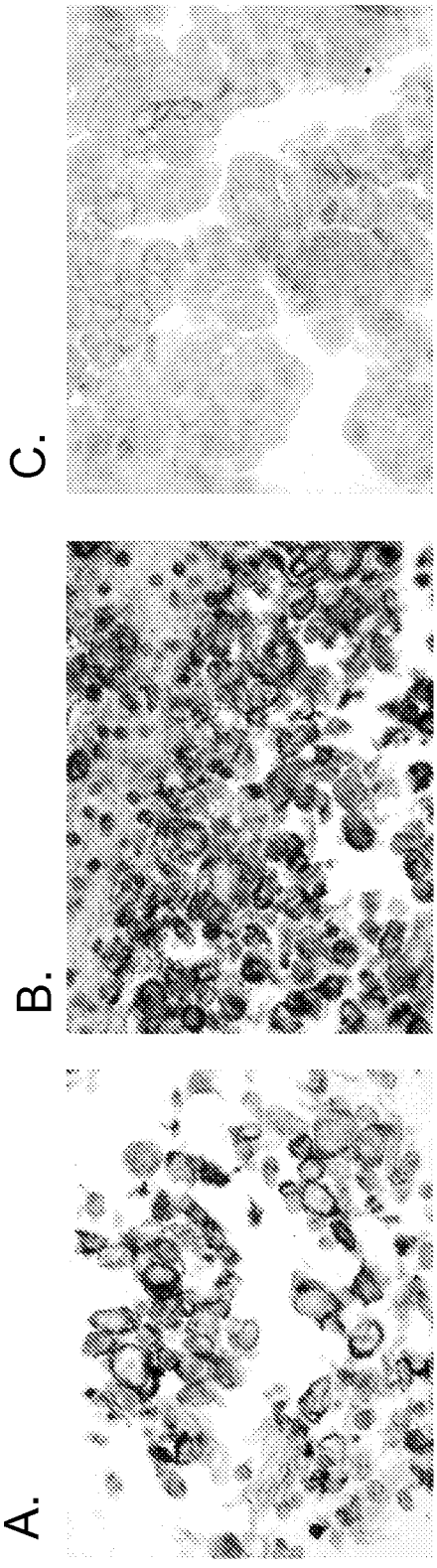


Figure 10

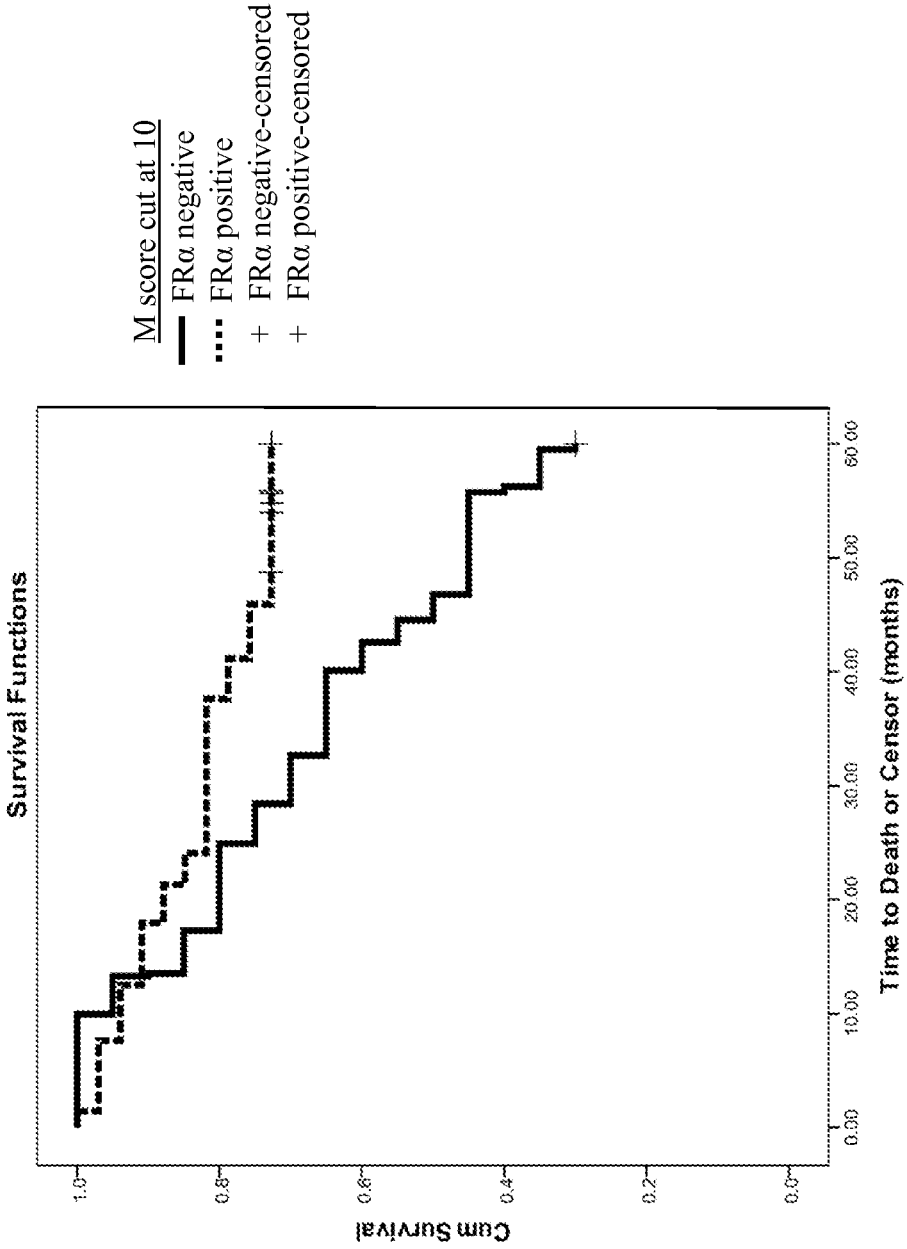

M score cut at 10
FRα negative
FRα positive
FRα negative-censored
FRα positive-censored
Survival Functions
Cum Survival
1.0
0.8
0.6
0.4
0.2
0.0
0.00
10.00
20.00
30.00
40.00
50.00
60.00
Time to Death or Censor (months)

Figure 11

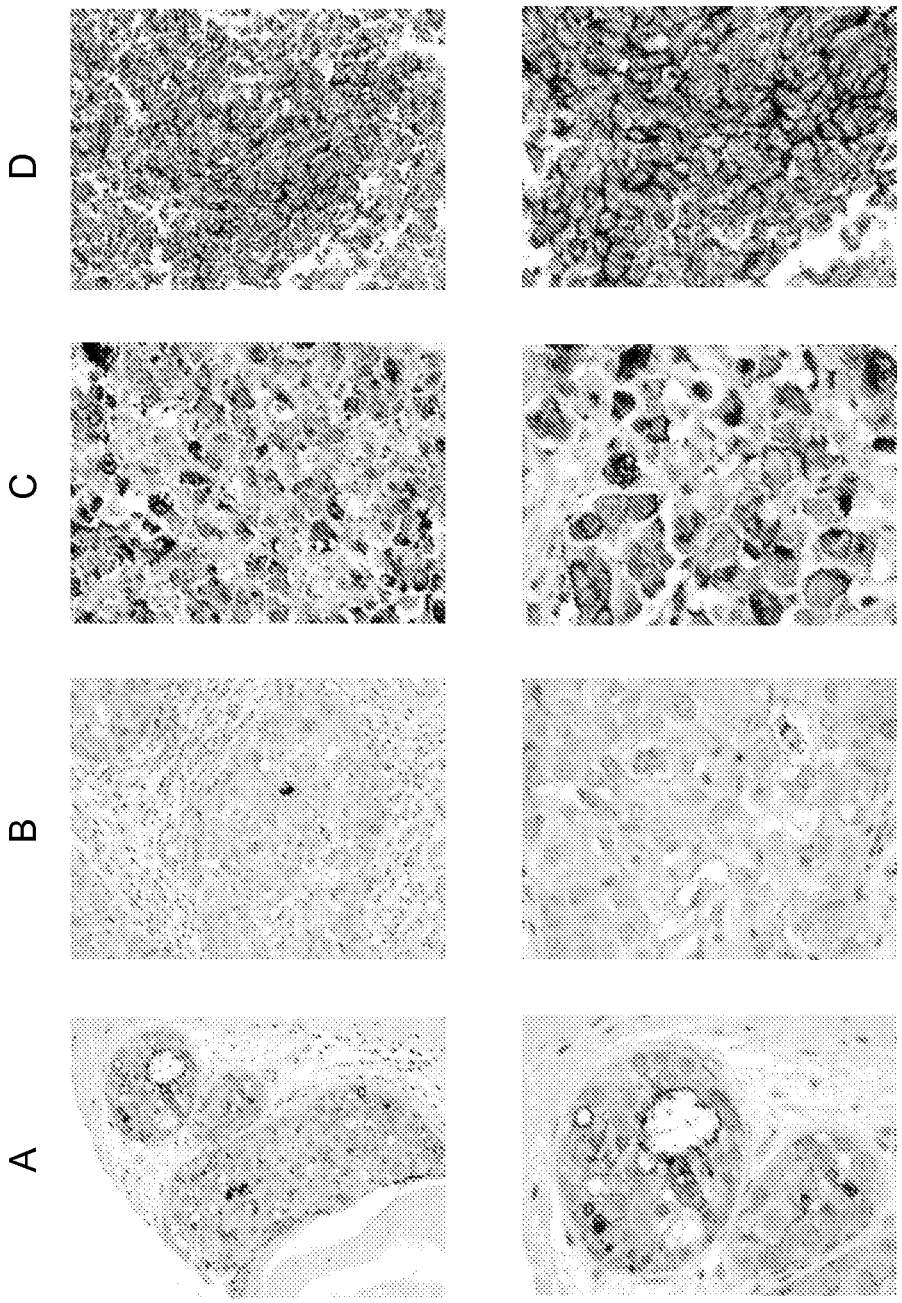


Figure 12

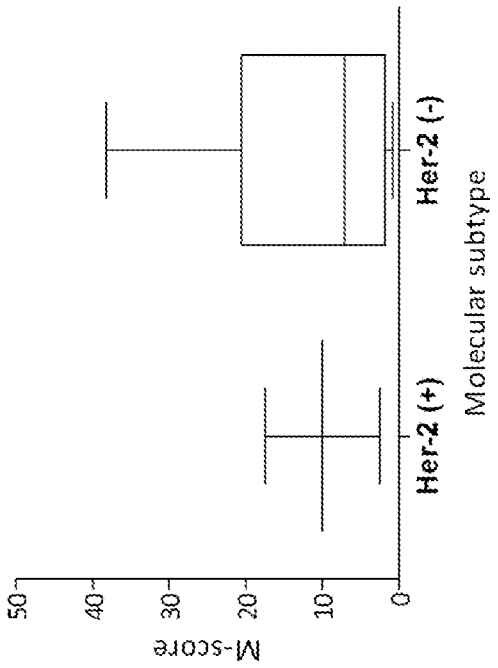


Figure 13

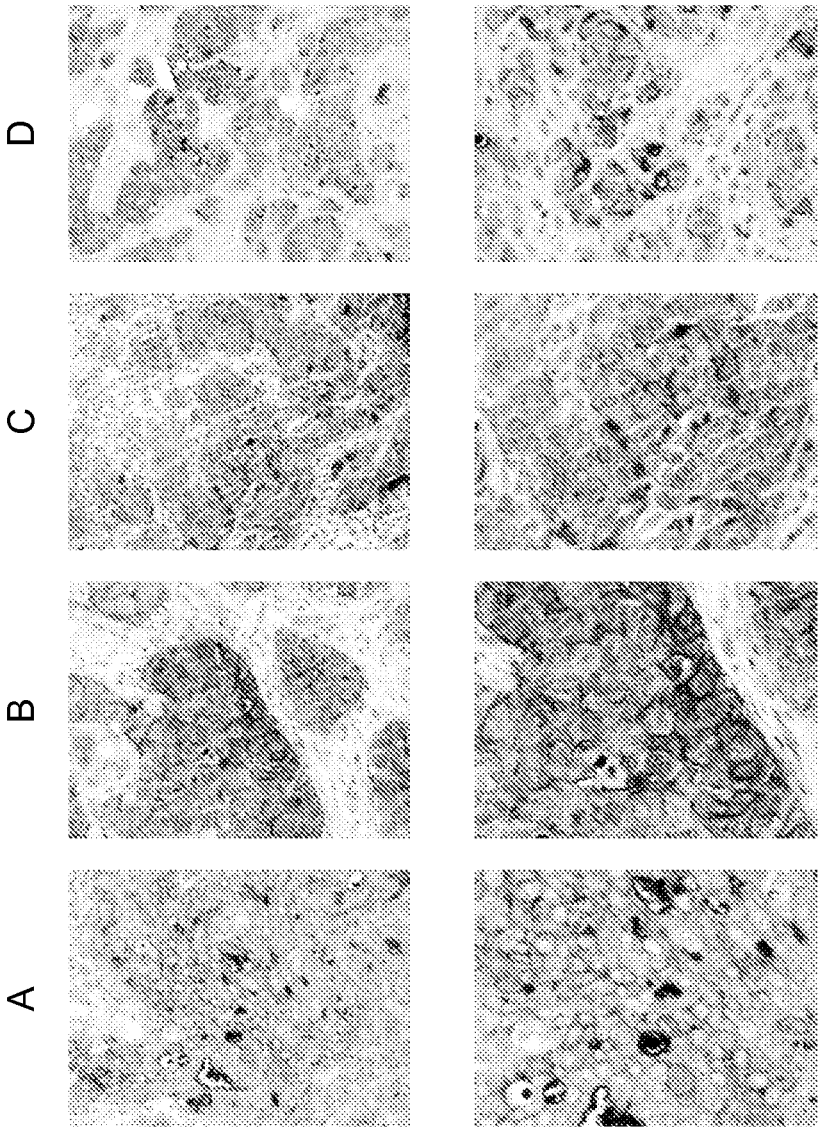


Figure 14

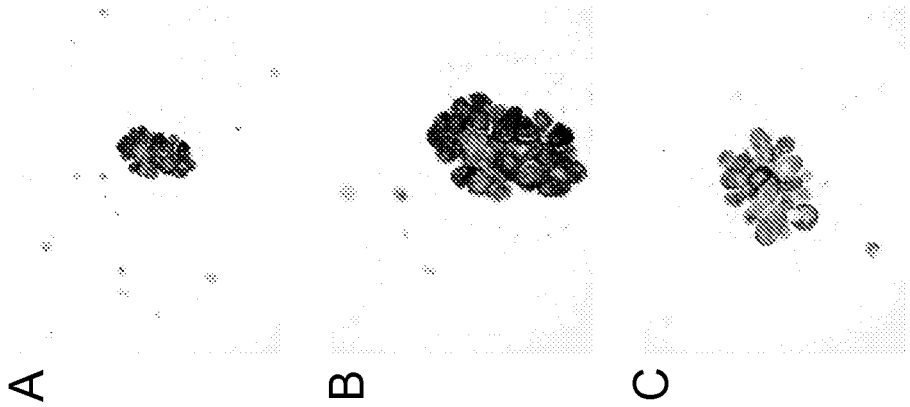


Figure 15

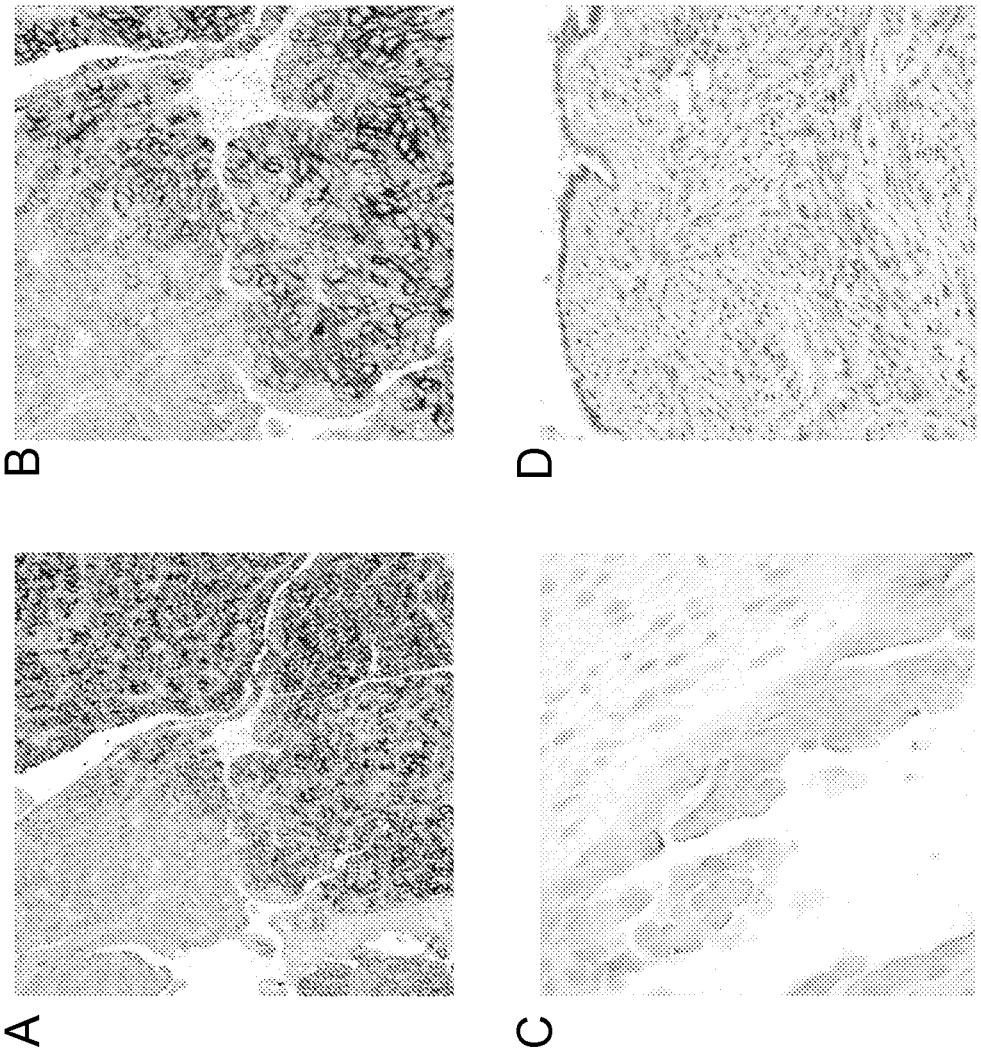


Figure 16

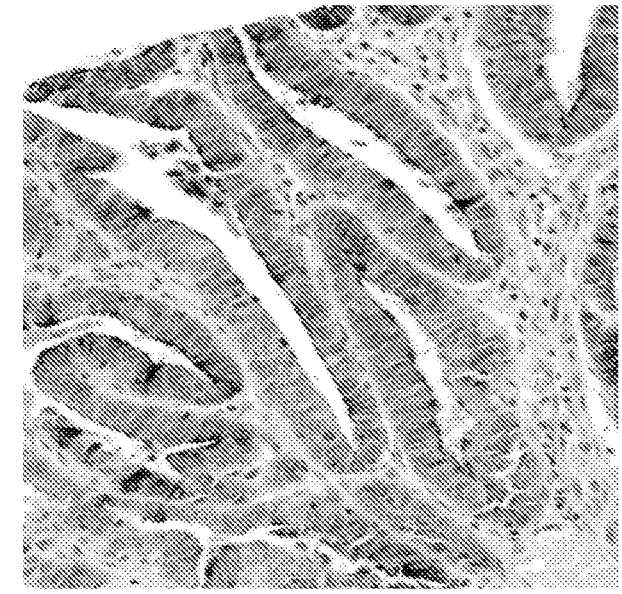




Figure 17

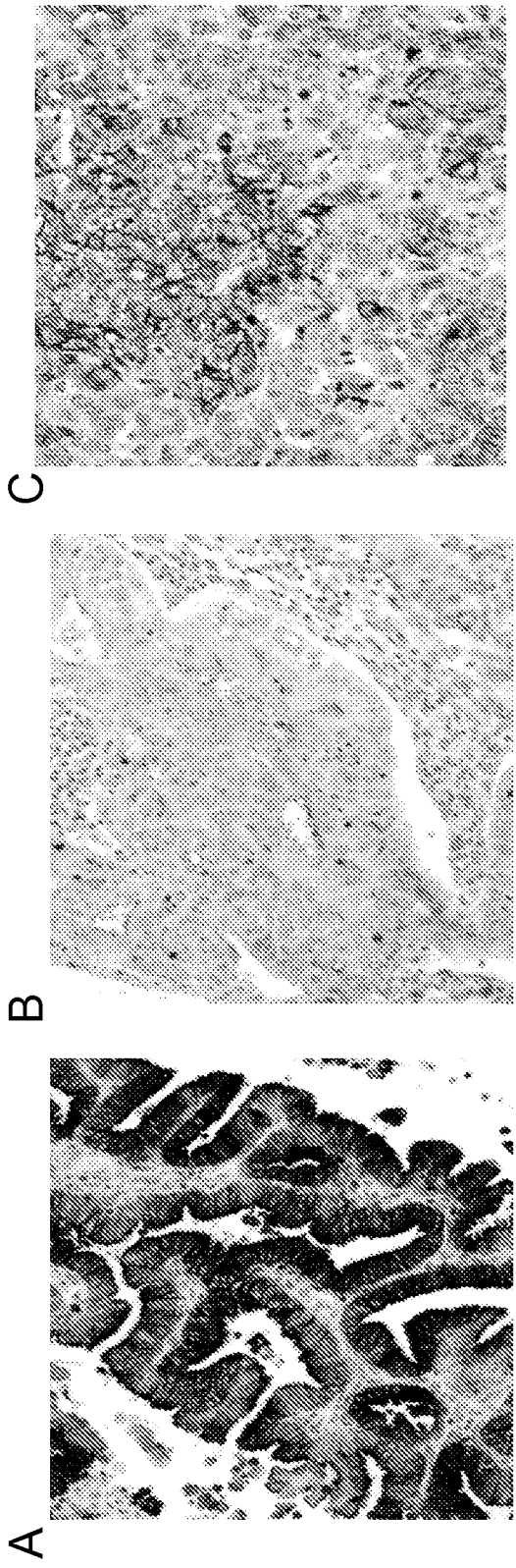


Figure 18

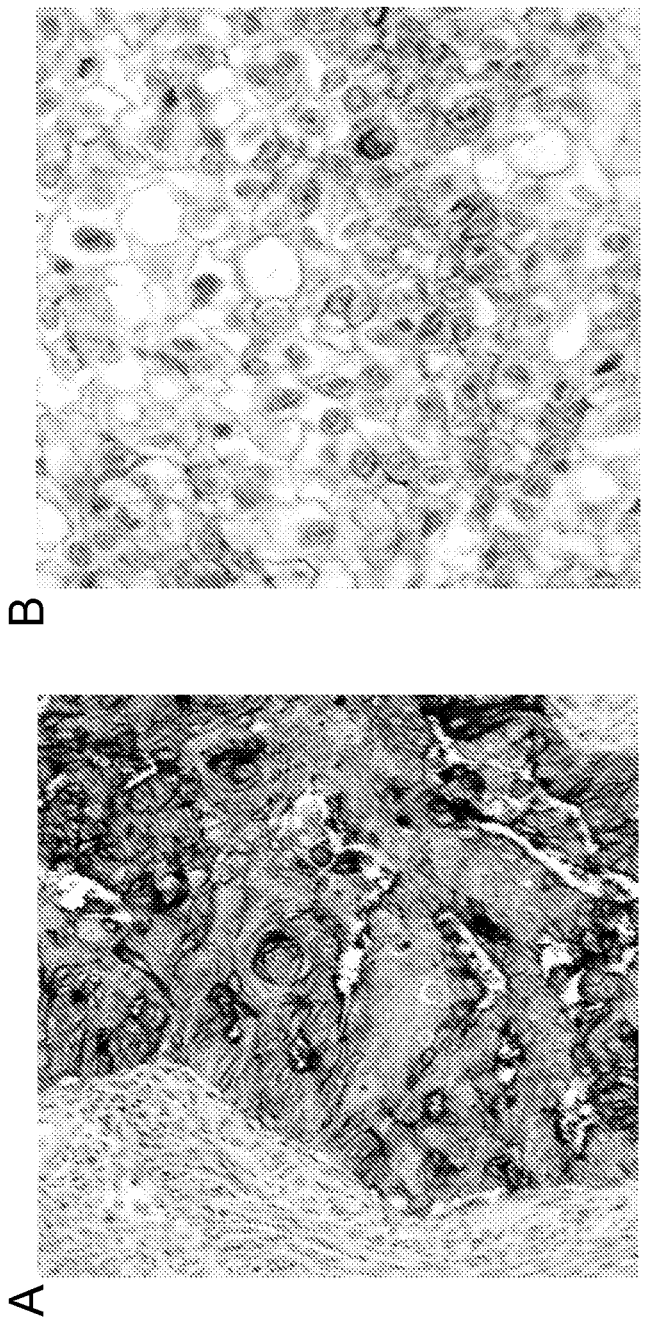


Figure 19

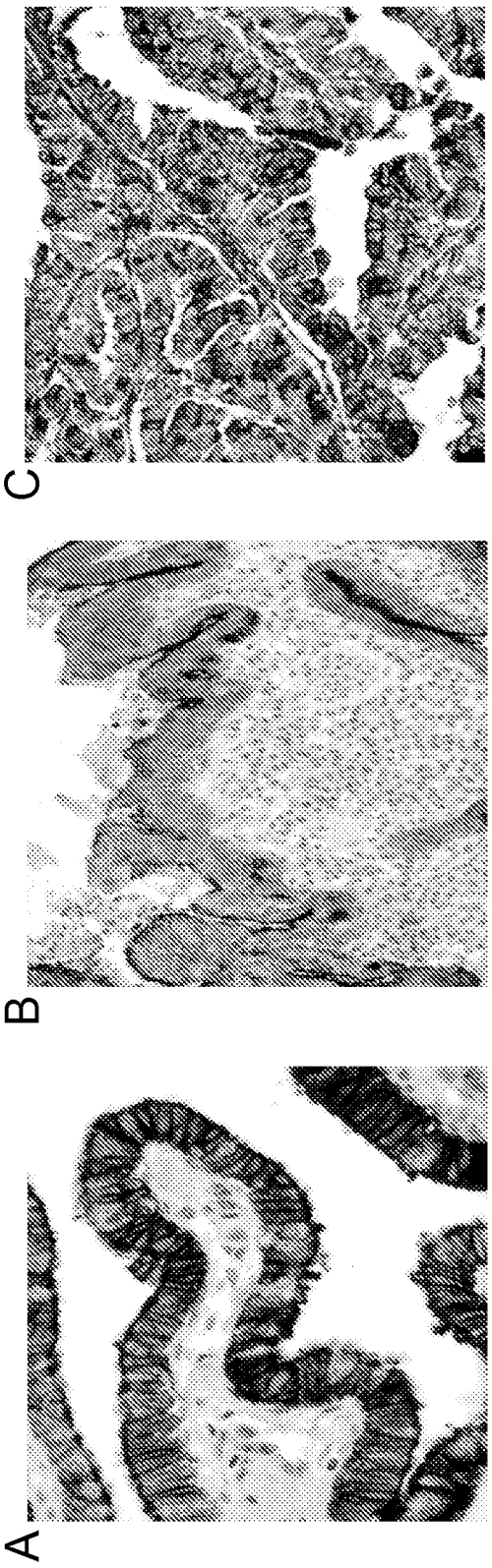


Figure 20

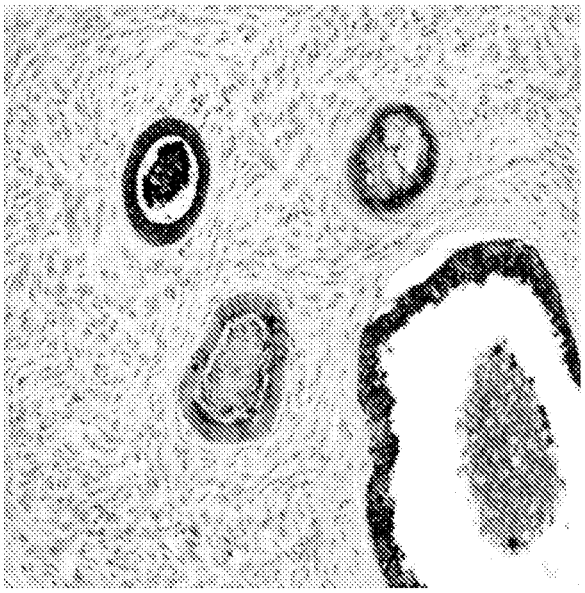


Figure 21