FIGURE 1

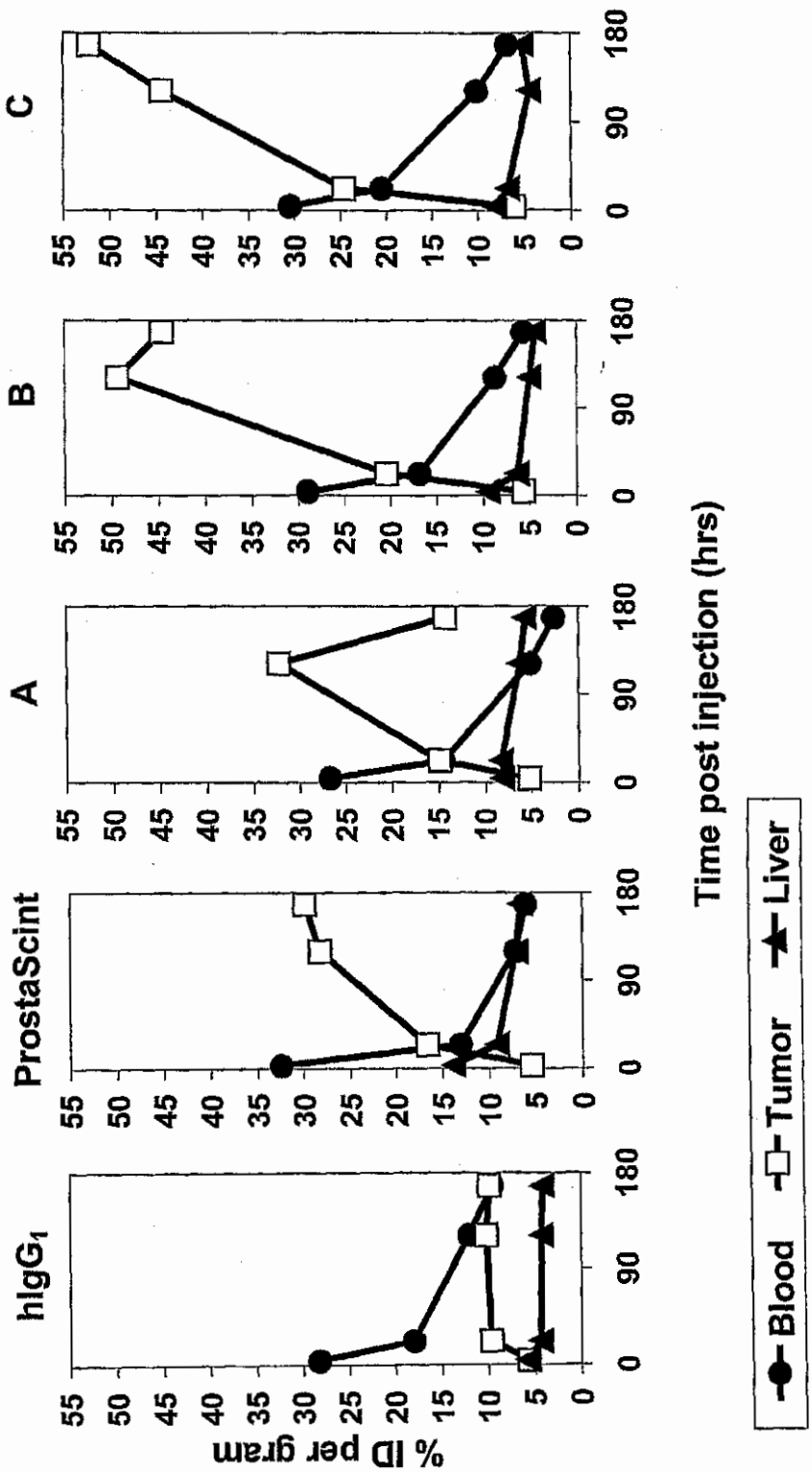

hIgG1
ProstaScint
A
B
C
% ID per gram
Time post injection (hrs)
Blood
Tumor
Liver

FIGURE 2

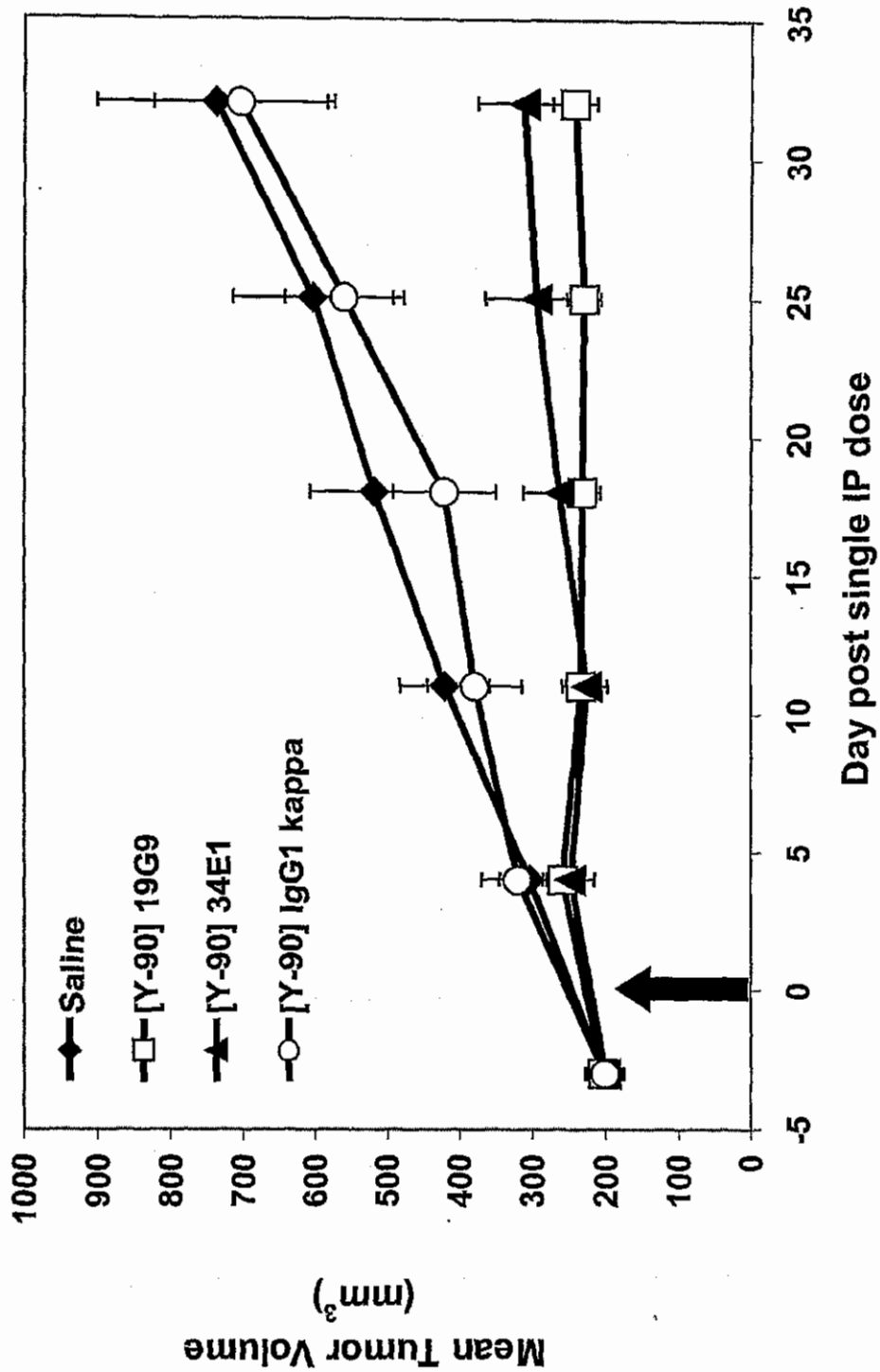

Mean Tumor Volume
(mm3)
1000
900
800
700
600
500
400
300
200
100
0
-5
0
5
10
15
20
25
30
35
Day post single IP dose
Saline
[Y-90] 19G9
[Y-90] 34E1
[Y-90] IgG1 kappa

FIGURE 3

huMAb B Variable Region Sequences

V_L

```
  1 METPAQLLFLLLLWLPDTTGEIVLTQSPGTLSLSPGERATLSCRASQSVS 50
 51 SSYLAWYQQKPGQAPRLLIYGASSRATGIPDRFSGSGSGTDFTLTISRLE 100
101 PEDFAVYYCQQYSSSLTFGGGTKVEIK                         150
```

V_H

```
  1 MEFVLSWVFLVAILKGVQCEVQLVQSGGGLVHPGGSLRLSCAGSGFTFSS 50
 51 YVMHWLRQAPGKGLEWVSVIGTGGVTHYADSVKGRFTISRDNAKNSLYLQ 100
101 MNSLRAEDMAMYYCARWGYYGSGSYENDAFDIWGQGTMVTVSSASTK     150
```

B_3M, V_H (mutations in bold)

```
  1 MEFVLSWVFLVAILKGVQCEVQLVQSGGGLVQPGGSLRLSCAGSGFTFSS 50
 51 YVMHWLRQAPGKGLEWVSVIGTGGVTHYADSVKGRFTISRDNAKNSLYLQ 100
101 MNSLRAEDTAVYYCARWGYYGSGSYENDAFDIWGQGTMVTVSSASTK     150
```

CDR sequences (1,2,and 3)for each variable region are underlined

FIGURE 4

HuMAb C Variable Region Sequences

V_L

```
  1 METPAQLLFLLLLWLPDTTGEIVLTQSPGTLSLSPGERATLSCRASQSVS 50
  51 SSYLAWYQQKPGQAPRLLIYGASSRATGIPDRFSGSGSGTDFTLTISRLE 100
 101 PEDFAVYYCQQYGSSLTFGGGTKVEIK                        150
```

V_H

```
  1 MEFVLSWVFLVAILKGVQCEVQLVQSGGGLVHPGGSLRLSCAGSGFTFSS 50
  51 YVMHWVRQAPGKGLEWVSVIGTGGVTNYADSVKGRFTISRDNAKNSLYLQ 100
 101 MNSLRAEDMAVYYCARWGDWDDAFDIWGQGTMVTVSSASTK          144
```

C_2m, V_H (mutations in bold)

```
  1 MEFVLSWVFLVAILKGVQCEVQLVQSGGGLVQPGGSLRLSCAGSGFTFSS 50
  51 YVMHWVRQAPGKGLEWVSVIGTGGVTNYADSVKGRFTISRDNAKNSLYLQ 100
 101 MNSLRAEDTAVYYCARWGDWDDAFDIWGQGTMVTVSSASTK          144
```

CDR sequences (CDR1,2,3) for each variable region are underlined