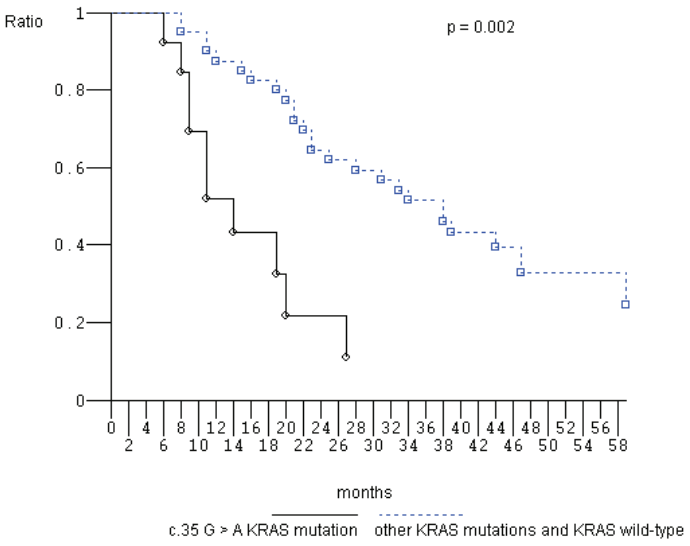
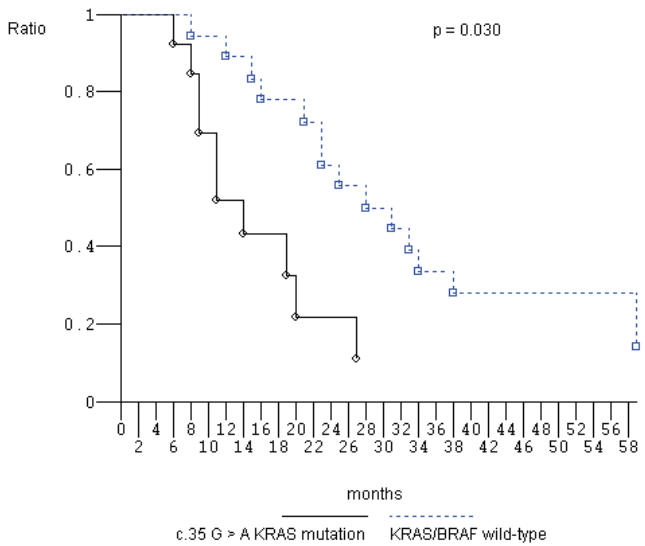
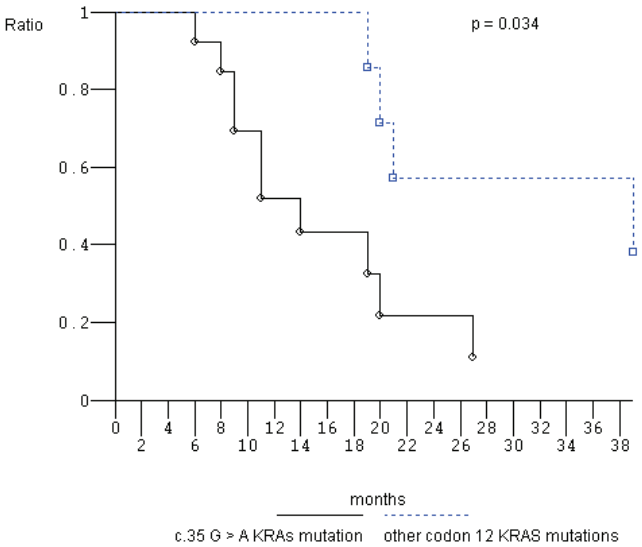

Ratio
p = 0.002
1
0.8
0.6
0.4
0.2
0
0 2 4 6 8 10 12 14 16 18 20 22 24 26 28 30 32 34 36 38 40 42 44 46 48 50 52 54 56 58
months
c.35 G > A KRAS mutation
other KRAS mutations and KRAS wild-type

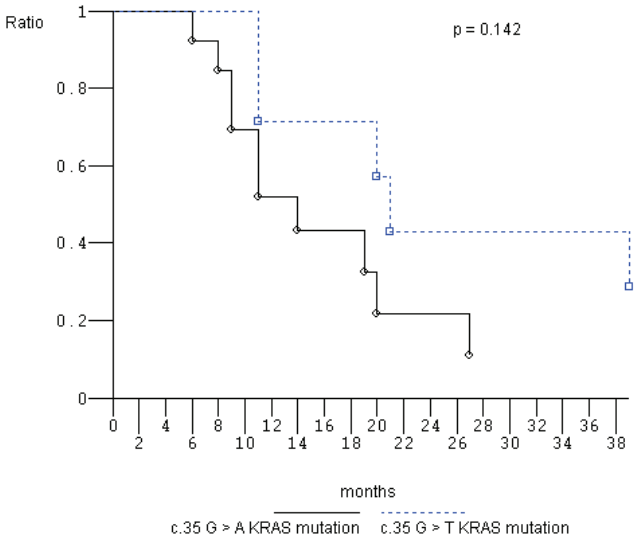
(A)


Ratio
p = 0.030
1
0.8
0.6
0.4
0.2
0
0 2 4 6 8 10 12 14 16 18 20 22 24 26 28 30 32 34 36 38 40 42 44 46 48 50 52 54 56 58
months
c.35 G > A KRAS mutation
KRAS/BRAF wild-type

(B)


Ratio
p = 0.034
1
0.8
0.6
0.4
0.2
0
0 2 4 6 8 10 12 14 16 18 20 22 24 26 28 30 32 34 36 38
months
c.35 G > A KRAs mutation
other codon 12 KRAS mutations

(C)


Ratio
p = 0.142
1
0.8
0.6
0.4
0.2
0
0 2 4 6 8 10 12 14 16 18 20 22 24 26 28 30 32 34 36 38
months
c.35 G > A KRAS mutation
c.35 G > T KRAS mutation

(D)