

Table 1. Distinct prognostic values of the clonal and subclonal events

Variable	Mut vs. Wild	Clonal Mut vs. Wild	Subclonal Mut vs. Wild
<i>KRAS</i>	N = 15	N = 15	NA
(Univariate analysis)	HR = 2.31 (1.22 ~ 4.38) P = 0.01	HR = 2.31 (1.22 ~ 4.38) P = 0.01	
(Multivariate analysis: Age, Gender, TNM stage)	HR = 2.91 (1.52 ~ 5.59) P = 0.001	HR = 2.91 (1.52 ~ 5.59) P = 0.001	NA
<i>Del18p</i>	N = 182	N = 119	N = 63
(Univariate analysis)	HR = 1.50 (1.1 ~ 2.05) P = 0.01	HR = 1.34 (0.949 ~ 1.89) P = 0.10	HR = 1.91 (1.27 ~ 2.86) P = 0.002
(Multivariate analysis: Age, Gender, TNM stage)	HR = 1.35 (0.99 ~ 1.85) P = 0.06	HR = 1.21 (0.856 ~ 1.72) P = 0.28	HR = 1.7 (1.12 ~ 2.58) P = 0.01
<i>Del21q</i>	N = 167	N = 103	N = 64
(Univariate analysis)	HR = 1.42 (1.04 ~ 1.92) P = 0.03	HR = 1.45 (1.03 ~ 2.05) P = 0.03	HR = 1.36 (0.896 ~ 2.05) P = 0.15
(Multivariate analysis: Age, Gender, TNM stage)	HR = 1.29 (0.949 ~ 1.76) P = 0.10	HR = 1.35 (0.954 ~ 1.91) P = 0.09	HR = 1.2 (0.788 ~ 1.82) P = 0.40
<i>Del12p</i>	N = 156	N = 114	N = 42
(Univariate analysis)	HR = 1.35 (0.993 ~ 1.83) P = 0.06	HR = 1.55 (1.12 ~ 2.14) P = 0.009	HR = 0.881 (0.516 ~ 1.5) P = 0.64
(Multivariate analysis: Age, Gender, TNM stage)	HR = 1.19 (0.877 ~ 1.63) P = 0.26	HR = 1.35 (0.972 ~ 1.89) P = 0.07	HR = 0.824 (0.481 ~ 1.41) P = 0.48