OS MM–BO (model 2)

OLD	(N=513)	2.61 (1.139 – 6.0)		0.023 *
SEX	F (N=211)	reference		
	M (N=302)	1.17 (0.784 – 1.8)		0.438
ISS	2 (N=191)	reference		
	1 (N=192)	0.87 (0.546 – 1.4)		0.57
	3 (N=123)	2.07 (1.307 – 3.3)		0.002 **
HB_m_105	(N=513)	1.00 (0.651 – 1.5)		0.987
PLT_m_150	(N=513)	2.07 (1.339 – 3.2)		0.001 **
MMrisk_class_t_CCND2	1q&13– (N=212)	reference		
	1q&13+_pure (N=66)	1.56 (0.854 – 2.9)		0.147
	del 13q only (N=124)	1.18 (0.689 – 2.0)		0.552
	gain 1q only (N=46)	1.12 (0.527 – 2.4)		0.771
	t&1q&13+ (N=62)	4.63 (2.497 – 8.6)		<0.001 ***
FISH_Del_17p	(N=513)	3.09 (1.959 – 4.9)		<0.001 ***
FISH_Del_1p36	(N=513)	1.50 (0.897 – 2.5)		0.121
HyperDiploidy	(N=513)	1.22 (0.787 – 1.9)		0.369
FISH_T_4_14_pure	(N=513)	1.46 (0.729 – 2.9)		0.287
FISH_T_14_16_pure	(N=513)	5.97 (1.863 – 19.1)		0.003 **
FISH_T_14_20_pure	(N=513)	5.17 (1.106 – 24.2)		0.037 *
FISH_T_11_14	(N=513)	1.55 (0.854 – 2.8)		0.15
FISH_T_6_14	(N=513)	0.76 (0.097 – 6.0)		0.798

 # Events: 126; Global p–value (Log–Rank): 0.00000000000016678
AIC: 1210.21; Concordance Index: 0.76

0.1   0.2   0.5   1   2   5   10   20