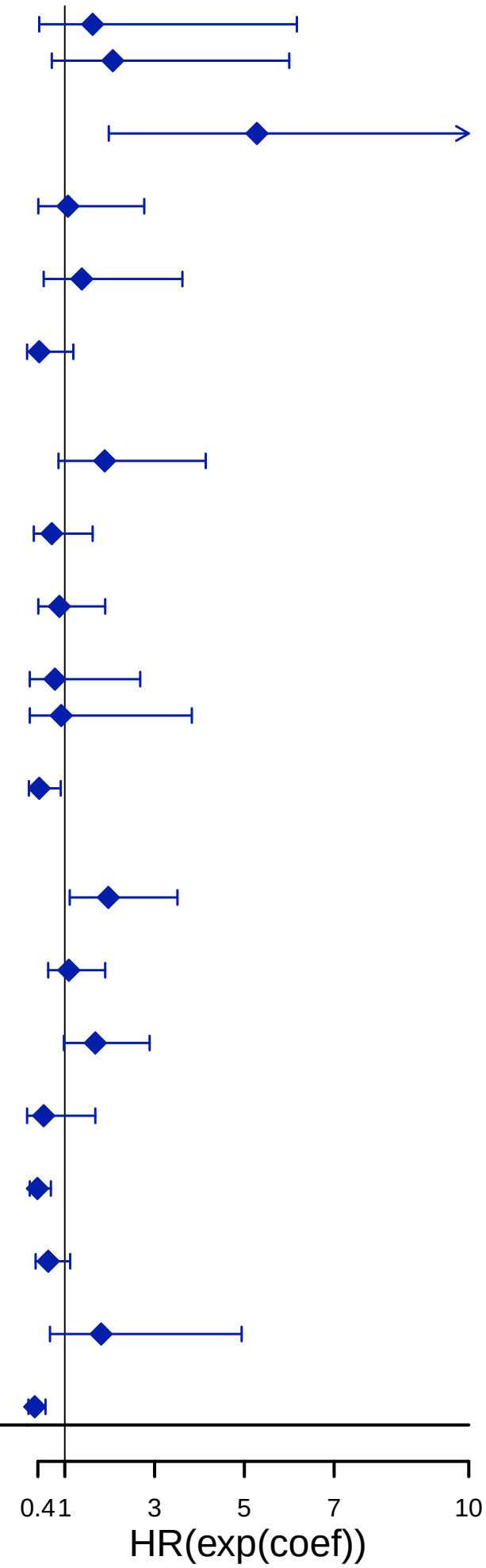
Multi−variables forest plot

Features	HR(95%CI)	p−value
GSE14333		
Location(Left/Rectum/Right)		
Rectum versus Left	1.62 [0.43, 6.17]	0.479
Right versus Left	2.07 [0.71, 6.00]	0.181
Duck Stage		
StageIII versus StageII	5.28 [1.98, 14.09]	0.001
Age		
Age>=70 versus Age<70	1.07 [0.41, 2.77]	0.891
Gender		
Male versus Female	1.38 [0.53, 3.62]	0.513
Signature Label		
Low−risk versus High−risk	0.43 [0.16, 1.19]	0.104
GSE17536		
AJCC Stage		
StageIII versus StageII	1.89 [0.86, 4.14]	0.114
Age		
Age>=70 versus Age<70	0.71 [0.31, 1.62]	0.422
Gender		
Male versus Female	0.88 [0.41, 1.90]	0.749
Histologic Grade		
2−MD(Moderately differentiated) versus 1−WD(Well differentiated)	0.78 [0.22, 2.68]	0.689
3−PD(Poorly differentiated) versus 1−WD(Well differentiated)	0.92 [0.22, 3.83]	0.912
Signature Label		
Low−risk versus High−risk	0.43 [0.20, 0.91]	0.028
GSE39582		
TNM Stage		
StageIII versus StageII	1.97 [1.11, 3.51]	0.021
Age		
Age>=70 versus Age<70	1.09 [0.63, 1.90]	0.755
Gender		
Male versus Female	1.68 [0.98, 2.89]	0.060
BRAF Mutation		
BRAF Mutation versus Wild−type	0.53 [0.16, 1.68]	0.279
KRAS Mutation		
KRAS Mutation versus Wild−type	0.39 [0.22, 0.69]	0.001
TP53 Mutation		
TP53 versus Wild−type	0.63 [0.35, 1.12]	0.113
MMR status		
pMMR versus Wild−dMMR	1.81 [0.67, 4.94]	0.243
Signature Label		
Low−risk versus High−risk	0.33 [0.19, 0.57]	<0.001