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Supplementary Material

Identification of Immune-Related LncRNA for Predicting Survival in Skin Cutaneous Melanoma

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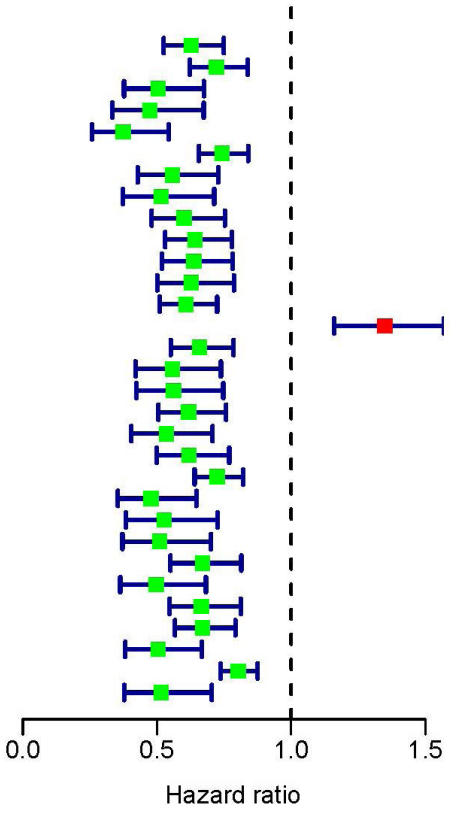
	pvalue	Hazard ratio
HLA-DQB1-AS1	<0.001	0.628(0.525-0.749)
PCED1B-AS1	<0.001	0.722(0.622-0.838)
LINC01943	<0.001	0.504(0.377-0.675)
AC011899.2	<0.001	0.474(0.333-0.674)
C5orf56	<0.001	0.374(0.258-0.543)
PSMB8-AS1	<0.001	0.742(0.656-0.840)
AL133371.2	<0.001	0.558(0.428-0.728)
MMP25-AS1	<0.001	0.515(0.372-0.713)
AC242842.1	<0.001	0.601(0.480-0.753)
AC093726.1	<0.001	0.642(0.530-0.779)
AC243960.1	<0.001	0.637(0.519-0.782)
WAC-AS1	<0.001	0.628(0.501-0.787)
USP30-AS1	<0.001	0.608(0.510-0.724)
MIR205HG	<0.001	1.349(1.160-1.569)
MIR155HG	<0.001	0.658(0.551-0.785)
MIAT	<0.001	0.557(0.420-0.738)
U62317.1	<0.001	0.562(0.423-0.747)
AC083799.1	<0.001	0.618(0.505-0.757)
ZEB1-AS1	<0.001	0.535(0.404-0.707)
AL365361.1	<0.001	0.619(0.498-0.769)
LINC01871	<0.001	0.725(0.640-0.821)
LINC00324	<0.001	0.478(0.354-0.647)
NCK1-DT	<0.001	0.527(0.383-0.726)
AC098613.1	<0.001	0.510(0.371-0.701)
EBLN3P	<0.001	0.669(0.550-0.814)
VIM-AS1	<0.001	0.497(0.363-0.682)
AC136475.3	<0.001	0.666(0.547-0.812)
LINC02446	<0.001	0.670(0.567-0.792)
AC015911.3	<0.001	0.505(0.382-0.667)
HCP5	<0.001	0.803(0.737-0.875)
AL590764.1	<0.001	0.516(0.378-0.704)



Supplementary Fig. S1. A Forest plot of 31 candidates immune-related lncRNAs selected by univariate Cox regression analysis associated with SKCM survival.

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