



Supplementary Material

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

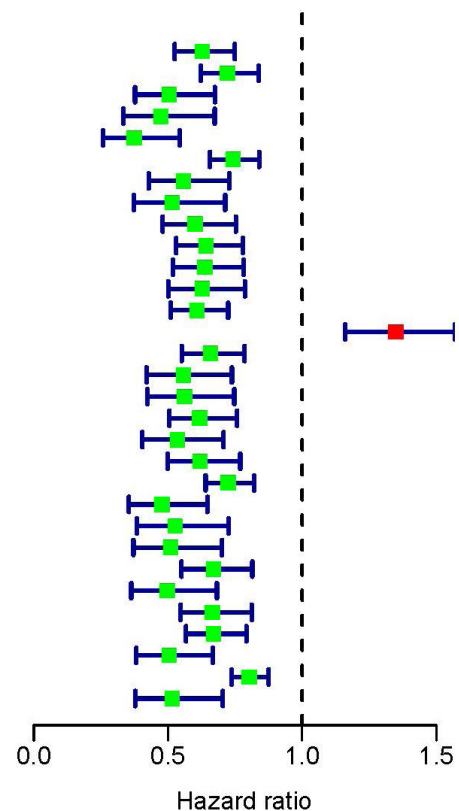
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| | pvalue | Hazard ratio |
|--------------|--------|--------------------|
| HLA-DQB1-AS1 | <0.001 | 0.628(0.525-0.749) |
| PCED1B-AS1 | <0.001 | 0.722(0.622-0.838) |
| LINC01943 | <0.001 | 0.504(0.377-0.675) |
| AC011899.2 | <0.001 | 0.474(0.333-0.674) |
| 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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