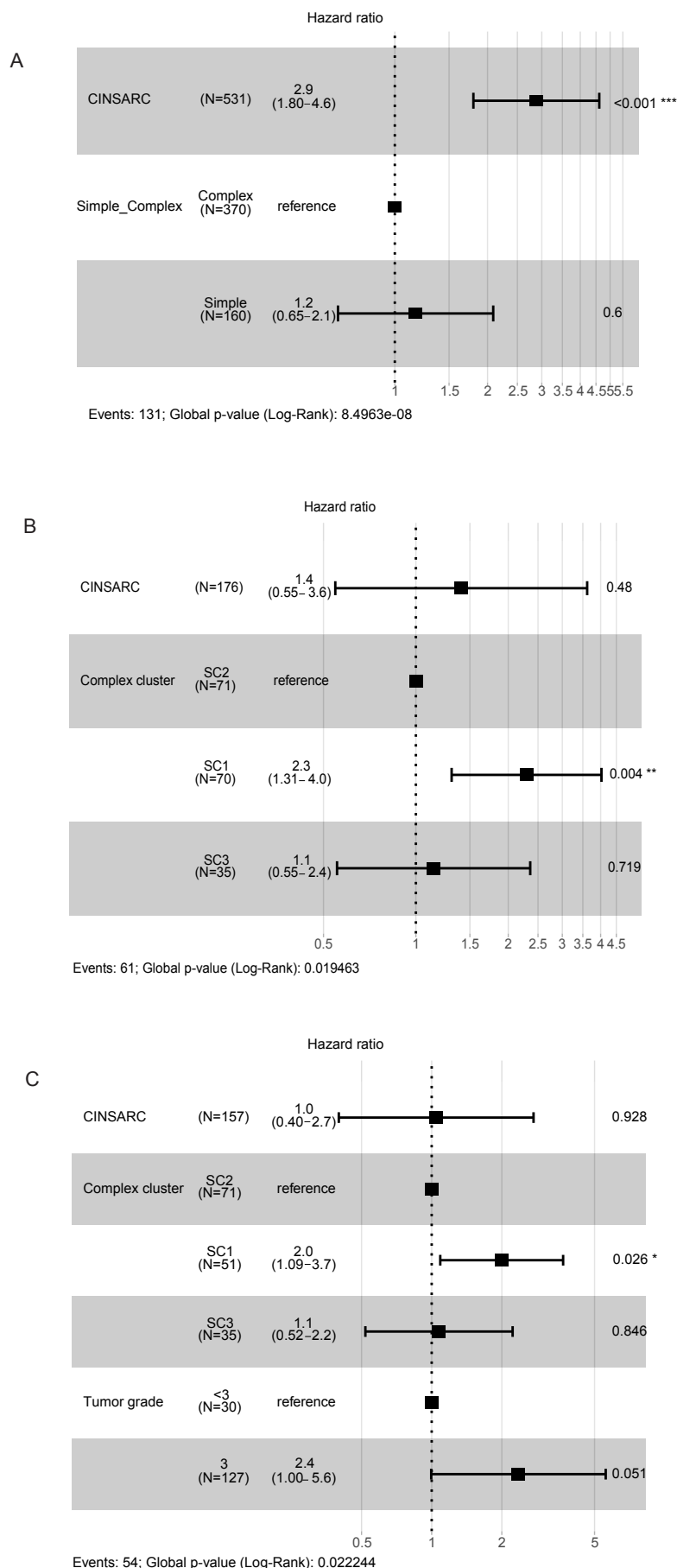




Supplementary Figure S8: Survival analysis with CINSARC enrichment

Forest plots displaying the output from multivariate cox regression analysis of malignant and intermediate tumors, including the hazard ratio and 95% confidence interval. The CINSARC enrichment score was used as a continuous variable and analyzed with (A), tumors having simple or complex genomes (n = 531) (B), the transcriptomic subgroups; Complex SC1-3 (n = 176) or (C), transcriptomic subgroups; Complex SC1-3 and tumor grade (n = 157).