Multivariate Cox Regression: Full Model

Age (per year) — Significant (p<0.05) — 1.03, p=0.000

Male (vs Female) — Not significant — 1.08, p=0.545

TP53 mutation — 2.96, p=0.000

RUNX1 mutation — 1.13, p=0.518

ASXL1 mutation — 1.21, p=0.331
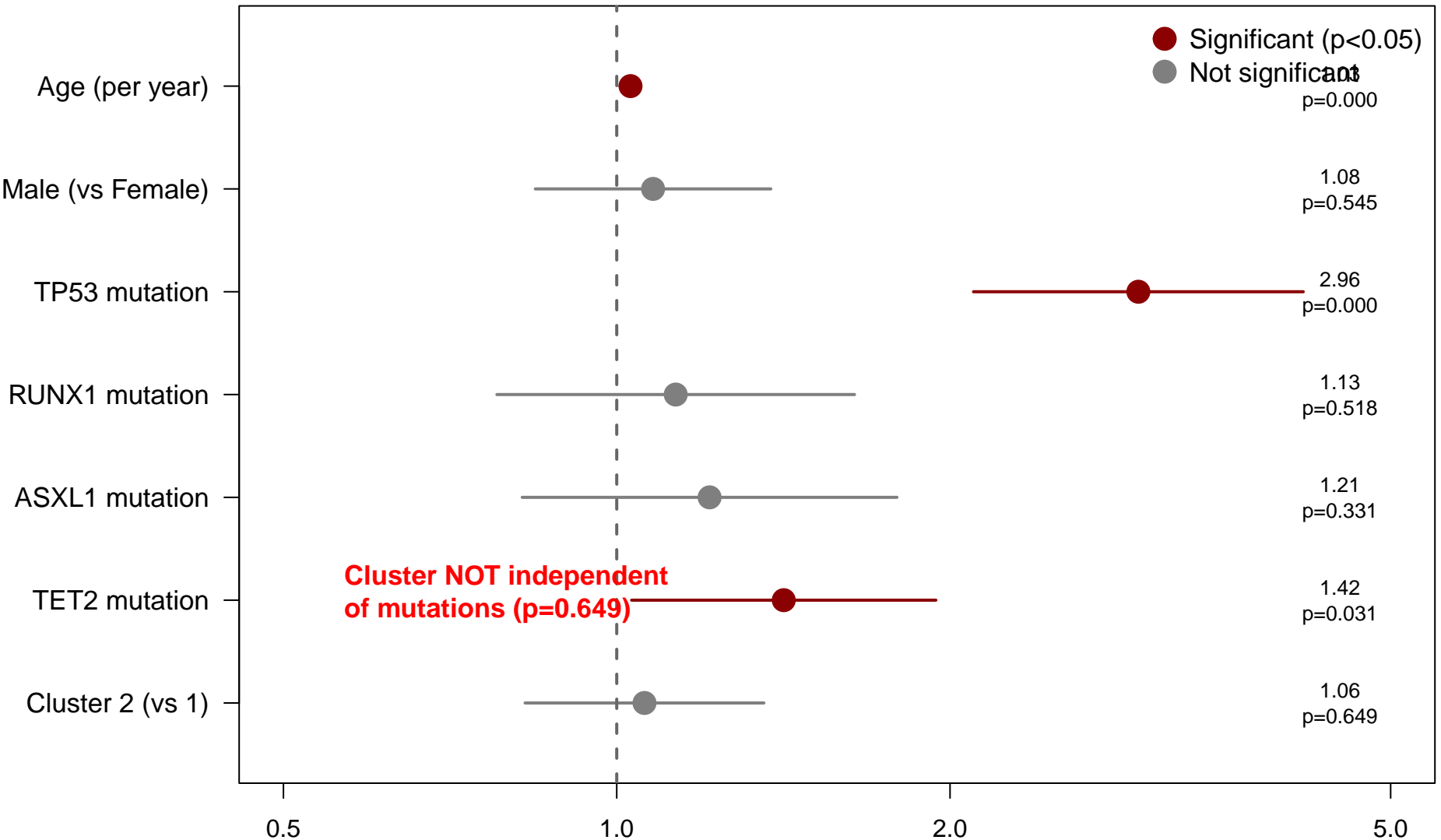
TET2 mutation — **Cluster NOT independent of mutations (p=0.649)** — 1.42, p=0.031

Cluster 2 (vs 1) — 1.06, p=0.649

Hazard Ratio (95% CI)

n=459, 282 events | Concordance=0.685