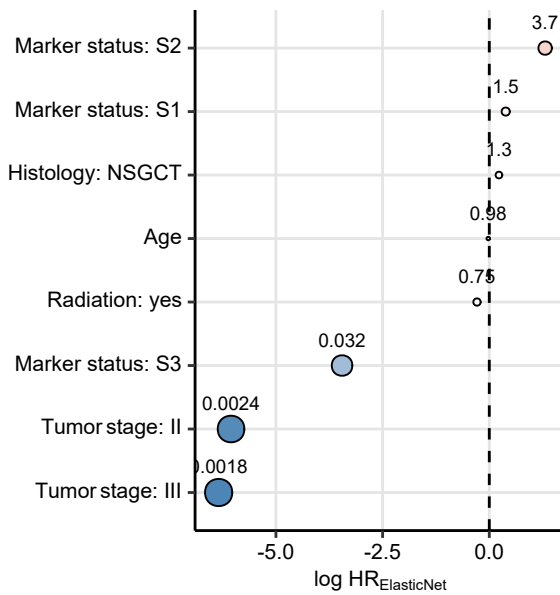
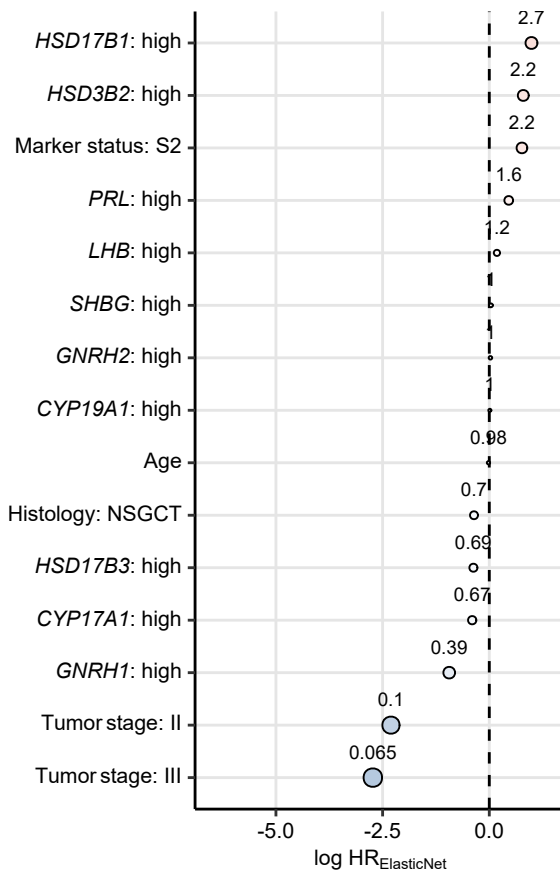


**A****Clinical model**

total: n = 107, events: n = 30

**B****Clinical + genes**

total: n = 107, events: n = 30

**C****Clinical + hormonal subsets**

total: n = 107, events: n = 30

