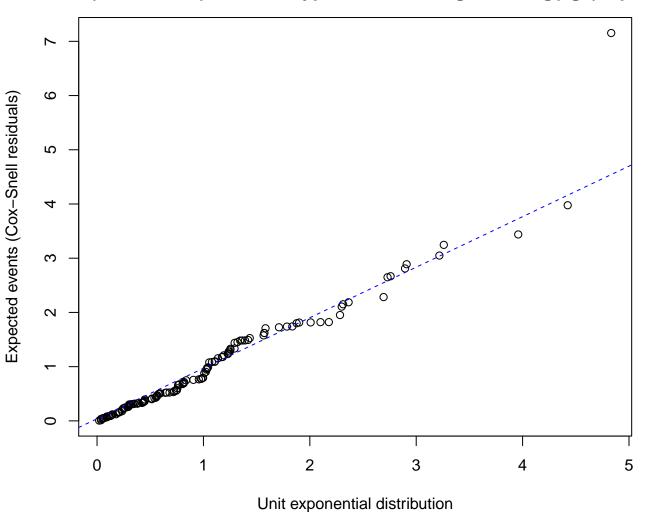
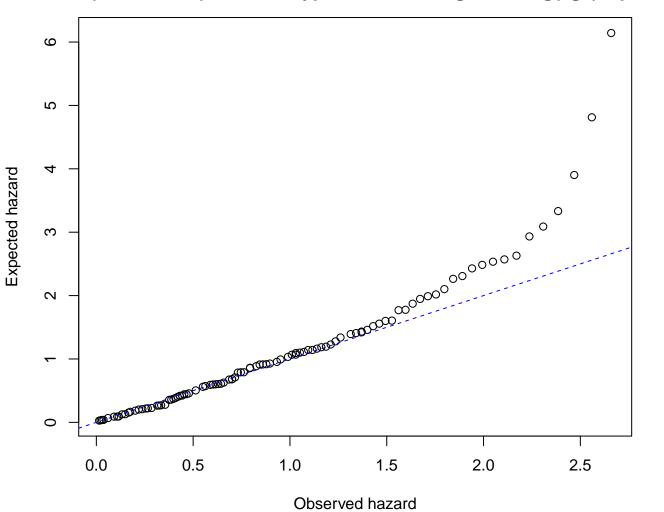
Quantile–quantile plot.
Unit exponential distribution vs. expected events (Cox–Snell residuals).
Should follow line through origin at 45 degrees (blue) if well fit.

Complete model:
Surv(time, status) ~ trt \* celltype + karno + diagtime + log(age) + prior



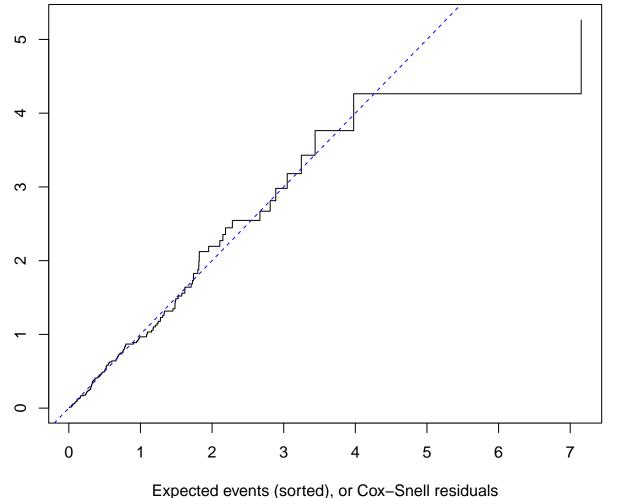
Observed vs. expected hazard. Should follow line through origin at 45 degrees (blue) if well fit.

Complete model:
Surv(time, status) ~ trt \* celltype + karno + diagtime + log(age) + prior



Expected events vs. hazard based on sorted expected events or Cox–Snell residuals vs. cumulative hazard of these residuals. Should follow line through origin at 45 degrees (blue) if well fit.

# Complete model: Surv(time, status) ~ trt \* celltype + karno + diagtime + log(age) + prior

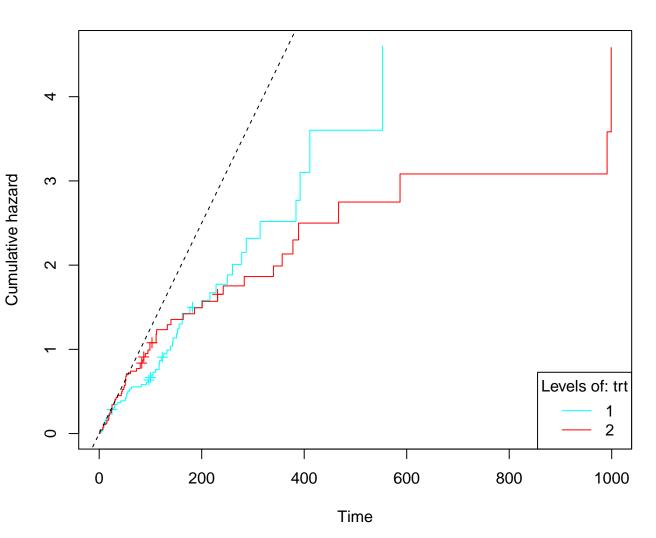


Time vs. hazard, per predictor.

If hazards proportional then curves should be constant multiples of a baseline.

Reference (black) line is 45 degrees.

**Predictor: trt** 

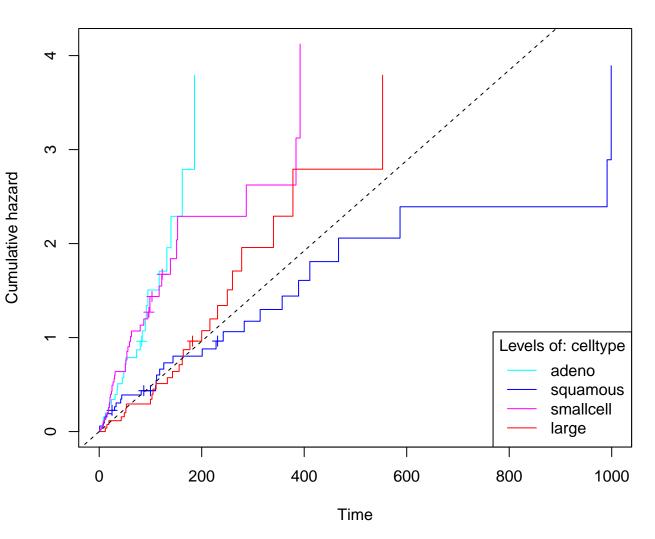


Time vs. hazard, per predictor.

If hazards proportional then curves should be constant multiples of a baseline.

Reference (black) line is 45 degrees.

## **Predictor: celltype**

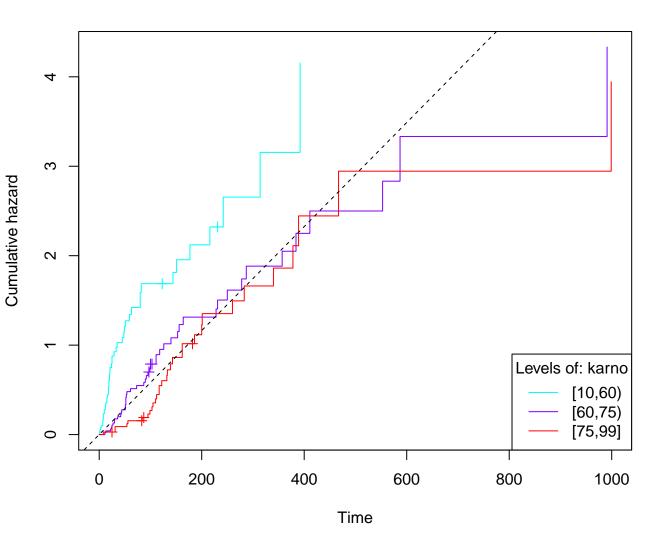


Time vs. hazard, per predictor.

If hazards proportional then curves should be constant multiples of a baseline.

Reference (black) line is 45 degrees.

**Predictor: karno** 

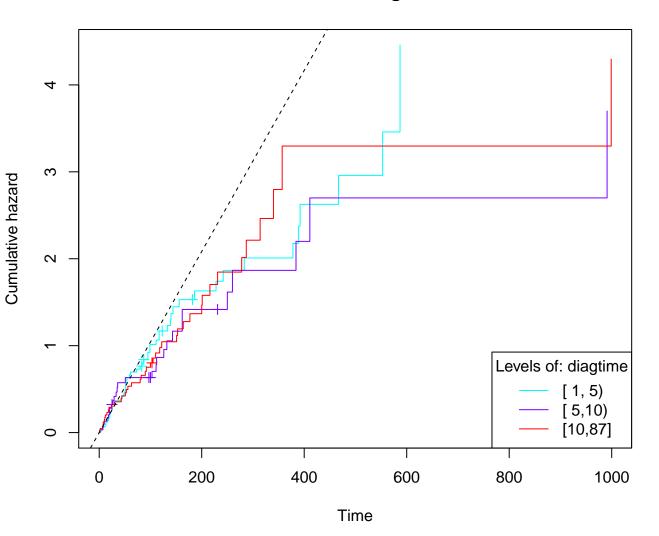


Time vs. hazard, per predictor.

If hazards proportional then curves should be constant multiples of a baseline.

Reference (black) line is 45 degrees.

#### **Predictor: diagtime**

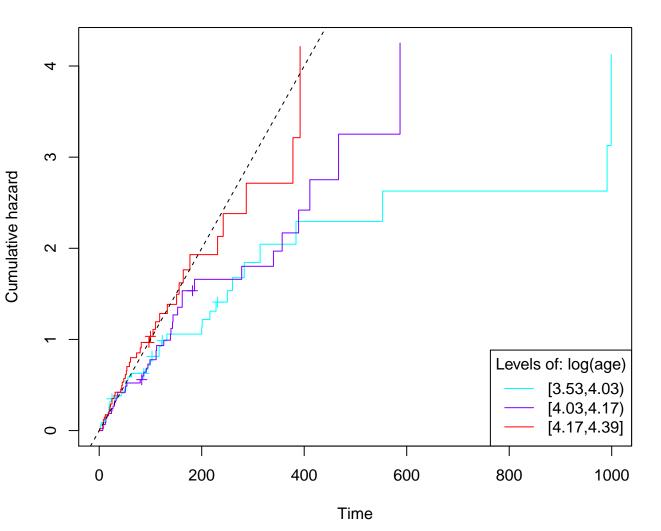


Time vs. hazard, per predictor.

If hazards proportional then curves should be constant multiples of a baseline.

Reference (black) line is 45 degrees.

**Predictor:** log(age)

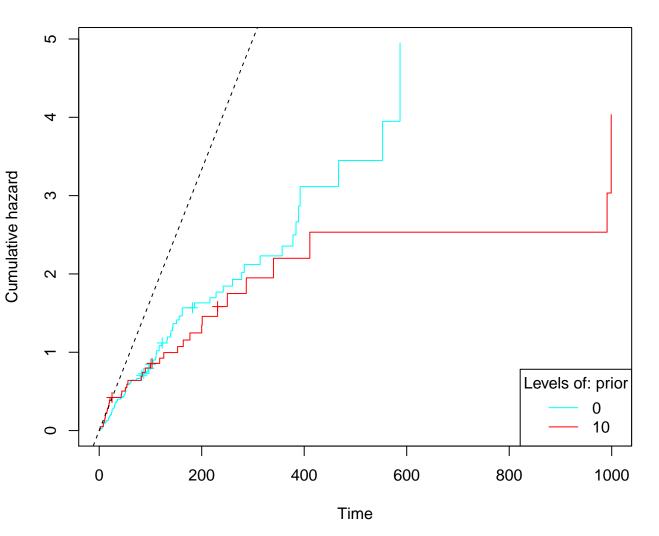


Time vs. hazard, per predictor.

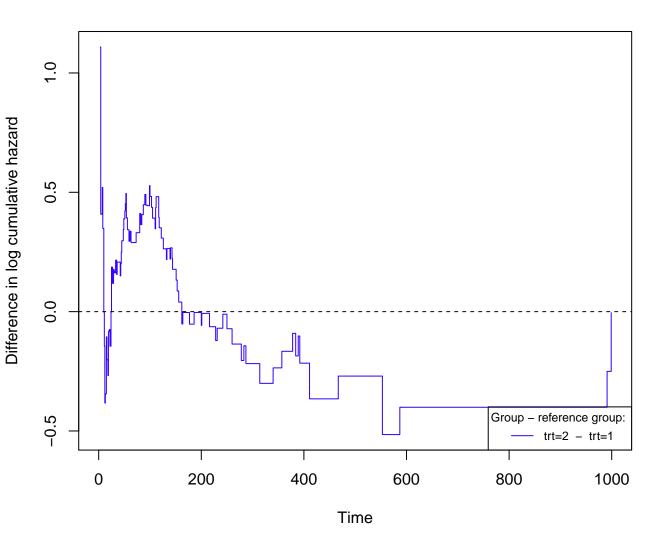
If hazards proportional then curves should be constant multiples of a baseline.

Reference (black) line is 45 degrees.

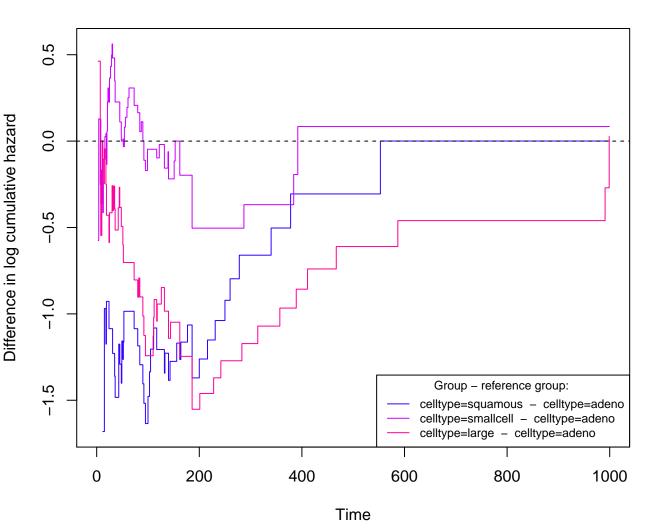




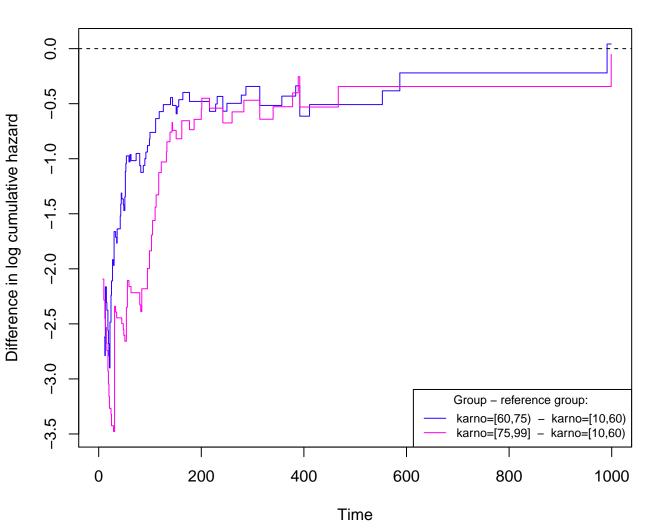
Reference: trt = 1



### Reference: celltype = adeno

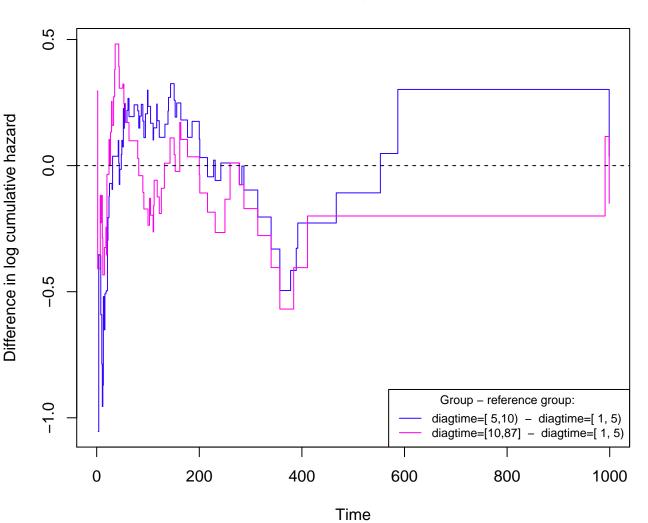


Reference: karno = [10,60)

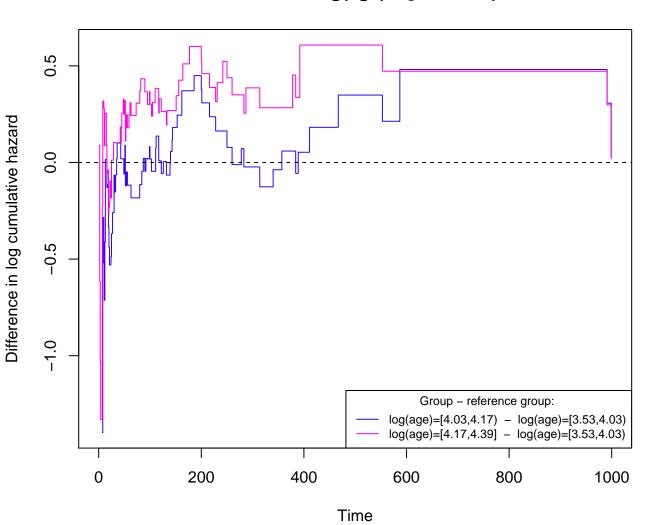


Time vs. difference in log hazards, per predictor.
Should be constant over time.
If >0 (black line) shows survival advantage for reference group.

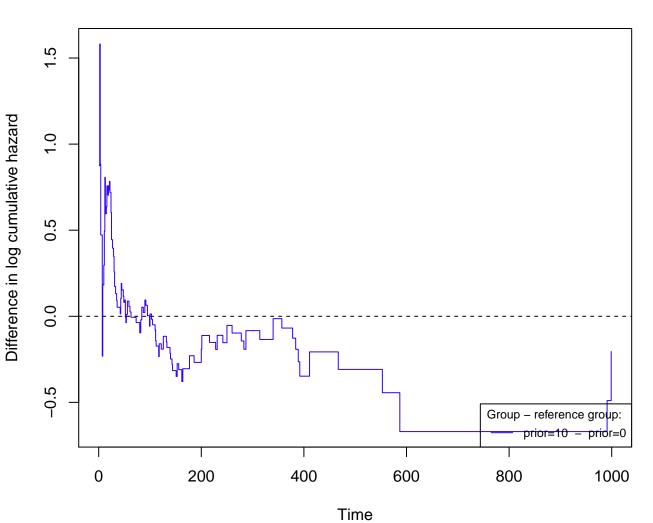
Reference: diagtime = [1, 5)



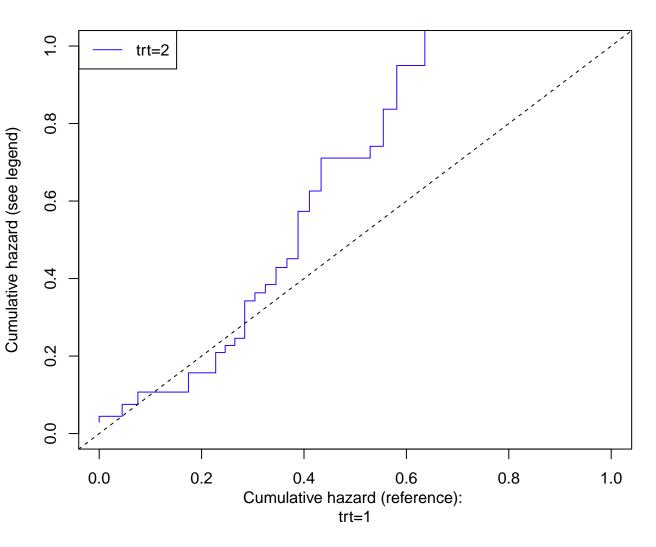
### Reference: log(age) = [3.53,4.03)



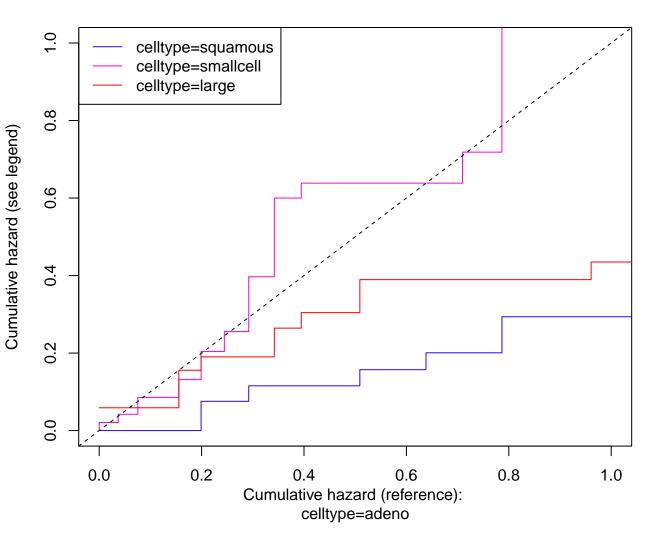
Reference: prior = 0



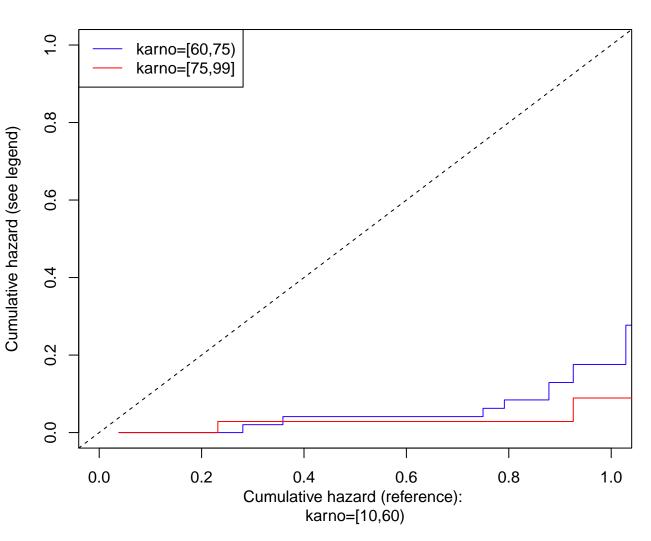




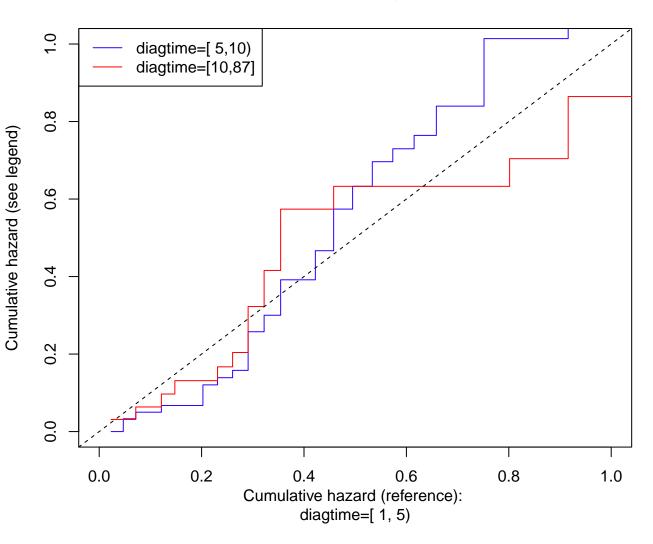
## Predictor: celltype



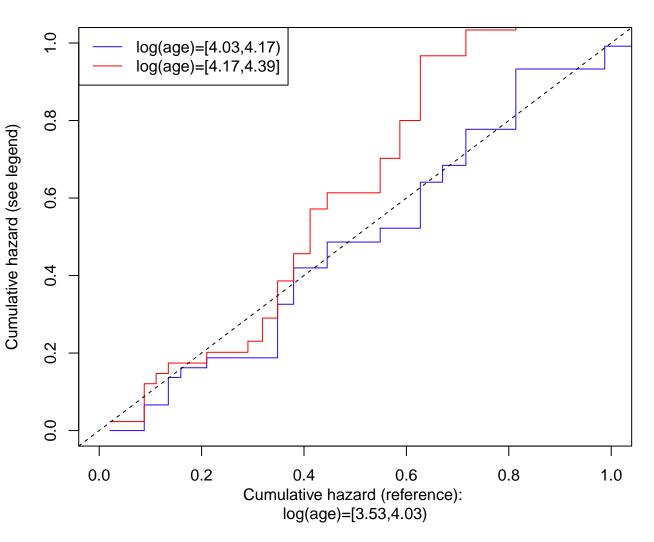
#### **Predictor: karno**



## Predictor: diagtime



## **Predictor:** log(age)



## **Predictor: prior**

