Plots that could not fit in the report
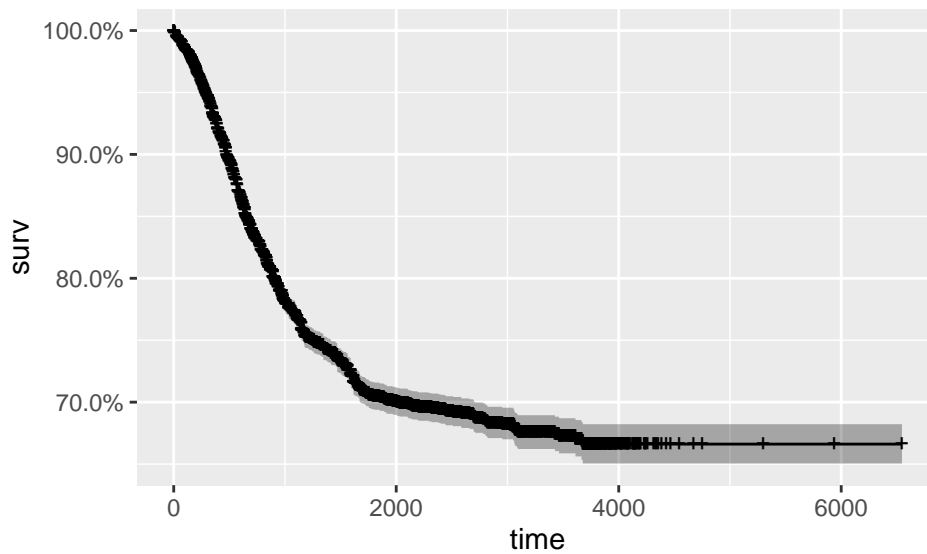
```
#generating KM plots
msurv<-survfit(Surv(days_to_last_follow_up,vital_status)~1, data = data)

autoplot(msurv)
```
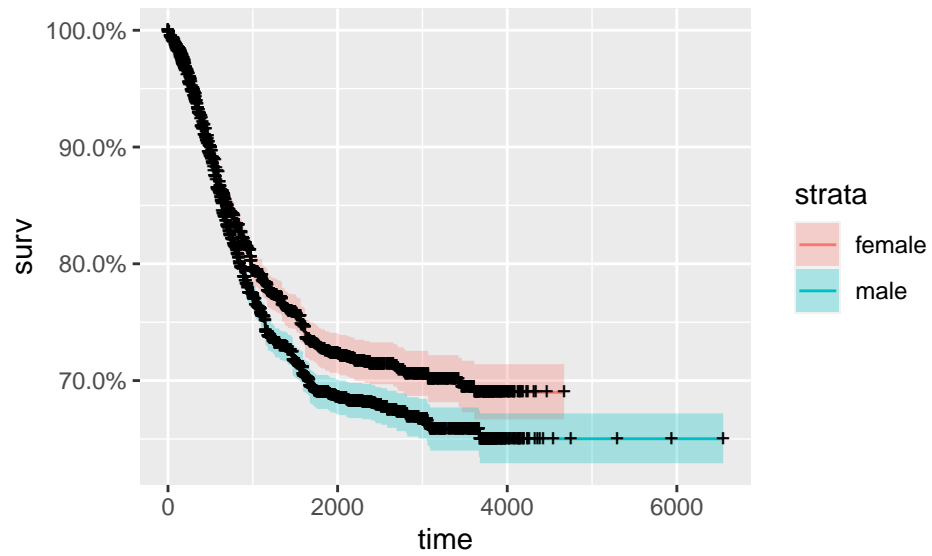


```
###############
msurv2<-survfit(Surv(days_to_last_follow_up,vital_status)~gender, data = data)

autoplot(msurv2)
```
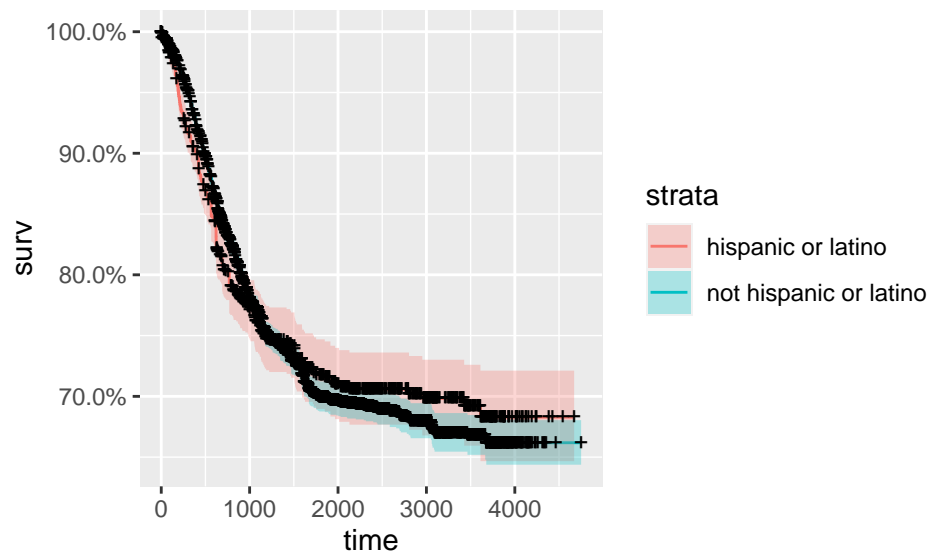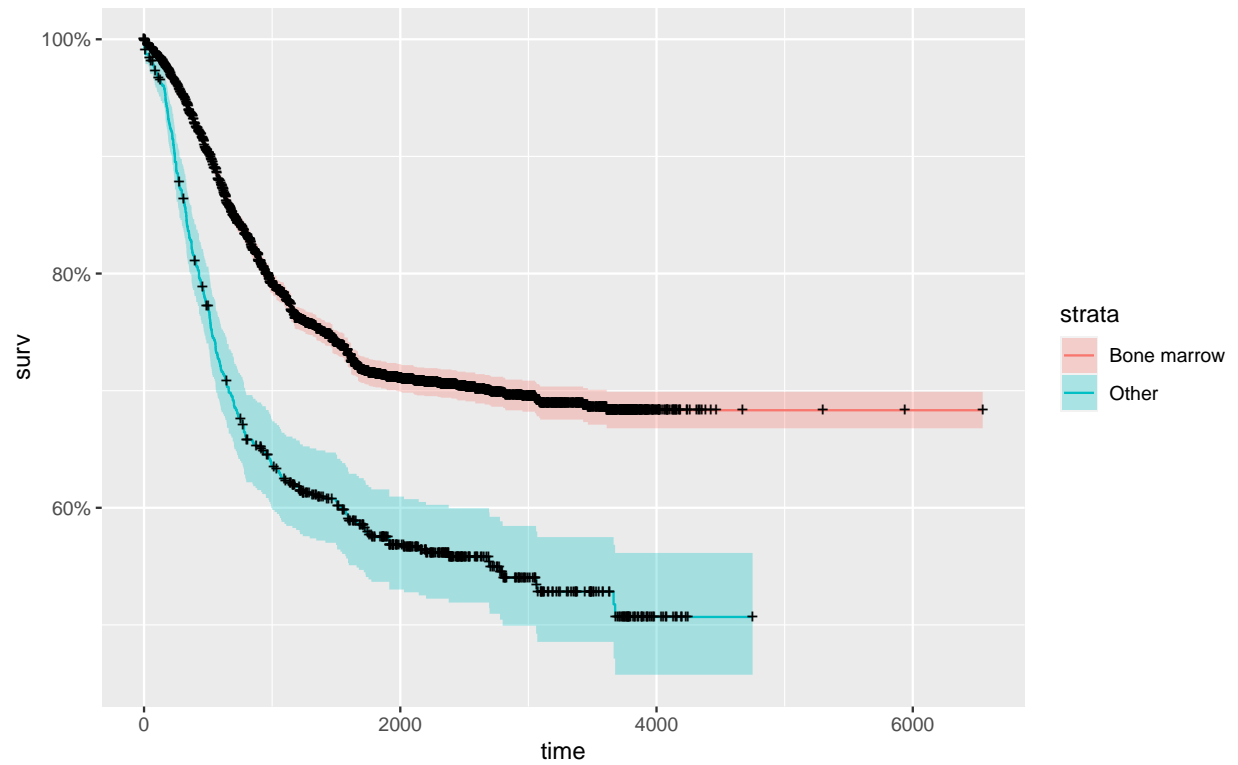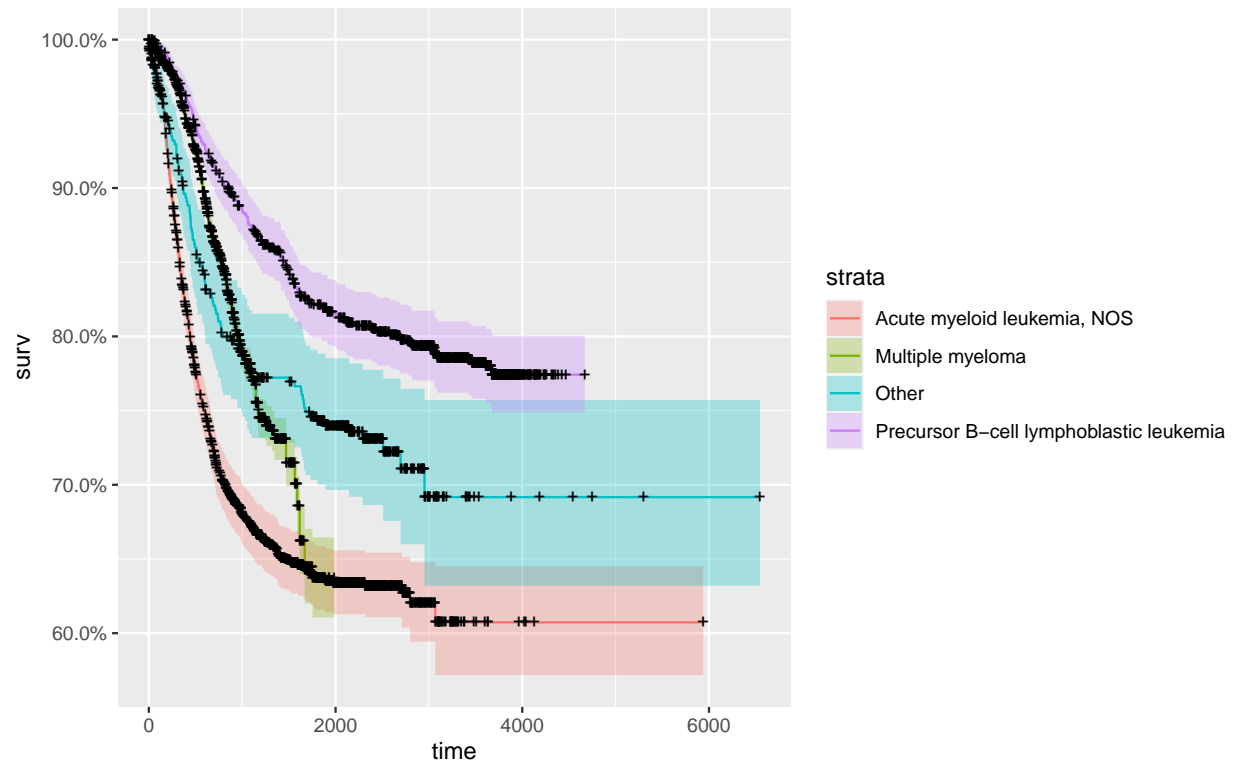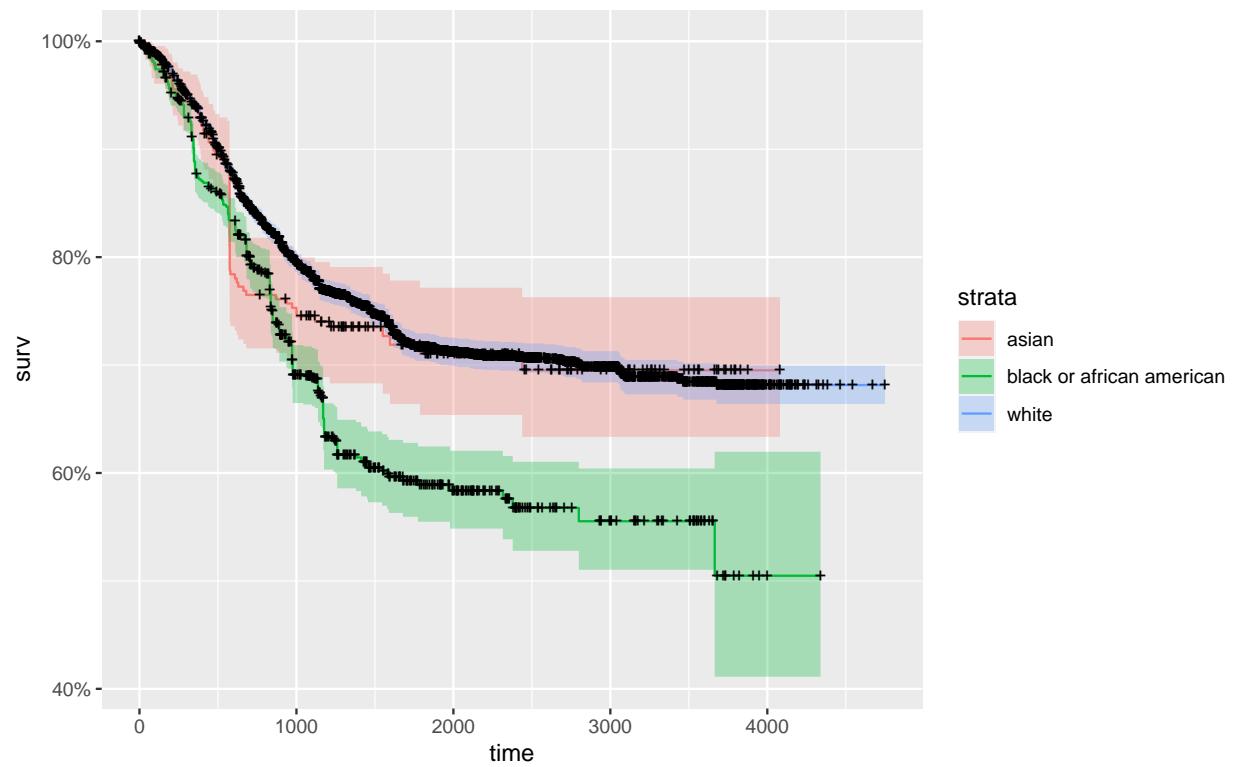
```
msurv3<-survfit(Surv(days_to_last_follow_up,vital_status)~ ethnicity, data = data)

autoplot(msurv3)
```
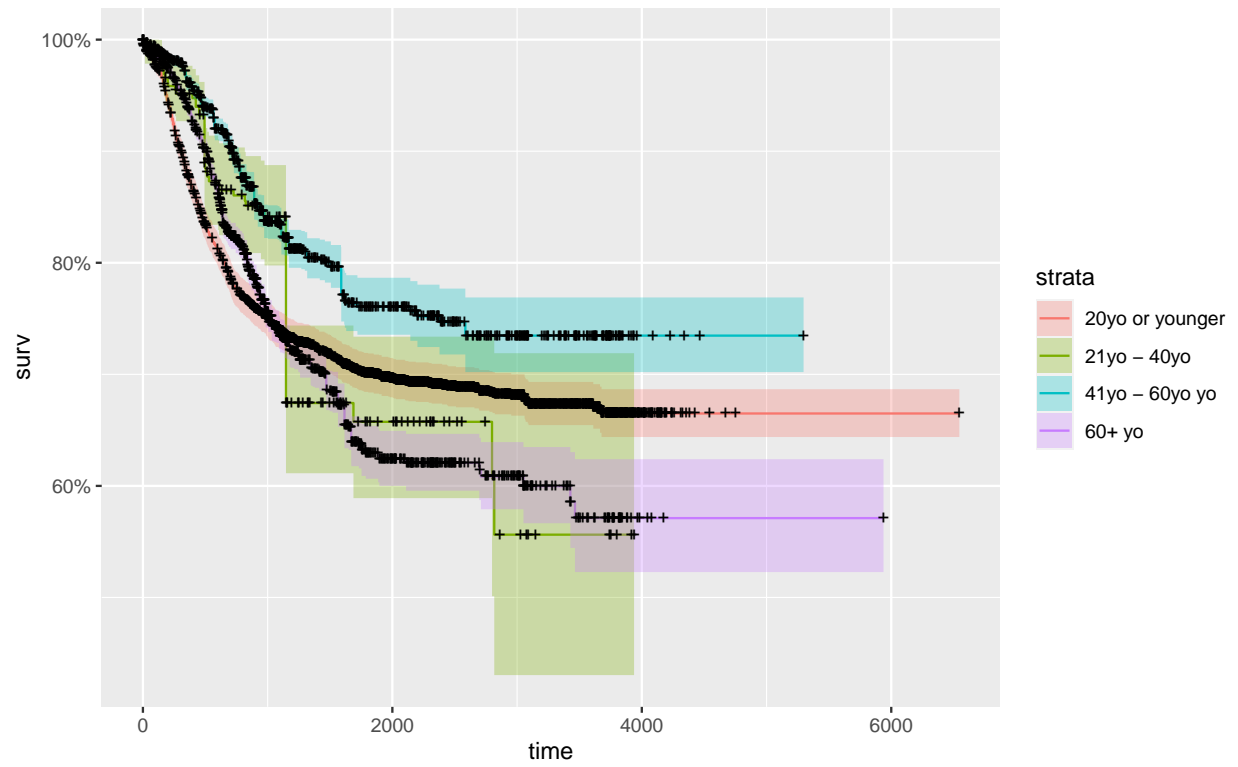
Hazard ratio

ethnicity	hispanic or latino (N=1284)	reference		
	not hispanic or latino (N=11080)	0.94 (0.81 – 1.08)	0.369	
gender	female (N=6467)	reference		
	male (N=9424)	1.22 (1.12 – 1.33)	<0.001	***
agecat	20yo or younger (N=3114)	reference		
	21yo – 40yo (N=411)	0.83 (0.63 – 1.09)	0.175	
	41yo – 60yo yo (N=3664)	0.55 (0.45 – 0.67)	<0.001	***
	60+ yo (N=5190)	0.93 (0.78 – 1.11)	0.411	
race	asian (N=351)	reference		
	black or african american (N=2200)	1.37 (1.06 – 1.76)	0.014	*
	white (N=9870)	0.88 (0.70 – 1.12)	0.309	
tissue_or_organ_of_origin	Bone marrow (N=12212)	reference		
	Hematopoietic system, NOS (N=3808)	reference		
site_of_resection_or_biopsy	Bone marrow (N=10820)	reference		
	Other (N=2841)	1.65 (1.41 – 1.93)	<0.001	***
primary_diagnosis	Acute myeloid leukemia, NOS (N=2739)	reference		
	Multiple myeloma (N=7255)	0.96 (0.81 – 1.14)	0.648	
	Other (N=4583)	0.12 (0.07 – 0.22)	<0.001	***
	Precursor B-cell lymphoblastic leukemia (N=1334)	0.42 (0.36 – 0.49)	<0.001	***

 # Events: 2188; Global p–value (Log–Rank): 1.8452e–85

AIC: 37822.78; Concordance Index: 0.63

0.05   0.1   0.2   0.5   1   2