

```
summary(sem.fit, rsq = T)
```

```
summary(sem.fit) # look for Variances:
```

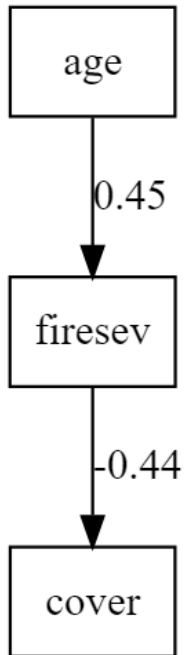


Fig. 1

```
> anova(sem_fit1, sem_fit2)
Chi-Squared Difference Test
```

	Df	AIC	BIC	Chisq	Chisq diff	Df diff	Pr(>Chisq)
sem_fit2	0	359.4	371.9	0.0000			
sem_fit1	1	360.7	370.7	3.2974	3.2974	1	0.06939 .

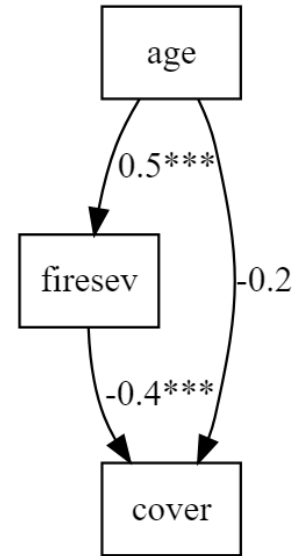


Fig. 2

We select **model 1** (simpler model) because adding link to the model (as shown on fig 2) did not significantly improved the fit. This means that the more complex model (sem_fit2) is NOT significantly more accurate than the simpler model sem_fit1 as our **Pr(>Chisq)** is larger .05