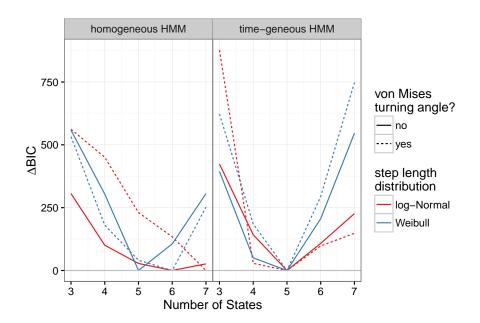
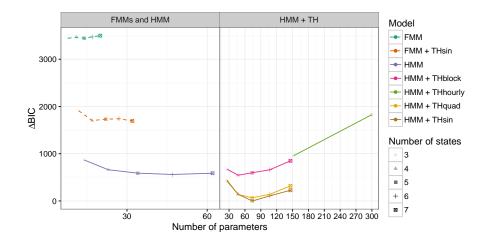
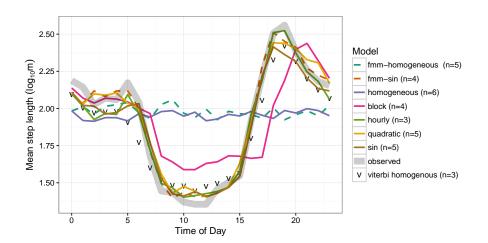
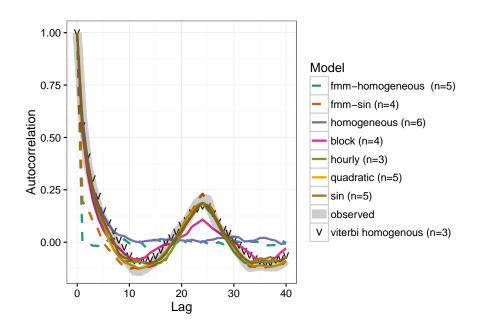


Figure 1: Simulation test for BIC-optimal approach

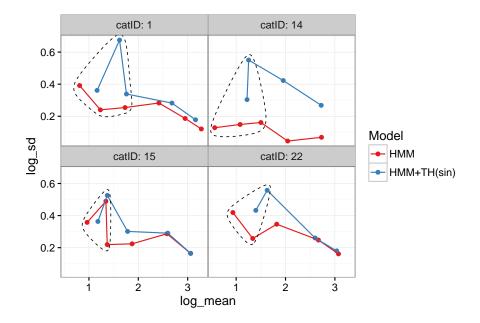




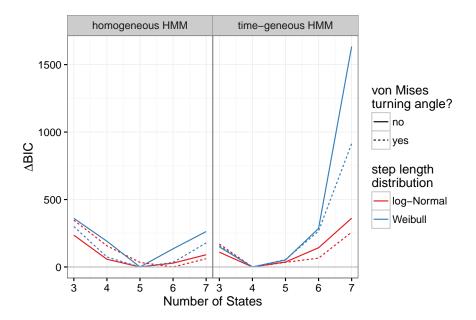


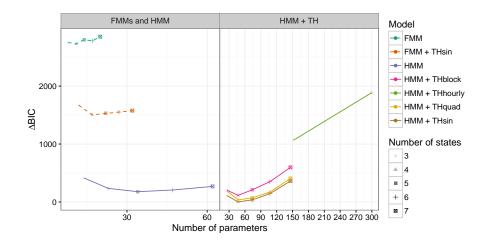


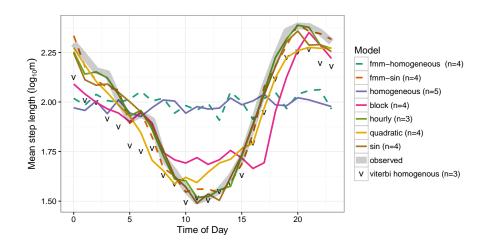
```
get_cat_pars2 <- function(catid){</pre>
  templist <- readRDS(paste("../summary_stats/cat.",catid,".df.RDS",sep=""))</pre>
  moddf <- templist$msdlist</pre>
  df <- data.frame(catID = catid</pre>
           , log_mean = c(unlist(moddf[1]),unlist(moddf[3]))
           , log_sd = c(unlist(moddf[2]),unlist(moddf[4]))
           , Model = c(rep("HMM",length(unlist(moddf[1]))),rep("HMM+TH(sin)",length(unlist(moddf[1])))
  return(df)
all_cat_pars2 <- ldply(lme4:::namedList(1,22,14,15),get_cat_pars2)</pre>
g0 <- ggplot(all_cat_pars2,aes(log_mean,log_sd,color=Model))+</pre>
     geom_point()+geom_line() + facet_wrap(~catID,labeller=label_both)+zmargin
 (g0c \leftarrow g0 +
     geom_encircle(data=subset(all_cat_pars2,
                                 (catID=="15" & log_mean<1.5) |
                                  (catID!="15" & log_mean<1.8)),
                    colour="black",linetype=2,expand=0.05))
```

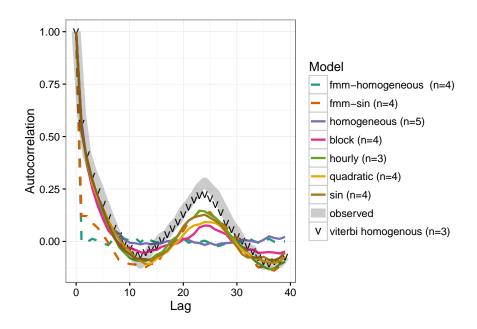


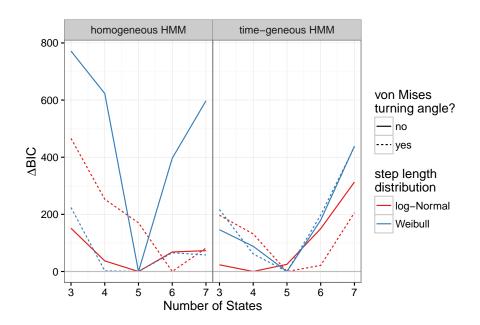
## 1 sup

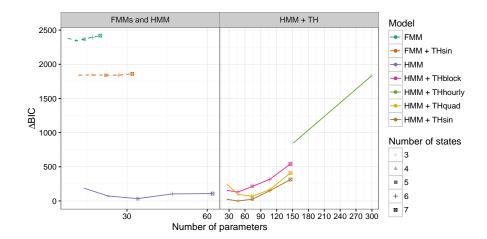












```
## Error in gzfile(file, "rb"): cannot open the connection
## Error in gzfile(file, "rb"): cannot open the connection
```

