

# Mazurka paper figures

*DJM*

8/20/2018

## Suggested order

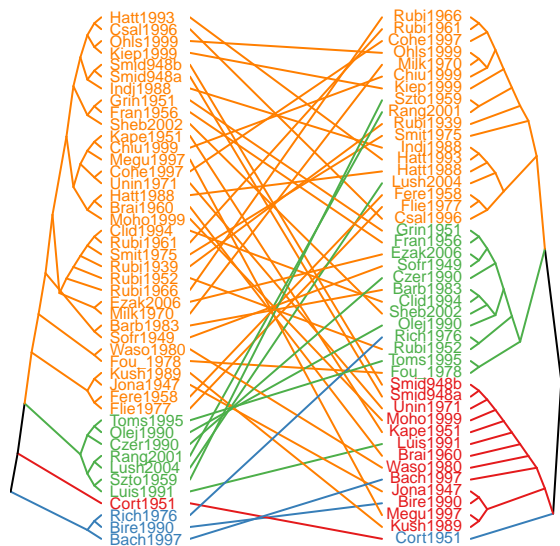
1. Parameter interpretation in Fliere
2. Using parameters to examine two different performances
3. Clustering performances (compare the clusters)
  - a. what can we say about the parameters of each cluster? what is different about them?
4. Similar performances (Rubinstein)
5. Model issues

## Comparing clusters

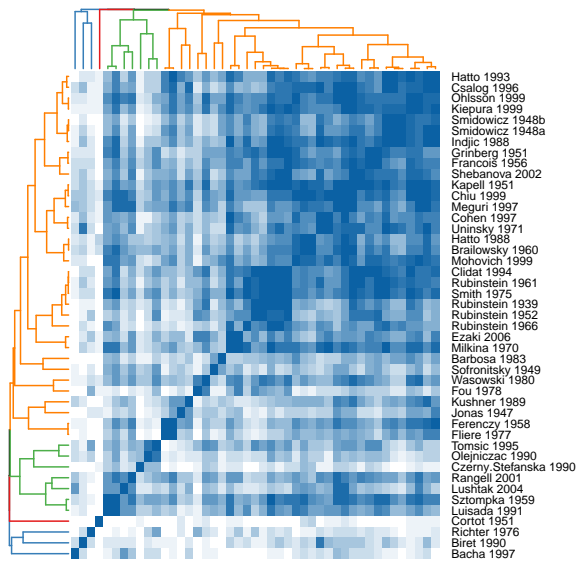
```
perfs = tempos[,-c(1:3)] %>% as.matrix %>% t
bad_perf = grep('Block',rownames(pvec_ml))
hc_parm = pvec_ml[-bad_perf,] %>% dist %>% percentize %>% hclust
hc_perf = perfs[-bad_perf,] %>% dist %>% percentize %>% hclust
short_labs = rownames(perfs)[-bad_perf]
lens = nchar(short_labs)
short_labs = paste0(substr(short_labs,1,4), substr(short_labs,lens-3,lens))
hc_parm$labels = short_labs
hc_perf$labels = short_labs
dend_parm = hc_parm %>% as.dendrogram
dend_perf = hc_perf %>% as.dendrogram

dend_parm = dend_parm %>% set('labels_col', value=fivecolors[1:4], k=4) %>%
  set('branches_lty', 1) %>%
  set('branches_k_color', value=fivecolors[1:4], k=4)
dend_perf = dend_perf %>% set('labels_col', value=fivecolors[1:4], k=4) %>%
  set('branches_lty', 1) %>%
  set('branches_k_color', value=fivecolors[1:4], k=4)
col_lines_by_left_groups <- fivecolors[cutree(dend_parm, 4, order_clusters_as_data=FALSE)]

tanglegram(dend_parm,dend_perf, color_lines = col_lines_by_left_groups,
  columns_width = c(1,1,1), axes=FALSE, rank_branches = TRUE, type='t',
  # left_dendo_mar = c(0,1,0,8), right_dendo_mar = c(0,8,0,1),
  margin_top = 0,
  margin_bottom = 0, margin_inner = 3.5,
  #remove_nodePar = TRUE,
  lab.cex=.75, lwd=1, edge.lwd=1)
```



```
heatmap.2(as.matrix(percentize(dist(pvec_ml[-bad_perf,]))),
  Rowv = dend_parm, Colv = dend_parm,
  symm=TRUE,
  density.info = 'none', trace='none',
  labRow = sub('_', ' ', row.names(pvec_ml)[-bad_perf]),
  labCol = NA,
  key.title = NA,
  col=colorRampPalette(c('#0b61a4','white')),
  key.xlab = NA,
  margins = c(1,6),
  cexRow = .6,
  cexCol = .6,
  lhei=c(1,8),
  lwid=c(1,8),
  offsetCol = 0, offsetRow = 0,
  key=FALSE
)
```

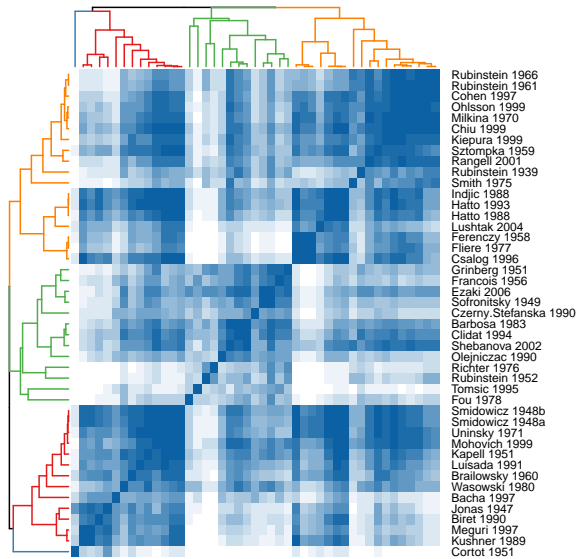


```

clusts = cutree(as.hclust(dend_perm), k = 4)
save(clusts, file = '../extras/ClusterLabels.Rdata')

heatmap.2(as.matrix(percentize(dist(perfs[~bad_perf]))),
  Rowv = dend_perm, Colv = dend_perm,
  symm=TRUE,
  density.info = 'none', trace='none',
  labRow = sub('_', ' ', row.names(pvec_ml)[~bad_perf]),
  labCol = NA,
  key.title = NA,
  col=colorRampPalette(c('#0b61a4', 'white')),
  key.xlab = NA,
  margins = c(1,6),
  cexRow = .6,
  cexCol = .6,
  lhei=c(1,8),
  lwid=c(1,8),
  offsetCol = 0, offsetRow = 0,
  key=FALSE
)

```



## Interpreting parameters

```
plotStates <- function(performers, pars, tempos,
                      noplot=FALSE,
                      particleNumber = 200,
                      initialMean = c(132,0),
                      initialVariance = c(400,10)){
  lt = diff(c(tempos$note_onset, 61))
  alldfs = NULL
  for(perf in performers){
    params = unlist(pars[row.names(pars)==perf,])
    y = matrix(tempos[[perf]], nrow = 1)
    mats = yupengMats(lt, params[1], params[2:4], params[5:8],
                      params[9:12], initialMean, initialVariance)
    bs = beamSearch(mats$a0, mats$P0, c(1,0,0,0,0,0,0,0), mats$dt,
                    mats$ct, mats$Tt, mats$Zt,
                    mats$Rt, mats$Qt, mats$GGt, y, mats$transMat, particleNumber)
    bestpath = bs$paths[which.max(bs$weights),]
    kal = kalman(mats, bestpath, y)
    df = data.frame(performer=perf, measure = tempos$note_onset, tempo = c(y),
                    inferred = c(kal$ests), state = convert8to4(bestpath))
    alldfs = rbind(alldfs, df)
  }
  if(noplot) return(alldfs)
  ggplot(alldfs, aes(x=measure, y=tempo)) + ylim(0, max(df$tempo)) +
    annotate('rect',xmin = 33, xmax = 45, ymin = -Inf, ymax = Inf,
            alpha=.2) +
    theme_minimal(base_family = 'Times') +
    geom_line(aes(y=tempo), color='black')+
    geom_point(aes(y=inferred),color=fivecolors[alldfs$state])+
    facet_wrap(~performer, labeller = as_labeller(function(x) gsub('_', ' ', x))) +
    theme(legend.position = 'none', legend.title = element_blank())+
    ylab('tempo (beats/minute)') + xlab('measure number') +
```

```

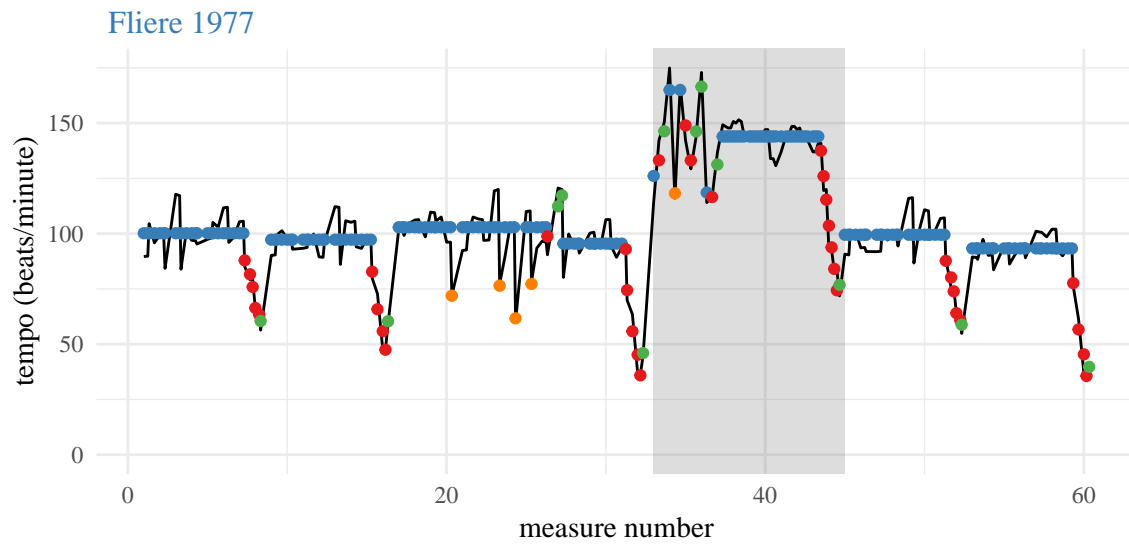
theme(strip.text = element_text(size=12, hjust=0,color = fivecolors[1]))
      #strip.background = element_rect(fill='grey90',linetype = 'blank'))
}

```

```

remove(pvec_ml)
load("mazurka2results.Rdata")
perfs = c('Fliere_1977','Tomsic_1995')
plotStates(perfs[1], pvec_ml, tempos)

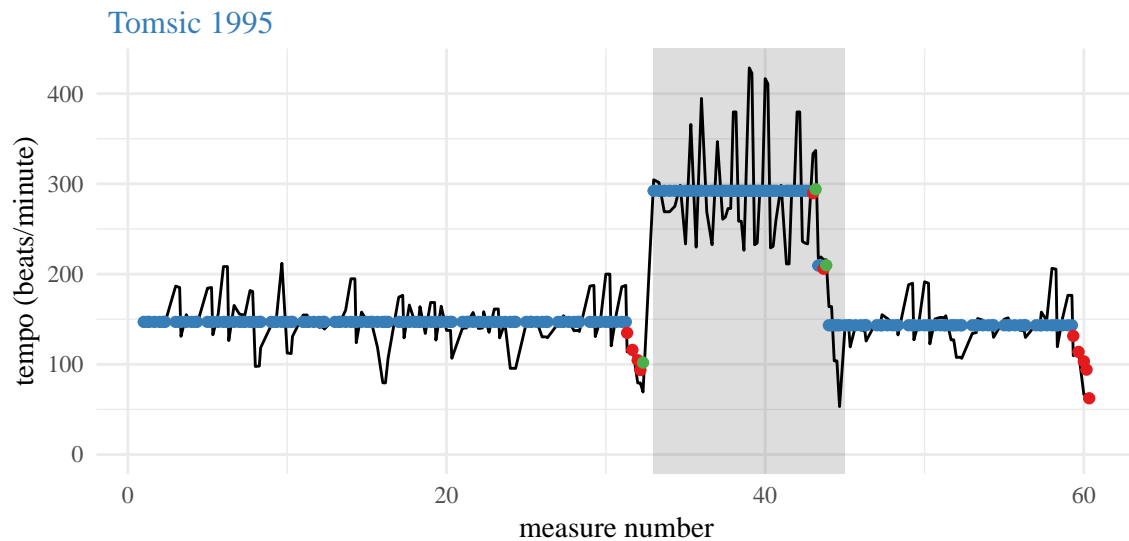
```



```

plotStates(perfs[2], pvec_ml, tempos)

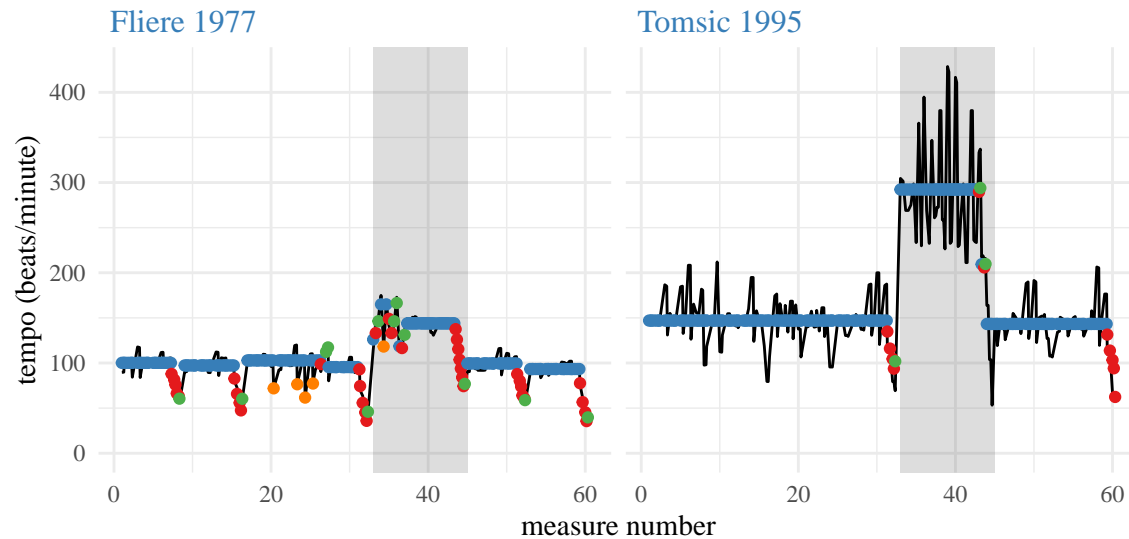
```



```

plotStates(perfs, pvec_ml, tempos)

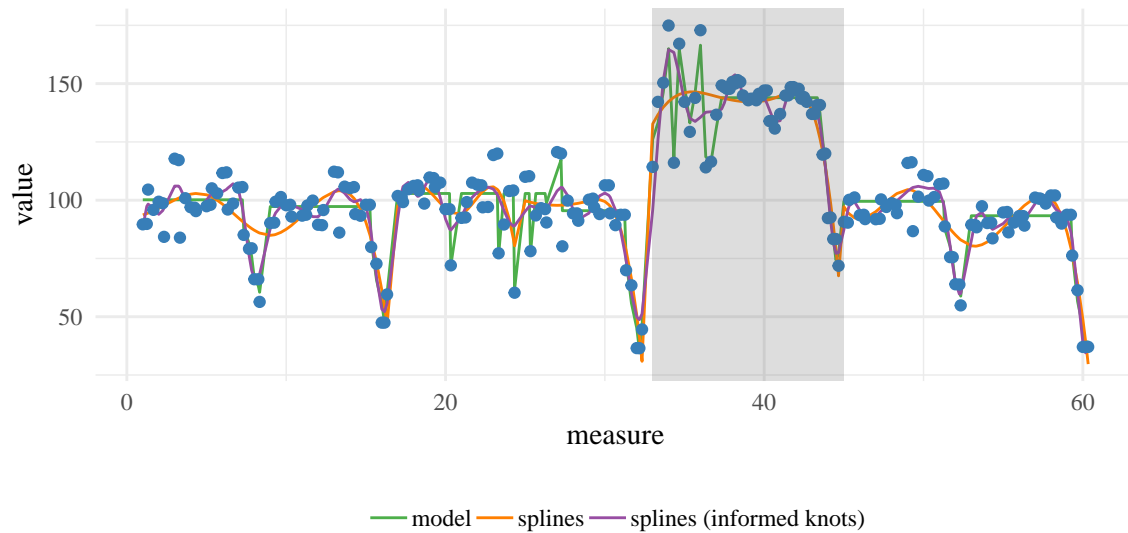
```



## Different smoothing

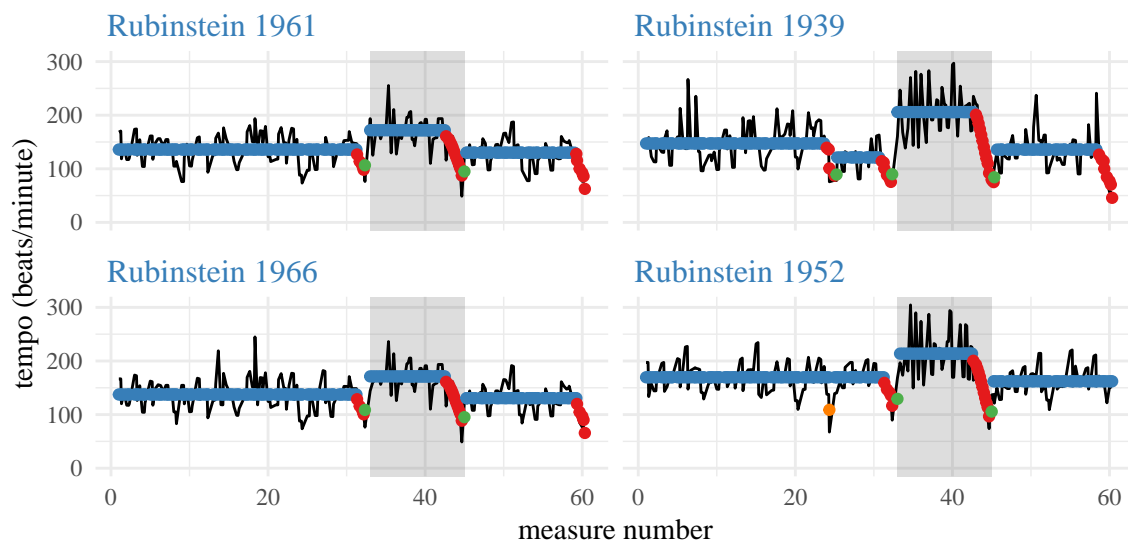
Try splines, replicating knots, lltf?

```
nsplines = 64 # 1 knot per bar plus boundary
B = bs(tempos$note_onset, df=nsplines, intercept = TRUE)
preds_smooth = fitted(lm(tempos[[perfs[1]]]~B-1))
single.knots = match(seq(4,56,by=4)+1,tempos$meas_num)
double.knots = match(c(16,24,32,44)+1, tempos$meas_num)
triple.knots = match(c(16,24,32,44)+1, tempos$meas_num)
quad.knots = match(c(16,24,32,44)+1, tempos$meas_num)
all.knots = tempos$note_onset[
  sort(c(single.knots,double.knots,triple.knots,quad.knots))]
B1 = bs(tempos$note_onset, knots = all.knots, intercept = TRUE,Boundary.knots = c(1,61))
preds_music = fitted(lm(tempos[[perfs[1]]]~B1-1))
extras=data.frame(x=tempos$note_onset,y1=preds_smooth,y2=preds_music)
perf1 = plotStates(perfs[1], pvec_ml, tempos, noplot = TRUE)
perf1$ss = preds_smooth
perf1$ms = preds_music
perf1 %>% select(measure, ss, ms, inferred) %>%
  gather(key='key',value='value',-measure) %>%
  ggplot(aes(x=measure)) + geom_line(aes(y=value, color=key)) +
  scale_color_manual(values = fivecolors[c(3,4,5)],
    labels = c('model','splines','splines (informed knots)')) +
  geom_point(data=perf1, aes(x=measure, y=tempo), color=fivecolors[1]) +
  annotate('rect',xmin = 33, xmax = 45, ymin = -Inf, ymax = Inf,
    alpha=.2) +
  theme_minimal(base_family = 'Times') +
  theme(legend.position = 'bottom',legend.title = element_blank())
```



## Similar performances

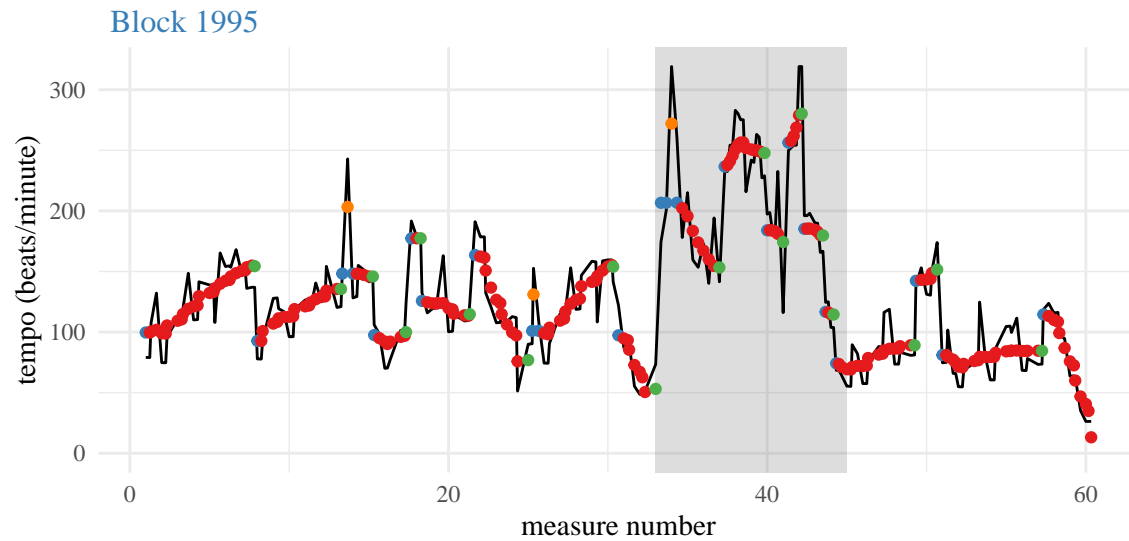
```
rubins = row.names(pvec_ml)[grep('Rubinstein', row.names(pvec_ml))]
plotStates(rubins, pvec_ml, tempos)
```



*# note that the 1939 recording is the only one in a different cluster*

## Bad estimation

```
plotStates('Block_1995', pvec_ml, tempos)
```



## Problems with the model

- Problem with retransitioning to state 1
- states 2 and 3 aren't constrained to always decrease/increase, only in mean
- state 4 may not always emphasize a slow down
- previous 2 have to do with Gaussian assumptions
- necessity for strong priors
- but priors are on parameters, not on path (how would we want this to change?)