

```

#
# Implements procedureWithGrowthPattern() which allows a general
# thresholding procedure to be run with a growth pattern where
# when one location finishes, it is used to seed any of its
# immediate neighbours that have a 'wave number' one greater
# than itself.
# Note that this is different from some growth patterns which allow
# for the starting guess to be the average of several neighbours,
# thus requiring all neighbours to complete before that location
# can commence (eg Turpin et al's implementation of HFA growth pattern).
#
# The procedure is controlled by a growth pattern matrix, and
# requires a matrix of guesses for the first subset of locations,
# and functions to start, step, test-for-stop, and get-final-threshold
# for the procedure. See comments and Zest242.r for an example.
# The grid must have a 'chain' of immediate neighbours to locations
# not numbered 1 (the initial points) as they can only be opened for
# testing if an immediate 8-neighbour is complete.
#
# This version also has allowance for catch trials and an adaptive response
# window.
#
# WARNING - the function setResponseWindow assume that the responseWindow
#           is in position 8 of the 3rd line of the makeStim function
#           This is truly disgusting! It needs to be fixed in the future.
#
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# Date: Sun 17 May 2015 09:26:05 AEST
#

#####
# Use startFun, stepFun and stopFun to run all locations with number
# 1 in the growthPattern. When one finishes it opens up any locations
# numbered 2 in the immediate 8-neighbourhood. When one of those finishes
# a 3 is opened if abutting, and so on.
# After each location finishes, use finalFun to get threshold and num pres.
# Within each subset of possible locations,
# locations are presented in random order.
#
# INPUTS:
# gp - growth pattern matrix (min(gp) == 1). See 'chain' comment above.
# gn - growth next. lookup table of next locations to open up in the growth pattern
# statrs - matrix same size as gp giving start values for gp==1
# startFun - function(guess, rw, cl) that creates a state for
#           location (rw, cl) in gp using guess as start guess
# stepFun - function(state) returns a new state after one presentation
# stopFun - function(state) returns TRUE if location finished
# finalFun - function(state) returns c(final threshold, number presentations)
# FPTrials - perform an FP trial every time the presentation counter hits this value
# FNTrials - perform an FN trial every time the presentation counter hits this value
# FPLevel - level of stimuli for false positive in cd/m^2
# FNPause - duration of pause (in ms) after a FN presentation, to allow for recovery after bright stimulus.
# FNDelta - the degree of brightness relative to current threshold at that location (in dB)
#           e.g. if FNDelta = 10, FN stimulus of 10 dB brighter than current estimated threshold is presented.
# FNLocationThreshold - locations with a final threshold of at least this value (cd/m^2)
#                       are candidates for a FN presentation.
# FPSize - size of FP stimulus (diameter in mm)
# FNSize - size of FN stimulus (diameter in mm)
# initialRespWin - starting response window in ms
# respWinBuffer - time added to mean of respWin to determine final response Window
# catchTrialLoadFreq - Frequency of catch trials in the first minute. Usually more frequent in order to front load.
# catchTrialFreq - Frequency of catch trials for the remainder of the test
# catchTrialMax - Maximum number of FP and FN catch trials for the test (e.g. if catchTrialMax = 5, will result in 5 FP and 5 FN catch trials)
#
# RETURNS: list of two matrices, each with same dimensions as gp
#          t is final threshold at each location
#          n is number of presentations at each location
#####
source("testStatusOutput.r")
if(!('audio' %in% installed.packages()))
  install.packages("audio")
library("audio")

procedureSuprathreshold <- function(
  startTime,
  locations, #previously 'gp'
  # gn,
  # starts,
  startFun,
  stepFun,
  stopFun,
  finalFun,
  gridPat,
  catchTrialLoadFreq=6,
  catchTrialFreq=20,
  catchTrialMax=14,
  FPLevel=dbToacd(60, 4000/pi),
  FNDelta=10,
  FNPause=300,
  FNLocationThreshold=20,
  FPSize=1.72,
  FNSize=1.72,
  initialRespWin=1200,
  respWinBuffer=250,
  moveProj = T,
  minInterStimInt = 0) {

  #####
  ## Return any locations that are in the immediate 8 neighbours
  ## of [rw,cl] and have a wave number == gp[rw,cl]+1.
  ##
  ## INPUTS
  ## gp - growth Pattern matrix
  ## rw - row of location
  ## cl - column of location
  ##
  ## RETURNS: matrix of locations where column 1 = row index, column 2 = column index
  #####
  openUP <- function(gp, gn, rw, cl, states) {
    wave <- gp[rw,cl]
    locations <- NULL
    x <- states[[rw,cl]]$x
    y <- states[[rw,cl]]$y
    #
    nextLocs <- gn[[wave]][[paste(x,y,sep=" ")]]
    #
    if (!is.null(unlist(nextLocs))) {
    #
      locations <- matrix(nextLocs,length(unlist(nextLocs))/2,2) #make sure locations are in matrix form (becomes numeric if there is only one row)
    #
      for (row in 1:nrow(locations)) { # convert back to row and column
        locations[row,1] <- 91 + locations[row,1]
        locations[row,2] <- 55 - locations[row,2]
      }
      locations <- t(apply(locations,1,rev)) # flip so that row index is in first column and col index in second column
    # locations <- locations[apply(locations,1,function (x) {is.null(states[[x[1],x[2]]]}),)] # remove locations that have already been opened up
    # locations <- matrix(locations,length(unlist(locations))/2,2)
    # if (nrow(locations) == 0) {locations <- NULL}
    #
    }
    return(locations)
  }

  #####

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# Present FP or FN trial. Return TRUE(seen) or FALSE (not seen)
#####
# NEED TO REDO FOR NEG CATCH TRIALS
presentCatch <- function(posOrNeg, responseWindow, currentThresholds, states,index) {
  if (posOrNeg == "POS") {
    s <- list(x=9, y=9, level=FPLevel, size=FPSize, duration=200,
              responseWindow=responseWindow)
  } else {

    k <- which(currentThresholds > FNLocationThreshold,arr.ind=TRUE) ## find rw and cl of eligible locations
    l <- k[sample(nrow(k),size=1),] # choose a location at random: l[rw,cl]

    s <- list(x=states[l[1],l[2]]$x, y=states[l[1],l[2]]$y,
              level=dbTocd(round(currentThresholds[l[1],l[2]] - FNDelta),4000/pi),
              size=FNSize, duration=200,
              responseWindow=responseWindow)
  }
  class(s) <- "opiStaticStimulus"

  if (moveProj) {
    showStim <- opiPresent(stim=s,states[[index[[1]][1],index[[1]][2]]]$makeStim(0,0))
  } else {
    showStim <- opiPresent(stim=s)
  }

  return (c(showStim,list(stimulus=s$level)))
}

#####
# If there is only one location remaining, present a random stimulus
# so that that particular location is not tested in a row
#
#####
# NEED TO REDO
presentDummy <- function (grid,responseWindow,dummyState,states) {
  locIndex <- which(!is.na(grid),arr.ind=TRUE)
  loc <- locIndex[round(runif(1,1,nrow(locIndex))),] # choose a location at random
  stimulus <- round(runif(1,0,30)) # choose a stimulus intensity at random
  s <- list(x=dummyState(30,loc[1],loc[2])$x,y=dummyState(30,loc[1],loc[2])$y,
            level=dbTocd(stimulus, 4000/pi),size=FNSize,duration=200,responseWindow=responseWindow)

  class(s) <- "opiStaticStimulus"

  if (moveProj) {
    showStim <- c(opiPresent(stim=s,states[[index[[1]][1],index[[1]][2]]]$makeStim(0,0)),list(x=s$x,y=s$y,stimulus=stimulus))
  } else {
    showStim <- c(opiPresent(stim=s),list(x=s$x,y=s$y,stimulus=stimulus))
  }
  return (showStim)
}

#####
# Function to alter the response window of a state
# Note does not allow responseWindow to be less than respWinBuffer
# Returns the state after alteration
#####
setResponseWindow <- function(state,respWinBuffer,responseWindow) {

  m <- state$makeStim
  body(m)[[2]][[3]][[8]] <- respWinBuffer + responseWindow
  state$makeStim <- m

  return(state)
}

#####
# Function to remove previous presentation response information
# if the observer made a known false response
#
#####
#NEED TO FIX
applyUndos <- function () {
  #print(locsPresented)
  myEnv <- parent.env(environment())
  delLocs <- NULL
  while (gUndos > 0 && nrow(locsPresented) >= gUndos) {
    rr <- locsPresented[nrow(locsPresented),1]
    cc <- locsPresented[nrow(locsPresented),2]

    #delLocs - table of locations to be deleted.
    #Purpose: to identify locations that need to be deleted twice.
    delLocs <- rbind(delLocs,c(rr,cc))

    z <- unlist(lapply(locs, function(x) all(x == c(rr,cc))))

    if (any(z)) {
      print('removing an unterminated location')
    } else {
      print('removing a terminated location')
      myEnv$locs <- c(locs,list(c(rr,cc)))
      myEnv$finished_counter <- finished_counter - 1
    }

    # check for locations with repeated deletions.
    # If there has been a repeat, need to look up second last stim value rather than last.
    lookupIndex <- sum(apply(delLocs,1,function (x) ((x[1] == rr) && x[2] == cc)))
    prevStimVal <- tail(states[[rr,cc]]$stimuli,lookupIndex)[1]
    prevStimVal <- which(prevStimVal == states[[rr,cc]]$domain)

    if (tail(states[[rr,cc]]$responses,lookupIndex)[1] == FALSE) {
      states[[rr,cc]]$pdf <- states[[rr,cc]]$pdf / (1-states[[rr,cc]]$likelihood[prevStimVal,])
      myEnv$states[[rr,cc]]$pdf <- states[[rr,cc]]$pdf / sum(states[[rr,cc]]$pdf)
    } else {
      states[[rr,cc]]$pdf <- states[[rr,cc]]$pdf / states[[rr,cc]]$likelihood[prevStimVal,]
      myEnv$states[[rr,cc]]$pdf <- states[[rr,cc]]$pdf / sum(states[[rr,cc]]$pdf)
    }
    myEnv$locsPresented <- locsPresented[-nrow(locsPresented),]
    gUndos <- gUndos - 1

    if (details$gridType != "practice") {
      cat(file=paste(details$dx,"/",details$gridType,"
",details$stimSizeRoman,"/",details$name,"_",details$dx,"_",details$grid,"_",details$stimSizeRoman,"_",details$eye,"Eye_",details$date,"_",details$startTime,"_stimResponses.txt",sep=""),
          append=TRUE,paste("Presentation at location x =",states[[rr,cc]]$x,"y =",states[[rr,cc]]$y,"was deleted\n",sep=" "))
    }
  }
}

#####
# Function for the adaptive interstimulus interval
#
# INPUTS
# responseTime - vector of total response times throughout test
# minISI - the minimum inter-stimulus interval
# interStimMultiplier - multiplier of the mean response time, which determines the max interstim interval
#
#####
interStimInt <- function (responseTime = respTime,minISI,interStimMultiplier = 1) {
  if (!is.null(respTime)) {
    Sys.sleep(runif(1, min=minISI, max= max(minISI,mean(responseTime) * interStimMultiplier)/1000) # pause before presenting next stimulus
  ) else {
    Sys.sleep(200/1000) #If there have been no response times recorded yet, make interstim interval 200 ms
  }
}

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}
}

#####
# set up answers and loop vars
#####

currentThresholds <- starts
finishedThresholds <- matrix(NA, nrow(gp), ncol(gp))
currentNumPres <- matrix(NA, nrow(gp), ncol(gp))

states <- array(list(), dim=c(nrow(gp), ncol(gp)))

# Set up locations whose wave == 1
locs <- which(gp == 1, arr.ind=TRUE)
locs <- split(locs, 1:nrow(locs))
for (i in 1:length(locs)) {
  rc <- locs[[i]]
  states[[rc[1], rc[2]]] <- startFun(starts[[rc[1],rc[2]]], rc[1], rc[2])
}

respWin <- rep(initialRespWin,5) ## set up adaptive response window

respTime <- NULL # vector of response times
fp_counter <- NULL # vector of responses for FP
fn_counter <- NULL # vector of responses for FN
finished_counter <- 0 # number of terminated locations
counter <- 1 # number of presentations
locsPresented <- NULL # vector of locations presented in order
gUndos <- 0
index <- locs[sample(1:length(locs))] # choose random location to test first

#####
# loop while still some unterminated locations
#####
# what is gRunning?
while (length(locs) > 0 && gRunning) {
  start_time <- Sys.time()
  applyUndos()

  if ((counter <= 60 && length(fp_counter) < catchTrialMax && (counter %% catchTrialLoadFreq == 0) && ((counter/catchTrialLoadFreq) %% 2 != 0)) ||
      (counter > 60 && length(fp_counter) < catchTrialMax && (counter %% catchTrialFreq == 0) && ((counter/catchTrialFreq) %% 2 != 0))) {

    result <- presentCatch("POS", mean(respWin) + respWinBuffer, currentThresholds, states,index) #adaptive response window
    #result <- presentCatch("POS", initialRespWin, currentThresholds, states,index) # fixed response window

    if (result$seen) {
      for (i in 1:2) {
        wait((play(sin(1:8000/20)))) ## play 2 beeps if FP error is made
      }
    }

    fp_counter <- c(fp_counter, min(1,result$seen))

    testStatus(result$seen,currentNumPres,currentThresholds,finishedThresholds,finished_counter,gp,fp_counter,fn_counter,stateInfo=states[[rw,cl]],respTime,testGrid = gridPat)
    if (details$gridType != "practice") {
      cat(file=paste(details$dx,"/",details$gridType,"
",details$stimSizeRoman,"/",details$name,"_",details$dx,"_",details$grid,"_",details$stimSizeRoman,"_",details$eye,"Eye_",details$date,"_",details$startTime,"_stimResponses.txt",sep=""),
          append=TRUE,sprintf("Location: %5s Stim: %2g dB Seen: %5s Resp Time: %5.2f Trial Time: %.0f\n", "FPCatch",cdTodb(FPLevel,4000/pi), result$seen, result$time,
diffTime(Sys.time(),start_time,units = "secs") * 1000))
    }
    counter <- counter + 1
    start_time <- Sys.time() #reset start_time counter for trial time
  }

  if ((counter <= 60 && length(fn_counter) < catchTrialMax && ((counter %% catchTrialLoadFreq == 0) && ((counter/catchTrialLoadFreq) %% 2 == 0)) && any(currentThresholds >
FNLocationThreshold,na.rm=TRUE)) ||
      (counter > 60 && length(fn_counter) < catchTrialMax && ((counter %% catchTrialFreq == 0) && ((counter/catchTrialFreq) %% 2 == 0)) && any(currentThresholds >
FNLocationThreshold,na.rm=TRUE))) {

    result <- presentCatch("NEG", mean(respWin) + respWinBuffer, currentThresholds, states,index)
    #result <- presentCatch("NEG", initialRespWin, currentThresholds, states,index) #fixed response window

    Sys.sleep(FNPause/1000)
    fn_counter <- c(fn_counter, result$seen == FALSE)
    testStatus(result$seen,currentNumPres,currentThresholds,finishedThresholds,finished_counter,gp,fp_counter,fn_counter,stateInfo=states[[rw,cl]],respTime, testGrid = gridPat)
    if (details$gridType != "practice") {
      cat(file=paste(details$dx,"/",details$gridType,"
",details$stimSizeRoman,"/",details$name,"_",details$dx,"_",details$grid,"_",details$stimSizeRoman,"_",details$eye,"Eye_",details$date,"_",details$startTime,"_stimResponses.txt",sep=""),
          append=TRUE,sprintf("Location: %5s Stim: %2g dB Seen: %5s Resp Time: %5.2f\n Trial Time: %.0f\n", "FNCatch",cdTodb(result$stimulus,4000/pi), result$seen,
result$time,diffTime(Sys.time(),start_time,units = "secs") * 1000))
    }

    counter <- counter + 1
    start_time <- Sys.time() #reset start_time counter for trial time
  }

  counter <- counter + 1

  # weight stimulus choice by growth pattern wave
  getWave <- lapply(locs, function (x) {gp[x[1],x[2]]})
  weight <- sapply(getWave, function (x) {1/(x^2)})

  if (length(locs) > 1) {
    index[2] <- locs[sample(1:length(locs),1,prob=weight)]
    while (all(index[[1]] == index[[2]])) {
      index[2] <- locs[sample(1:length(locs),1,prob=weight)]
    }
  } else {
    index[[1]] <- locs[[1]]
    index[[2]] <- locs[[1]]
  }

  rw <- index[[1]][1]
  cl <- index[[1]][2]
  rw2 <- index[[2]][1]
  cl2 <- index[[2]][2]

  locsPresented <- rbind(locsPresented,c(rw,cl))

  states[[rw,cl]] <- setResponseWindow(states[[rw,cl]],respWinBuffer,mean(respWin)) #Updates response window

  if (length(locs) > 1 && moveProj == TRUE) {
    states[[rw,cl]] <- stepFun(states[[rw,cl]],nextStimState=states[[rw2,cl2]])
  } else {
    states[[rw,cl]] <- stepFun(states[[rw,cl]])
  }

  if (all(details$gridType != c("Peripheral","P-Peripheral","P-Edge"))) {interStimInt(respTime,minInterStimInt)}
  currentThresholds[[rw,cl]] <- sum(states[[rw,cl]]$pdf*states[[rw,cl]]$domain) # update currentThresholds
  currentNumPres[[rw,cl]] <- states[[rw,cl]]$numPresentations

  testStatus(tail(states[[rw,cl]]$responses,1),currentNumPres,currentThresholds,finishedThresholds,finished_counter,gp,fp_counter,fn_counter,stateInfo=states[[rw,cl]],respTime,
testGrid = gridPat)
  if (details$gridType != "practice") {
    cat(file=paste(details$dx,"/",details$gridType,"
",details$stimSizeRoman,"/",details$name,"_",details$dx,"_",details$grid,"_",details$stimSizeRoman,"_",details$eye,"Eye_",details$date,"_",details$startTime,"_stimResponses.txt",sep=""),
        append=TRUE, sprintf("Location: x=%3g, y=%3g Stim: %2g dB Seen: %5s Resp Time: %5.2f Trial Time: %.0f\n", states[[rw,cl]]$x, states[[rw,cl]]$y,
tail(states[[rw,cl]]$stimuli,1), tail(states[[rw,cl]]$responses,1), tail(states[[rw,cl]]$responseTimes,1),diffTime(Sys.time(),start_time,units = "secs") * 1000))
  }
}

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}

if (length(locs) == 1) {
  dummy_start_time <- Sys.time()
  result <- presentDummy (gridPat,mean(respWin) + respWinBuffer,startFun,states)
  #result <- presentDummy (gridPat,initialRespWin,startFun,states) #fixed response window

  if (all(details$gridType != c("Peripheral","P-Peripheral","P-Edge"))) {interStimInt(respTime,minInterStimInt)}

  if (details$gridType != "practice") {
    testStatus(result$seen,currentNumPres,currentThresholds,finishedThresholds,finished_counter,gp,fp_counter,fn_counter,stateInfo=list(x=result$x,y=result$y),respTime, testGrid
= gridPat)
    cat(file=paste(details$dx,"/",details$gridType,"
",details$stimSizeRoman,"/",details$name,"_",details$dx,"_",details$grid,"_",details$stimSizeRoman,"_",details$eye,"Eye_",details$date,"_",details$startTime,"_stimResponses.txt",sep=""),
        append=TRUE,sprintf("Location: x=%3g, y=%3g Stim: %2g dB Seen: %5s Resp Time: %5.2f Trial Time: %.0f %5s\n",result$x,result$y,result$stimulus, result$seen,
result$time,difftime(Sys.time(),dummy_start_time,units = "secs") * 1000,"(Dummy Trial)"))
  }
}

if (tail(states[[rw,cl]]$responses,1)) {
  respWin <- c(tail(states[[rw,cl]]$responseTimes,1),respWin[-5])
  respTime <- c(tail(states[[rw,cl]]$responseTimes,1),respTime)
}

if (stopFun(states[[rw,cl]])) {
  # fill in finishedThresholds and remove from locs
  finishedThresholds[rw,cl] <- finalFun(states[[rw,cl]])[1]
  finished_counter <- finished_counter + 1

  locs <- locs[-which(sapply(locs,function(x) {all(x == index[[1]]))})]

  # look around for neighbours that can be opened
  newLocs <- openUP(gp, gn, rw, cl,states)

  if (!is.null(newLocs)) {
    for (i in 1:nrow(newLocs)) {
      rc <- newLocs[i,]
      states[[rc[1], rc[2]]] <- startFun(finishedThresholds[rw,cl], rc[1], rc[2])
      locs <- c(locs, list(rc))
    }
  }
  index[[1]] <- index[[2]] #move next stimulus to current stimulus before next presentation sequence
}
if (gRunning) {
  currentThresholds[currentThresholds < states[[rw,cl]]$domain[6]] <- -1 #Set censored thresholds to -1
  finishedThresholds[finishedThresholds < states[[rw,cl]]$domain[6]] <- -1 #Set censored thresholds to -1
}
### NEED TO INCORPORATE PARENT FILE FOR THIS

testStatus(result$seen,currentNumPres,currentThresholds,finishedThresholds,finished_counter,gp,fp_counter,fn_counter,stateInfo=states[[rw,cl]],respTime,plotStimResponse=FALSE,testGrid =
gridPat)
  return(list(t=currentThresholds, n=currentNumPres,fpc=fp_counter,fnc=fn_counter,rt=respTime))
}

```