

Anova

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1 Statistical analysis of Line Task 2014 data

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Author: Seb James seb.james@sheffield.ac.uk

This is a companion analysis notebook for the paper Target-distractor Synchrony Affects Performance in a Novel Motor Task for Studying Action Selection

This notebook uses data which was generated by the Octave/Matlab script `lt_analyse_all.m`.

It does some analysis itself, but it also passes much analysis off to R scripts, the output of which it then presents. It is rather unparsemonious to have used three coding environments to have analysed the data from the line task; in retrospect I would probably have limited this to two (Octave and R).

1.1 Data import

Import latency and error data which is computed from the raw data by the script `lt_analyse_all.m`. This script saves the resulting data into a Matlab v7 file.

See the `Analysing Data.ipynb` notebook for details about how the data is organised. This notebook contains just the analysis.

This script requires the python modules `scipy` and `statsmodels`. It also makes external system calls to evaluate R scripts and so R must also be installed together with the R modules `effsize` and `nlme`.

```
In [12]: # Import the data, which should be available in Matlab v7 format.  
         # fnames is used throughout the rest of this notebook, so this section needs to be evaluated!  
         import scipy.io as sio  
         mat_workspace = sio.loadmat('AllData/fnames.mat')  
         fnames = mat_workspace['fnames']  
         print 'fnames.mat has been imported.'  
         # We need Image from IPython.display to view graphs  
         from IPython.display import Image
```

`fnames.mat` has been imported.

1.1.1 On subject omission

Omitted subjects, with reasons:

EM1, KW, JD, LC, YC - Experimenter chose the wrong jump time on one or more conditions.

AB2_ - Experimenter did not carry out a Synchronous Distractor trial

is132 - Age not recorded for this participant - could not match data to participant in Spreadsheet.

After omitting these subjects, 55 remained.

1.1.2 On event omission

The script `lt_analyse_latency.m` will try to produce accurate latencies to first movement, along with direction errors from the time series data in the 2014*.txt files. It omits some events for a set of reasons which are

described in the paper (grep for omit_reason in the .m code to find them and see the omit_reasons() function in the analysis code below).

Later, when the individual analysis is called, a standard approach to outlier removal is taken (see individual.excludeOutliers()).

1.2 Count the number of target events

This next section of code tallies up the number of target events in each task. This is not the number of latency measurements obtained from the data by the Octave pre-processing script.

```
In [13]: from __future__ import division
import numpy as np

# Find event counts
import scipy.io as sio
mat_workspace2 = sio.loadmat('AllData/eventcounts.mat')
evcounts = mat_workspace2['fnames']

num_targs = []
nd_num_targs = []
sd_num_targs = []
ad_num_targs = []
min_nd_num_targ = 1000
min_sd_num_targ = 1000
min_ad_num_targ = 1000
min_nd_num_targ_idx = ''
min_sd_num_targ_idx = ''
min_ad_num_targ_idx = ''
max_nd_num_targ = 0
max_sd_num_targ = 0
max_ad_num_targ = 0
max_nd_num_targ_idx = ''
max_sd_num_targ_idx = ''
max_ad_num_targ_idx = ''

allevents = dict()

# Note: This code written before individual.num_target_events() and friends were implemented.
for task in zip(*evcounts):

    num_targ = task[1][0][0]
    cond = task[2][0]
    subj = task[0][0].split('/',2)[1]
    #print 'Subj',subj,'Num target events:',num_targ, 'Condition:',cond

    if cond == "No Distractor":
        nd_num_targs = np.append(nd_num_targs, num_targ)
        allevents[subj+"nd"] = num_targ
        if num_targ < min_nd_num_targ:
            min_nd_num_targ = num_targ
            min_nd_num_targ_idx = task[0][0]
        if num_targ > max_nd_num_targ:
            max_nd_num_targ = num_targ
            max_nd_num_targ_idx = task[0][0]
```

```

elif cond == "Synchronous Distractor":
    sd_num_targs = np.append(sd_num_targs, num_targ)
    allevents[subj+"sd"] = num_targ
    if num_targ < min_sd_num_targ:
        min_sd_num_targ = num_targ
        min_sd_num_targ_idx = task[0][0]
    if num_targ > max_sd_num_targ:
        max_sd_num_targ = num_targ
        max_sd_num_targ_idx = task[0][0]
elif cond == "Asynchronous Distractor":
    ad_num_targs = np.append(ad_num_targs, num_targ)
    allevents[subj+"ad"] = num_targ
    if num_targ < min_ad_num_targ:
        min_ad_num_targ = num_targ
        min_ad_num_targ_idx = task[0][0]
    if num_targ > max_ad_num_targ:
        max_ad_num_targ = num_targ
        max_ad_num_targ_idx = task[0][0]
num_targs = np.append(num_targs, num_targ)

print 'Mean number of target events:'
print 'Overall:', num_targs.mean(), '(' , num_targs.std(), ')', 'ND:', nd_num_targs.mean(), 'SD:', sd,
print 'Min nums:'
print 'ND:', min_nd_num_targ, min_nd_num_targ_idx, 'SD:', min_sd_num_targ, min_sd_num_targ_idx, 'AD'
print 'Max nums:'
print 'ND:', max_nd_num_targ, max_nd_num_targ_idx, 'SD:', max_sd_num_targ, max_sd_num_targ_idx, 'AD'

```

Mean number of target events:

Overall: 64.703030303 (6.8721874621) ND: 65.5636363636 SD: 64.8545454545 AD: 63.6909090909

Min nums:

ND: 56.0 Aizat/EB1/line/20141204160911.txt SD: 42.0 Katie/JS/line/20141117163927.txt AD 37.0 Rachel/AW1

Max nums:

ND: 75.0 Rachel/AW1/line/20141118163434.txt SD: 75.0 Rachel/EB2/line/20141118165737.txt AD 73.0 Jon/SY

1.3 Analysis code

The following code block provides a number of global functions and a class which are used in the rest of this analysis. It's the majority of the code used in the analysis here.

The class called individual stores the latency data for each individual and has a set of methods for analysing the data.

In [14]: `from __future__ import division`

```

import numpy as np
import random

def getfnameid (filename):
    # idarr[1] is the ID, idarr[0] is the experimenter, idarr[3] is the dated filename.
    idarr = filename.split('/')
    return idarr[1]

# From the condition string, return an index for the condition. 0 is
# "No Distractor trial", 1 is "Synchronous Distractor trail", 2 is
# "Asynchronous Distractor trial".
def getcondition (condition_string):

```

```

condition_index = -1
if 'No Dist' in condition_string:
    condition_index = 0
elif 'Synchro' in condition_string:
    condition_index = 1
elif 'Asynchr' in condition_string:
    condition_index = 2
return condition_index

# A Single Factor ANOVA calculation for three datasets
def group_anova(nodist_latencies, sync_latencies, async_latencies):

    all_latencies = np.concatenate((nodist_latencies, sync_latencies, async_latencies))

    # Compute grand mean
    grand_mean = all_latencies.mean()
    #print 'Grand mean:', grand_mean, '(', all_latencies.var(ddof=1), ') not ', all_latencies.var(d

    # Compute within-group variance
    tmp1 = all_latencies
    np.power(tmp1, 2)
    within_group_dof = all_latencies.size-3
    within_group_variance = tmp1.sum()/within_group_dof
    #print 'within_group_variance', within_group_variance

    nodist_mean = nodist_latencies.mean()
    sync_mean = sync_latencies.mean()
    async_mean = async_latencies.mean()

    # Compute among-group variance
    tmp1 = np.power (grand_mean - nodist_mean, 2)*nodist_latencies.size
    tmp2 = np.power (grand_mean - sync_mean, 2)*sync_latencies.size
    tmp3 = np.power (grand_mean - async_mean, 2)*async_latencies.size
    sosquares = tmp1 + tmp2 + tmp3
    between_group_dof = 2 # 3 conditions => 3 groups, so 3-1 DOF
    between_group_variance = sosquares / between_group_dof
    #print 'between_group_variance', between_group_variance

    # Now compute the F ratio
    F = between_group_variance/within_group_variance

    # Lastly, what's the probability for this?
    P = 1-special.fdftr(between_group_dof, within_group_dof, F)

    return (F, between_group_dof, within_group_dof, P)

# Convert an omit reason id to a reason string
def omit_reason (or_id):
    reason_str = ""
    if or_id == 0:
        reason_str = "0 not omitted"
    if or_id == 1:
        reason_str = "1 target posn change insignificant"
    elif or_id == 2:

```

```

        reason_str = "2 target posn held less than min. duration"
    elif or_id == 3:
        reason_str = "3 target posn change less than min. jump size"
    elif or_id == 4:
        reason_str = "4 Stable position later than event onset"
    elif or_id == 5:
        reason_str = "5 Another event caused this movement"
    elif or_id == 6:
        reason_str = "6 Not distracted"
    elif or_id == 7:
        reason_str = "7 Too fast (targ)" # faster than A.fastest_brain_decision
    elif or_id == 8:
        reason_str = "8 Too fast (distractor)" # faster than A.fastest_brain_decision
    elif or_id == 9:
        reason_str = "9 No movement detected"
    elif or_id == 10:
        reason_str = "10 Failed to find stable stylus posn"
    elif or_id == 11:
        reason_str = "11 Stable stylus period too short"
    elif or_id == 12:
        reason_str = "12 Drift too great during stable stylus period"
    elif or_id == 13:
        reason_str = "13 Drift too great during stable period (avg)"
    elif or_id == 14:
        reason_str = "14 Stylus moving at event onset"
    elif or_id == 15:
        reason_str = "15 Stylus didn't move away from target"
    elif or_id == 16:
        reason_str = "16 Movement occurs beyond next target"
    elif or_id == 17:
        reason_str = "17 Subject was distracted by closely previous distractor"
    elif or_id == 18:
        reason_str = "18 Incorrect move was recorded in previous distractor event"
    elif or_id == 19:
        reason_str = "19 This distractor event did not distract the stylus movement"
    elif or_id == 20:
        reason_str = "20 Recorded this stylus movement as a distraction towards the next distractor"

    return reason_str

# Take n sub-samples from distn
def subsample (distn, n):
    distn_cp = distn
    np.random.seed(19742016)
    counter = 0
    subsamp = []
    if n > len(distn_cp):
        print 'No possible to get',n,'samples from this distribution!'
        return subsamp
    while counter < n:
        i = 0
        for s in distn_cp:
            if np.random.uniform()>0.5:
                np.delete(distn_cp, i)

```

```

        i -= 1
        subsamp = np.append(subsamp, s)
        counter = counter + 1
    if counter >= n:
        break
    i += 1
return subsamp

def compute_mad_outliers (points, thresh=3.5):
    """
    Returns a boolean array with True if points are outliers and False
    otherwise.

    Parameters:
    -----
        points : An numobservations by numdimensions array of observations
        thresh : The modified z-score to use as a threshold. Observations with
            a modified z-score (based on the median absolute deviation) greater
            than this value will be classified as outliers.

    Returns:
    -----
        mask : A numobservations-length boolean array.

    References:
    -----
        Boris Iglewicz and David Hoaglin (1993), "Volume 16: How to Detect and
        Handle Outliers", The ASQC Basic References in Quality Control:
        Statistical Techniques, Edward F. Mykytka, Ph.D., Editor.
    """
    if len(points.shape) == 1:
        points = points[:,None]
    median = np.median(points, axis=0)
    # Compute root mean square deviation from median:
    diff = np.sum((points - median)**2, axis=-1)
    diff = np.sqrt(diff)
    # The median of this is the median abs. deviation from the median.
    med_abs_deviation = np.median(diff)
    # A modified Z score is analogous to the Z score for number of SDs from the mean.
    modified_z_score = 0.6745 * diff / med_abs_deviation

    return modified_z_score > thresh

def compute_mad (points):
    """
    Returns the median abs deviation value for the points

    Parameters:
    -----
        points : An numobservations by numdimensions array of observations

    Returns:
    -----
        med_abs_deviation: The median absolute deviation

```

References:

Boris Iglewicz and David Hoaglin (1993), "Volume 16: How to Detect and Handle Outliers", The ASQC Basic References in Quality Control: Statistical Techniques, Edward F. Mykytka, Ph.D., Editor.

"""

```
if len(points.shape) == 1:
    points = points[:,None]
median = np.median(points, axis=0)
# Compute root mean square deviation from median:
diff = np.sum((points - median)**2, axis=-1)
diff = np.sqrt(diff)
# The median of this is the median abs. deviation from the median.
med_abs_deviation = np.median(diff)
return med_abs_deviation
```

Libs used in class individual

```
from scipy import special
from scipy import stats
import statsmodels.api as sm
from matplotlib import pyplot as plt
import matplotlib.lines as mlines
```

A class for an individual's data.

```
class individual:
```

```
    def __init__(self, subj_id):
```

Hindsight note: I should have made an "event" class.

```
    self.subj_id = subj_id;
    self.idnum = -1
    self.filename = ""
```

Default number of subsamples to take from a distribution

```
    self.numSubsamples = 20
```

All data, errorred and ok, for each condition.

```
    self.alldata_nd = np.ndarray(0)
    self.alldata_sd = np.ndarray(0)
    self.alldata_ad = np.ndarray(0)
```

These are the "non-movement-error target latencies"

```
    self.nodist_latencies = np.ndarray(0)
    self.sync_latencies = np.ndarray(0)
    self.async_latencies = np.ndarray(0)
    self.nodist_latencies_rank = np.ndarray(0)
    self.sync_latencies_rank = np.ndarray(0)
    self.async_latencies_rank = np.ndarray(0)
```

These are the "non-movement-error distractor latencies"

```
    self.nodist_dgood_latencies = np.ndarray(0)
    self.sync_dgood_latencies = np.ndarray(0)
    self.async_dgood_latencies = np.ndarray(0)
```

```

self.nodist_dgood_latencies_rank = np.ndarray(0)
self.sync_dgood_latencies_rank = np.ndarray(0)
self.async_dgood_latencies_rank = np.ndarray(0)

# These are the "target event error latencies"
self.nodist_err_latencies = np.ndarray(0)
self.sync_err_latencies = np.ndarray(0)
self.async_err_latencies = np.ndarray(0)
self.nodist_err_latencies_rank = np.ndarray(0)
self.sync_err_latencies_rank = np.ndarray(0)
self.async_err_latencies_rank = np.ndarray(0)

# These are the "distractor event error latencies" (N/A for ND or for Sync, where targ
self.nodist_derr_latencies = np.ndarray(0)
self.sync_derr_latencies = np.ndarray(0)
self.async_derr_latencies = np.ndarray(0)
self.nodist_derr_latencies_rank = np.ndarray(0)
self.sync_derr_latencies_rank = np.ndarray(0)
self.async_derr_latencies_rank = np.ndarray(0)

self.async_derr_timesince = np.ndarray(0)
self.async_tnoerr_timesince = np.ndarray(0)

self.nodist_dirns = np.ndarray(0)
self.sync_dirns = np.ndarray(0)
self.async_dirns = np.ndarray(0)

self.n_errors_per_target_nd = 0
self.n_errors_per_distractor_sync = 0
self.n_errors_per_distractor_async = 0

# Number of distractor events in the asynchronous case
self.n_async_distractors = 0

# Some constants
self.DIST_EVENT = 0
self.TARG_EVENT = 1

def csvheader(self):
    hl = 'Experimenter,Subject,Distractor type,RT (M),RT (SD),N correct,RT incorrect (M),'
    hl += 'RT incorrect (SD),N incorrect>Error rate,D1 (-ve=>error)'
    for i in range(2,65):
        hl += ',D{0}'.format(i)
    hl += '\n'
    return hl

def ad_omit_reasons (self, event_type):
    event_indices = [i for i,elem in enumerate(self.alldata_ad[:,1]) if elem == event_type]
    # Last col gives omit reason
    return np.bincount(self.alldata_ad[event_indices,6].astype(int),minlength=21)

def sd_omit_reasons (self, event_type):
    event_indices = [i for i,elem in enumerate(self.alldata_sd[:,1]) if elem == event_type]
    # Last col gives omit reason

```



```

        return np.bincount(self.alldata_sd[event_indices,6].astype(int),minlength=21)

def nd_omit_reasons (self, event_type):
    event_indices = [i for i,elem in enumerate(self.alldata_nd[:,1]) if elem == event_type]
    # Last col gives omit reason
    return np.bincount(self.alldata_nd[event_indices,6].astype(int),minlength=21)

# Experimenter,Subject,Distractor type,RT (M),RT (SD),N correct,RT incorrect (M),
# RT incorrect (SD),N incorrect>Error rate,Delays (negative = incorrect)
def csvlineset(self):
    line = self.filename_nd + ',' + self.subj_id + ',0,'
    # ND mean/SD. NB: /1000 to output numbers as seconds, like Mauro reports.
    line += '{0},{1}'.format(self.nodist_mean()/1000,self.nodist_latencies.std()/1000)
    # ND RT incorrect mean/SD
    line += '{0},{1},{2}'.format(self.num_tgood(0),self.nodist_err_mean()/1000,self.nodis
    # ND num incorrect, error rate
    line += '{0},{1}'.format(self.num_tmoveerrors(0),ind.num_tmoveerrors_percent(0))
    # ND delays
    #for i in self.nodist_latencies[:,1]:
    #    line += ',{0}'.format(i/1000)
    for i in self.alldata_nd:
        if i[5]==1:
            line += ',OR{0}'.format(omit_reason(i[6]))
        else:
            if i[2]==0: # ND and no error
                line += ',{0}'.format(i[4]/1000)
            elif i[2]==1: # ND and error
                # make errorred ones -ve as Mauro does.
                line += ',{0}'.format(-i[4]/1000)

    line += '\n'

    line += self.filename_sd + ',' + self.subj_id + ',1,'
    # SD mean/SD
    line += '{0},{1}'.format(self.sync_mean()/1000,self.sync_latencies.std()/1000)
    # SD RT incorrect mean/SD
    line += '{0},{1},{2}'.format(self.num_tgood(1),self.sync_err_mean()/1000,self.sync_er
    # SD num incorrect, error rate
    line += '{0},{1}'.format(self.num_tmoveerrors(1),ind.num_tmoveerrors_percent(1))
    # SD delays
    for i in self.alldata_sd:
        if i[5]==1:
            line += ',OR{0}'.format(omit_reason(i[6]))
        else:
            if i[2]==0: # SD and no error
                line += ',{0}'.format(i[4]/1000)
            elif i[2]==1: # SD and error
                # make errorred ones -ve as Mauro does.
                line += ',{0}'.format(-i[4]/1000)

    line += '\n'

    line += self.filename_ad + ',' + self.subj_id + ',2,'
    # AD mean/SD
    line += '{0},{1}'.format(self.async_latencies.mean()/1000,self.async_latencies.std()/

```

```

# AD RT incorrect mean/SD
line += '{0},{1},{2},'.format(self.num_tgood(2),self.async_err_mean()/1000,self.async_
# AD num incorrect, error rate
line += '{0},{1}'.format(self.num_tmoveerrors(2),ind.num_tmoveerrors_percent(2))
# AD delays
for i in self.alldata_ad:
    if i[5]==1:
        line += ',OR{0}'.format(omit_reason(i[6]))
    else:
        if i[1]==1 and i[2]==0: # TARG_EVENT and AD and no error
            line += ',{0}'.format(i[4]/1000)
        elif i[1]==1 and i[2]==1: # TARG_EVENT and AD and error
            # make errorred ones -ve as Mauro does.
            line += ',{0}'.format(-i[4]/1000)
line += '\n'

return line

# Compute ANOVA for this individual
def anova(self):
    F, between_group_dof, within_group_dof, P = group_anova (self.nodist_latencies[:,1], s
    return (F, between_group_dof, within_group_dof, P)

def reportmeans (self):
    print "Mean(SD): No distr: {0:.2f} ({1:.2f}) Sync: {2:.2f} ({3:.2f}) Async: {4:.2f} ({

# Batch up all data in a form suitable for statsmodel's MultiComparison class. This means
# concatenating the ND, SD & AD data into a single array, and making a "label" array to mat
def getMultiComparisonData (self):
    d = np.hstack((self.nodist_latencies[:,1],self.sync_latencies[:,1],self.async_latencies

    nd_labels = np.ndarray(shape=(self.num_tgood(0),), dtype=object)
    nd_labels.fill('ND')

    sd_labels = np.ndarray(shape=(self.num_tgood(1),), dtype=object)
    sd_labels.fill('SD')

    ad_labels = np.ndarray(shape=(self.num_tgood(2),), dtype=object)
    ad_labels.fill('AD')

    l = np.hstack((nd_labels,sd_labels,ad_labels))

    # d is the data array, l is the label array.
    return (d, l)

# Do a full set of graphs to show the normality of the data. Show QQ plots,
# histograms of the distributions and results of Shapiro-Wilks tests for comparison.
# Pass in the significance level for the S-W test.
def shownormality (self, alpha):
    f, axarr = plt.subplots(3, 2)

    #ax1.set_title('QQ plots')
    fig1 = sm.qqplot(self.nodist_latencies[:,1], fit=True, line='45',ax=axarr[0,0])
    fig2 = sm.qqplot(self.sync_latencies[:,1], fit=True, line='45',ax=axarr[1,0])

```

```

fig3 = sm.qqplot(self.async_latencies[:,1], fit=True, line='45', ax=axarr[2,0])
axarr[0,0].set_title('QQ Plots')

W, p = stats.shapiro (subsample(self.nodist_latencies[:,1], self.numSubsamples))
isNormal = False
if p > alpha:
    isNormal = True
sw = 'ND. Mean/SD:{2:.2f}/{3:.2f} W={0:.2f}, p={1:.2f} (Normal:{4})'.format(W,p,self.nodist_latencies[:,1])
axarr[0,1].hist(self.nodist_latencies[:,1], bins=20, label=sw)
axarr[0,1].legend(prop={'size':9})
axarr[0,1].set_title('Dist\'ns with Shapiro-Wilks stats')

W, p = stats.shapiro (subsample(self.sync_latencies[:,1], self.numSubsamples))
isNormal = False
if p > alpha:
    isNormal = True
sw = 'SD. Mean/SD:{2:.2f}/{3:.2f} W={0:.2f}, p={1:.2f} (Normal:{4})'.format(W,p,self.sync_latencies[:,1])
axarr[1,1].hist(self.sync_latencies[:,1], bins=20, label=sw)
axarr[1,1].legend(prop={'size':9})

W, p = stats.shapiro (subsample(self.async_latencies[:,1], self.numSubsamples))
isNormal = False
if p > alpha:
    isNormal = True
sw = 'AD. Mean/SD:{2:.2f}/{3:.2f} W={0:.2f}, p={1:.2f} (Normal:{4})'.format(W,p,self.async_latencies[:,1])
axarr[2,1].hist(self.async_latencies[:,1], bins=20, label=sw)
axarr[2,1].legend(prop={'size':9})

# Fine-tune figure; make subplots close to each other and hide x ticks for
# all but bottom plot.
f.subplots_adjust(hspace=0)
plt.setp([a.get_xticklabels() for a in f.axes[:-1]], visible=False)

savename = 'images/' + self.subj_id + 'normplot.png'
plt.savefig(savename)
plt.show()

# Apply Shapiro-Wilk test. Null hypothesis is that the data are normally
# distributed. If p < alpha then null hypothesis must be rejected and data
# cannot be considered to be normally distributed.
def shapiroWilk (self, condition, alpha):

    W = -1
    p = -1
    isNormal = False

    if condition == 0:
        W, p = stats.shapiro (subsample(self.nodist_latencies[:,1], self.numSubsamples))
    elif condition == 1:
        ss = subsample(self.sync_latencies[:,1], self.numSubsamples)
        W, p = stats.shapiro (ss)
    elif condition == 2:
        W, p = stats.shapiro (subsample(self.async_latencies[:,1], self.numSubsamples))
    # else leave W,p,isNormal with default values

```

```

        if p > alpha:
            isNormal = True

    return W, p, isNormal

# Do a Quantile-Quantile plot to compare against normal distribution
def qqplot (self):
    f, (ax1, ax2, ax3) = plt.subplots(3, sharex=True, sharey=True)
    ax1.set_title('QQ plots',fontsize=18)
    fig1 = sm.qqplot(self.nodist_latencies[:,1], fit=True, line='45',ax=ax1)
    fig2 = sm.qqplot(self.sync_latencies[:,1], fit=True, line='45',ax=ax2)
    fig3 = sm.qqplot(self.async_latencies[:,1], fit=True, line='45', ax=ax3)
    # Fine-tune figure; make subplots close to each other and hide x ticks for
    # all but bottom plot.
    f.subplots_adjust(hspace=0)
    plt.setp([a.get_xticklabels() for a in f.axes[:-1]], visible=False, fontsize=18)
    plt.show()
    return f

def boxplot (self):
    data = [self.nodist_latencies[:,1], self.sync_latencies[:,1], self.async_latencies[:,1]]
    plt.figure()
    plt.boxplot(data, 0, 'gD')
    nodist_cond_x = np.ones(self.nodist_latencies[:,1].size)-0.1
    sync_cond_x = 2*np.ones(self.sync_latencies[:,1].size)-0.1
    async_cond_x = 3*np.ones(self.async_latencies[:,1].size)-0.1
    nodist_pts = plt.scatter(nodist_cond_x, self.nodist_latencies[:,1])
    sync_pts = plt.scatter(sync_cond_x, self.sync_latencies[:,1])
    async_pts = plt.scatter(async_cond_x, self.async_latencies[:,1])
    plt.xlabel('Condition 1:ND 2:S 3:AS',fontsize=18)
    plt.ylabel('Latency (ms)',fontsize=18)
    savename = 'images/' + self.subj_id + 'boxplot.png'
    plt.savefig(savename)
    plt.show()

# A standard method for excluding outliers
def excludeOutliers (self, show_excluded=0):
    # Exclude from latency data
    if show_excluded:
        print 'Excluding good movement outliers...'
        self.excludeOutliers_mad_based(3.5,show_excluded)
    # Exclude from error data:
    if show_excluded:
        print 'Excluding error outliers...'
        self.excludeErrOutliers(3.5,show_excluded)# also mad based

# subroutine of excludeOutliers_mad_based()
def excludeOutliers_mad_based(self, latencies, alldata, thresh, show_excluded=0):
    nd_outliers = compute_mad_outliers (latencies,thresh)
    # nd_outliers is a large array of True/False.
    # print 'nd_outliers:',nd_outliers
    del_indices = [i for i,elem in enumerate(nd_outliers) if elem == True]
    # del_indices are the indices of the True values in nd_outliers

```

```

if not del_indices:
    return latencies, alldata

# The event number is not the same as the index into latencies
event_indices = latencies[[del_indices]][:,0]

if show_excluded:
    values_to_delete = latencies[[del_indices]][:,1]
    print 'deleting the following events', event_indices, 'which have values', values_to_delete

keepmask_bool = np.invert(np.in1d(latencies[:,0], event_indices, assume_unique=True))
latencies = latencies[keepmask_bool]
# Remove same indices from self.alldata:
keepmask_bool = np.invert(np.in1d(alldata[:,0], event_indices, assume_unique=True))
alldata = alldata[keepmask_bool]
return latencies, alldata

# Exclude outliers from each latency set based on median absolute deviation method.
def excludeOutliers_mad_based(self, thresh=3.5, show_excluded=0):
    if show_excluded:
        print 'exclude nodist latencies for subject ', self.subj_id, 'filename', self.filename
        self.nodist_latencies, self.alldata_nd = self.excludeOutliers_mad_based(self.nodist_latencies, self.alldata_nd)
    if show_excluded:
        print 'exclude sync latencies for subject ', self.subj_id, 'filename', self.filename
        self.sync_latencies, self.alldata_sd = self.excludeOutliers_mad_based(self.sync_latencies, self.alldata_sd)
    if show_excluded:
        print 'exclude async latencies for subject ', self.subj_id, 'filename', self.filename
        self.async_latencies, self.alldata_ad = self.excludeOutliers_mad_based(self.async_latencies, self.alldata_ad)

def excludeErrOutliers (self, num_sds, show_excluded):
    if show_excluded:
        print 'exclude nodist latencies for subject ', self.subj_id, 'filename', self.filename
        self.nodist_err_latencies, self.alldata_nd = self.excludeOutliers_mad_based(self.nodist_latencies, self.alldata_nd)
    if show_excluded:
        print 'exclude sync latencies for subject ', self.subj_id, 'filename', self.filename
        self.sync_err_latencies, self.alldata_sd = self.excludeOutliers_mad_based(self.sync_latencies, self.alldata_sd)
    if show_excluded:
        print 'exclude async latencies for subject ', self.subj_id, 'filename', self.filename
        self.async_err_latencies, self.alldata_ad = self.excludeOutliers_mad_based(self.async_latencies, self.alldata_ad)

def randomly_subsample_data (self, num_data):
    random.seed(19742016)
    while len(self.nodist_latencies[:,1]) > num_data:
        remove_this = random.randint (0, len(self.nodist_latencies[:,1])-1)
        self.nodist_latencies = np.delete(self.nodist_latencies, remove_this, axis=0)
    while len(self.sync_latencies[:,1]) > num_data:
        remove_this = random.randint (0, len(self.sync_latencies[:,1])-1)
        self.sync_latencies = np.delete(self.sync_latencies, remove_this, axis=0)
    while len(self.async_latencies[:,1]) > num_data:
        remove_this = random.randint (0, len(self.async_latencies[:,1])-1)
        self.async_latencies = np.delete(self.async_latencies, remove_this, axis=0)

def graph1(self):
    print 'Showing graph for ', ind.subj_id

```

```

means = (self.nodist_mean(), self.sync_mean(), self.async_mean())
stds = (self.nodist_std(), self.sync_std(), self.async_std())
index = np.arange(3)
opacity = 0.4
error_config = {'ecolor': '0.3'}
rects1 = plt.bar(index, means, 0.2,
                  alpha=opacity,
                  color='b',
                  yerr=stds,
                  error_kw=error_config,
                  label=ind.subj_id)
# Now draw the points on a scatter graph
nodist_cond_x = np.zeros(self.nodist_latencies[:,1].size)
sync_cond_x = np.ones(self.sync_latencies[:,1].size)
async_cond_x = 2*np.ones(self.async_latencies[:,1].size)
nodist_pts = plt.scatter(nodist_cond_x, self.nodist_latencies)
sync_pts = plt.scatter(sync_cond_x, self.sync_latencies)
async_pts = plt.scatter(async_cond_x, self.async_latencies)
plt.xlabel('Condition 0:ND 1:S 2:AS')
plt.ylabel('Latency (ms)')
plt.title(self.subj_id)
plt.show()
return

# Compute the sum of the squared displacements from the mean for all three conditions
def sumofsquare_displacements_all_from_value(self, value):
    sos = self.sumofsquare_displacements_from_value(0,value) + self.sumofsquare_displacements(1) + self.sumofsquare_displacements(2)
    return sos

# Compute the sum of the squared displacements from the mean for all three conditions
def sumofsquare_displacements_all(self):
    sos = self.sumofsquare_displacements(0) + self.sumofsquare_displacements(1) + self.sumofsquare_displacements(2)
    return sos

# Compute the sum of the squared displacements from the mean for the given condition
def sumofsquare_displacements(self, condition):
    if condition == 0:
        mn = self.nodist_mean()
        squares = np.power((self.nodist_latencies[:,1] - mn), 2)
    elif condition == 1:
        mn = self.sync_mean()
        squares = np.power((self.sync_latencies[:,1] - mn), 2)
    else: # condition 2
        mn = self.async_mean()
        squares = np.power((self.async_latencies[:,1] - mn), 2)
    sos = np.sum(squares)
    return sos

# Compute the sum of the squared displacements from the mean for the given condition
def sumofsquare_displacements_from_value(self, condition, value):
    if condition == 0:
        squares = np.power((self.nodist_latencies[:,1] - value), 2)
        # Verification of this method:
        #squares_alt = 0

```

```

        #for i in self.nodist_latencies:
        #    imv = i - value
        #    squares_alt += imv*imv
        #print 'nd sum of squares:',np.sum(squares),'squares_alt:',squares_alt
    elif condition == 1:
        squares = np.power((self.sync_latencies[:,1] - value), 2)
    else: # condition 2
        squares = np.power((self.async_latencies[:,1] - value), 2)
    sos = np.sum(squares)
    return sos

def alldata_for_condition(self, condition):
    if condition == 0:
        thedata = self.alldata_nd
    elif condition == 1:
        thedata = self.alldata_sd
    else: # condition 2
        thedata = self.alldata_ad
    return thedata

def num_target_omissions(self, condition):
    thedata = self.alldata_for_condition(condition)
    # Select rows for which omit==1:
    omit_indices = [i for i,elem in enumerate(thedata[:,5]) if elem == 1]
    num_omit = 0
    if omit_indices:
        # From those rows, select those for which event type is target (1)
        omit_target_indices = [i for i,elem in enumerate(thedata[omit_indices,1]) if elem == 1]
        if omit_target_indices:
            num_omit = thedata[[omit_target_indices]][:,4].size

    return num_omit

def num_target_non_omissions(self, condition):
    thedata = self.alldata_for_condition(condition)
    # Select rows for which omit==0:
    nomit_indices = [i for i,elem in enumerate(thedata[:,5]) if elem == 0]
    num_nomit = 0
    if nomit_indices:
        # From those rows, select those for which event type is target (1)
        nomit_target_indices = [i for i,elem in enumerate(thedata[nomit_indices,1]) if elem == 1]
        if nomit_target_indices:
            num_nomit = thedata[[nomit_target_indices]][:,4].size

    return num_nomit

def num_distractor_events(self, condition):
    # In ND case, num_distractor_events == 0 by definition
    num = 0
    if condition == 0:
        return num

    # In SD case, num_distractor_events == num_target_events

```

```

        if condition == 1:
            return self.num_target_events(condition)

        thedata = self.alldata_for_condition(condition)
        distractor_indices = [i for i,elem in enumerate(thedata[:,1]) if elem == 0]
        if distractor_indices:
            num = thedata[[distractor_indices]][:,4].size
        return num

def num_target_events(self, condition):
    thedata = self.alldata_for_condition(condition)
    distractor_indices = [i for i,elem in enumerate(thedata[:,1]) if elem == 1]
    num = 0
    if distractor_indices:
        num = thedata[[distractor_indices]][:,4].size
    return num

def num_distractor_omissions(self, condition):
    thedata = self.alldata_for_condition(condition)

    # Select rows for which omit==1:
    omit_indices = [i for i,elem in enumerate(thedata[:,5]) if elem == 1]
    num_omit = 0
    if omit_indices:
        # From those rows, select those for which event type is distractor (0)
        omit_distractor_indices = [i for i,elem in enumerate(thedata[omit_indices,1]) if elem == 0]
        if omit_distractor_indices:
            num_omit = thedata[[omit_distractor_indices]][:,4].size

    return num_omit

def num_distractor_non_omissions(self, condition):
    thedata = self.alldata_for_condition(condition)

    # Select rows for which omit==0:
    nomit_indices = [i for i,elem in enumerate(thedata[:,5]) if elem == 0]
    num_nomit = 0
    if nomit_indices:
        # From those rows, select those for which event type is distractor (0)
        nomit_distractor_indices = [i for i,elem in enumerate(thedata[nomit_indices,1]) if elem == 0]
        if nomit_distractor_indices:
            num_nomit = thedata[[nomit_distractor_indices]][:,4].size

    return num_nomit

# Movements following target events which are good and without error
def num_tgood(self, condition):
    if condition == 0:
        if self.nodist_latencies.size<2:
            return 0
        return self.nodist_latencies[:,1].size
    elif condition == 1:
        if self.sync_latencies.size<2:
            return 0

```



```

        return self.sync_latencies[:,1].size
    else: # condition 2
        if self.async_latencies.size<2:
            return 0
        return self.async_latencies[:,1].size

def num_tgood_all(self):
    n = self.num_tgood(0) + self.num_tgood(1) + self.num_tgood(2)
    return n

# num_tgood_percent. Number of good movements as a percentage of number of target events
def num_tgood_percent(self, condition):
    if condition == 0:
        if self.nodist_latencies.size<2:
            return 0
        return 100*self.nodist_latencies[:,1].size/self.num_target_events(condition)
    elif condition == 1:
        if self.sync_latencies.size<2:
            return 0
        return 100*self.sync_latencies[:,1].size/self.num_target_events(condition)
    else: # condition 2
        if self.async_latencies.size<2:
            return 0
        #prop_targets_non_omitted = self.num_target_non_omissions(condition)/self.num_targ
        #representative_num_distractors = prop_targets_non_omitted * self.num_distractor_e
        return 100*self.async_latencies[:,1].size/self.num_target_events(condition)

# moveerrors are "errors for target movements"
def num_tmoveerrors_all(self):
    n = self.num_tmoveerrors(0) + self.num_tmoveerrors(1) + self.num_tmoveerrors(2)
    return n

# Number of target movement errors
def num_tmoveerrors(self, condition):

    #print 'num_tmoveerrors(',condition,') called'
    if condition == 0:
        #print 'nodist_err_latencies.size:',self.nodist_err_latencies.size
        if self.nodist_err_latencies.size<2:
            #print 'Returning 0 for nodist_err_latencies:',self.nodist_err_latencies
            return 0
        elif self.nodist_err_latencies.size == 2:
            #print 'Returning 1 for nodist_err_latencies:',self.nodist_err_latencies
            return 1
        #print 'Returning[:,1].size for nodist_err_latencies'
        return self.nodist_err_latencies[:,1].size
    elif condition == 1:
        #print 'sync_err_latencies.size:',self.sync_err_latencies.size
        if self.sync_err_latencies.size<2:
            return 0
        elif self.sync_err_latencies.size == 2:
            return 1
        return self.sync_err_latencies[:,1].size
    else: # condition 2

```

```

        #print 'async_err_latencies.size:',self.async_err_latencies.size
        if self.async_err_latencies.size<2:
            return 0
        elif self.async_err_latencies.size == 2:
            return 1
        return self.async_err_latencies[:,1].size

# Number of target movement errors as a percentage of number of target events
def num_tmoveerrors_percent(self, condition):
    if condition == 0:
        if self.nodist_err_latencies.size<2:
            return 0
        return 100*self.nodist_err_latencies[:,1].size/self.num_target_non_omissions(condition)
    elif condition == 1:
        if self.sync_err_latencies.size<2:
            return 0
        return 100*self.sync_err_latencies[:,1].size/self.num_target_non_omissions(condition)
    else: # condition 2
        if self.async_err_latencies.size<2:
            return 0
        return 100*self.async_err_latencies[:,1].size/self.num_target_non_omissions(condition)

# dmoveerrors are : Distractor move errors
def num_dmoveerrors_all(self):
    n = self.num_dmoveerrors(0) + self.num_dmoveerrors(1) + self.num_dmoveerrors(2)
    return n

def num_dmoveerrors(self, condition):
    if condition == 0:
        if self.nodist_derr_latencies.size<2:
            return 0
        return self.nodist_derr_latencies[:,1].size
    elif condition == 1:
        if self.sync_derr_latencies.size<2:
            return 0
        return self.sync_derr_latencies[:,1].size
    else: # condition 2
        if self.async_derr_latencies.size<2:
            return 0
        return self.async_derr_latencies[:,1].size

# number of detected distractor errors as a percentage of the
# total number of distractor events
def num_dmoveerrors_percent(self, condition):
    if condition == 0:
        return 0 # 0 by definition
    elif condition == 1:
        if self.sync_derr_latencies.size<2:
            return 0
        return 100*self.sync_derr_latencies[:,1].size/self.num_distractor_events(condition)
    else: # condition 2
        if self.async_derr_latencies.size<2:
            return 0
        return 100*self.async_derr_latencies[:,1].size/self.num_distractor_events(condition)

```

```

# Total number of movement errors (target and distractor) as a
# percentage of the number of target events not omitted
def num_moveerrors_per_target_percent(self, condition):
    moveerrors_percent = 0
    if condition == 0:
        # same as num_tmoveerrors_percent:
        moveerrors_percent = self.num_tmoveerrors_percent(condition)
    elif condition == 1:
        numerr = self.num_tmoveerrors(condition)+self.num_dmoveerrors(condition)
        moveerrors_percent = 100 * numerr / self.num_target_non_omissions(condition)
    else: # condition == 2
        numerr = self.num_tmoveerrors(condition)+self.num_dmoveerrors(condition)
        moveerrors_percent = 100 * numerr / self.num_target_non_omissions(condition)
    return moveerrors_percent

# as above, but not as a percentage
def num_moveerrors_per_target(self, condition):
    moveerrors_ = 0
    if condition == 0 or condition == 1:
        # same as num_tmoveerrors_ for the ND and SD cases
        moveerrors_ = self.num_tmoveerrors_percent(condition)/100
    else: # condition == 2
        numerr = self.num_tmoveerrors(condition)+self.num_dmoveerrors(condition)
        moveerrors_ = numerr / self.num_target_non_omissions(condition)
    return moveerrors_

# Total number of movement errors as a percentage of "the
# proportion of all distractor events matching the proportion of
# target events not omitted from latency measurement"
def num_moveerrors_per_distractor_percent(self, condition):
    moveerrors_percent = 0
    prop_target_events = self.num_target_non_omissions(condition)/self.num_target_events(c
    if condition == 0:
        moveerrors_percent = -1 # nan really
    elif condition == 1:
        # For the SD condition, target moveerrors == distractor moveerrors
        moveerrors_percent = self.num_moveerrors_per_target_percent(condition)
    else: # condition == 2
        numerr = self.num_tmoveerrors(condition)+self.num_dmoveerrors(condition)
        moveerrors_percent = 100 * numerr / self.num_distractor_events(condition)*prop_tar
    return moveerrors_percent

# dgood : Distractor events without a move error
def num_dgood_all(self):
    n = self.num_dgood(0) + self.num_dgood(1) + self.num_dgood(2)
    return n

def num_dgood(self, condition):
    if condition == 0:
        if self.nodist_dgood_latencies.size<2:
            return 0
        return self.nodist_dgood_latencies[:,1].size
    elif condition == 1:

```

```

        if self.sync_dgood_latencies.size<2:
            return 0
        return self.sync_dgood_latencies[:,1].size
    else: # condition 2
        if self.async_dgood_latencies.size<2:
            return 0
        return self.async_dgood_latencies[:,1].size

def num_dgood_percent(self, condition):
    if condition == 0:
        if self.nodist_dgood_latencies.size<2:
            return 0
        return 100*self.nodist_dgood_latencies[:,1].size/(self.num_tgood(condition)+self.num)
    elif condition == 1:
        if self.sync_dgood_latencies.size<2:
            return 0
        return 100*self.sync_dgood_latencies[:,1].size/(self.num_tgood(condition)+self.num)
    else: # condition 2
        if self.async_dgood_latencies.size<2:
            return 0
        return 100*self.async_dgood_latencies[:,1].size/(self.num_tgood(condition)+self.num)

def nodist_mean(self):
    if self.nodist_latencies.size<2:
        return 0
    return self.nodist_latencies[:,1].mean()

def sync_mean(self):
    if self.sync_latencies.size<2:
        return 0
    return self.sync_latencies[:,1].mean()

def async_mean(self):
    if self.async_latencies.size<2:
        return 0
    return self.async_latencies[:,1].mean()

def overall_mean(self):
    all_latencies = np.concatenate((self.nodist_latencies, self.sync_latencies, self.async_latencies))
    return all_latencies[:,1].mean()

def overall_std(self):
    all_latencies = np.concatenate((self.nodist_latencies, self.sync_latencies, self.async_latencies))
    return all_latencies[:,1].std()

def nodist_std(self):
    if self.nodist_latencies.size<2:
        return 0
    return self.nodist_latencies[:,1].std()

def sync_std(self):
    if self.sync_latencies.size<2:
        return 0
    return self.sync_latencies[:,1].std()

```

```

def async_std(self):
    if self.async_latencies.size<2:
        return 0
    return self.async_latencies[:,1].std()

# mean/std accessors for the error latencies:
def nodist_err_mean(self):
    if self.nodist_err_latencies.size<2:
        return 0
    return self.nodist_err_latencies.mean()

def sync_err_mean(self):
    if self.sync_err_latencies.size<2:
        return 0
    return self.sync_err_latencies.mean()

def async_err_mean(self):
    if self.async_err_latencies.size<2:
        return 0
    return self.async_err_latencies.mean()

def overall_err_mean(self):
    all_err_latencies = np.concatenate((self.nodist_err_latencies, self.sync_err_latencies)
    return all_err_latencies[:,1].mean()

def overall_err_std(self):
    all_err_latencies = np.concatenate((self.nodist_err_latencies, self.sync_err_latencies)
    return all_err_latencies[:,1].std()

def nodist_err_std(self):
    if self.nodist_err_latencies.size<2:
        return 0
    return self.nodist_err_latencies.std()

def sync_err_std(self):
    if self.sync_err_latencies.size<2:
        return 0
    return self.sync_err_latencies.std()

def async_err_std(self):
    if self.async_err_latencies.size<2:
        return 0
    return self.async_err_latencies.std()

# mean/std accessors for the distractor error latencies:
def nodist_derr_mean(self):
    if self.nodist_derr_latencies.size<2:
        return 0
    return self.nodist_derr_latencies.mean()

def sync_derr_mean(self):
    if self.sync_derr_latencies.size<2:
        return 0

```

```

        return self.sync_derr_latencies.mean()

def async_derr_mean(self):
    if self.async_derr_latencies.size<2:
        return 0
    return self.async_derr_latencies.mean()

def overall_derr_mean(self):
    all_derr_latencies = np.concatenate((self.nodist_derr_latencies, self.sync_derr_latencies))
    return all_derr_latencies[:,1].mean()

def overall_derr_std(self):
    all_derr_latencies = np.concatenate((self.nodist_derr_latencies, self.sync_derr_latencies))
    return all_derr_latencies[:,1].std()

def nodist_derr_std(self):
    if self.nodist_derr_latencies.size<2:
        return 0
    return self.nodist_derr_latencies.std()

def sync_derr_std(self):
    if self.sync_derr_latencies.size<2:
        return 0
    return self.sync_derr_latencies.std()

def async_derr_std(self):
    if self.async_derr_latencies.size<2:
        return 0
    return self.async_derr_latencies.std()

# mean/std accessors for the distractor non-movement-error latencies:
def nodist_dgood_mean(self):
    if self.nodist_dgood_latencies.size<2:
        return 0
    return self.nodist_dgood_latencies.mean()

def sync_dgood_mean(self):
    if self.sync_dgood_latencies.size<2:
        return 0
    return self.sync_dgood_latencies.mean()

def async_dgood_mean(self):
    if self.async_dgood_latencies.size<2:
        return 0
    return self.async_dgood_latencies.mean()

def overall_dgood_mean(self):
    all_dgood_latencies = np.concatenate((self.nodist_dgood_latencies, self.sync_dgood_latencies))
    return all_dgood_latencies[:,1].mean()

def overall_dgood_std(self):
    all_dgood_latencies = np.concatenate((self.nodist_dgood_latencies, self.sync_dgood_latencies))
    return all_dgood_latencies[:,1].std()

```

```

def nodist_dgood_std(self):
    if self.nodist_dgood_latencies.size<2:
        return 0
    return self.nodist_dgood_latencies.std()

def sync_dgood_std(self):
    if self.sync_dgood_latencies.size<2:
        return 0
    return self.sync_dgood_latencies.std()

def async_dgood_std(self):
    if self.async_dgood_latencies.size<2:
        return 0
    return self.async_dgood_latencies.std()

def report_movements(self,cond):
    if cond==0:
        str = '    ND'
    elif cond==1:
        str = '    Sync'
    else:
        print 'cond=',cond
        str = '    Async'

    print str,'non-error target movements',ind.num_tgood(cond),'-',100*ind.num_tgood(cond)
    print str,'non-error distractor events',ind.num_dgood(cond),'-',100*ind.num_dgood(cond)
    print str,'target movement errors:',ind.num_tmoveerrors(cond),'-',100*ind.num_tmoveerrors(cond)
    print str,'distractor movement errors:',ind.num_dmoveerrors(cond),'-',100*ind.num_dmoveerrors(cond)
    print str,'total events:',ind.num_tgood(cond)+ind.num_tmoveerrors(cond)+ind.num_dmoveerrors(cond)
    print ''

def __str__(self):
    return "Data container for subject {0}".format(self.subj_id)

# Output latency means for the three conditions as three lines for a csv file.
def outputDataForR(self):
    # R doesn't like 'NA' in a field - it reads it as "not available" so substitute "NA_" for "NA" here.
    if self.subj_id == 'NA':
        self.subj_id = 'NA_';
    line = '{2},{1},0,ND,{0}\n'.format(self.subj_id,self.nodist_latencies[:,1].mean(),self.nodist_latencies[:,1].size)
    line += '{2},{1},1,SD,{0}\n'.format(self.subj_id,self.sync_latencies[:,1].mean(),self.sync_latencies[:,1].size)
    line += '{2},{1},2,AD,{0}\n'.format(self.subj_id,self.async_latencies[:,1].mean(),self.async_latencies[:,1].size)
    return line

# Write out per-individual data for analysis in R into a file.
def writeDataForR(self):
    dfname = 'IndDat' + self.subj_id + '.csv'

    theheader = "latency,condition_str\n"
    minsize = np.min([self.nodist_latencies[:,1].size,self.sync_latencies[:,1].size,self.async_latencies[:,1].size])
    _nd = subsample(self.nodist_latencies[:,1], minsize)
    _sd = subsample(self.sync_latencies[:,1], minsize)
    _ad = subsample(self.async_latencies[:,1], minsize)

```

```

f = open(dfname, 'w')
f.write(theheader)
# allaltlines is the latency trials all together
allaltlines = ''
for i in range(1,minsize):
    line = "{0},ND\n{1},SD\n{2},AD\n".format(_nd[i],_sd[i],_ad[i])
    altline = "{3},{0},ND\n{3},{1},SD\n{3},{2},AD\n".format(_nd[i],_sd[i],_ad[i],self)
    allaltlines += altline
    f.write(line)
f.close()
return allaltlines

def writeNoDistDataForR(self):
    all_lines = ''
    if self.subj_id == 'NA':
        self.subj_id = 'NA_';
    theheader = "subj_id,num,type,error,correctmove,latency,omit,direction\n"
    dfname = 'NoDistDat' + self.subj_id + '.csv'
    f = open(dfname, 'w')
    f.write(theheader)
    minsize = self.alldata_nd[:,1].size
    for i in range(1,minsize):
        line = "{0},{1},{2},{3},{4},{5},{6},{7}\n".format(
            self.subj_id,
            self.alldata_nd[i,0],
            self.alldata_nd[i,1],
            self.alldata_nd[i,2],
            self.alldata_nd[i,3],
            self.alldata_nd[i,4],
            self.alldata_nd[i,5],#omit
            self.alldata_nd[i,8]
        )
        f.write(line)
        all_lines += line
    f.close()
    return all_lines

def writeSyncDataForR(self):
    all_lines = ''
    if self.subj_id == 'NA':
        self.subj_id = 'NA_';
    theheader = "subj_id,num,type,error,correctmove,latency,omit,direction\n"
    dfname = 'SyncDat' + self.subj_id + '.csv'
    f = open(dfname, 'w')
    f.write(theheader)
    minsize = self.alldata_sd[:,1].size
    for i in range(1,minsize):
        line = "{0},{1},{2},{3},{4},{5},{6},{7}\n".format(
            self.subj_id,
            self.alldata_sd[i,0],#evnum
            self.alldata_sd[i,1],#evtype
            self.alldata_sd[i,2],#everror
            self.alldata_sd[i,3],#correct_move

```



```

        self.alldata_sd[i,4],#latency
        self.alldata_sd[i,5],#omit
        self.alldata_sd[i,8]#direction
    )
    f.write(line)
    all_lines += line
f.close()
return all_lines

# Write out asynchronous data including both distractors and targets and the times of each
def writeAsyncDataForR(self):
    # We have alldata_ad which contains columns: evnum, evtype, everror, evcorrectmove, evlatency
    # The simplest approach is probably to simply write this out, along with the subj_id,
    # manipulations in R.
    all_lines = ''
    if self.subj_id == 'NA':
        self.subj_id = 'NA_';
    theheader = "subj_id,num,type,error,correctmove,latency,omit,oreas,timesincelast,direction"
    dfname = 'AsyncDat' + self.subj_id + '.csv'
    f = open(dfname, 'w')
    f.write(theheader)
    minsize = self.alldata_ad[:,1].size
    for i in range(1,minsize):
        line = "{0},{1},{2},{3},{4},{5},{6},{7},{8},{9},{10}\n".format(
            self.subj_id,
            self.alldata_ad[i,0],#evnum
            self.alldata_ad[i,1],#evtype
            self.alldata_ad[i,2],#everror
            self.alldata_ad[i,3],#correct_move
            self.alldata_ad[i,4],#latency
            self.alldata_ad[i,5],#omit
            self.alldata_ad[i,6],#oreas
            self.alldata_ad[i,7],#time_since_last
            self.alldata_ad[i,8],#direction
            self.alldata_ad[i,9] #destination
        )
        f.write(line)
        all_lines += line
    f.close()
    return all_lines

# Output for two factor anova analysis in SPSS
def outputDataForSPSS(self):
    for lat in self.nodist_latencies[:,1]:
        print '{0},ND,{1}'.format(self.subj_id,lat)
    for lat in self.sync_latencies[:,1]:
        print '{0},SD,{1}'.format(self.subj_id,lat)
    for lat in self.async_latencies[:,1]:
        print '{0},AD,{1}'.format(self.subj_id,lat)

# Output for univariate or multivariate repeated measures analysis
def outputDataForSPSS_MV(self):
    print '{0},ND,{1},{2}'.format(self.subj_id,self.nodist_latencies[:,1].mean(),self.n_err)

```

```

        print '{0},SD,{1},{2}'.format(self.subj_id,self.sync_latencies[:,1].mean(),self.n_errors)
        print '{0},AD,{1},{2}'.format(self.subj_id,self.async_latencies[:,1].mean(),self.n_errors)

    # Output for repeated measures analysis of latency
    def outputDataForSPSS_RepMeasLat(self):
        print '{0},{1},{2},{3}'.format(self.subj_id,self.nodist_latencies.mean(),self.sync_latencies.mean(),self.n_errors)

    # Output for repeated measures analysis of error rate
    def outputDataForSPSS_RepMeasError(self):
        print '{0},{1},{2},{3}'.format(self.subj_id,self.n_errors_per_target_nd,self.n_errors_per_target_ad,self.n_errors_per_target_sync)

### END CLASS individual ###

# This function sets up a list of individuals and populates the latencies.
def readIndividuals():
    individuals = dict()
    idnum_counter = 1
    # Extract the data from the raw format and collate it into individual
    # data containers, one per subject.
    for fname in zip(*fnames):
        # I'll use the subject ID as a key into output data structures
        subj_id = getfnameid(fname[0][0])

        # condition index is for the no distractor/sync distractor/async distractor
        condition_index = getcondition(fname[1][0,0][36][0])

        # Need ONE individual object for each subj_id.
        if subj_id not in individuals:
            individuals[subj_id] = individual(subj_id) # or if imported individual.individual()
            individuals[subj_id].idnum = idnum_counter
            idnum_counter += 1

        # latencies for all events
        #
        # All the fname[n] contain:
        # ev.number, ev.type, ev.error, ev.correct_move, ev.latency. Then add on the condition
        #
        # fname[2] additionally has: ev.omit and omit_reason (as a
        # numeric code) cols. So can obtain a value for "proportion of
        # targets omitted".
        #
        #if individuals[subj_id].alldata.size < 2:
        #    individuals[subj_id].alldata = np.append(fname[2], condition_index*np.ones(fname[2].shape[0]),axis=1)
        #else:
        #    newdata = np.append(fname[2], condition_index*np.ones(fname[2].shape[0])[:,None],axis=1)
        #    individuals[subj_id].alldata = np.concatenate((individuals[subj_id].alldata,newdata),axis=1)

        # latencies for non-movement-error target events
        tnoerr_latencies = fname[4] # R.latency_noerror_target - Use table with python index 4

        # latencies for non-movement-error distractor events
        dnoerr_latencies = fname[5]

        # err_latencies = fname[6] # ALL error events, distractor and target.

```

```

# latencies for target movement error events
terr_latencies = fname[7] # R.latency_error_target

# latencies for distractor movement error events
derr_latencies = fname[8] # R.latency_error_distractor

#omitted = fname[1] # omitted events?

if condition_index == 0:
    individuals[subj_id].alldata_nd = fname[2] # contains dirn information
    #print 'alldata_nd has shape', fname[2].shape
    individuals[subj_id].filename_nd = fname[0][0]

    #print err_latencies
    individuals[subj_id].nodist_latencies = tnoerr_latencies[:, [0,4]]
    individuals[subj_id].nodist_latencies_rank = stats.rankdata(individuals[subj_id].nodist_latencies)
    if terr_latencies.size > 0:
        individuals[subj_id].nodist_err_latencies = terr_latencies[:, [0,4]]
        individuals[subj_id].nodist_err_latencies_rank = stats.rankdata(individuals[subj_id].nodist_err_latencies)
    # We'll have an error rate for nodist - it should be very low. Did it get passed i
    nerrs = 0
    ntargets = 0
    for d in individuals[subj_id].alldata_nd:
        ntargets += 1 # "ndistractors" is really "ntargets" for the ND condition.
        if d[2] > 0.0:
            nerrs += 1
    individuals[subj_id].n_errors_per_target_nd = (float(nerrs) / float(ntargets))

elif condition_index == 1:
    individuals[subj_id].alldata_sd = fname[2]
    individuals[subj_id].filename_sd = fname[0][0]
    #print latencies
    individuals[subj_id].sync_latencies = tnoerr_latencies[:, [0,4]]
    individuals[subj_id].sync_latencies_rank = stats.rankdata(individuals[subj_id].sync_latencies)
    if terr_latencies.size > 0:
        individuals[subj_id].sync_err_latencies = terr_latencies[:, [0,4]]
        individuals[subj_id].sync_err_latencies_rank = stats.rankdata(individuals[subj_id].sync_err_latencies)
    # For sync, also read errors
    nerrs = 0
    ndistractors = 0
    #if subj_id == 'JS':
    #    print 'Sync'
    for d in individuals[subj_id].alldata_sd:
        #if subj_id == 'JS' and d[2] > 0.0:
        #    print d
        ndistractors += 1
        if d[2] > 0.0:
            nerrs += 1
    individuals[subj_id].n_errors_per_distractor_sync = (float(nerrs) / float(ndistractors))

elif condition_index == 2:
    individuals[subj_id].alldata_ad = fname[2]
    individuals[subj_id].filename_ad = fname[0][0]

```

```

individuals[subj_id].async_latencies = tnoerr_latencies[:, [0,4]]
individuals[subj_id].async_latencies_rank = stats.rankdata(individuals[subj_id].as
#test[test[:, 1] == 4]
if terr_latencies.size > 0:
    individuals[subj_id].async_err_latencies = terr_latencies[:, [0,4]]
    individuals[subj_id].async_err_latencies_rank = stats.rankdata(individuals[sub
if derr_latencies.size > 0:
    individuals[subj_id].async_derr_latencies = derr_latencies[:, [0,4]]
    individuals[subj_id].async_derr_latencies_rank = stats.rankdata(individuals[su

individuals[subj_id].async_tnoerr_timesince = tnoerr_latencies[:, [0,5]]
individuals[subj_id].async_derr_timesince = derr_latencies[:, [0,5]]
# plus derr_latencies[:, []]

# For async, also read errors
nerrs = 0
ndistractors = 0
#if subj_id == 'JS':
#    print 'Async'
for d in individuals[subj_id].alldata_ad:
    #if subj_id == 'JS' and d[2] > 0.0:
    #    print d
    if d[2] > 0.0: # Count all errors for async
        nerrs += 1
    if d[1] < 1.0:
        ndistractors += 1
#if subj_id == 'JS':
#    print 'Num movement errors:', nerrs, 'Num distractor events:', ndistractors
individuals[subj_id].n_errors_per_distractor_async = (float(nerrs) / float(ndistrac
individuals[subj_id].n_async_distractors = ndistractors

return individuals

def equaliseReplicates(individuals):
    # Find smallest number of cell replications and reduce all cells to this number.
    smallest_n = 1000000
    for i in individuals:
        individuals[i].excludeOutliers()
        if individuals[i].num_tgood(0) < smallest_n:
            smallest_n = individuals[i].num_tgood(0)
        if individuals[i].num_tgood(1) < smallest_n:
            smallest_n = individuals[i].num_tgood(1)
        if individuals[i].num_tgood(2) < smallest_n:
            smallest_n = individuals[i].num_tgood(2)

    # Randomly select smallest_n data from each condition in each individual
    for i in individuals:
        individuals[i].randomly_subsample_data(smallest_n)

return individuals

```

1.4 Numbers of errors and omitted events

This is the definitive example of how to find out the number of target & distractor errors and the number of omitted events.

```
In [15]: for i,ind in readIndividuals().iteritems():
    print 'Subject ID: ', ind.subj_id

    print 'ND'
    print 'num target events:', ind.num_target_events(0)
    print 'num distractor events:', ind.num_distractor_events(0)
    print 'num_target_omissions:', ind.num_target_omissions(0)
    print 'num_tgood:', ind.num_tgood(0)
    print 'num_tmoveerrors', ind.num_tmoveerrors(0)
    print 'num_target_non_omissions (should equal num_tgood + num_tmoveerrors):', ind.num_target_non_omissions(0)
    print 'tmoveerrors percent', ind.num_tmoveerrors_percent(0)
    print 'dmoveerrors percent', ind.num_dmoveerrors_percent(0)
    print 'moveerrors per target percent', ind.num_moveerrors_per_target_percent(0)
    print 'moveerrors per distractor percent', ind.num_moveerrors_per_distractor_percent(0)
    print 'num_target_non_omissions', ind.num_target_non_omissions(0)
    print 'num_target_omissions', ind.num_target_omissions(0)
    print 'num_distractor_non_omissions', ind.num_distractor_non_omissions(0)
    print 'num_distractor_omissions', ind.num_distractor_omissions(0)

    print 'SD'
    print 'num target events:', ind.num_target_events(1)
    print 'num distractor events:', ind.num_distractor_events(1)
    print 'num_target_omissions:', ind.num_target_omissions(1)
    print 'num_tgood:', ind.num_tgood(1)
    print 'num_tmoveerrors', ind.num_tmoveerrors(1)
    print 'num_target_non_omissions (should equal num_tgood + num_tmoveerrors):', ind.num_target_non_omissions(1)
    ##
    print 'tmoveerrors percent', ind.num_tmoveerrors_percent(1)
    print 'dmoveerrors percent', ind.num_dmoveerrors_percent(1)
    print 'moveerrors per target percent', ind.num_moveerrors_per_target_percent(1)
    print 'moveerrors per distractor percent', ind.num_moveerrors_per_distractor_percent(1)
    print 'num_target_non_omissions', ind.num_target_non_omissions(1)
    print 'num_target_omissions', ind.num_target_omissions(1)
    print 'num_distractor_omissions:', ind.num_distractor_omissions(1)
    print 'num_distractor_non_omissions:', ind.num_distractor_non_omissions(1)

    print 'AD'
    print 'num target events:', ind.num_target_events(2)
    print 'num distractor events:', ind.num_distractor_events(2)
    print 'num_target_omissions:', ind.num_target_omissions(2)
    print 'num_tgood:', ind.num_tgood(2)
    print 'num_tmoveerrors', ind.num_tmoveerrors(2)
    print 'num_target_non_omissions (num_tgood+num_tmoveerrors):', ind.num_target_non_omissions(2)
    print 'num_dgood:', ind.num_dgood(2)
    print 'num_dmoveerrors', ind.num_dmoveerrors(2)
    ##
    print 'tmoveerrors percent', ind.num_tmoveerrors_percent(2)
    print 'dmoveerrors percent', ind.num_dmoveerrors_percent(2)
    print 'moveerrors per target percent', ind.num_moveerrors_per_target_percent(2)
```

```

print 'moveerrors per distractor percent', ind.num_moveerrors_per_distractor_percent(2)
print 'num_target_non_omissions', ind.num_target_non_omissions(2)
print 'num_target_omissions', ind.num_target_omissions(2)
print 'num_distractor_omissions:', ind.num_distractor_omissions(2)
print 'num_distractor_non_omissions:', ind.num_distractor_non_omissions(2)

break # So that we only output the information for one individual

```

Subject ID: SB2

ND

```

num target events: 68
num distractor events: 0
num_target_omissions: 31
num_tgood: 35
num_tmoveerrors 2
num_target_non_omissions (should equal num_tgood + num_tmoveerrors): 37
tmoveerrors percent 5.40540540541
dmoveerrors percent 0
moveerrors per target percent 5.40540540541
moveerrors per distractor percent -1
num_target_non_omissions 37
num_target_omissions 31
num_distractor_non_omissions 0
num_distractor_omissions 0

```

SD

```

num target events: 66
num distractor events: 66
num_target_omissions: 23
num_tgood: 32
num_tmoveerrors 11
num_target_non_omissions (should equal num_tgood + num_tmoveerrors): 43
tmoveerrors percent 25.5813953488
dmoveerrors percent 0
moveerrors per target percent 25.5813953488
moveerrors per distractor percent 25.5813953488
num_target_non_omissions 43
num_target_omissions 23
num_distractor_omissions: 0
num_distractor_non_omissions: 0

```

AD

```

num target events: 70
num distractor events: 118
num_target_omissions: 33
num_tgood: 33
num_tmoveerrors 4
num_target_non_omissions (num_tgood+tmoveerrors): 37
num_dgood: 0
num_dmoveerrors 4
tmoveerrors percent 10.8108108108
dmoveerrors percent 3.38983050847
moveerrors per target percent 21.6216216216
moveerrors per distractor percent 3.58353510896
num_target_non_omissions 37

```

```

num.target_omissions 33
num.distractor_omissions: 89
num.distractor_non_omissions: 29

```

1.5 Breakdown of omit reasons

I want to count up the number of different omit reasons for various event types. To do this, need to add any of the new omit reasons in `lt_analyse.latency.m`.

In the octave script “omit” means “omit from latency measurements”. That means that an omitted distractor movement may be a correct, error-free event; it’s just that because the user did not move in response to the distractor, no latency measurement could be measured.

So, when counting omit reasons to see how many events were omitted and how many considered for errors, if the reason is “16 - Movement occurs beyond next target”, that’s no error and it goes in the “kept” pile.

```

In [16]: # Count omit reasons...
         from __future__ import division
         import numpy as np
         DIST_EVENT=0
         TARG_EVENT=1

         def omit_reason_summary (omit_reason_count_arr):
             iterator = 0 # into omit_reason_count_arr
             latmeas_count = 0 # "latency measurement count"
             nolatmeas_count = 0 # "omitted from latency measurement count"
             omit_count = 0 # "omitted from error counting"
             nomit_count = 0 # "included in error counting"
             for thecount in omit_reason_count_arr:
                 if thecount > 0:
                     print 'omit reason',omit_reason(iterator),'count:',thecount

                     # How many events considered for latency measurements?
                     if iterator == 0:
                         latmeas_count += thecount
                     else:
                         nolatmeas_count += thecount

                     # How many considered for error counting?
                     if iterator == 0 or iterator == 16:
                         # Note - 0 means no omission at all. 16 is special - it is
                         # where the distractor event was omitted because it didn't
                         # distract the subject, so it should be counted as no
                         # omission.
                         nomit_count += thecount
                     else:
                         omit_count += thecount

                 iterator += 1
             print nomit_count,'kept for error-counting,',omit_count,'omitted from error-counting'
             print latmeas_count,'kept for latency measurements,',nolatmeas_count,'couldnt provide later'
             print 'There were ', nomit_count+omit_count, 'events in total.'
             print ''
             return nomit_count+omit_count

```

```

# To count up total events:
targ_event_count = 0
dist_event_count = 0

omit_reason_count_arr = np.zeros(21).astype(int)
for i,ind in readIndividuals().iteritems():
    ador = ind.ad_omit_reasons(TARG_EVENT)
    omit_reason_count_arr += ador
    #print 'Omit reason summary for Async condition, target events, INDIVIDUAL:',ind.subj_id
    #omit_reason_summary (ador)
print 'Omit reason summary for Async condition, target events'
print '-----',
targ_event_count += omit_reason_summary (omit_reason_count_arr)

omit_reason_count_arr = np.zeros(21).astype(int)
for i,ind in readIndividuals().iteritems():
    ador = ind.ad_omit_reasons(DIST_EVENT)
    omit_reason_count_arr += ador
    #print 'Omit reason summary for Async condition, distractor events, INDIVIDUAL:',ind.subj_
    #omit_reason_summary (ador)
print 'Omit reason summary for Async condition, distractor events'
print '-----',
dist_event_count += omit_reason_summary (omit_reason_count_arr)

omit_reason_count_arr = np.zeros(21).astype(int)
for i,ind in readIndividuals().iteritems():
    ador = ind.sd_omit_reasons(TARG_EVENT)
    omit_reason_count_arr += ador
print 'Omit reason summary for Sync condition, target events'
print '-----',
targ_event_count += omit_reason_summary (omit_reason_count_arr)

omit_reason_count_arr = np.zeros(21).astype(int)
for i,ind in readIndividuals().iteritems():
    ador = ind.nd_omit_reasons(TARG_EVENT)
    omit_reason_count_arr += ador
    #print 'Omit reason summary for ND condition, target events, INDIVIDUAL:',ind.subj_id
    #omit_reason_summary (ador)
    #print ador
    #print omit_reason_count_arr
print 'Omit reason summary for ND condition, target events'
print '-----',
targ_event_count += omit_reason_summary (omit_reason_count_arr)

print ''
print 'Total target events:',targ_event_count,'Total distractor events:',dist_event_count
print 'Grand total events:',targ_event_count+dist_event_count

```

Omit reason summary for Async condition, target events

```

omit reason 0 not omitted count: 2313
omit reason 3 target posn change less than min. jump size count: 91
omit reason 7 Too fast (targ) count: 62
omit reason 9 No movement detected count: 8

```


omit reason 11 Stable stylus period too short count: 351
omit reason 12 Drift too great during stable stylus period count: 6
omit reason 13 Drift too great during stable period (avg) count: 661
omit reason 17 Subject was distracted by closely previous distractor count: 4
omit reason 18 Incorrect move was recorded in previous distractor event count: 7
2313 kept for error-counting, 1190 omitted from error-counting
2313 kept for latency measurements, 1190 couldnt provide latency measurements
There were 3503 events in total.

Omit reason summary for Async condition, distractor events

omit reason 0 not omitted count: 1692
omit reason 3 target posn change less than min. jump size count: 98
omit reason 4 Stable position later than event onset count: 123
omit reason 8 Too fast (distractor) count: 20
omit reason 11 Stable stylus period too short count: 1743
omit reason 12 Drift too great during stable stylus period count: 18
omit reason 13 Drift too great during stable period (avg) count: 957
omit reason 15 Stylus didn't move away from target count: 1326
omit reason 16 Movement occurs beyond next target count: 896
omit reason 19 This distractor event did not distract the stylus movement count: 64
omit reason 20 Recorded this stylus movement as a distraction towards the next distractor count: 4
2588 kept for error-counting, 4353 omitted from error-counting
1692 kept for latency measurements, 5249 couldnt provide latency measurements
There were 6941 events in total.

Omit reason summary for Sync condition, target events

omit reason 0 not omitted count: 2623
omit reason 3 target posn change less than min. jump size count: 119
omit reason 7 Too fast (targ) count: 32
omit reason 9 No movement detected count: 8
omit reason 11 Stable stylus period too short count: 214
omit reason 12 Drift too great during stable stylus period count: 7
omit reason 13 Drift too great during stable period (avg) count: 564
2623 kept for error-counting, 944 omitted from error-counting
2623 kept for latency measurements, 944 couldnt provide latency measurements
There were 3567 events in total.

Omit reason summary for ND condition, target events

omit reason 0 not omitted count: 2684
omit reason 3 target posn change less than min. jump size count: 102
omit reason 7 Too fast (targ) count: 46
omit reason 9 No movement detected count: 5
omit reason 11 Stable stylus period too short count: 166
omit reason 12 Drift too great during stable stylus period count: 8
omit reason 13 Drift too great during stable period (avg) count: 595
2684 kept for error-counting, 922 omitted from error-counting
2684 kept for latency measurements, 922 couldnt provide latency measurements
There were 3606 events in total.

Total target events: 10676 Total distractor events: 6941

Grand total events: 17617

1.6 Show which events are outliers

This code runs through each individual showing which events are deleted from each dataset by passing 1 to the `individuals.excludeOutliers()` method.

```
In [17]: print 'Showing latency outlier event numbers and values'
         print '-----'
         individuals = readIndividuals()
         show_excluded = 0
         if show_excluded:
             for i,ind in individuals.iteritems():
                 ind.excludeOutliers(show_excluded)
         else:
             print '[Set show_excluded = 1 to see the (rather verbose) output of this block.]'
```

Showing latency outlier event numbers and values

[Set show_excluded = 1 to see the (rather verbose) output of this block.]

1.7 Mean number of latency-to-first-movement values obtained by Octave pre-processing

Mean numbers of trials overall and for the 3 different conditions. Note that these are the mean numbers of trials which the Octave pre-processing script was able to turn into latency-to-first-movement values, rather than the mean number of target events, which is determined separately in a previous block called **Count number of target events**.

```
In [18]: from __future__ import division
         import numpy as np

         num_trials = []
         nd_num_trials = []
         sd_num_trials = []
         ad_num_trials = []
         min_nd_num_trials = 1000
         min_sd_num_trials = 1000
         min_ad_num_trials = 1000
         min_nd_num_trials_idx = ''
         min_sd_num_trials_idx = ''
         min_ad_num_trials_idx = ''
         max_nd_num_trials = 0
         max_sd_num_trials = 0
         max_ad_num_trials = 0
         max_nd_num_trials_idx = ''
         max_sd_num_trials_idx = ''
         max_ad_num_trials_idx = ''
         ind_ids = []
         nd_percent = []
         sd_percent = []
         ad_percent = []

         for i,ind in readIndividuals().iteritems():
             ind_ids.append(ind.subj_id)
```


SD Percent min: 46.1538461538 % max: 88.0952380952 % mean/sd 64.7803976178 10.5121907677
AD Percent (of targ events) min: 30.5555555556 % max: 97.5609756098 % mean/sd 65.0555862042 13.78277843

1.7.1 Summary of the above

The block above gives (for each condition) the minimum and maximum percentages of target events for which a latency-to-first-movement value was found by the Octave preprocessing scripts. Also given is the mean percentage success rate and associated SD and the minimum absolute number of latencies found.

1.8 Repeated measures ANOVA

If we collapse the replication of latency measurements into a mean latency for each condition/individual combination, then it's possible to perform a repeated measures ANOVA analysis. This was the original design of the experiment and is the primary analysis reported in the paper, however there an ANOVA on ranked data is reported as the Shapiro Wilks tests on normality of the mean latencies fails.

The QQ plot from this code block is reported in the paper.

```
In [19]: from __future__ import division
         # This repmeasures_anova works on the mean latency for each individual.
         # This is effectively a "two factor anova without replication"
         import numpy as np

         def repmeasures_anova(individuals):

             # A one-way repeated measures anova is computed like a "two-factor anova without replicati
             # for a random block design. That means sums of squares are computed like a two factor ano
             # This example follows McKillup Section 15.5 (p216).

             # Containers to fill
             nodist_latencies = [];
             sync_latencies = [];
             async_latencies = [];

             num_individuals = len(individuals)
             num_conditions = 3

             # Zeroth, exclude outliers (assumed already to have been done before this method was calle

             # In this loop extract the latencies into external containers:
             for i,ind in individuals.iteritems():
                 # NB: Assume here that any outlier excluding has ALREADY been carried out.
                 nodist_latencies = np.concatenate((nodist_latencies, [ind.nodist_mean()]))
                 sync_latencies = np.concatenate((sync_latencies, [ind.sync_mean()]))
                 async_latencies = np.concatenate((async_latencies, [ind.async_mean()]))

             # Compute grand mean
             all_latencies = np.concatenate((nodist_latencies, sync_latencies, async_latencies))
             grand_mean = all_latencies.mean()
             print 'Grand mean:',grand_mean

             # Displacement due to ALL sources of variation in the experiment. Factor A, Factor B, inte
             # total_variance. This is displacement from the grand mean.
             total_sos = 0
             total_dof = -1 # dof is number of replicates -1
             for i in individuals:
```

```

    ind = individuals[i]
    total_sos += np.power(grand_mean - ind.nodist_mean(), 2)
    total_sos += np.power(grand_mean - ind.sync_mean(), 2)
    total_sos += np.power(grand_mean - ind.async_mean(), 2)

total_dof = (num_conditions*num_individuals)-1

# So here's the total variance:
total_variance = total_sos/total_dof
print 'total sos:',total_sos,'total dof:',total_dof,'total variance:',total_variance
# Could also get total sos with:
# total_variance = all_latencies.var(ddof=1) # but it doesn't work as above.

# Now consider the data in relation to each of the two factors (individual and condition)

# Compute condition_plus_error_sos (Factor A - equiv to Temperature in p177 of McKillup)
nd_treatment_mean = 0
sd_treatment_mean = 0
ad_treatment_mean = 0

for i,ind in individuals.iteritems():
    nd_treatment_mean += ind.nodist_mean()
    sd_treatment_mean += ind.sync_mean()
    ad_treatment_mean += ind.async_mean()

nd_treatment_mean = nd_treatment_mean / num_individuals
sd_treatment_mean = sd_treatment_mean / num_individuals
ad_treatment_mean = ad_treatment_mean / num_individuals

# Now we have treatment means for each condition.

condition_sos = num_individuals*(np.power(nd_treatment_mean - grand_mean,2) + np.power(sd_
condition_dof = num_conditions-1 # 3 conditions - 1 = 2

condition_variance = condition_sos / condition_dof

# Compute individual_plus_error_sos (p178).
individual_sos = 0
for i,ind in individuals.iteritems():
    ind_mean = (ind.nodist_mean() + ind.sync_mean() + ind.async_mean()) / 3
    individual_sos += num_conditions*np.power(ind_mean-grand_mean,2)
individual_dof = (num_individuals-1)

individual_variance = individual_sos / individual_dof

# remainder variance computed by subtraction
##remainder_variance = total_variance - condition_variance - individual_variance
remainder_sos = total_sos - condition_sos - individual_sos
print 'total_sos:', total_sos
print 'condition_sos:', condition_sos
print 'individual_sos:', individual_sos
print 'remainder_sos:', remainder_sos
remainder_dof = total_dof - condition_dof - individual_dof
# This is the best estimate for error in a two factor ANOVA without replication:

```

```

remainder_variance = remainder_sos/remainder_dof
print 'remainder_variance:',remainder_variance,'remainder_sos:',remainder_sos

print 'total_sos:',total_sos,'condition_sos:',condition_sos,'indiv_sos:',individual_sos,'e

# Now compute the F ratios
F_cond = condition_variance / remainder_variance # fixed. condition + error
F_ind = individual_variance / remainder_variance # random. individual + error

# Lastly, what's the probability for this?
P_cond = 1-special.fdtr(condition_dof,remainder_dof,F_cond)
P_ind = 1-special.fdtr(individual_dof,remainder_dof,F_ind)

print '\nF_cond:{0} P_cond:{1}'.format(F_cond,P_cond)
print 'F_ind:{0} P_ind:{1}'.format(F_ind,P_ind)
#print 'F_interaction:{0} P_interaction:{1}'.format(F_interaction,P_interaction)

print '\nCondition:\nF({0},{1})={2}, p={3}'.format(condition_dof,remainder_dof,F_cond,P_cond)
print '\nBetween individuals:\nF({0},{1})={2}, p={3}'.format(individual_dof,remainder_dof,F_ind,P_ind)

print '\n'
print 'Source of var, Sumof Squares, df, Mean square , F ratio          ', 'P'
print '-----'
print '{0}, {1}, {2}, {3}, {4}, {5}'.format('Individual', individual_sos, individual_dof, F_ind, P_ind)
print '{0}, {1}, {2}, {3}, {4}, {5}'.format('Condition', condition_sos, condition_dof, F_cond, P_cond)
print '{0}, {1}, {2}, {3}, {4}, {5}'.format('Remainder', remainder_sos, remainder_dof, F_ind, P_ind)
print '{0}, {1}, {2}, {3}, {4}, {5}'.format('Total', total_sos, total_dof, F_ind, P_ind)

def normality_test(individuals):
    print '\nNormality of data:'

    nd_latency_means = []
    sd_latency_means = []
    ad_latency_means = []
    for i, ind in individuals.iteritems():
        nd_latency_means = np.append(nd_latency_means, ind.nodist_mean())
        sd_latency_means = np.append(sd_latency_means, ind.sync_mean())
        ad_latency_means = np.append(ad_latency_means, ind.async_mean())

    print ' Shapiro-Wilks:'
    W, p = stats.shapiro (subsample(nd_latency_means, 25))
    print ' No Distractor, 25 sub-samples: W', W, 'p-value', p, '(Reject Normal NULL hypothesis if p<0.05)'
    W, p = stats.shapiro (subsample(sd_latency_means, 25))
    print ' Sync Distractor, 25 sub-samples: W', W, 'p-value', p
    W, p = stats.shapiro (subsample(ad_latency_means, 25))
    print ' Async Distractor, 25 sub-samples: W', W, 'p-value', p

    print '\nSee also QQ Plots (set show_graph to 1)'
    # Quantile-Quantile Plot to show normality
    show_graph = 1
    if show_graph==1:
        %matplotlib inline
        f, (ax1, ax2, ax3) = plt.subplots(3)

```

```

fig1 = sm.qqplot(nd_latency_means, fit=True, line='45', ax=ax1)
ax1.set_title('ND')
fig2 = sm.qqplot(sd_latency_means, fit=True, line='45', ax=ax2)
ax2.set_title('SD')
fig3 = sm.qqplot(ad_latency_means, fit=True, line='45', ax=ax3)
ax3.set_title('AD')
savename = 'images/latency_means_normplot.png'
plt.savefig(savename)
plt.show()

# Call the repeated measures ANOVA
individuals = readIndividuals()

# Could equalise replicants, but this would be entirely optional for repeated measures as the
# individuals = equaliseReplicates(individuals)
# If we don't call equalise, then must exclude outliers anyway:
for i, ind in individuals.iteritems():
    ind.excludeOutliers()

normality_test(individuals)

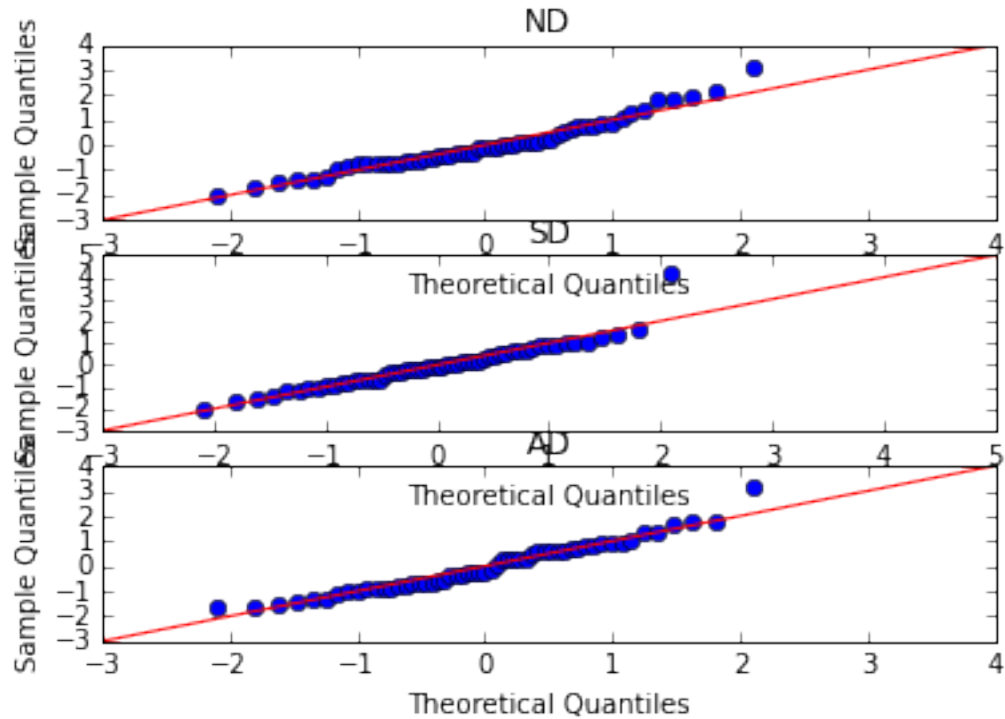
repmeasures_anova(individuals)

Normality of data:
Shapiro-Wilks:
No Distractor, 25 sub-samples: W 0.914870500565 p-value 0.0391994863749 (Reject Normal NULL hypothesis)
Sync Distractor, 25 sub-samples: W 0.845669388771 p-value 0.00145912088919
Async Distractor, 25 sub-samples: W 0.962160229683 p-value 0.459158211946

See also QQ Plots (set show_graph to 1)

/usr/local/lib/python2.7/dist-packages/numpy/core/_methods.py:59: RuntimeWarning: Mean of empty slice.
  warnings.warn("Mean of empty slice.", RuntimeWarning)

```



Grand mean: 327.018604194
total sos: 215033.599942 total dof: 164 total variance: 1311.18048745
total_sos: 215033.599942
condition_sos: 80551.4780108
individual_sos: 99090.3959277
remainder_sos: 35391.726004
remainder_variance: 327.701166704 remainder_sos: 35391.726004
total_sos: 215033.599942 condition_sos: 80551.4780108 indiv_sos: 99090.3959277 error (i.e. remainder): 3

F_cond:122.903862109 P_cond:1.11022302463e-16
F_ind:5.59963624925 P_ind:1.62092561595e-14

Condition:
F(2,108)=122.903862109, p=1.11022302463e-16

Between individuals:
F(54,108)=5.59963624925, p=1.62092561595e-14

Source of var,	Sumof Squares,	df,	Mean square ,	F ratio	P
Individual	, 99090.3959277,	54,	1835.00733199,	5.59963624925,	1.62092561595e-14
Condition	, 80551.4780108,	2,	40275.7390054,	122.903862109,	1.11022302463e-16
Remainder	, 35391.726004,	108,	327.701166704,	-, -	
Total	, 215033.599942,	164,	-, -, -		

1.8.1 Repeated measures ANOVA summary

The null hypotheses are that “condition does not change latency” and “individual does not affect latency”.

A repeated measures anova, where I use the mean latency for each condition as a single data point for each individual shows a significant effect both for condition and for individual - the null hypotheses would be rejected in each case if the data fulfilled normality.

1.9 Output data for analysis in R

How many different stats systems can we use for one piece of data analysis? See various .r scripts which carry out much of the analysis reported in the paper.

This code block generates AnovaR.csv which contains the latencies and error_rates.csv which contains the error rates.

```
In [20]: # Output data in format suitable for R
         individuals = readIndividuals()

         # Optional:
         #individuals = equaliseReplicates(individuals)
         # or
         # If we don't call equalise, exclude outliers anyway:
         for i,ind in individuals.iteritems():
             ind.excludeOutliers()

         async_lines = 'subj_id,num,type,error,correctmove,latency,omit,oreas,timesincelast,direction,d
         sync_lines = 'subj_id,num,type,error,correctmove,latency,omit,direction\n'
         nodist_lines = 'subj_id,num,type,error,correctmove,latency,omit,direction\n'
         csvdata = 'idnum,latency,condition,condition_str,subj_id\n'
         csvtrialdata = 'subj_id,latency,condition_str\n'
         for i,ind in individuals.iteritems():
             csvdata += ind.outputDataForR()
             # Save this particular individuals latency values
             csvtrialdata += ind.writeDataForR()
             async_lines += ind.writeAsyncDataForR()
             sync_lines += ind.writeSyncDataForR()
             nodist_lines += ind.writeNoDistDataForR()

         print 'Writing AnovaR.csv...'
         f = open('AnovaR.csv', 'w')
         f.write(csvdata)
         f.close()

         print 'Writing AllTrials.csv...'
         f = open('AllTrials.csv', 'w')
         f.write(csvtrialdata)
         f.close()

         print 'Writing AsyncTrials.csv...'
         f = open('AsyncTrials.csv', 'w')
         f.write(async_lines)
         f.close()

         print 'Writing SyncTrials.csv...'
         f = open('SyncTrials.csv', 'w')
         f.write(sync_lines)
```

```

f.close()

print 'Writing NoDistTrials.csv...'
f = open('NoDistTrials.csv', 'w')
f.write(nodist_lines)
f.close()

print 'Writing error_rates.csv...'
f = open('error_rates.csv', 'w')
f.write ('idnum,error_rate,condition,condition_str,subj_id\n')
for i,ind in readIndividuals().iteritems():
    subj_id = ind.subj_id
    if subj_id == 'NA':
        subj_id = 'NA_'
    f.write ('{0},{1},1,ND,{2}\n'.format(ind.idnum,ind.num_moveerrors_per_target(0),subj_id))
    f.write ('{0},{1},2,SD,{2}\n'.format(ind.idnum,ind.num_moveerrors_per_target(1),subj_id))
    f.write ('{0},{1},3,AD,{2}\n'.format(ind.idnum,ind.num_moveerrors_per_target(2),subj_id))
f.close()

Writing AnovaR.csv...
Writing AllTrials.csv...
Writing AsyncTrials.csv...
Writing SyncTrials.csv...
Writing NoDistTrials.csv...
Writing error_rates.csv...

```

1.10 ANOVA on latency ranks

This code block calls the R script **RankedAnova.r** and outputs the result. It means you have to have R installed (version 3.0.2 will match my system) with the packages nlme and effsize.

```

In [21]: from subprocess import CalledProcessError, check_output
        try:
            out = check_output(["R", "-q --file=RankedAnova.r"])
        except CalledProcessError as err:
            print 'Error for:',err.cmd,'with error code:',err.returncode
            print "\nOutput:\n\n",err.output
        else:
            for line in out.splitlines():
                print line

> # I found guidance on Anovas in R here:
> # https://gribblelab.wordpress.com/2009/03/09/repeated-measures-anova-using-r/
> # and here:
> # https://seriousstats.wordpress.com/tag/rank-transformation/
>
> # Set factor contrasts option, important for aov() function.
> options(contrasts=c("contr.treatment","contr.treatment"))
>
> latdat <- read.csv('AnovaR.csv')
>
> # Using Linear Mixed Effects models
> require(nlme)
>
> # Compute ranks to carry out ANOVA on the ranks

```

```

> rlatency <- rank(latdat$latency)
> #
> # The formulae here are:
> # fixed: rlatency "is predicted by" condition_str
> # random: "is predicted by" "the mean" "given" "condition_str nested within subj_id"
> anova_ranked_cond <- lme(fixed = rlatency ~ condition_str, random = ~1|subj_id/condition_str, data=latdat)
> # -----
> print(summary(anova_ranked_cond))
Linear mixed-effects model fit by REML
Data: latdat
      AIC      BIC    logLik
1598.539 1617.065 -793.2697

Random effects:
Formula: ~1 | subj_id
      (Intercept)
StdDev:      28.38689

Formula: ~1 | condition_str %in% subj_id
      (Intercept) Residual
StdDev:      21.43293 9.895613

Fixed effects: rlatency ~ condition_str
              Value Std.Error  DF    t-value p-value
(Intercept)  109.89091  4.978334 108   22.073834  0.000
condition_strND -69.54545  4.501691 108  -15.448740  0.000
condition_strSD -11.12727  4.501691 108   -2.471798  0.015
Correlation:
      (Intr) cnd_ND
condition_strND -0.452
condition_strSD -0.452  0.500

Standardized Within-Group Residuals:
      Min      Q1      Med      Q3      Max
-0.86143597 -0.25401473  0.02105463  0.24965003  0.90271124

Number of Observations: 165
Number of Groups:
      subj_id condition_str %in% subj_id
      55              165

> # -----
> print(anova(anova_ranked_cond))
      numDF denDF  F-value p-value
(Intercept)      1   108 382.1123 <.0001
condition_str      2   108 137.7248 <.0001
> # -----
> #
> # Showing residuals of ANOVA on ranks:
>
> setEPS()
> postscript(file='../paper/figures/anova_ranked_cond_resid.eps')
> plot(fitted(anova_ranked_cond), residuals(anova_ranked_cond))
> dev.off()
null device

```

```

1
>
> png(filename='r_images/anova_ranked_cond_resid.png')
> plot(fitted(anova_ranked_cond), residuals(anova_ranked_cond))
> dev.off()
null device
1
>
> # Carry out the LME on subj_id - the inverse of the above.
> anova_ranked_indiv <- lme(fixed = rlatency ~ subj_id, random = ~1|condition_str/subj_id, data=latdat)
> # -----
> print(anova(anova_ranked_indiv))
              numDF denDF    F-value p-value
(Intercept)      1   108 14.809621  2e-04
subj_id          54   108  5.337834 <.0001
> # -----
> setEPS()
> postscript(file='../paper/figures/anova_ranked_indiv_resid.eps')
> plot (fitted(anova_ranked_indiv), residuals(anova_ranked_indiv))
> dev.off()
null device
1
>
> png (filename='r_images/anova_ranked_indiv_resid.png')
> plot (fitted(anova_ranked_indiv), residuals(anova_ranked_indiv))
> dev.off()
null device
1
>
> #
> # Apply Wilcoxon Signed-Rank test for pairwise comparisons
> #
> # individual latencies
> nd <- latdat[latdat$condition_str == "ND",]$latency
> sd <- latdat[latdat$condition_str == "SD",]$latency
> ad <- latdat[latdat$condition_str == "AD",]$latency
> #
> # paired=TRUE indicates signed rank test
> # Note in wilcox.test, V is what R calls W.
> print(wilcox.test(nd, sd, paired=TRUE))

```

Wilcoxon signed rank test with continuity correction

```

data: nd and sd
V = 3, p-value = 1.344e-10
alternative hypothesis: true location shift is not equal to 0

> print(wilcox.test(nd, ad, paired=TRUE))

```

Wilcoxon signed rank test with continuity correction

```

data: nd and ad
V = 1, p-value = 1.203e-10
alternative hypothesis: true location shift is not equal to 0

```

```

> print(wilcox.test(ad, sd, paired=TRUE, exact=T))

      Wilcoxon signed rank test

data:  ad and sd
V = 1042, p-value = 0.02208
alternative hypothesis: true location shift is not equal to 0

>
> #
> # Cliff's delta
> require(effsize)
> #
> # Between No Distractor and Synchronous Distractor:
> cliff.delta(nd,sd)

Cliff's Delta

delta estimate: -0.7500826 (large)
95 percent confidence interval:
      inf      sup
-0.8601172 -0.5733459
> #
> # Between No Distractor and Asynchronous Distractor:
> cliff.delta(nd,ad)

Cliff's Delta

delta estimate: -0.8009917 (large)
95 percent confidence interval:
      inf      sup
-0.9070022 -0.5995156
> #
> # Between Asynchronous Distractor and Synchronous Distractor:
> cliff.delta(ad,sd)

Cliff's Delta

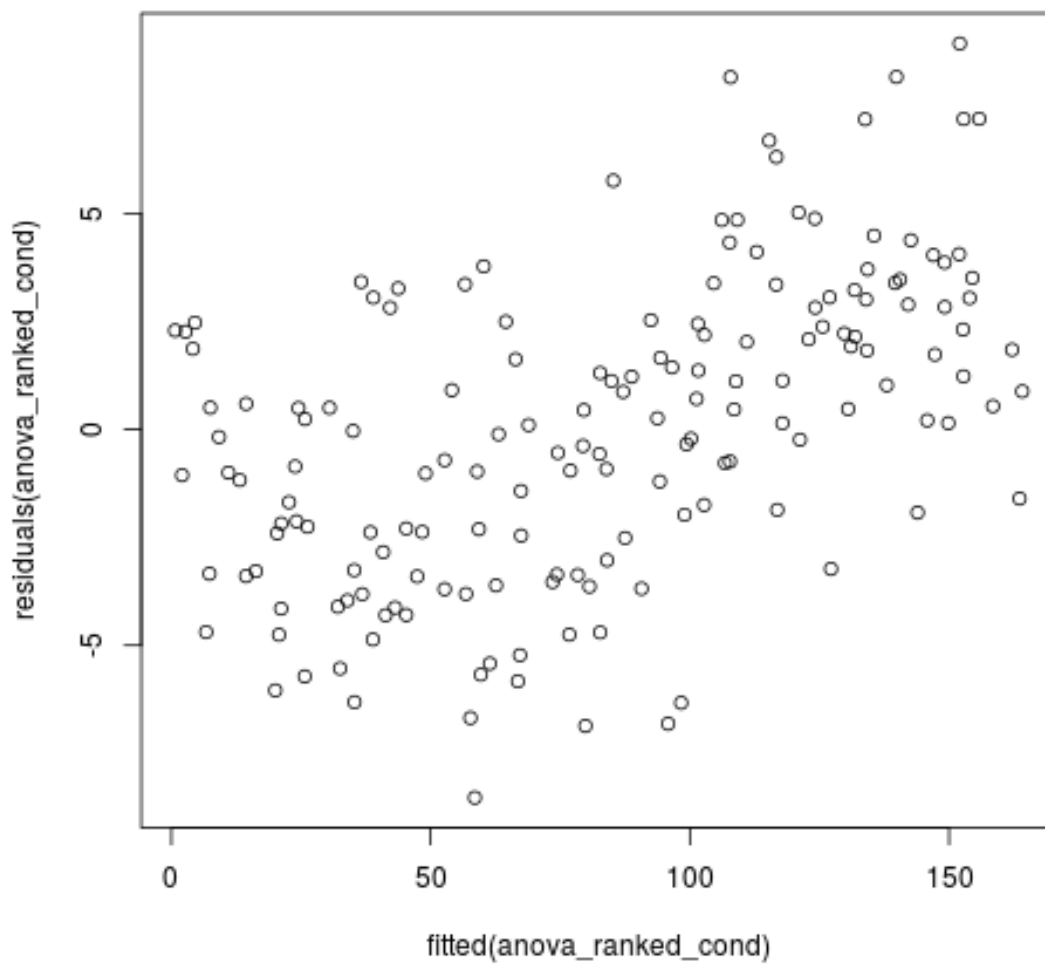
delta estimate: 0.1768595 (small)
95 percent confidence interval:
      inf      sup
-0.03903052  0.37697444
>

In [22]: print 'Residuals (Condition):'
         Image(filename='r_images/anova_ranked_cond_resid.png')

Residuals (Condition):

Out[22]:

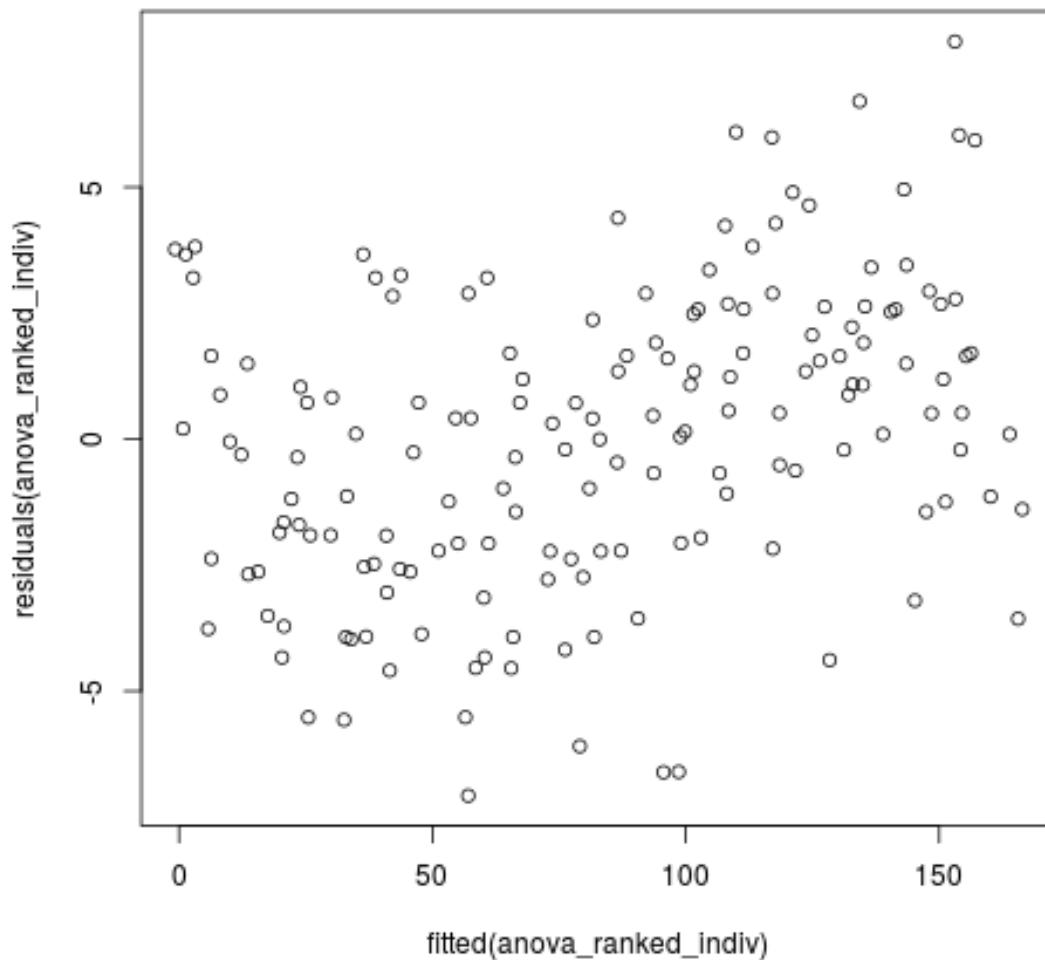
```



```
In [23]: print 'Residuals (Individual)'  
         Image(filename='r_images/anova_ranked_indiv_resid.png')
```

Residuals (Individual)

Out[23]:



1.11 Individual analysis of latencies

Now run **Bootstrap_indiv.r** to compute per-individual results.

```
In [24]: from subprocess import CalledProcessError, check_output
try:
    out = check_output(["R", "-q --file=Bootstrap_indiv.r"])
except CalledProcessError as err:
    print 'Error for:',err.cmd,'with error code:',err.returncode
    print "\nOutput:\n\n",err.output
else:
    for line in out.splitlines():
        #if line[0] != ">" and line[0] != "+":
        print line
```

```

> # Here, I'm going to bootstrap the latencies of each individual
>
> set.seed(19742016)
>
> # This loads individual latencies. each file contains two columns,
> # latency and condition_str. The latencies are all from non-error
> # events.
> fns <- list.files(pattern="IndDat*")
>
> fasterSlower <- c()
>
> sdadNoDiff<-0
> adFaster<-1
> sdFaster<-2
>
> ndall <- c()
> sdall <- c()
> adall <- c()
> for (indiv in fns) {
+
+   latdat <- read.csv(indiv)
+
+   # Extract latencies for the three conditions
+   nd <- latdat[latdat$condition_str == "ND",]$latency
+   sd <- latdat[latdat$condition_str == "SD",]$latency
+   ad <- latdat[latdat$condition_str == "AD",]$latency
+
+   # Make vectors of all the latencies together, for all individuals
+   ndall <-c(ndall, nd)
+   sdall <-c(sdall, sd)
+   adall <-c(adall, ad)
+
+   lat.n <- 1000
+
+   # Plot will show that 87% of the possible means that I could have
+   # measured for SD and AD were different with
+   lat.highconf <- 0.9
+   lat.lowconf <- 1-lat.highconf
+   lat.confint <- 1 - 2*lat.lowconf
+
+   # Colour scheme
+   lat.ndcol <- "black"
+   lat.sdcol <- "steelblue"
+   lat.adcol <- "red"
+
+   lat.minx <- min(c(nd,sd,ad))
+   lat.maxx <- max(c(nd,sd,ad))
+   lat.miny <- 0
+   lat.maxy <- 0.15
+
+   # ND is blue
+   lat.nd.mean <- numeric(lat.n) # numeric vector lat.n long
+   # This is a bootstrap loop:
+   for(i in 1:lat.n) {

```



```

+       this.samp <- nd[ sample(length(nd), length(nd), replace=TRUE) ]
+       lat.nd.mean[i] <- mean(this.samp)
+     }
+   df <- density(lat.nd.mean, n=1024)
+
+   setEPS()
+   # This fails to make the eps files (dev.copy fails). However, we don't really need those for now.
+   postscript(file=sprintf('../paper/figures/bootstrap_indiv_%s.eps', indiv))
+   dev.copy (png, filename=sprintf('r_images/bootstrap_indiv_%s.png', indiv))
+   plot(df, lwd=3, col=lat.ndcol, ylim=range(lat.miny,lat.maxy), xlim=range(lat.minx:lat.maxx), main=)
+   abline(v=mean(nd), lty=2, lwd=2, col=lat.ndcol)
+   q5nd <- quantile(lat.nd.mean,lat.lowconf)
+   abline(v=q5nd[1], lty=6, lwd=2, col=lat.ndcol)
+   q95nd <- quantile(lat.nd.mean,lat.highconf)
+   abline(v=q95nd[1], lty=6, lwd=2, col=lat.ndcol)
+
+   # SD is dark red
+   lat.sd.mean <- numeric(lat.n) # numeric vector lat.n long
+   for(i in 1:lat.n) {
+     this.samp <- sd[ sample(length(sd), length(sd), replace=TRUE) ]
+     lat.sd.mean[i] <- mean(this.samp)
+   }
+   df <- density(lat.sd.mean, n=1024)
+   lines(df, lwd=3, col=lat.sdcol, xlim=range(280:365))
+   abline(v=mean(sd), lty=2, lwd=2, col=lat.sdcol)
+   q5sd <- quantile(lat.sd.mean,lat.lowconf)
+   abline(v=q5sd[1], lty=6, lwd=2, col=lat.sdcol)
+   q95sd <- quantile(lat.sd.mean,lat.highconf)
+   abline(v=q95sd[1], lty=6, lwd=2, col=lat.sdcol)
+
+   # AD is black
+   lat.ad.mean <- numeric(lat.n) # numeric vector lat.n long
+   for(i in 1:lat.n) {
+     this.samp <- ad[ sample(length(ad), length(ad), replace=TRUE) ]
+     lat.ad.mean[i] <- mean(this.samp)
+   }
+   df <- density(lat.ad.mean, n=1024)
+   lines(df, lwd=3, col=lat.adcol, xlim=range(280:365))
+   abline(v=mean(ad), lty=2, lwd=2, col=lat.adcol)
+   q5ad <- quantile(lat.ad.mean,lat.lowconf)
+   abline(v=q5ad[1], lty=6, lwd=2, col=lat.adcol)
+   q95ad <- quantile(lat.ad.mean,lat.highconf)
+   abline(v=q95ad[1], lty=6, lwd=2, col=lat.adcol)
+
+   # Stick in a legend
+   legend (320,0.15, bg="white",
+          c("PDF, ND resampled latencies","ND mean latency",
+            paste("ND latency conf. interval",lat.confint),"ibid, SD","ibid, AD"),
+          lty=c(1,2,6,1,1),
+          lwd=c(3,2,2,3,3),
+          col=c(lat.ndcol,lat.ndcol,lat.ndcol,lat.sdcol,lat.adcol))

```

```

+
+   # Close plotting devices
+   dev.off(dev.prev())
+   dev.off()
+
+   if (lat.sd.mean < lat.ad.mean) {
+     if (q5ad < q95sd) {
+       # overlap SD==AD
+       fasterSlower[indiv] <- sdadNoDiff
+     } else {
+       # no overlap, SD faster
+       fasterSlower[indiv] <- sdFaster
+     }
+   } else {
+     if (q5sd < q95ad) {
+       # overlap SD==AD
+       fasterSlower[indiv] <- sdadNoDiff
+     } else {
+       # no overlap AD faster
+       fasterSlower[indiv] <- adFaster
+     }
+   }
+ }
+ }
+
>
>
> print (sprintf ("%d individuals were faster in the AD condition", length(fasterSlower[fasterSlower=="adFaster"])
[1] "6 individuals were faster in the AD condition"
> print (sprintf ("%d individuals were faster in the SD condition", length(fasterSlower[fasterSlower=="sdFaster"])
[1] "13 individuals were faster in the SD condition"
> print (sprintf ("%d individuals were no faster in either AD or SD", length(fasterSlower[fasterSlower=="sdadNoDiff"])
[1] "36 individuals were no faster in either AD or SD"
>
> # Output the individuals from fasterSlower in a form suitable for copy
> # & paste into python, which is formatted like this:
> #
> # fasterSyncSubjects = ['CD1','CD2','CP','EC1','EC2','EF','IR','RF','RQ','SB2']
> fssNames <- c()
> fss <- "fasterSyncSubjects = ["
> for (s in names(fasterSlower[fasterSlower=="sdFaster"])) {
+   # Process name, remove "IndDat" and ".csv":
+   s <- sub ("IndDat", "", s)
+   s <- sub (".csv", "", s)
+   fssNames <- c(fssNames, s)
+   # Switch the R-form of "NA" back to the original as used in python:
+   s <- sub ("NA_", "NA", s)
+   fss <- sprintf ("%s '%s'", fss, s)
+ }
> fss <- sprintf ("%s]", fss)
> # Get rid of last ', '
> fss <- sub (",]", " ]", fss)
> print (sprintf("fasterSyncSubjects (N=%d) python list code:", length(fssNames)))
[1] "fasterSyncSubjects (N=13) python list code:"
> print (fss)
[1] "fasterSyncSubjects = [ 'AB1_', 'CD1', 'CD2', 'CH', 'CP', 'EC1', 'EC2', 'EF', 'IR', 'RF', 'RQ', 'SB1'

```

```

>
>
> fasNames <- c()
> fas <- "fasterAsyncSubjects = ["
> for (s in names(fasterSlower[fasterSlower==adFaster])) {
+   s <- sub ("IndDat", "", s)
+   s <- sub (".csv", "", s)
+   fasNames <- c(fasNames, s)
+   s <- sub ("NA_", "NA", s)
+   fas <- sprintf ("%s '%s'", fas, s)
+ }
> fas <- sprintf ("%s]", fas)
> fas <- sub (",]", " ]", fas)
> print (sprintf("fasterAsyncSubjects (N=%d) python list code:", length(fasNames)))
[1] "fasterAsyncSubjects (N=6) python list code:"
> print (fas)
[1] "fasterAsyncSubjects = [ 'AW3_', 'EJ', 'LH1', 'NA', 'PO', 'SF' ]"
>
>
> ndNames <- c()
> nds <- "noDiffSubjects = ["
> for (s in names(fasterSlower[fasterSlower==sdadNoDiff])) {
+   s <- sub ("IndDat", "", s)
+   s <- sub (".csv", "", s)
+   ndNames <- c(ndNames, s)
+   s <- sub ("NA_", "NA", s)
+   nds <- sprintf ("%s '%s'", nds, s)
+ }
> nds <- sprintf ("%s]", nds)
> nds <- sub (",]", " ]", nds)
> print (sprintf("noDiffSubjects (N=%d) python list code:", length(ndNames)))
[1] "noDiffSubjects (N=36) python list code:"
> print (nds)
[1] "noDiffSubjects = [ 'AL_', 'AM', 'AS', 'AW1_', 'AW2_', 'BG', 'CM', 'EB1', 'EB2', 'EK', 'EM2', 'EW1_',
>
>
> #
> # Now do analysis on ndall etc.
> #
>
> # Save some variables that are used in Bootstrap_all and Err.r
> save (ndall, sdall, adall, fssNames, fasNames, ndNames, file="all_latencies.rdat", ascii=TRUE)
>
> print ('You can now call Bootstrap_all.r for "the groupstrap"')
[1] "You can now call Bootstrap_all.r for \"the groupstrap\""
> print ('You can also call Err.r for the per-group error analysis')
[1] "You can also call Err.r for the per-group error analysis"
>

```

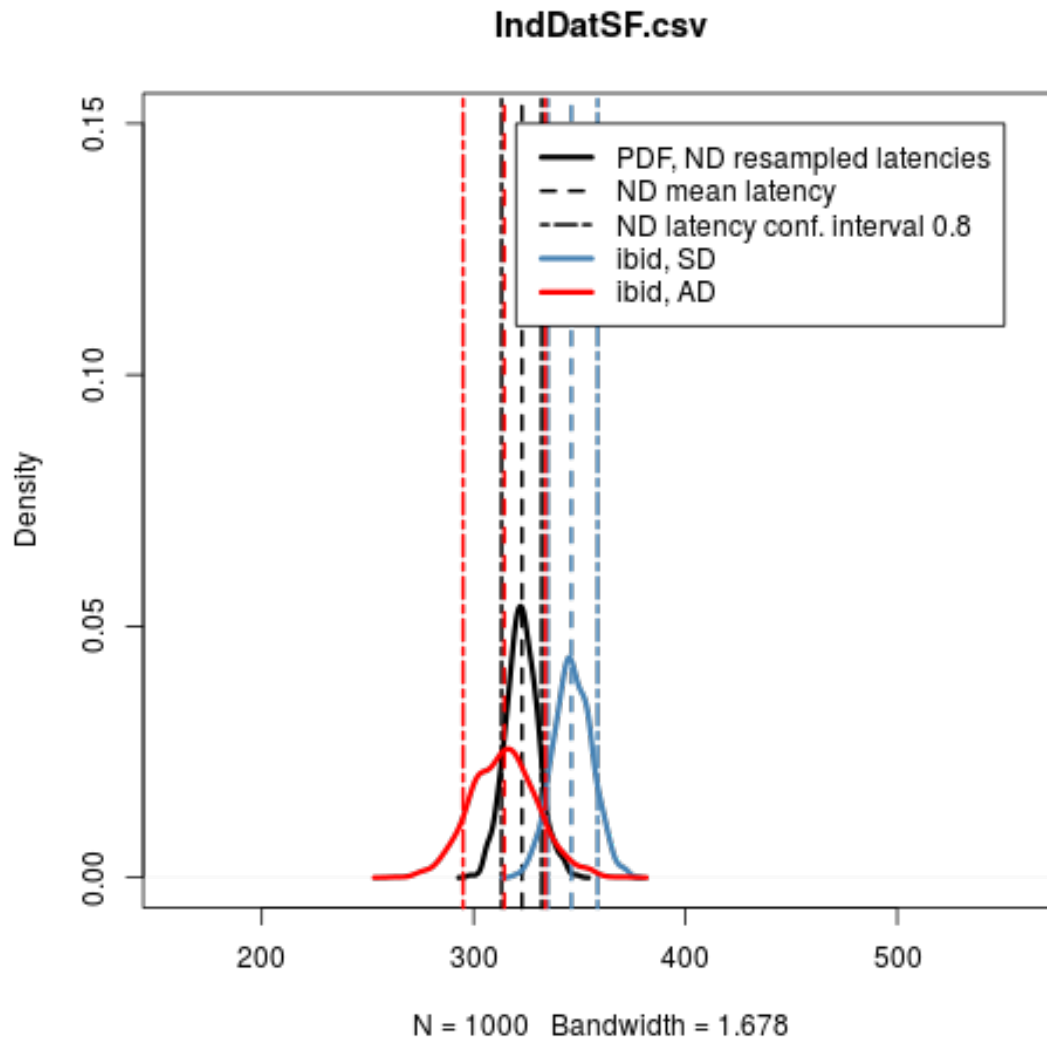
```

In [25]: print 'An example individual set of latency bootstrap means'
         Image(filename='r_images/bootstrap_indiv_IndDatSF.csv.png')

```

An example individual set of latency bootstrap means

```
Out[25]:
```



1.12 Bootstrap analysis of latencies

After calling **Bootstrap_indiv.r**, **Bootstrap_all.r** can be called.

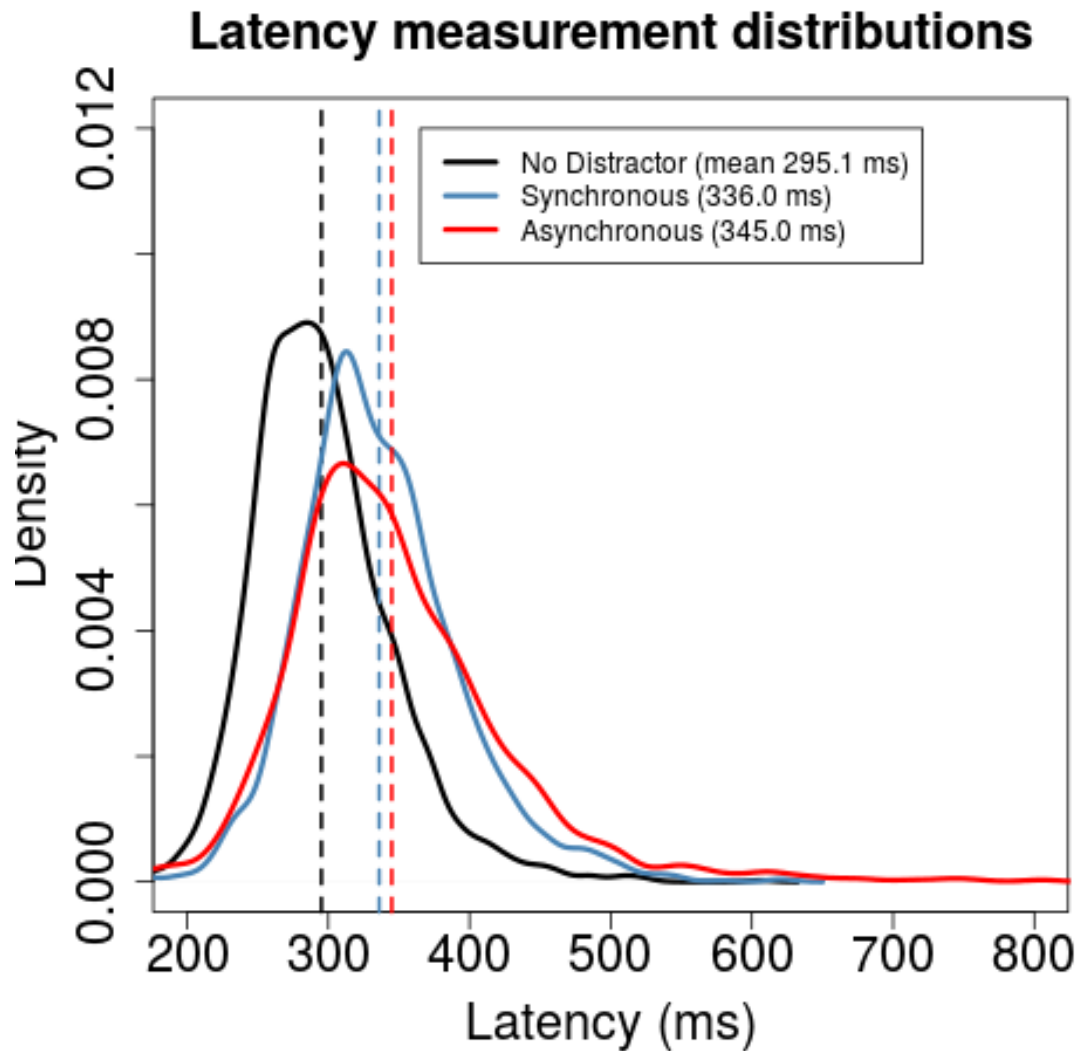
```
In [26]: from subprocess import CalledProcessError, check_output
try:
    out = check_output(["R", "-q --file=Bootstrap_all.r"])
except CalledProcessError as err:
    print 'Error for:',err.cmd,'with error code:',err.returncode
    print "\nOutput:\n\n",err.output
else:
    print 'Success'
    #for line in out.splitlines():
    #    print line
```

Success

```
In [27]: print 'Latency measurement distributions, all subjects'
         Image(filename='r_images/data_density_all.png')
```

Latency measurement distributions, all subjects

Out[27]:

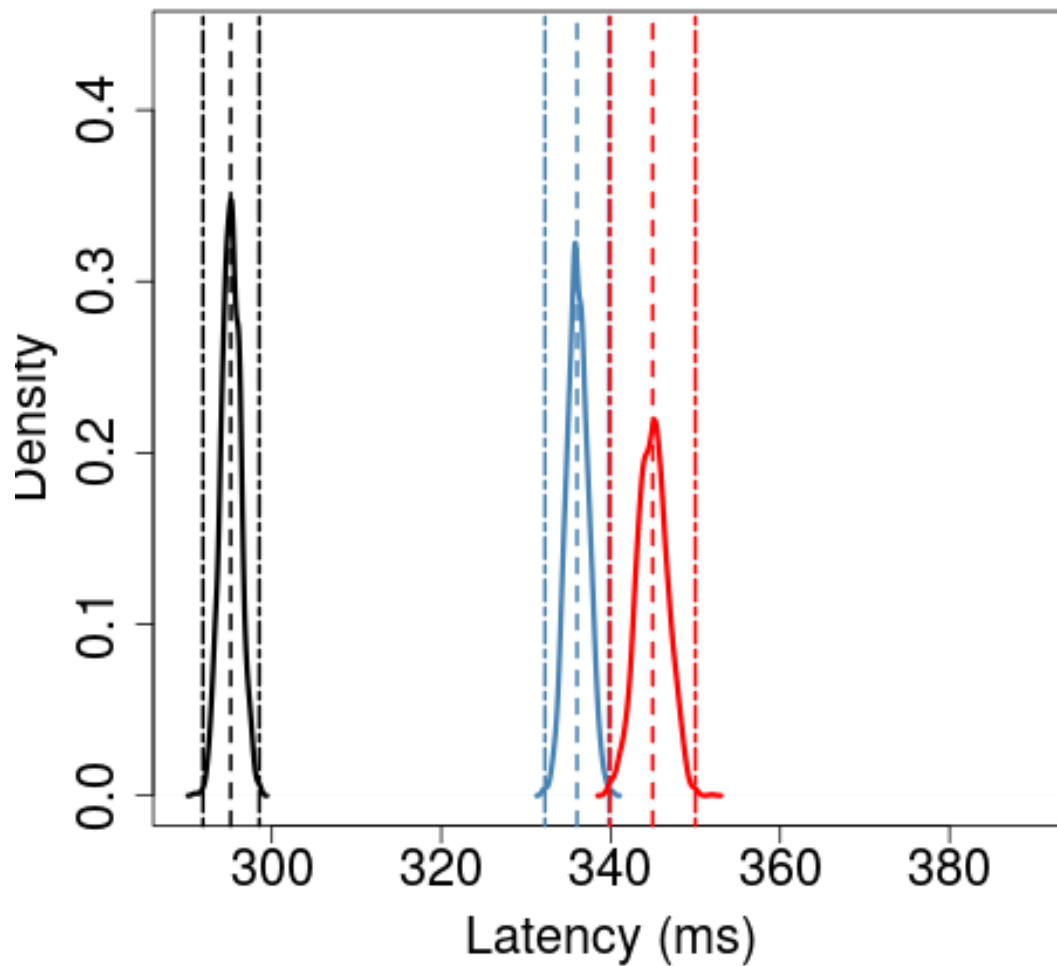


```
In [28]: print 'Bootstrapped latency means, all subjects'
         Image(filename='r_images/bootstrapped_means_all.png')
```

Bootstrapped latency means, all subjects

Out[28]:

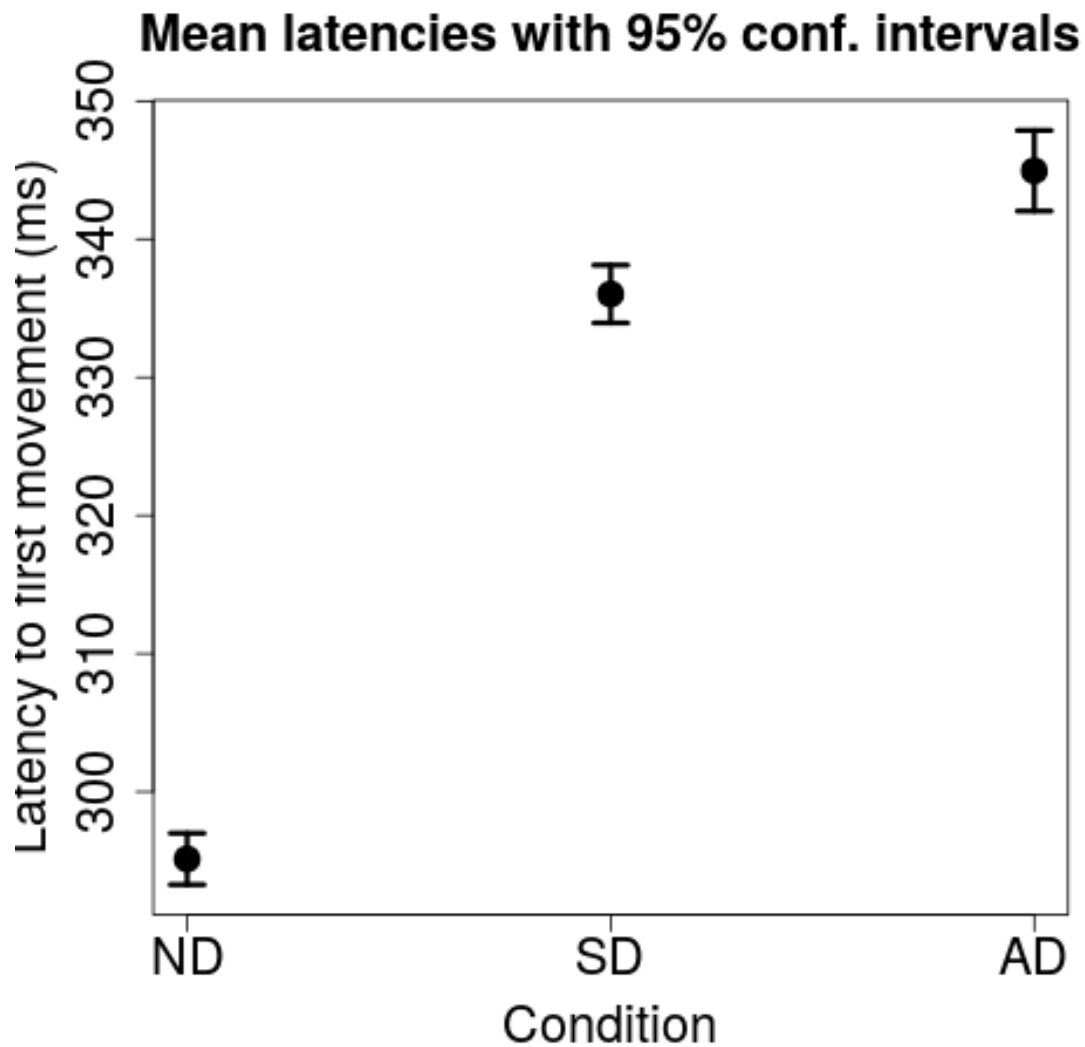
Distributions of bootstrapped means



```
In [29]: print 'Means again, this time with 95% confidence intervals as error bars'  
         Image(filename='r_images/meanmad_all.png')
```

Means again, this time with 95% confidence intervals as error bars

Out[29]:



1.13 Analysis of Movement Errors

It's possible to count the number of errors which the subject makes when following the target. In some trials, the subject will mistakenly begin to move towards the distractor line, before correcting (sometimes after a very short period of time) to move towards the target. I wanted to see whether the synchronous distractor or the asynchronous distractor produce more errors, but it is hard to compare. In the synchronous distractor condition, there are always the same number of distractor events as target events. In the asynchronous distractor condition, there may be more distractor events (or fewer) than target events. I concluded that it's not possible to compare SD and AD conditions.

1.13.1 Overall and Group-based movement error analysis

This R code block computes mean error rates and their MAD and StDev statistics, and it also then computes Wilcoxon signed rank test and Cliff's Delta for the ND and SD error rates. There is a bootstrap analysis of the error rates and then, in the same file, a bootstrap analysis of the fasterSync/fasterAsync sub-groups.

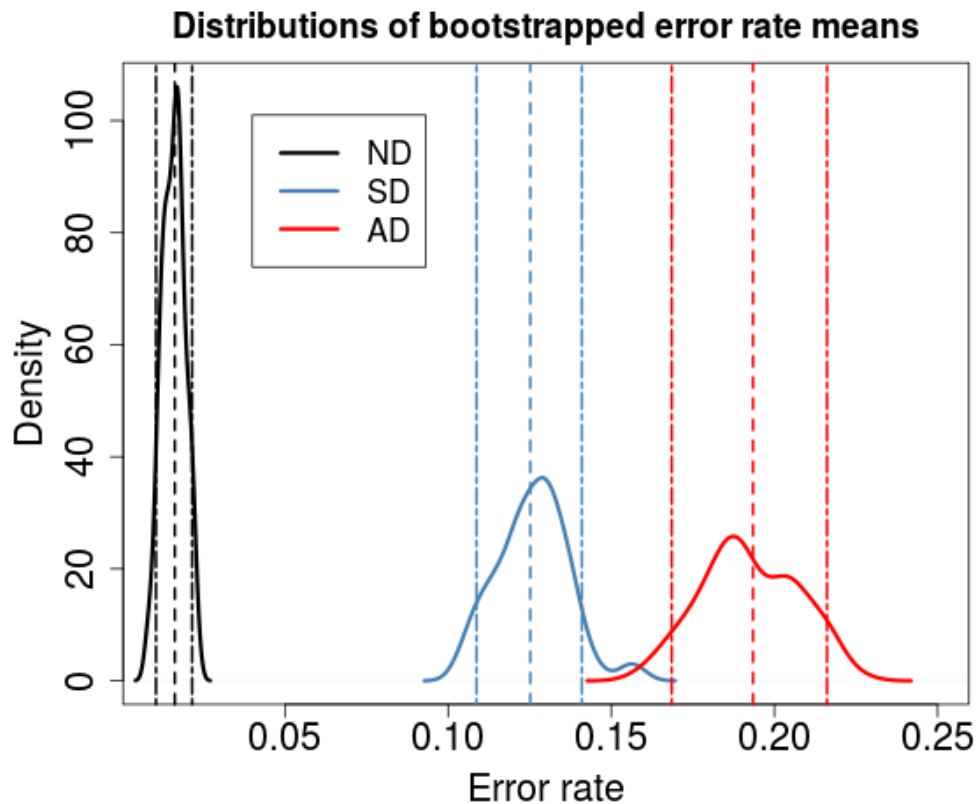
```
In [30]: from subprocess import CalledProcessError, check_output
try:
    out = check_output(["R", "-q --file=Err.r"])
except CalledProcessError as err:
    print 'Error for:',err.cmd,'with error code:',err.returncode
    print "\nOutput:\n\n",err.output
else:
    print 'Success'
    #for line in out.splitlines():
    #    print line
```

Success

```
In [31]: print 'Bootstrapped error means'
Image(filename='r_images/bootstrap_errors.png')
```

Bootstrapped error means

Out[31]:



```
In [32]: from subprocess import CalledProcessError, check_output
try:
    out = check_output(["R", "-q --file=Bootstrap_async.r"])
except CalledProcessError as err:
    print 'Error for:',err.cmd,'with error code:',err.returncode
```



```

        print "\nOutput:\n\n",err.output
    else:
        for line in out.splitlines():
            print line

> #
> # Analysis of Async-specific statistics. Especially:
> #
> # 1) Does last event recency affect latency of distractor movements? (Ans: No)
> # 2) Does distractor recency affect latency of target movements? (Ans: No)
> # 3) Does distractor in same direction as target reduce latency?
> # 4) What's the reaction time for distractors cf. targets?
> #
>
> maxlatency <- 1000
>
> # a function which will bootstrap the standard error of the mean
> bs.mean <- function(data, num) {
+   resamples <- lapply(1:num, function(i) sample(data, replace=T))
+   r.mean <- sapply(resamples, mean)
+   std.err <- sqrt(var(r.mean))
+   list(std.err=std.err, resamples=resamples, means=r.mean)
+ }
> bs.median <- function(data, num) {
+   resamples <- lapply(1:num, function(i) sample(data, replace=T))
+   r.median <- sapply(resamples, median)
+   std.err <- sqrt(var(r.median))
+   list(std.err=std.err, resamples=resamples, medians=r.median)
+ }
>
> d <- read.csv('AsyncTrials.csv')
>
> set.seed(197420162)
>
> # This takes a 2-D data set of latency vs time since last event. It
> # collects "bins" of data by dividing time since last event into 30
> # bins, then computes bootstrap mean & median values for each bin.
> binned <- function (data, fname) {
+   listPoints <- list()
+   listBStrap <- list()
+   dfBStrap <- data.frame()
+   nBreaks <- 30
+   iter <- 1
+   nResamples <- 200
+   h <- hist(data$timesincelast, breaks=nBreaks, plot=F)
+   for (b in h$breaks) {
+     if (b==0) {
+       bLast = b
+       next
+     }
+
+     points <- data[which(data$timesincelast >= bLast & data$timesincelast < b),]
+     listPoints[[iter]] <- points
+
+   }
+ }

```

```

+         # Now bootstrap each member of listPoints, compute mean & std err of mean
+         bsmed <- bs.median(points$latency, nResamples)
+         bsmean <- bs.mean(points$latency, nResamples)
+         if (nrow(points)) {
+             dfBStrap <- rbind (dfBStrap, c((b-bLast/2), median(points$latency),
+                                     bsmed$std.err, mean(points$latency),
+                                     bsmean$std.err))
+         }
+
+         iter <- iter + 1
+         bLast = b
+     }
+     names(dfBStrap) <- c("timesincelast","median","med.stderr","mean","mean.stderr")
+
+     png (filename=fname)
+     plot (dfBStrap$timesincelast,dfBStrap$median, xlim=c(0,1500),# ylim=c(0,400),
+          pch=19, cex=1, cex.lab=1.5, cex.axis=1.5, cex.main=1.5, cex.sub=1.5,
+          xlab="Time since last event (ms)",ylab="Median latency (ms)", col="black")
+     lines(dfBStrap$timesincelast,dfBStrap$median, lty=1, lwd=2)
+     arrows(dfBStrap$timesincelast,dfBStrap$median-dfBStrap$med.stderr,dfBStrap$timesincelast,dfBStrap$
+     dev.off()
+ }
>
> #
> # 1) Does last event recency affect latency of distractor movements?
> #
>
> # This selects out distractor latencies and allows me to plot the
> # latencies vs. time since last event.
> # Note: 15 is "stylus didn't move away from target" 16 is "movement occurs beyond next target"
> dlt <- d[which (d$type == 0 & d$correctmove == 0 & d$latency<maxlatency & !(d$omit == 1 & (d$reas ==
> png (filename='./r_images/async_dist.timesince_vs_latency.png')
> plot (dlt$timesincelast,dlt$latency,xlab="Time since last event (ms)",ylab="Latency to distracted mot.
> dev.off()
null device
      1
> binned (dlt, './r_images/async_dist.binned_latencies.png')
null device
      1
>
> #
> # 2) Does distractor recency affect latency of target movements?
> #
> tlt <- d[which (d$type == 1 & d$correctmove == 1 & d$latency<maxlatency & d$omit == 0),c("latency","t.
> png (filename='./r_images/async_targ.timesince_vs_latency.png')
> plot (tlt$timesincelast,tlt$latency)
> dev.off()
null device
      1
> binned (tlt, './r_images/async_targ.binned_latencies.png')
null device
      1
>
> # Moving bin approach for timesincelast (Question 2)

```

```

> movingbinwidth <- 80 # ms
> movingbin <- function (data) {
+   # First sort data wrt direction
+   data <- data[order(data$timesincelast),]
+   listPoints <- list()
+   listBStrap <- list()
+   dfBStrap <- data.frame()
+   iter <- 1
+   nResamples <- 200
+   maxtime <- 1000 # ms
+   bLast <- -1
+   for (b in unique(data$timesincelast)) {
+     if (b < movingbinwidth/2) {
+       next
+     }
+     if (maxtime - b < movingbinwidth/2) {
+       next
+     }
+     points <- data[which(data$timesincelast >= b-movingbinwidth/2 & data$timesincelast < b+movingbinwidth/2),]
+     listPoints[[iter]] <- points
+     bsmed <- bs.median(points$latency, length(points$latency))
+     bsmean <- bs.mean(points$latency, length(points$latency))
+     if (nrow(points)) {
+       dfBStrap <- rbind (dfBStrap, c(b, median(points$latency),
+                                       bsmed$std.err, mean(points$latency),
+                                       bsmean$std.err))
+     }
+     iter <- iter + 1
+     bLast = b
+   }
+   names(dfBStrap) <- c("distance", "median", "med.stderr", "mean", "mean.stderr")
+   return (dfBStrap)
+ }
>
> amb <- movingbin (tlt)
> png (filename='./r_images/async_targ.timesince_vs_latency_movingbin.png')
> plot (amb$distance, amb$mean, xlim=c(0,1000), ylim=c(270,405),
+       pch=19, cex=0.2, cex.lab=1.5, cex.axis=1.5, cex.main=1.5, cex.sub=1.5,
+       xlab="Time since last event (ms)", ylab="Mean latency (ms)", col="white")
> lines(amb$distance, amb$mean, lty=1, lwd=3, col="red")
> lines(amb$distance, amb$mean-1.96*amb$mean.stderr, lty=1, lwd=1, col="red")
> lines(amb$distance, amb$mean+1.96*amb$mean.stderr, lty=1, lwd=1, col="red")
> dev.off()
null device
    1
>
> # A linear model shows that timesincelast does not predict latency:
> lmt <- lm(latency ~ timesincelast, data = tlt)
> print(summary(lmt))

```

Call:

```
lm(formula = latency ~ timesincelast, data = tlt)
```

Residuals:

Min	1Q	Median	3Q	Max
-246.30	-50.14	-9.82	37.51	459.08

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	3.433e+02	2.787e+00	123.141	<2e-16 ***
timesincelast	2.950e-03	3.821e-03	0.772	0.44

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 81.31 on 2230 degrees of freedom

Multiple R-squared: 0.0002673, Adjusted R-squared: -0.000181

F-statistic: 0.5962 on 1 and 2230 DF, p-value: 0.4401

```
>
> #
> # 3) Does distractor in same direction as target reduce latency?
> #
>
> # A graphing function for target data and distractor data, both
> # assumed to be simple vectors of data
> bs_graph <- function(dat1, dat2, dat1label, dat2label, xrange, yrange, alpha, mean=FALSE) {
+
+   densityN <- 256
+   nResamples <- 1024
+   dat1Col <- "steelblue"
+   dat2Col <- "red"
+
+   # Raw distributions
+   dat1dens = density(dat1, n=512)
+   dat1densScale = max(dat1dens$y)
+   dat2dens = density(dat2, n=512)
+   dat2densScale = max(dat2dens$y)
+   xmax = max(dat1dens$x, dat2dens$x)
+   xmin = min(dat1dens$x, dat2dens$x)
+
+   if (xrange[2]==0) {
+     xrange <- c(xmin, xmax)
+   }
+
+   if (mean == TRUE) {
+     bsdat1 <- bs.mean(dat1, nResamples)
+     q5 <- quantile(bsdat1$means, alpha)
+     q95 <- quantile(bsdat1$means, 1-alpha)
+     df <- density(bsdat1$means, n=densityN)
+     cent <- mean(dat1)
+     maintitle <- sprintf("Distns of bootstrapped means. alpha=%.3f", alpha)
+     bsdat2 <- bs.mean(dat2, nResamples)
+     df_ <- density(bsdat2$means, n=densityN)
+     centtype <- 'mean'
+   } else {
+     bsdat1 <- bs.median(dat1, nResamples)
+     q5 <- quantile(bsdat1$medians, alpha)
+     q95 <- quantile(bsdat1$medians, 1-alpha)
+   }
+ }
```

```

+     df <- density(bsdat1$medians, n=densityN)
+     cent <- median(dat1)
+     maintitle <- sprintf("Distns of bootstrapped medians. alpha=%.3f", alpha)
+     bsdat2 <- bs.median(dat2, nResamples)
+     df_ <- density(bsdat2$medians, n=densityN)
+     centtype <- 'median'
+   }
+
+   print (sprintf('%s: %s = %f +- %f', dat1label, centtype, cent, q95-cent))
+   print (sprintf('%s: std %f, mad: %f', dat1label, sd(dat1), mad(dat1)))
+
+   if (yrange[2]==0) {
+     yrange <- c(0,max(df$y,df_$y))
+   }
+
+   # Plot dat1 first
+   plot(df, lwd=2, col=dat1Col, xlim=xrange, ylim=yrange,
+        cex.lab=1.5, cex.axis=1.5, cex.main=1.5, cex.sub=1.5,
+        main=maintitle, xlab="Latency (ms)")
+   abline(v=cent, lty=2, lwd=2, col=dat1Col)
+   abline(v=q5[1], lty=6, lwd=2, col=dat1Col)
+   abline(v=q95[1], lty=6, lwd=2, col=dat1Col)
+
+   dat1densScale = max(df$y) / dat1densScale
+   print (dat1densScale)
+   lines(dat1dens$x, dat1dens$y * dat1densScale, lwd=2, lty=5, col=dat1Col)
+
+
+   if (mean == TRUE) {
+     q5 <- quantile(bsdat2$means,alpha)
+     q95 <- quantile(bsdat2$means,1-alpha)
+     df <- density(bsdat2$means, n=densityN)
+     cent <- mean(dat2)
+   } else {
+     q5 <- quantile(bsdat2$medians,alpha)
+     q95 <- quantile(bsdat2$medians,1-alpha)
+     df <- density(bsdat2$medians, n=densityN)
+     cent <- median(dat2)
+   }
+
+   print (sprintf('%s: %s = %f +- %f', dat2label, centtype, cent, q95-cent))
+   print (sprintf('%s: std %f, mad: %f', dat2label, sd(dat2), mad(dat2)))
+
+   # Now plot dat2
+   lines(df, lwd=2, col=dat2Col,
+        cex.lab=1.5, cex.axis=1.5, cex.main=1.5, cex.sub=1.5,
+        main=maintitle, xlab="Latency (ms)")
+   abline(v=cent, lty=2, lwd=2, col=dat2Col)
+   abline(v=q5[1], lty=6, lwd=2, col=dat2Col)
+   abline(v=q95[1], lty=6, lwd=2, col=dat2Col)
+
+   dat2densScale = max(df$y) / dat2densScale
+   lines(dat2dens$x, dat2dens$y * dat2densScale, lwd=2, lty=5, col=dat2Col)

```

```

+
+   legend ("topleft", c(dat1label, dat2label),
+         lty=c(1,1),
+         lwd=c(2,2),
+         col=c(dat1Col,dat2Col)
+       )
+   legend ("topright", c('bootstrap distribution','raw distn (scaled)',centtype,'bootstrap conf. int
+         lty=c(1,5,2,6),
+         lwd=c(2,2,2,2),
+         col=c(dat1Col,dat1Col,dat1Col,dat1Col)
+       )
+
+ }
+
>
> # Get filenames:
> fns <- list.files(pattern="AsyncDat*")
> # Create container
> d2 <- data.frame()
> for (indiv in fns) {
+   dat <- read.csv(indiv)
+   # Add the 'opposite' column:
+   dat$opposite <- 0
+   # From this individual's data, find those targets for which prev. distractor is in same direction
+   for (i in 1:nrow(dat)) {
+     if (dat[i,]$type == 1.0) {
+       # 1 is TARG_EVENT
+       earlier <- dat[which (dat$type == 0 & dat$num < i),]
+       if (nrow(earlier)) {
+         # Got earlier events, now find most recent earlier
+         # event and extract its destination. Compare this with
+         # the target destination.
+         dest_d <- earlier[earlier$num==max(earlier$num),]$destination
+         dest_t <- dat[i,]$destination
+         dir_t <- dat[i,]$direction
+         start <- dest_t - dir_t
+         if ((dest_d < start & dest_t > start)
+             | (dest_d > start & dest_t < start)) {
+           # targ and dist in opposite directions
+           dat[i,]$opposite=1
+         } else {
+           # targ and dist NOT in opposite directions
+         }
+       }
+     }
+   }
+
+   # Lastly? combine data frame with the others to make up a return
+   # data frame from which I can determine if opposite makes a
+   # difference.
+   d2 <- rbind(d2,dat)
+ }
> # Extract opposite and same-side data:
> tlt_opp <- d2[which (d2$type == 1 & d2$opposite == 1 & d2$correctmove == 1 & d2$latency<maxlatency & c
> tlt_same <- d2[which (d2$type == 1 & d2$opposite == 0 & d2$correctmove == 1 & d2$latency<maxlatency &

```

```

> # Now make a nice graph:
> png(filename='./r_images/async_targ_vs_oppositeness_of_distractor.png')
> xrange <- c(325,360)
> yrange <- c(0,0)
> bs_graph(tlt_opp, tlt_same, "distractor opposite", "distractor same", xrange, yrange, 0.05, FALSE) # T
[1] "distractor opposite: median = 330.000000 +- 5.500000"
[1] "distractor opposite: std 83.191973, mad: 60.786600"
[1] 26.33083
[1] "distractor same: median = 336.000000 +- 4.000000"
[1] "distractor same: std 79.961187, mad: 65.234400"
> dev.off()
null device
      1
> # This appears to show that it is likely (0.79 probability) that the
> # "distractor opposite" does have a small effect, increasing the
> # latency by about 5 ms. However, there's a 0.1958 probability that
> # the "distractor opposite" makes *no difference* to the latency and a
> # 0.0121 probability that the "distractor opposite" actually decreases
> # the latency. A 0.2 probability of no effect/the means being opposite
> # doesn't pass the usual 0.05 alpha test, so conclude that this is
> # still non-significant.
>
>
> #
> # 4) What's the reaction time for distractors cf. targets?
> #
> png(filename='./r_images/async_targ_vs_dist.png')
> xrange <- c(180,500)
> yrange <- c(0,0)
> print('Async mean')
[1] "Async mean"
> bs_graph (tlt$latency, dlt$latency, "target latency","distractor latency", xrange, yrange, 0.001, TRUE)
[1] "target latency: mean = 344.943996 +- 5.098332"
[1] "target latency: std 81.303108, mad: 65.234400"
[1] 37.57107
[1] "distractor latency: mean = 270.267241 +- 16.447305"
[1] "distractor latency: std 95.227925, mad: 57.821400"
> dev.off()
null device
      1
> png(filename='./r_images/async_targ_vs_dist_median.png')
> xrange <- c(180,500)
> yrange <- c(0,0)
> print('Async median')
[1] "Async median"
> bs_graph (tlt$latency, dlt$latency, "target latency","distractor latency", xrange, yrange, 0.001, FALSE)
[1] "target latency: median = 335.000000 +- 5.000000"
[1] "target latency: std 81.303108, mad: 65.234400"
[1] 225.7886
[1] "distractor latency: median = 269.000000 +- 12.000000"
[1] "distractor latency: std 95.227925, mad: 57.821400"
> dev.off()
null device
      1

```

```

>
>
> #
> # Last - sync trials.
> #
> d <- read.csv('SyncTrials.csv')
> set.seed(197420163)
> # distracted latencies. correctmove is 0 when definitely incorrect, -1 when undetermined and 1 when d
> dlt <- d[which (d$correctmove == 0 & d$latency<maxlatency & d$latency>0),]$latency
> tlt <- d[which (d$correctmove == 1 & d$latency<maxlatency & d$latency>0),]$latency
> png(filename='./r_images/sync_targ_vs_dist.png')
> xrange <- c(150,550)
> yrange <- c(0,0)
> print('Sync mean')
[1] "Sync mean"
> bs_graph (tlt, dlt, "target latency","distractor latency", xrange, yrange, 0.001, TRUE)
[1] "target latency: mean = 334.104204 +- 4.165407"
[1] "target latency: std 60.492715, mad: 51.891000"
[1] 39.15872
[1] "distractor latency: mean = 263.697248 +- 11.959401"
[1] "distractor latency: std 78.343373, mad: 51.891000"
> dev.off()
null device
      1
>
> png(filename='./r_images/sync_targ_vs_dist_median.png')
> xrange <- c(150,550)
> yrange <- c(0,.33)
> print('Sync median')
[1] "Sync median"
> bs_graph (tlt, dlt, "target latency","distractor latency", xrange, yrange, 0.001, FALSE)
[1] "target latency: median = 330.000000 +- 5.000000"
[1] "target latency: std 60.492715, mad: 51.891000"
[1] 92.50053
[1] "distractor latency: median = 265.000000 +- 14.977000"
[1] "distractor latency: std 78.343373, mad: 51.891000"
> dev.off()
null device
      1
>
>
> #
> # REALLY Last - nodist trials.
> #
> d <- read.csv('NoDistTrials.csv')
> set.seed(197420166)
> # distracted latencies. correctmove is 0 when definitely incorrect, -1 when undetermined and 1 when d
> dlt <- d[which (d$correctmove == 0 & d$latency<maxlatency & d$latency>100),]$latency
> tlt <- d[which (d$correctmove == 1 & d$latency<maxlatency & d$latency>100),]$latency
> png(filename='./r_images/nodist_targ_vs_dist.png')
> xrange <- c(80,550)
> yrange <- c(0,0)
> print('Nodist mean')
[1] "Nodist mean"

```



```

> bs_graph (tlt, dlt, "target latency","distractor latency", xrange, yrange, 0.01, TRUE)
[1] "target latency: mean = 294.902810 +- 2.421714"
[1] "target latency: std 51.334525, mad: 44.478000"
[1] 41.66836
[1] "distractor latency: mean = 229.057143 +- 50.083714"
[1] "distractor latency: std 127.433707, mad: 106.747200"
> dev.off()
null device
      1
>
> png(filename='./r_images/nodist_targ_vs_dist_median.png')
> xrange <- c(80,550)
> yrange <- c(0,0)
> print('Nodist median')
[1] "Nodist median"
> bs_graph (tlt, dlt, "target latency","distractor latency", xrange, yrange, 0.01, FALSE)
[1] "target latency: median = 290.000000 +- 1.000000"
[1] "target latency: std 51.334525, mad: 44.478000"
[1] 221.3067
[1] "distractor latency: median = 204.000000 +- 44.250000"
[1] "distractor latency: std 127.433707, mad: 106.747200"
> dev.off()
null device
      1
>

```

```

In [33]: print '1) Does last event recency affect latency of DISTRACTOR movements? ANS: No.'
         print 'Raw distractor latency vs. time-since-last-event:'
         Image(filename='r_images/async_dist_timesince_vs_latency.png')

```

```

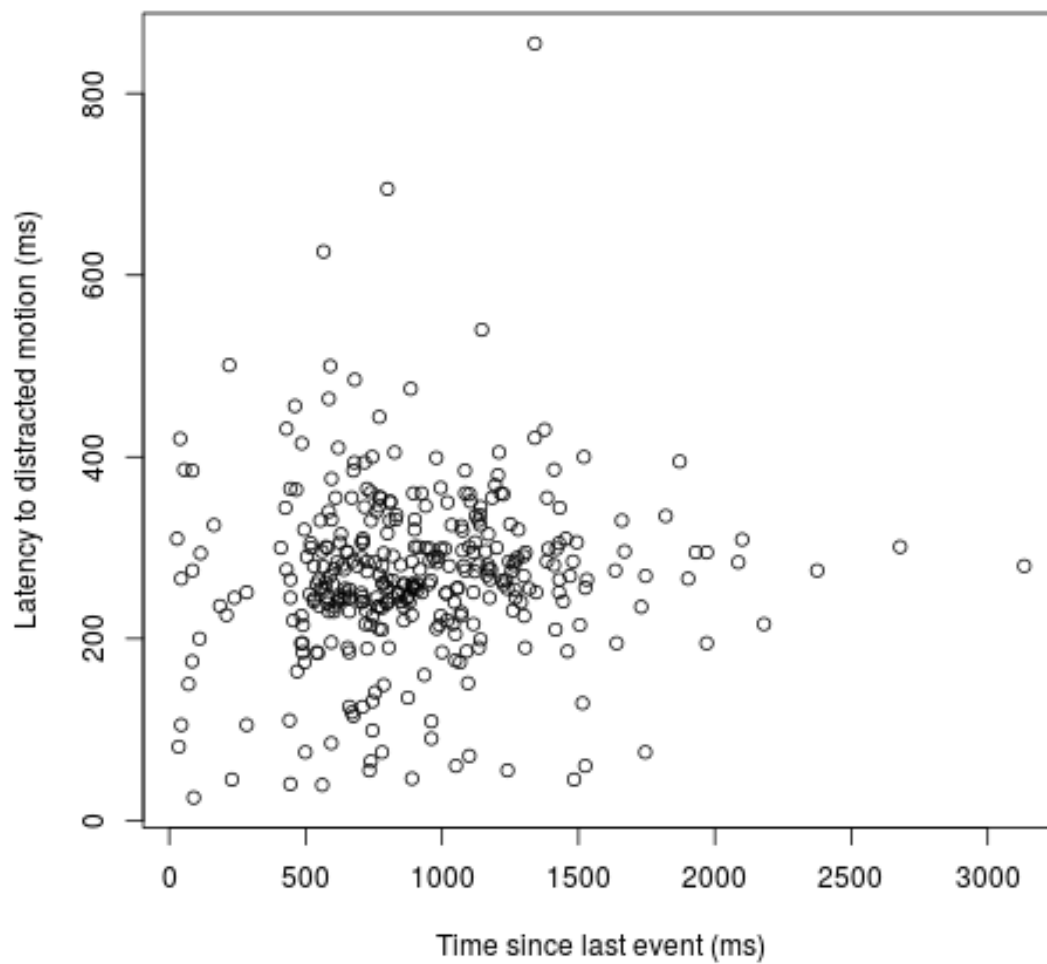
1) Does last event recency affect latency of DISTRACTOR movements? ANS: No.
Raw distractor latency vs. time-since-last-event:

```

```

Out[33]:

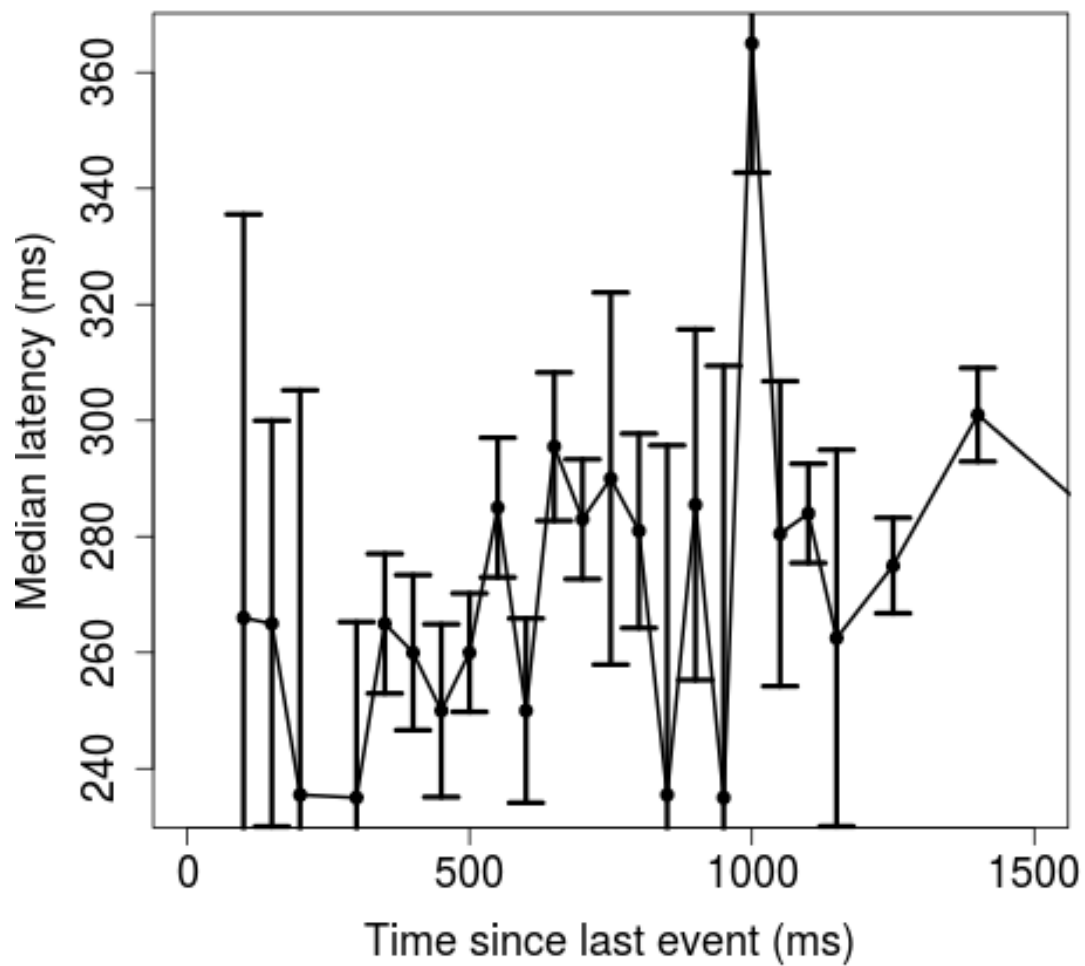
```



```
In [34]: print 'Binned median latencies'
         Image(filename='./r_images/async_dist_binned_latencies.png')
```

Binned median latencies

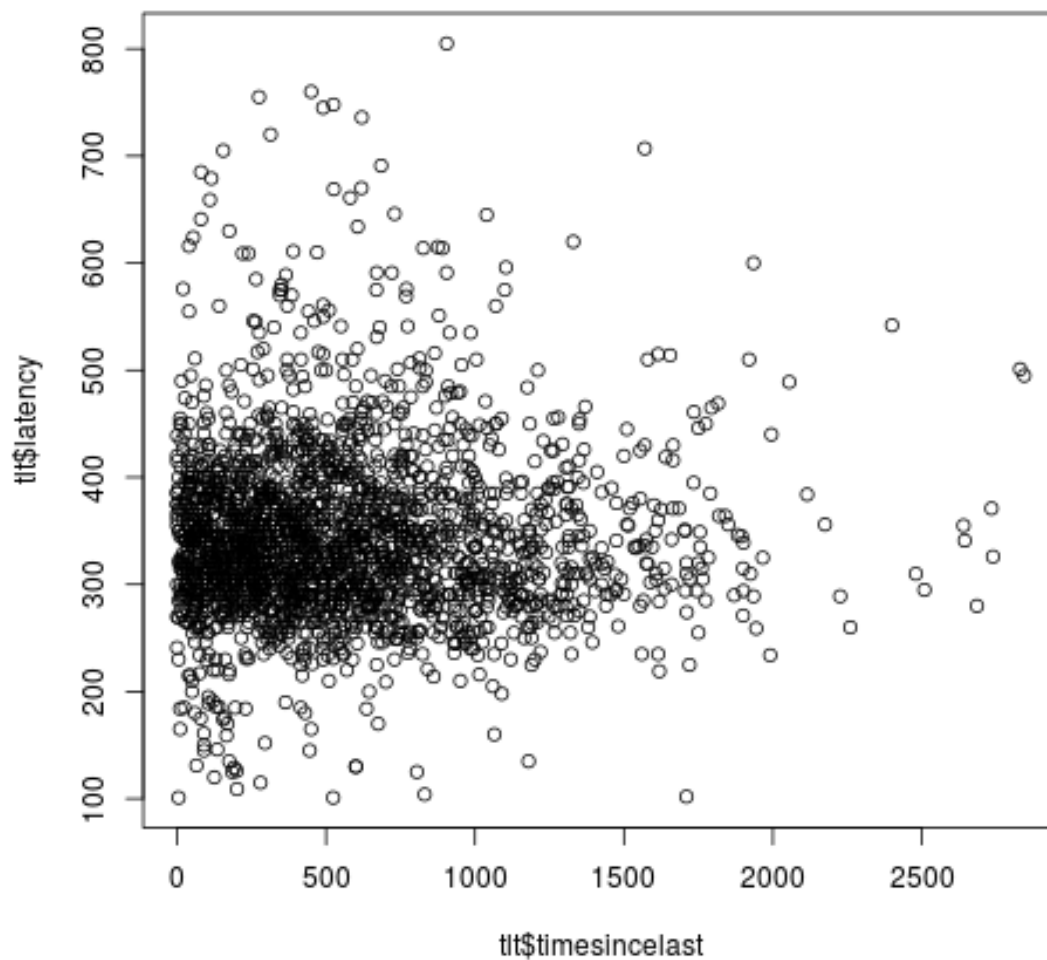
Out[34]:



```
In [35]: print '2) Does last event recency affect latency of TARGET movements? ANS: No.'
          print 'Raw target latency vs. time-since-last-event:'
          Image(filename='r_images/async_targ_timesince_vs_latency.png')
```

2) Does last event recency affect latency of TARGET movements? ANS: No.
Raw target latency vs. time-since-last-event:

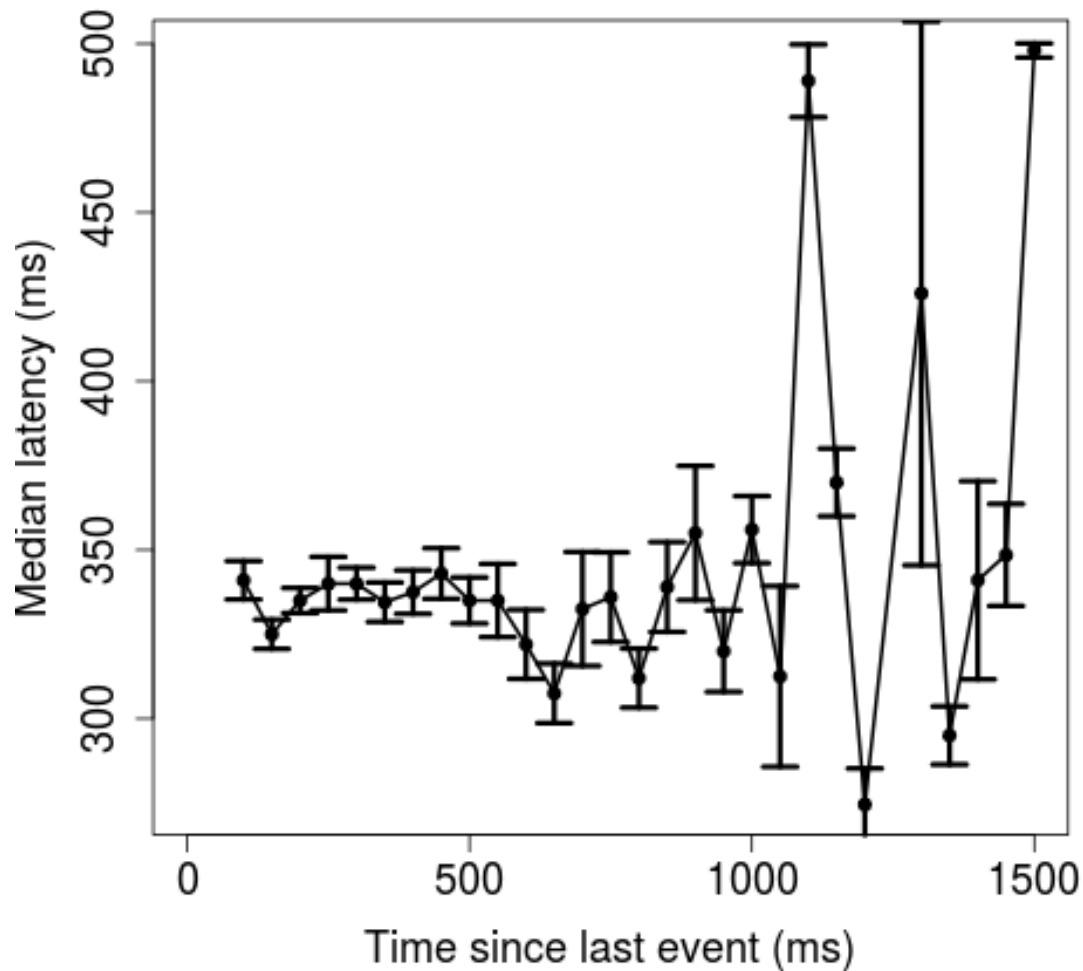
Out[35]:



```
In [36]: print 'Binned median latencies'  
         Image(filename='./r_images/async_targ_binned_latencies.png')
```

Binned median latencies

Out[36]:

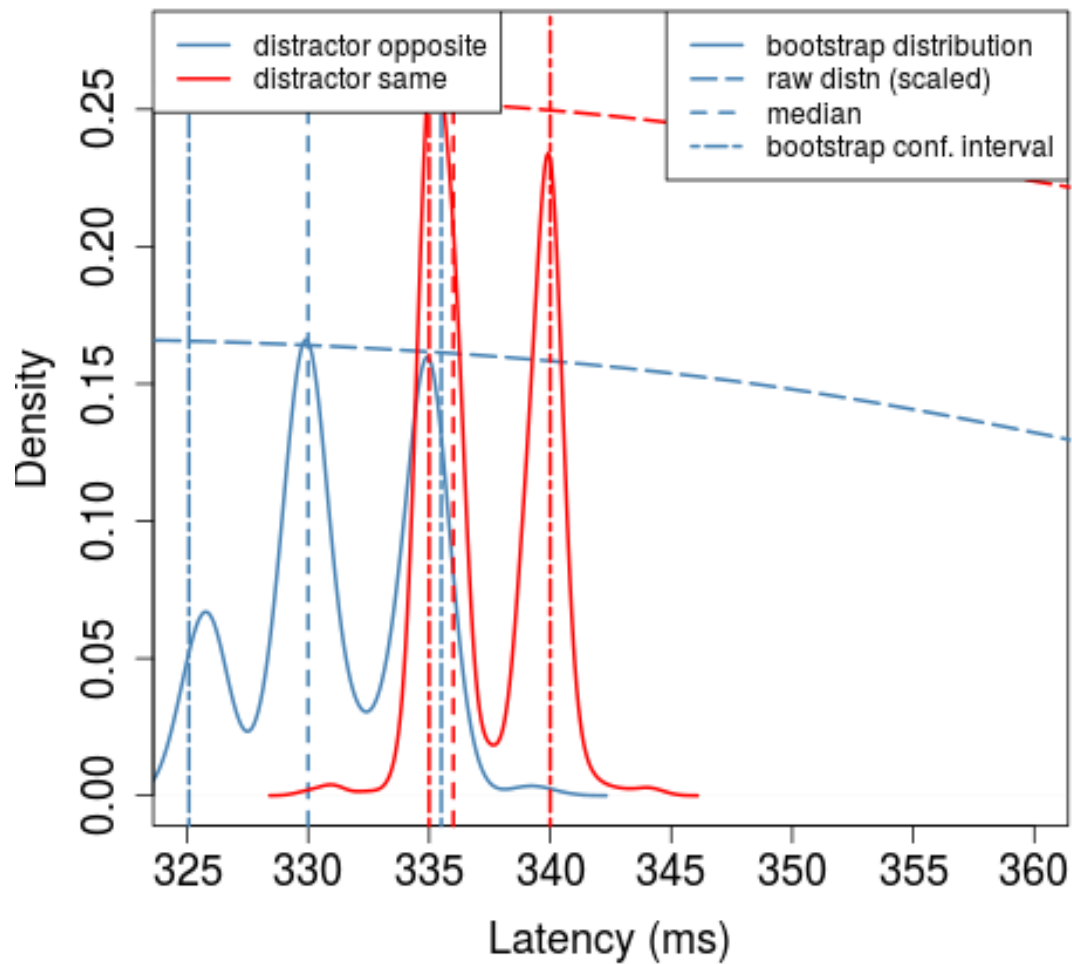


```
In [37]: print '3) Does distractor in same direction as target reduce latency? ANS: No (not significant)'
print 'Here "distractor opposite" means:\n"the existing distractor location was in the opposite direction compared with the new target location"'
print "Can say that: There's 0.7921 probability that lat_dist_same > lat_dist_opp; 0.1958 "
print "probability that lat_dist_same is not determinedly > or < lat_dist_opp. Finally, "
print "there's 0.0121 probability that lat_dist_same < lat_dist_opp."
Image(filename='./r_images/async_targ_vs_oppositeness_of_distractor.png')
```

3) Does distractor in same direction as target reduce latency? ANS: No (not significantly)
 Here "distractor opposite" means:
 "the existing distractor location was in the opposite direction compared with the new target location"
 Can say that: There's 0.7921 probability that lat_dist.same > lat_dist.opp; 0.1958
 probability that lat_dist.same is not determinedly > or < lat_dist.opp. Finally,
 there's 0.0121 probability that lat_dist.same < lat_dist.opp.

Out[37]:

Distsns of bootstrapped medians. alpha=0.050

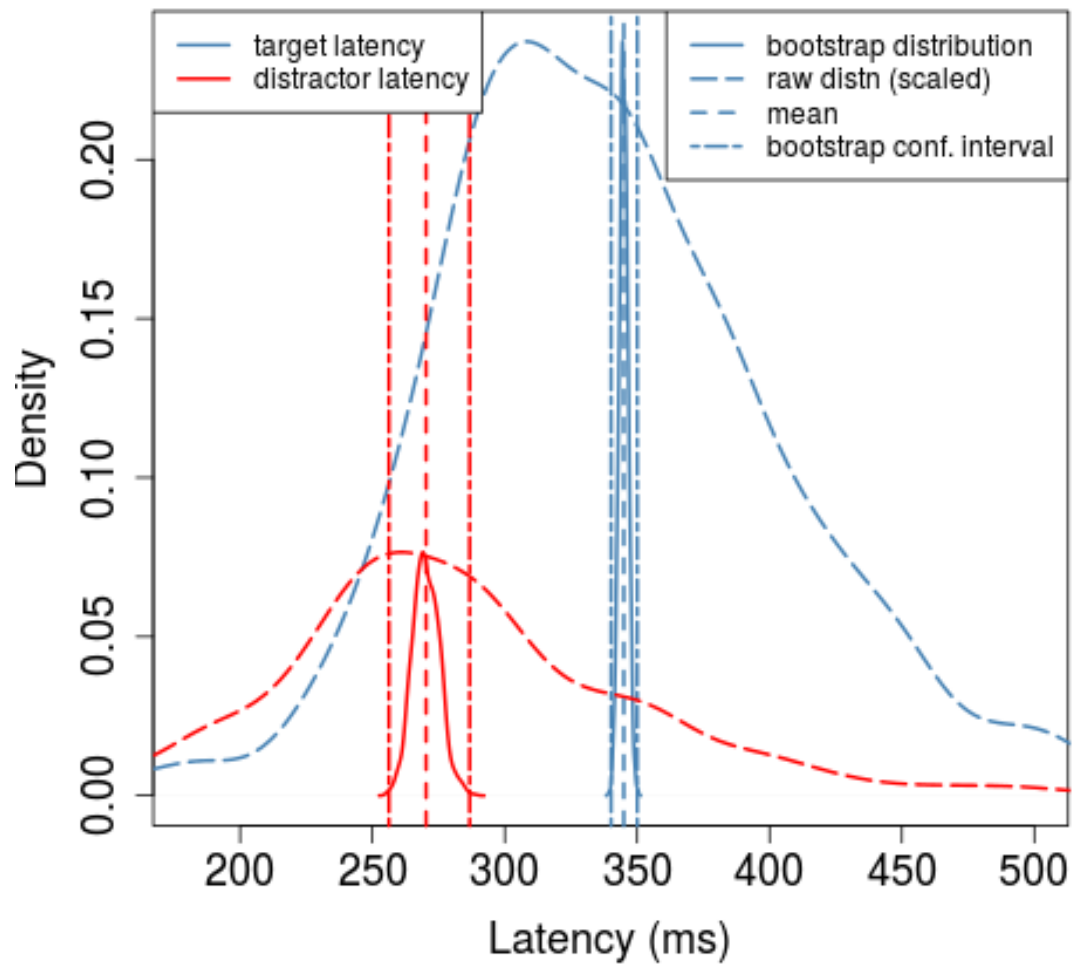


```
In [38]: print '4) What\'s the reaction time for distractors cf. targets *within the AD condition*?\nANS: Significantly different.'
          Image(filename='./r_images/async_targ_vs_dist.png')
```

4) What's the reaction time for distractors cf. targets *within the AD condition*?
ANS: Significantly different.

Out[38]:

Distsns of bootstrapped means. alpha=0.001

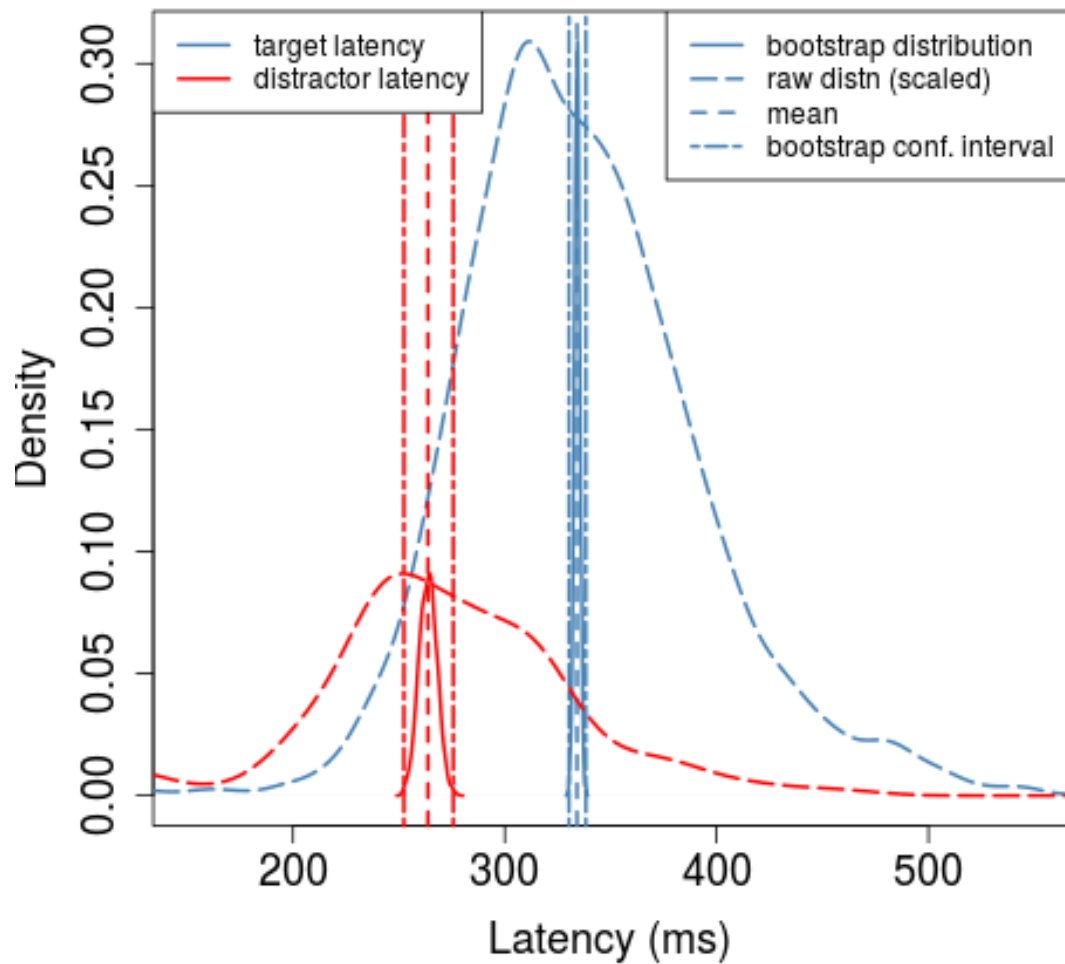


```
In [39]: print '5) What\'s the reaction time for distractors cf. targets *within the SD condition*?'
          Image(filename='./r_images/sync_targ_vs_dist.png')
```

5) What's the reaction time for distractors cf. targets *within the SD condition*?

Out[39]:

Distsns of bootstrapped means. alpha=0.001

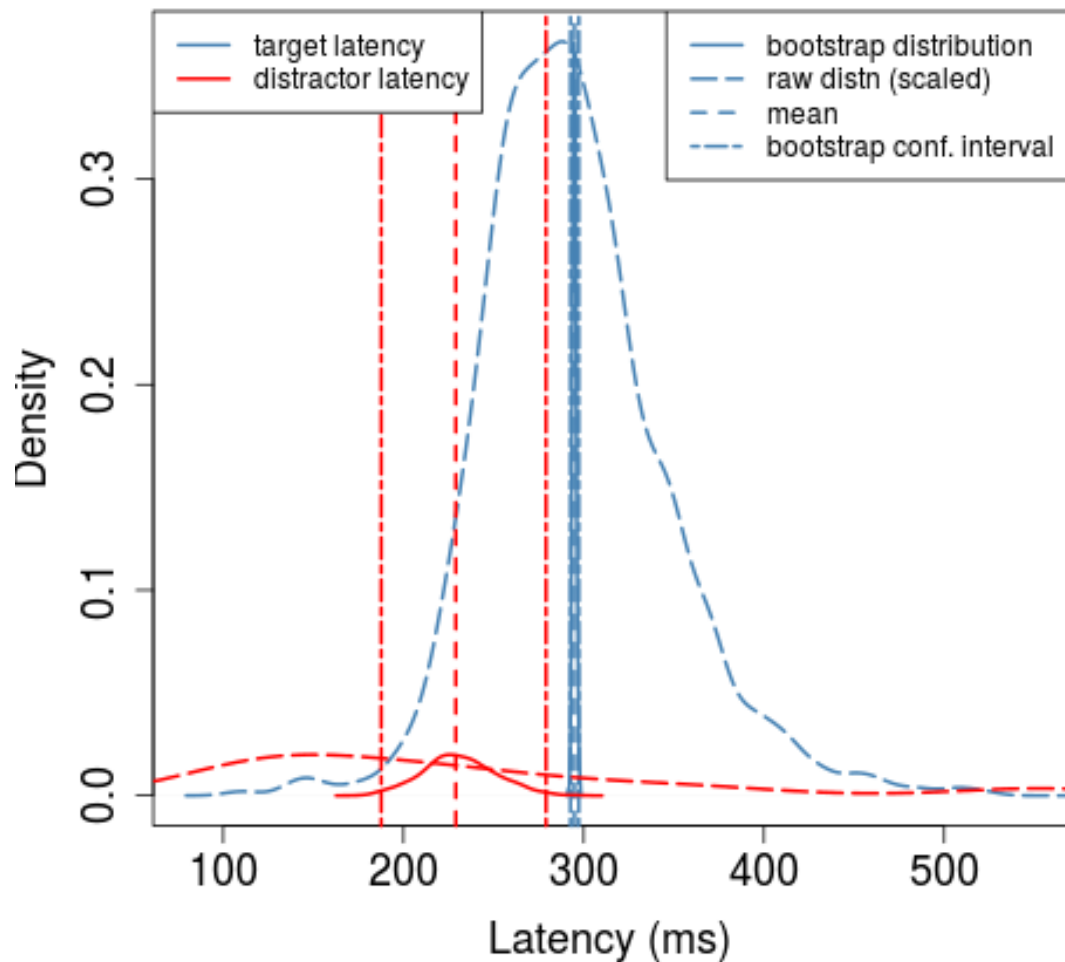


```
In [40]: print '6) What\'s the reaction time for error moves cf. correct, target movements *within the ND condition*?'
          Image(filename='./r_images/nodist_targ_vs_dist.png')
```

6) What's the reaction time for error moves cf. correct, target movements *within the ND condition*?

Out[40]:

Distsns of bootstrapped means. alpha=0.010



1.14 Analysis of latencies wrt target distance

This code (Bootstrap_targdist.r) looks at how the latency to first movement varies with distance to the target.

```
In [41]: from subprocess import CalledProcessError, check_output
try:
    out = check_output(["R", "-q --file=Bootstrap_targdist.r"])
except CalledProcessError as err:
    print 'Error for:',err.cmd,'with error code:',err.returncode
    print "\nOutput:\n\n",err.output
else:
    for line in out.splitlines():
        print line
```

```

> #
> # Analysis of distance-to-target statistics.
> #
> # 1) Does the distance to the target have an effect on the latency?
> # This has been reported by Meegan & Tipper 99, Pratt and Abrams 94,
> # Tipper et al 92 and 97.
> #
>
> maxlatency <- 1000
>
> set.seed(197420168)
>
> # a function which will bootstrap the standard error of the mean
> bs.mean <- function(data, num) {
+   resamples <- lapply(1:num, function(i) sample(data, replace=T))
+   r.mean <- sapply(resamples, mean)
+   std.err <- sqrt(var(r.mean))
+   list(std.err=std.err, resamples=resamples, means=r.mean)
+ }
> bs.median <- function(data, num) {
+   resamples <- lapply(1:num, function(i) sample(data, replace=T))
+   r.median <- sapply(resamples, median)
+   std.err <- sqrt(var(r.median))
+   list(std.err=std.err, resamples=resamples, medians=r.median)
+ }
>
> # This takes a 2-D data set of latency vs time since last event. It
> # collects "bins" of data by dividing time since last event into 30
> # bins, then computes bootstrap mean & median values for each bin.
> binned <- function (data, fname) {
+   listPoints <- list()
+   listBStrap <- list()
+   dfBStrap <- data.frame()
+   nBreaks <- 30
+   iter <- 1
+   nResamples <- 200
+   h <- hist(abs(data$direction), breaks=nBreaks, plot=F)
+   bLast <- -1
+   for (b in h$breaks) {
+     if (b==0) {
+       bLast = b
+       next
+     }
+
+     points <- data[which(abs(data$direction) >= bLast & abs(data$direction) < b),]
+     listPoints[[iter]] <- points
+
+     # Now bootstrap each member of listPoints, compute mean & std err of mean
+     bsmed <- bs.median(points$latency, nResamples)
+     bsmean <- bs.mean(points$latency, nResamples)
+
+     if (nrow(points)) {
+       dfBStrap <- rbind (dfBStrap, c((b-bLast/2), median(points$latency),
+                                     bsmed$std.err, mean(points$latency),

```

```

+             bsmean$std.err))
+         }
+
+         iter <- iter + 1
+         bLast = b
+     }
+     names(dfBStrap) <- c("distance","median","med.stderr","mean","mean.stderr")
+     return (dfBStrap)
+ }
+
+ >
+ > movingbinwidth <- 44 # 44 corresponds to about 5 mm of screen; 1/7 foveal width
+ >
+ > movingbin <- function (data, fname) {
+
+     # First sort data wrt direction
+     data$distance <- abs(data$direction)
+     data <- data[order(data$distance),]
+
+     listPoints <- list()
+     listBStrap <- list()
+     dfBStrap <- data.frame()
+     iter <- 1
+     nResamples <- 200
+     maxdist <- max(data$distance)
+     bLast <- -1
+     for (b in unique(data$distance)) {
+
+         if (b < movingbinwidth/2) {
+             next
+         }
+
+         if (maxdist - b < movingbinwidth/2) {
+             next
+         }
+
+         points <- data[which(data$distance >= b-movingbinwidth/2 & data$distance < b+movingbinwidth/2),]
+
+         listPoints[[iter]] <- points
+
+         # Now bootstrap each member of listPoints, compute mean & std err of mean
+         bsmed <- bs.median(points$latency, length(points$latency))
+         bsmean <- bs.mean(points$latency, length(points$latency))
+
+         if (nrow(points)) {
+             # 8.80734 mm per pixel, so this will make distance in mm.
+             dfBStrap <- rbind (dfBStrap, c(b/8.80734, median(points$latency),
+                                     bsmed$std.err, mean(points$latency),
+                                     bsmean$std.err))
+         }
+
+         iter <- iter + 1
+         bLast = b
+     }
+     names(dfBStrap) <- c("distance","median","med.stderr","mean","mean.stderr")

```

```

+
+   return (dfBStrap)
+ }
>
> movingbindir <- function (data, fname) {
+
+   # First sort data wrt direction
+   data <- data[order(data$direction),]
+
+   listPoints <- list()
+   listBStrap <- list()
+   dfBStrap <- data.frame()
+   iter <- 1
+   nResamples <- 200
+   maxdist <- max(data$direction)
+   mindist <- min(data$direction)
+   bLast <- -1
+   for (b in unique(data$direction)) {
+
+     if (b < mindist + movingbinwidth/2) {
+       next
+     }
+
+     if (maxdist - b < movingbinwidth/2) {
+       next
+     }
+
+     points <- data[which(data$direction >= b-movingbinwidth/2 & data$direction < b+movingbinwidth,)]
+
+     listPoints[[iter]] <- points
+
+     # Now bootstrap each member of listPoints, compute mean & std err of mean
+     bsmed <- bs.median(points$latency, length(points$latency))
+     bsmean <- bs.mean(points$latency, length(points$latency))
+
+     if (nrow(points)) {
+       # 8.80734 mm per pixel, so this will make distance in mm.
+       dfBStrap <- rbind (dfBStrap, c(b/8.80734, median(points$latency),
+                                     bsmed$std.err, mean(points$latency),
+                                     bsmean$std.err))
+     }
+
+     iter <- iter + 1
+     bLast = b
+   }
+   names(dfBStrap) <- c("distance", "median", "med.stderr", "mean", "mean.stderr")
+
+   return (dfBStrap)
+ }
>
>
> #
> # 1) Does event destination magnitude affect latency of target movements?
> #

```

```

>
> # This selects out target latencies and allows me to plot the
> # latencies vs. distance to target (magnitude of direction)
> d <- read.csv('AsyncTrials.csv')
> alt <- d[which (d$type == 1 & d$latency > 0 & d$correctmove == 1 & d$omit == 0 ),c("latency","direction")]
> png (filename='./r_images/async_latency_vs_dist.png')
> plot (abs(alt$direction),alt$latency,xlab="Distance to target",ylab="Latency to motion (ms)")
> dev.off()
null device
      1
>
> d <- read.csv('SyncTrials.csv')
> slt <- d[which (d$type == 1 & d$latency > 0 & d$correctmove == 1 & d$omit == 0 ),c("latency","direction")]
> png (filename='./r_images/sync_latency_vs_dist.png')
> plot (abs(slt$direction),slt$latency,xlab="Distance to target",ylab="Latency to motion (ms)")
> dev.off()
null device
      1
>
> d <- read.csv('NoDistTrials.csv')
> nlt <- d[which (d$type == 1 & d$latency > 0 & d$correctmove == 1 & d$omit == 0 ),c("latency","direction")]
> png (filename='./r_images/nodist_latency_vs_dist.png')
> plot (abs(nlt$direction),nlt$latency,xlab="Distance to target",ylab="Latency to motion (ms)")
> dev.off()
null device
      1
>
> # Compute moving bootstraps
> amb <- movingbin (alt, './r_images/async_targ_binned_latvsdist.png')
> smb <- movingbin (slt, './r_images/sync_targ_binned_latvsdist.png')
> nmb <- movingbin (nlt, './r_images/nodist_targ_binned_latvsdist.png')
>
> # A common plotting function
> plotfn <- function (nmb, smb, amb, xlimits) {
+   plot (amb$distance,amb$mean, xlim=xlimits, ylim=c(270,405),
+         pch=19, cex=0.2, cex.lab=1.5, cex.axis=1.5, cex.main=1.5, cex.sub=1.5,
+         xlab="Distance to target (mm)",ylab="Mean latency (ms)", col="white")
+
+   lines(amb$distance,amb$mean, lty=1, lwd=3, col="red")
+   lines(amb$distance,amb$mean-1.96*amb$mean.stderr, lty=1, lwd=1, col="red")
+   lines(amb$distance,amb$mean+1.96*amb$mean.stderr, lty=1, lwd=1, col="red")
+
+   lines(smb$distance,smb$mean, lty=1, lwd=3, col="blue")
+   lines(smb$distance,smb$mean-1.96*smb$mean.stderr, lty=1, lwd=1, col="blue")
+   lines(smb$distance,smb$mean+1.96*smb$mean.stderr, lty=1, lwd=1, col="blue")
+
+   lines(nmb$distance,nmb$mean, lty=1, lwd=3, col="black")
+   lines(nmb$distance,nmb$mean-1.96*nmb$mean.stderr, lty=1, lwd=1, col="black")
+   lines(nmb$distance,nmb$mean+1.96*nmb$mean.stderr, lty=1, lwd=1, col="black")
+
+   lines(c(11.9,11.9),c(270,400),lty=2, lwd=2, col='steelblue')
+
+   text (7,280, sprintf('Bin width: %.1f mm', movingbinwidth/8.807))
+
+

```

```

+     legend (20, 408, bg="white",
+             c("No distractor","Synchronous distractor","Asynchronous distractor","Approx. foveal exten
+             lty=c(1,1,1,2),
+             lwd=c(3,3,3,2),
+             cex=1.0,
+             col=c('black','blue','red','steelblue'))
+ }
>
> # Do the png plot
> png (filename='./r_images/movingbin_latvsdist.png')
> plotfn(nmb,smb,amb,c(0,55))
> dev.off()
null device
      1
>
> # And the eps plot
> setEPS()
> postscript(file='./paper/figures/movingbin_latvsdist.eps',width=8,height=7)
> plotfn(nmb,smb,amb,c(0,55))
> dev.off()
null device
      1
>
> ambdir <- movingbindir (alt, './r_images/async_targ_binned_latvsdir.png')
> smbdir <- movingbindir (slt, './r_images/sync_targ_binned_latvsdir.png')
> nmbdir <- movingbindir (nlt, './r_images/nodist_targ_binned_latvsdir.png')
> png (filename='./r_images/movingbin_latvsdir.png')
> plotfn(nmbdir,smbdir,ambdir,c(-55,55))
> dev.off()
null device
      1
>

```

```

In [42]: print 'Moving-bin mean & 95% confidence intervals wrt distance to target'
         Image(filename='r_images/movingbin_latvsdist.png')

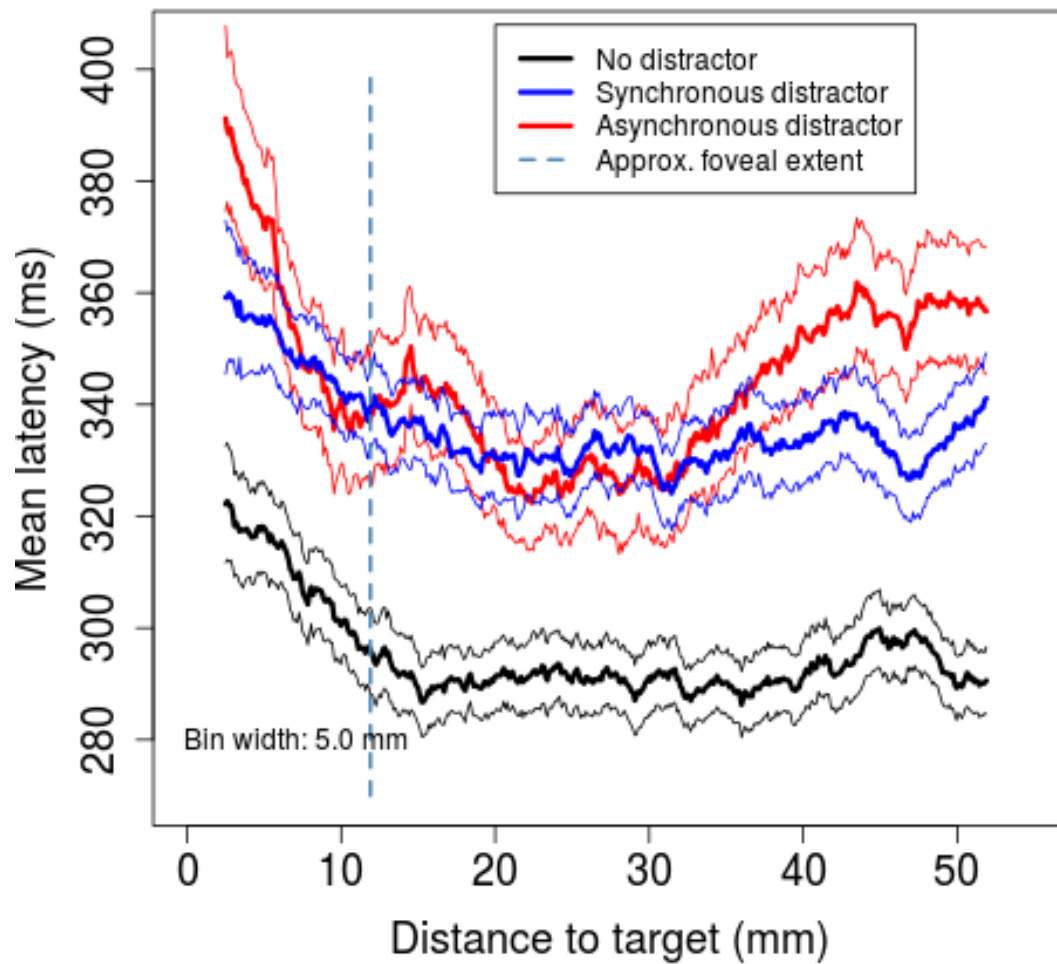
```

Moving-bin mean & 95% confidence intervals wrt distance to target

```

Out[42]:

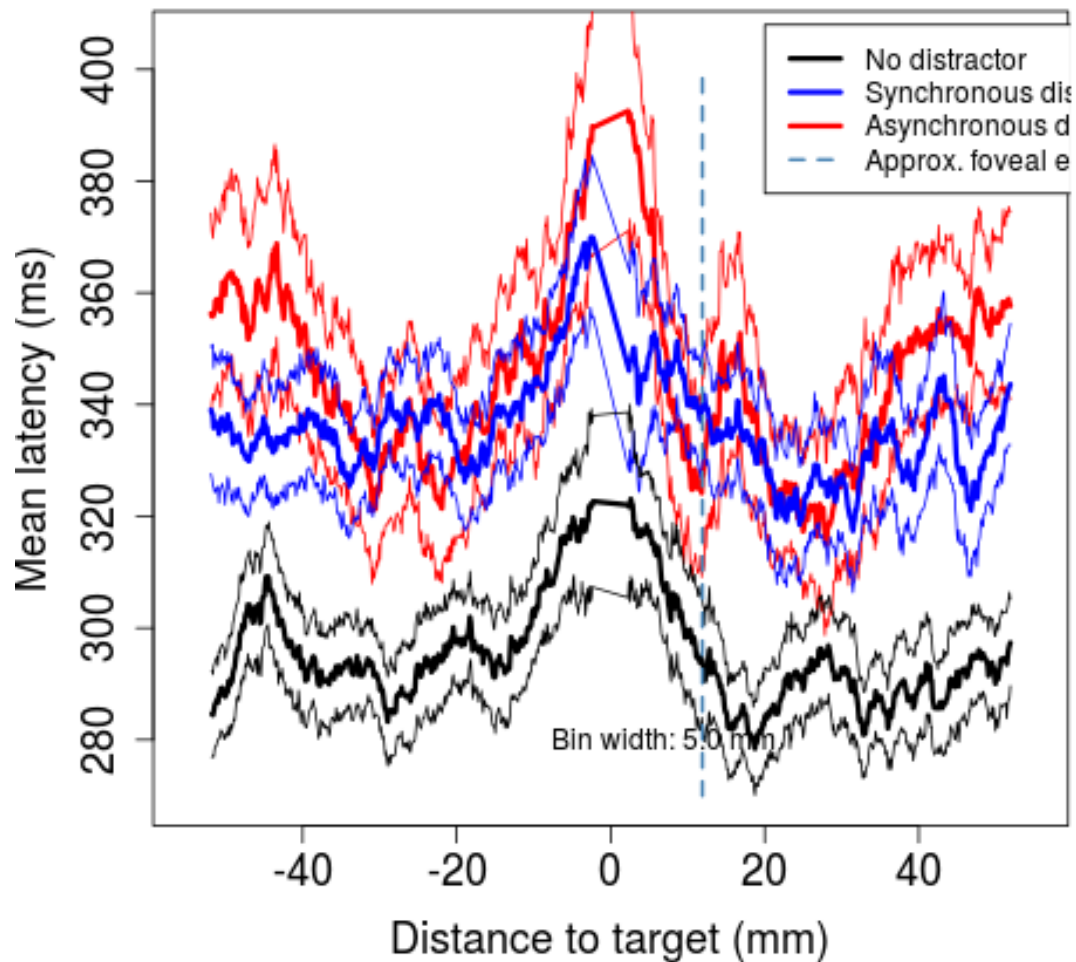
```



```
In [43]: print 'Directional version of the moving bin graph, above'
         Image(filename='r_images/movingbin_latvsdir.png')
```

Directional version of the moving bin graph, above

Out[43]:



2 Alternative movement analysis

Mauro da Lio developed an alternative approach to finding movement latencies and errors from the raw position data; an alternative to the Octave script which creates `fnames.mat`. Here are scripts to load in the `Summary.csv` that is provided by his analysis, and also a code block to re-create an equivalent `Summary.csv` file from the Octave-generated data.

2.1 Reading data

To read data in for the alternative movement analysis, replace calls to `readIndividuals()` in the code blocks above with this alternative function `readMauroIndividuals()`.

(I convert `Summary.xls` into `Summary.csv` before running this code.)

```
In [44]: # This function reads in data from Mauro's Summary.csv, producing the same list of individuals
        # as is generated by readIndividuals()
```



```

import csv
def readMauroIndividuals ():
    with open('Mauro_analysis/Summary.csv', 'rb') as csvfile:

        individuals = dict()
        idnum_counter = 1
        first = 1
        dreader = csv.reader(csvfile, delimiter=',', quotechar='"')

        for row in dreader:

            # Skip first row (the header)
            if first:
                first = 0
                continue

            # Can access rows by column number:
            #print row[3]
            #print row
            #print len(row)
            #break

            subj_id = row[1]
            exp_name = row[0]

            # condition index is for the no distractor/sync distractor/async distractor
            condition_index = int(row[2])

            # Need ONE individual object for each subj_id.
            if subj_id not in individuals:
                individuals[subj_id] = individual(subj_id)
                individuals[subj_id].idnum = idnum_counter
                individuals[subj_id].numSubsamples = 8
                idnum_counter += 1

            # No filename in Mauro's csv, just put name in each:
            individuals[subj_id].filename_nd = exp_name
            individuals[subj_id].filename_ad = exp_name
            individuals[subj_id].filename_sd = exp_name

            #individuals[subj_id].alldata_nd=np.ndarray(5)

            # latencies for non-movement-error target events
            ##tnoerr_latencies = fname[4] # R.latency_noerror_target - Use table with python i
            # latencies for non-movement-error distractor events
            ##dnoerr_latencies = fname[5]
            # err_latencies = fname[6] # ALL error events, distractor and target.
            # latencies for target movement error events
            ##terr_latencies = fname[7] # R.latency_error_target
            # latencies for distractor movement error events
            ##derr_latencies = fname[8] # R.latency_error_distractor
            if condition_index == 0:
                i=10 # starting col for the latencies in Mauro's spreadsheet
                while i < len(row):

```

```

        if len(row[i]) == 0:
            break
        rowval = float(row[i])
        # New latency:
        newlat = np.array([i+1,rowval])
        if (rowval >= 0):
            # number, type (TARG=1 here), error, correct_move, latency
            newalldata = np.array([i+1, 1, 0, 1, rowval])
            if individuals[subj_id].nodist_latencies.size == 0:
                individuals[subj_id].nodist_latencies = newlat
            else:
                individuals[subj_id].nodist_latencies = np.vstack((individuals[subj_id].nodist_latencies, newlat))
        else:
            newalldata = np.array([i+1, 1, 1, 0, rowval])
            if individuals[subj_id].nodist_err_latencies.size == 0:
                individuals[subj_id].nodist_err_latencies = -1*newlat
            else:
                individuals[subj_id].nodist_err_latencies = np.vstack((individuals[subj_id].nodist_err_latencies, -1*newlat))

        if individuals[subj_id].alldata_nd.size == 0:
            individuals[subj_id].alldata_nd = newalldata
        else:
            individuals[subj_id].alldata_nd = np.vstack((individuals[subj_id].alldata_nd, newalldata))

        #print 'alldata', individuals[subj_id].alldata_nd
        i+=1
    # Compute ranks
    #print 'nodist latencies:', individuals[subj_id].nodist_latencies
    #print 'nodist_err latencies:', individuals[subj_id].nodist_err_latencies
    if individuals[subj_id].nodist_latencies.size>0:
        individuals[subj_id].nodist_latencies_rank = stats.rankdata(individuals[subj_id].nodist_latencies)
    if individuals[subj_id].nodist_err_latencies.size>0:
        if individuals[subj_id].nodist_err_latencies.size==2:
            individuals[subj_id].nodist_err_latencies_rank = [1]
        else:
            individuals[subj_id].nodist_err_latencies_rank = stats.rankdata(individuals[subj_id].nodist_err_latencies)

    # nerrs code identical to that in readIndividuals
    nerrs = 0
    ntargets = 0
    for d in individuals[subj_id].alldata_nd:
        ntargets += 1 # "ndistractors" is really "ntargets" for the ND condition.
        if d[2] > 0.0:
            nerrs += 1
    individuals[subj_id].n_errors_per_target_nd = (float(nerrs) / float(ntargets))
    #print individuals[subj_id].n_errors_per_target_nd

if condition_index == 1:
    i=10 # starting col for the latencies in Mauro's spreadsheet
    while i < len(row):
        if len(row[i]) == 0:
            break
        rowval = float(row[i])
        # New latency:

```

```

newlat = np.array([i+1,rowval])
if (rowval >= 0):
    # number, type (TARG=1 here), error, correct_move, latency
    newalldata = np.array([i+1, 1, 0, 1, rowval])
    if individuals[subj_id].sync_latencies.size == 0:
        individuals[subj_id].sync_latencies = newlat
    else:
        individuals[subj_id].sync_latencies = np.vstack((individuals[subj_id].sync_latencies, newlat))
else:
    newalldata = np.array([i+1, 1, 1, 0, rowval])
    if individuals[subj_id].sync_err_latencies.size == 0:
        individuals[subj_id].sync_err_latencies = -1*newlat
    else:
        individuals[subj_id].sync_err_latencies = np.vstack((individuals[subj_id].sync_err_latencies, -1*newlat))

if individuals[subj_id].alldata_sd.size == 0:
    individuals[subj_id].alldata_sd = newalldata
else:
    individuals[subj_id].alldata_sd = np.vstack((individuals[subj_id].alldata_sd, newalldata))

# print 'alldata', individuals[subj_id].alldata_sd
i+=1
# print 'sync latencies:', individuals[subj_id].sync_latencies
# print 'sync_err latencies:', individuals[subj_id].sync_err_latencies
# Compute ranks
if individuals[subj_id].sync_latencies.size>0:
    individuals[subj_id].sync_latencies_rank = stats.rankdata(individuals[subj_id].sync_latencies)
if individuals[subj_id].sync_err_latencies.size>0:
    if individuals[subj_id].sync_err_latencies.size==2:
        individuals[subj_id].sync_err_latencies_rank = [1]
    else:
        individuals[subj_id].sync_err_latencies_rank = stats.rankdata(individuals[subj_id].sync_err_latencies)

# nerrs code identical to that in readIndividuals
nerrs = 0
ntargets = 0
for d in individuals[subj_id].alldata_sd:
    ntargets += 1 # "ndistractors" is really "ntargets" for the ND condition.
    if d[2] > 0.0:
        nerrs += 1
individuals[subj_id].n_errors_per_distractor_sync = (float(nerrs) / float(ntargets))
# print individuals[subj_id].n_errors_per_distractor_sync

if condition_index == 2:
    # FIXME: FINISH ME
    i=10 # starting col for the latencies in Mauro's spreadsheet
    while i < len(row):
        if len(row[i]) == 0:
            break
        rowval = float(row[i])
        # New latency:
        newlat = np.array([i+1,rowval])
        if (rowval >= 0):
            # number, type (TARG=1 here), error, correct_move, latency

```

```

        newalldata = np.array([i+1, 1, 0, 1, rowval])
        if individuals[subj_id].async_latencies.size == 0:
            individuals[subj_id].async_latencies = newlat
        else:
            individuals[subj_id].async_latencies = np.vstack((individuals[subj_id].async_latencies, newlat))
    else:
        newalldata = np.array([i+1, 1, 1, 0, rowval])
        if individuals[subj_id].async_err_latencies.size == 0:
            individuals[subj_id].async_err_latencies = -1*newlat
        else:
            individuals[subj_id].async_err_latencies = np.vstack((individuals[subj_id].async_err_latencies, -1*newlat))

    if individuals[subj_id].alldata_ad.size == 0:
        individuals[subj_id].alldata_ad = newalldata
    else:
        individuals[subj_id].alldata_ad = np.vstack((individuals[subj_id].alldata_ad, newalldata))

    #print 'alldata', individuals[subj_id].alldata_ad
    i+=1
    #print 'async latencies:', individuals[subj_id].async_latencies
    #print 'async_err latencies:', individuals[subj_id].async_err_latencies
    # Compute ranks
    if individuals[subj_id].async_latencies.size>0:
        individuals[subj_id].async_latencies_rank = stats.rankdata(individuals[subj_id].async_latencies)
    if individuals[subj_id].async_err_latencies.size>0:
        if individuals[subj_id].async_err_latencies.size==2:
            individuals[subj_id].async_err_latencies_rank = [1]
        else:
            individuals[subj_id].async_err_latencies_rank = stats.rankdata(individuals[subj_id].async_err_latencies)

    # nerrs code identical to that in readIndividuals
    nerrs = 0
    ntargets = 0
    for d in individuals[subj_id].alldata_ad:
        ntargets += 1 # "ndistractors" is really "ntargets" for the ND condition.
        if d[2] > 0.0:
            nerrs += 1
    individuals[subj_id].n_errors_per_distractor_async = (float(nerrs) / float(ntargets))
    #print individuals[subj_id].n_errors_per_distractor_async

    return individuals

```

2.2 Reproduce Summary Spreadsheet for comparison with Mauro's analysis

Mauro has Summary.xls, which lists the following columns:

Experimenter Subject Distractor type RT (M) RT (SD) N correct RT incorrect (M) RT incorrect (SD)
 N incorrect Error rate Delays (negative = incorrect)

The following code creates an equivalent Summary.csv from the data in frames.dat.

```

In [45]: from __future__ import division
import numpy as np
from statsmodels.stats.multicomp import MultiComparison

maxCount = 100 # 1 for debug or 100 for all

```

```

count = 1
csvdata = ''

individuals = readIndividuals()

for i in individuals:
    ind = individuals[i]

    if count == 1:
        csvdata = ind.csvheader()

    # Remove outliers in standard way
    ind.excludeOutliers()

    csvdata += ind.csvlineset()

    # For debugging, break after maxCount
    if count >= maxCount:
        break
    count += 1

print 'Writing Summary.csv...'
f = open('Summary.csv', 'w')
f.write(csvdata)
f.close()

```

Writing Summary.csv...