Brightness perception example

December 26, 2023

```
[1]: | ## Perception example of EZBHDDM
```

0.1 Perception example of EZBHDDM

Data from Ratcliff and Rouder (1998), experiment 1, participant N.H.:

subjects were asked to decide whether the overall brightness of pixel arrays displayed on a computer monitor was "high" or "low" (Fig. 3a). The brightness of a display was controlled by the proportion of the pixels that were white. For each trial, the proportion of white pixels was chosen from one of two distributions, a high distribution or a low distribution, each with fixed mean and standard deviation (Fig. 3b). Feedback was given after each trial to tell the subject whether his or her decision had correctly indicated the distribution

There are 66 cells in the design. 1-33 had an accuracy instruction, 34-66 a speed instruction. 1-16 and 34-49 had more black pixels and 18-33 and 51-66 had more white. 17 and 50 were ambiguous and will not be used here.

```
[2]: import pandas as pd
import numpy as np
import pyjags
import matplotlib.pyplot as plt
import seaborn as sns
```

	index	cond	response	RT
0	1	1	1	459
1	1	1	1	457
2	1	1	1	569
3	1	1	1	390
4	1	1	1	534
			•••	
7885	1	66	2	261
7886	1	66	2	333
7887	1	66	1	261

```
7888
                   66
                              2 351
              1
    7889
                              2 361
              1
                   66
    [7890 rows x 4 columns]
[4]: # Remove rows where the value in the 'RT' column is greater than 3000ms
     data = data[data['RT'] <= 3000]</pre>
     # Update the 'response' column based on conditions, so it reflects accuracy
     data['response'] = (
         ((data['cond'] > 0) & (data['cond'] < 17) & (data['response'] == 1)) |
         ((data['cond'] > 17) & (data['cond'] < 34) & (data['response'] == 2)) |
         ((data['cond'] > 33) & (data['cond'] < 50) & (data['response'] == 1)) |
         ((data['cond'] > 50) & (data['cond'] < 67) & (data['response'] == 2))
     ).astype(int)
[5]: # Initialize the output DataFrame
     df = pd.DataFrame(np.zeros((66, 6)),
                       columns=["nTrials",
                                "sum_accuracy",
                                "mean_rt_correct",
                                "variance_rt_correct",
                                "Xi",
                                "Xs"])
[6]: # Function to compute the summary statistics used by EZ
     def ez_summaries(subset):
         return [
             len(subset),
             subset['response'].sum(),
             subset['RT'].mean() / 1000,
             (subset['RT'] / 1000).var(),
             int(subset.iloc[0]['cond'] > 33)-0.5,
             ((subset.iloc[0]['cond'] - 1) % 33 - 16) / 5 # Arbitrary rescaling
         ]
[7]: # Populate the output DataFrame using the ez_summaries function
     for r in range(1, 67):
         df.iloc[r-1, :] = ez_summaries(data[data['cond'] == r])
     # Remove the two ambiguous conditions (17 and 50)
     df = df[df.iloc[:, -1] != 0]
     # PyJAGS format
     toJags = {
                 "nTrials": df.nTrials,
                 "meanRT": df.mean rt correct,
```

```
"varRT": df.variance_rt_correct,
    "correct": df.sum_accuracy,
    "Xi": df.Xi,
    "Xs": df.Xs,
}
```

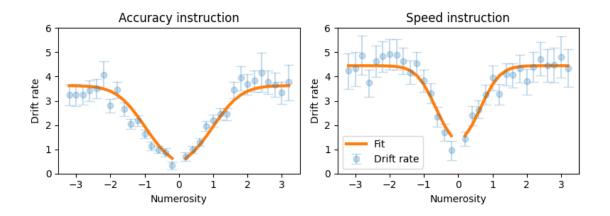
EZ JAGS code with an effect of instruction (Xi) on bound and a nonlinear regression using Xi and stimulus intensity Xs. Note the predictor is the absolute value of Xs to capture that the task gets easier as the black/white balance departs from 50%.

Beta3 and Beta4 capture effects of instruction on drift rate and so are of specific interest.

```
[8]: code = """
     # JAGS implementation of the EZ Bayesian
     # hierarchical drift diffusion model
     model {
       # Priors for hierarchical drift diffusion model
       # parameters
       betaweight ~ dnorm(0.00, 1.00)
       beta0 ~ dnorm(0.00, 1.00)
       beta1 ~ dnorm(0.00, 1.00)
       beta2 ~ dnorm(0.00, 1.00)
       beta3 ~ dnorm(0.00, 1.00)
       beta4 ~ dnorm(0.00, 1.00)
       bound_mean ~ dnorm(1.50, (0.20^-2))T( 0.10, 3.00)
       drift mean ~ dnorm(0.50, (0.50^-2))
       nondt_mean ~ dnorm(0.30, (0.06^-2))T( 0, )
       bound_sdev ~ dunif(0.01, 1.00)
       drift_sdev ~ dunif(0.01, 3.00)
       nondt_sdev ~ dunif(0.01, 0.50)
       for (p in 1:length(meanRT)) {
         # Hierarchical distributions of person-specific
         # diffusion model parameters. Here, drift rate
         # is the criterion.
         drift_pred[p] = beta0*phi(beta1 + beta2*abs(Xs[p]) + beta3*Xi[p]*abs(Xs[p]))
                       + beta4 * Xi[p] + drift_mean
         drift[p] ~ dnorm(drift_pred[p], (drift_sdev^-2))
         bound[p] ~ dnorm(bound_mean + betaweight * Xi[p],
                         (bound_sdev^-2))T( 0.10, 3.00)
         nondt[p] ~ dnorm(nondt mean, (nondt sdev^-2))
                                         T(0.05,)
         # Forward equations from EZ Diffusion
         ey[p] = exp(-bound[p] * drift[p])
         Pc[p] = 1 / (1 + ey[p])
         PRT[p] = 2 * pow(drift[p], 3) / bound[p] *
                  pow(ey[p] + 1, 2) / (2 * -bound[p] *
```

```
drift[p] * ey[p] - ey[p] * ey[p] + 1)
         MDT[p] = (bound[p] / (2 * drift[p])) *
                   (1 - ey[p]) / (1 + ey[p])
         MRT[p] = MDT[p] + nondt[p]
         # Sampling distributions for summary statistics
         correct[p] ~ dbin(Pc[p], nTrials[p])
         varRT[p] ~ dnorm(1/PRT[p], 0.5*(correct[p]-1)
                                       * PRT[p] * PRT[p])
         meanRT[p] ~ dnorm(MRT[p], PRT[p] * correct[p])
       }
      }
      0.00
 [9]: # Initial values for JAGS
      init = { "drift" : np.random.normal(0, 0.1, len(toJags['nTrials'])) }
      # Call JAGS and draw samples
      model = pyjags.Model(
                  progress_bar = True,
                  code
                       = code,
                  data
                        = toJags,
                  init = init,
                  adapt = 1000,
                  chains = 4,
                  threads = 4)
      # Overkill number of samples, takes a few seconds
      samples = model.sample(4000, vars = ['beta3', 'beta4', 'drift', 'drift_pred'])
     adapting: iterations 4000 of 4000, elapsed 0:00:02, remaining 0:00:00
     sampling: iterations 9858 of 16000, elapsed 0:00:09, remaining 0:00:05
     sampling: iterations 16000 of 16000, elapsed 0:00:09, remaining 0:00:00
[10]: # Extract samples for plotting the nonlinear regression
      drift = samples['drift']
      means = np.mean(drift, axis=(1, 2))
      preds = np.mean(samples['drift_pred'], axis=(1, 2))
      lower_percentiles = np.percentile(drift, 2.5, axis=(1, 2))
      upper_percentiles = np.percentile(drift, 97.5, axis=(1, 2))
      results = np.array([means, lower_percentiles, upper_percentiles, preds]).T
      # X-values
      x_values = df.Xs[df.Xi>0]
```

```
# Calculate the errors for the error bars (distance from mean to CI limits)
acccy = results[:32]
speed = results[-32:]
errors_acccy = [acccy[:, 0] - acccy[:, 1], acccy[:, 2] - acccy[:, 0]]
errors_speed = [speed[:, 0] - speed[:, 1], speed[:, 2] - speed[:, 0]]
# Fit line needs a break
fit_line_acccy = np.concatenate([acccy[:16, 3], [None], acccy[16:, 3]])
fit line speed = np.concatenate([speed[:16, 3], [None], speed[16:, 3]])
x_values_fit_line = np.concatenate([x_values[:16], [None], x_values[16:]])
# Plotting
plt.figure(figsize=(8, 3))
# Plot for the accuracy instruction condition
plt.subplot(1, 2, 1)
plt.errorbar(x_values, acccy[:, 0], yerr=errors_acccy,
             fmt='o', capsize=5, label='Drift rate', alpha=.25)
plt.plot(x_values_fit_line, fit_line_acccy, label='Fit', linewidth=3)
plt.title('Accuracy instruction')
plt.xlabel('Numerosity')
plt.ylabel('Drift rate')
plt.ylim((0,6))
# Plot for the speed instruction condition
plt.subplot(1, 2, 2)
plt.errorbar(x_values, speed[:, 0], yerr=errors_speed,
             fmt='o', capsize=5, label='Drift rate', alpha=.25)
plt.plot(x_values_fit_line, fit_line_speed, label='Fit', linewidth=3)
plt.title('Speed instruction')
plt.xlabel('Numerosity')
plt.ylabel('Drift rate')
plt.ylim((0,6))
plt.legend()
plt.tight_layout()
plt.show()
```



```
beta3 = samples['beta3'].flatten()

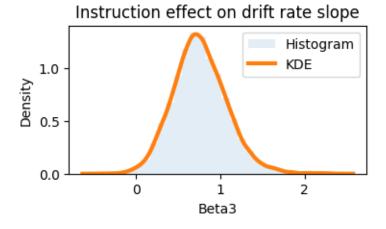
# Plotting
plt.figure(figsize=(4, 2))

# Histogram
plt.hist(beta3, bins=31, alpha=0.125, density=True, label='Histogram')

# Kernel density estimate
sns.kdeplot(beta3, linewidth=3, label='KDE')

plt.title('Instruction effect on drift rate slope')
plt.xlabel('Beta3')
plt.ylabel('Density')
plt.legend()

plt.show()
```



```
[12]: # Now focus on beta4
beta4 = samples['beta4'].flatten()

# Plotting
plt.figure(figsize=(4, 2))

# Histogram
plt.hist(beta4, bins=31, alpha=0.125, density=True, label='Histogram')

# Kernel density estimate
sns.kdeplot(beta4, linewidth=3, label='KDE')

plt.title('Instruction main effect on drift rate')
plt.xlabel('Beta4')
plt.ylabel('Density')
plt.legend()

plt.show()
```

