Semantic Networks Analysis

Abhilasha Kumar January 16, 2018

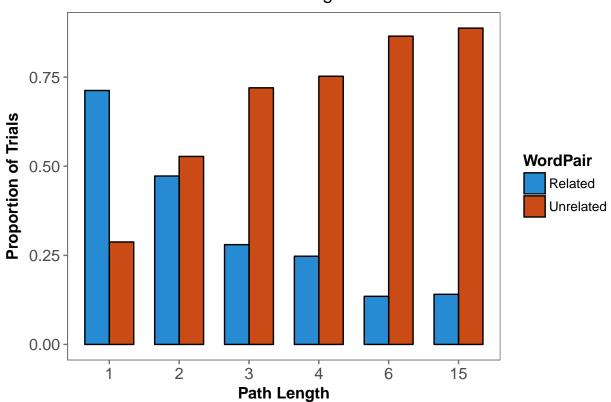
Reading the Data

Related-Unrelated Decisions

```
##
## Attaching package: 'dplyr'
## The following objects are masked from 'package:stats':
##
## filter, lag
## The following objects are masked from 'package:base':
##
## intersect, setdiff, setequal, union
```

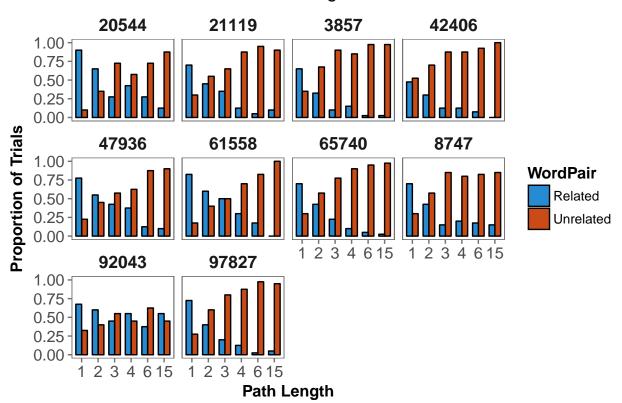
Plotting Proportions

Relatedness Judgments



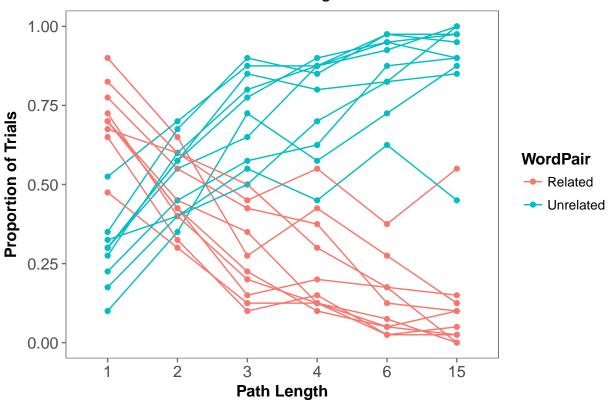
Subject-Wise

Relatedness Judgments



Line Plot Subject-Wise

Relatedness Judgments



ANOVA

```
relunrel_aov = aov(data = sem_decision, proportion ~ pathlengthfac*actual +
                     Error(subject/(pathlengthfac*actual)))
summary(relunrel_aov)
##
## Error: subject
                            Mean Sq F value Pr(>F)
##
            Df
                   Sum Sq
## Residuals 9 7.892e-31 8.769e-32
##
## Error: subject:pathlengthfac
                     Sum Sq Mean Sq F value Pr(>F)
## pathlengthfac 5 4.79e-31 9.577e-32
                                         1.122 0.362
                45 3.84e-30 8.533e-32
## Residuals
##
## Error: subject:actual
##
            Df Sum Sq Mean Sq F value
                                       Pr(>F)
## actual
             1 3.605
                        3.605
                                  24.3 0.000814 ***
## Residuals 9 1.336
                        0.148
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

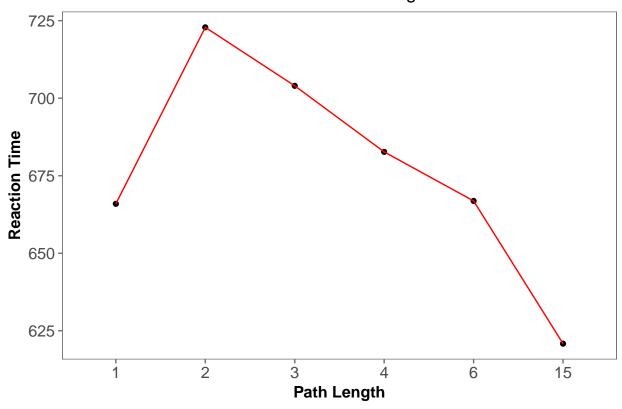
Raw Reaction Time

ANOVA

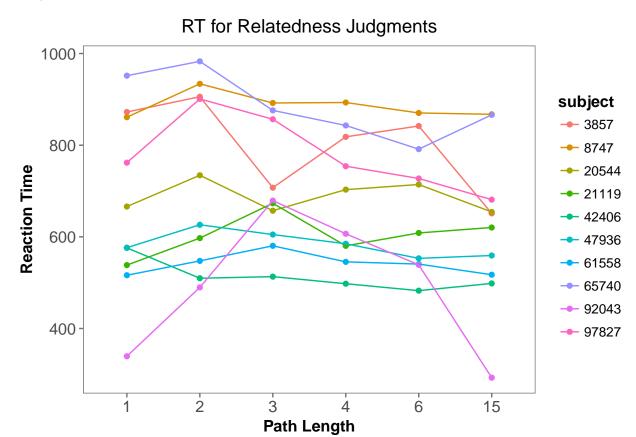
```
sem_rt$pathlengthfac = ordered(as.factor(as.character(sem_rt$pathlength)),
                           levels = c("1", "2", "3", "4", "6", "15"))
sem_rt$subject = as.factor(sem_rt$subject)
rt_aov = aov(data = sem_rt, rt ~ pathlengthfac +
                    Error(subject/(pathlengthfac)))
summary(rt_aov)
##
## Error: subject
            Df Sum Sq Mean Sq F value Pr(>F)
## Residuals 9 1241544 137949
## Error: subject:pathlengthfac
                Df Sum Sq Mean Sq F value Pr(>F)
## pathlengthfac 5 62403
                            12481
                                    2.95 0.0219 *
## Residuals
               45 190402
                             4231
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

Plotting RTs

RT for Relatedness Judgments



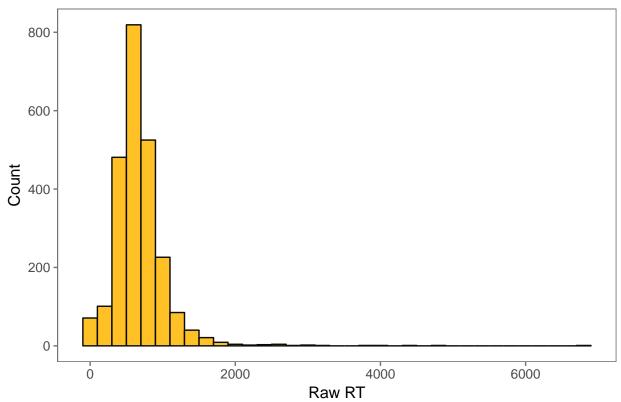
Subject-Wise



z-scored Reaction Time

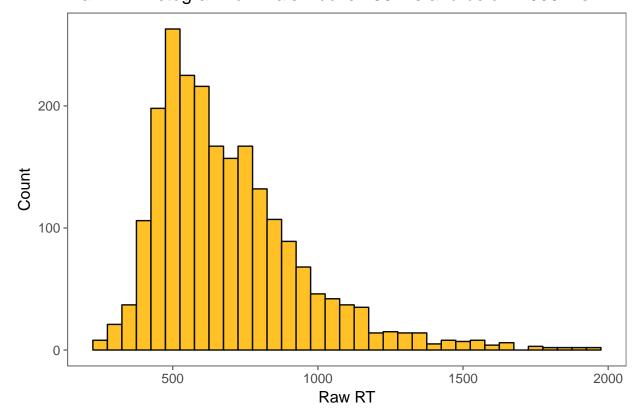
Histogram of RT

Raw RT Histogram for All Trials

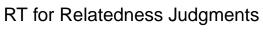


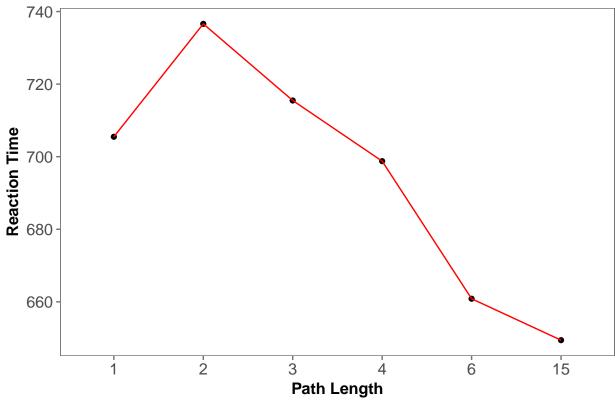
First Trim

Raw RT Histogram for Trials Above 250 ms and below 2000 ms

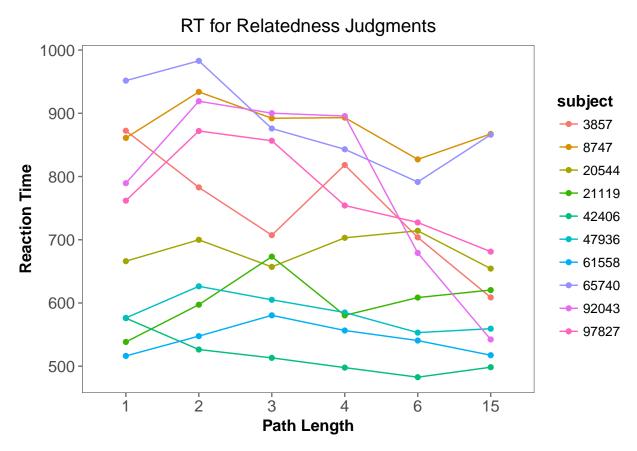


Raw RT aggregates After Trimming





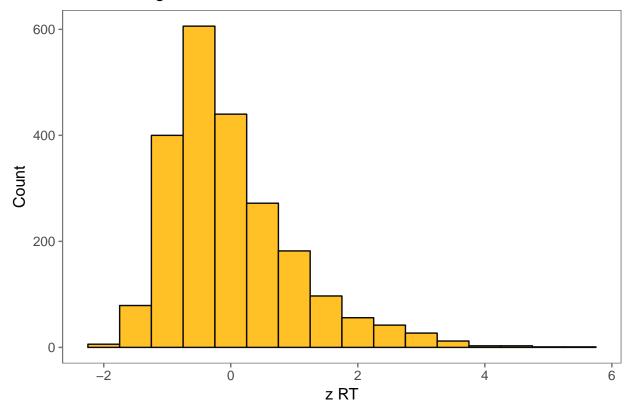
Subject Raw RT again



Making the z-scores

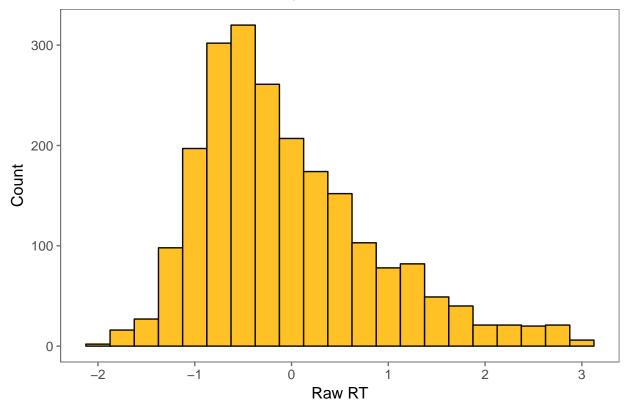
z-RT Distribution

z-RT Histogram for above 250 ms & <2s Trials



Trimming z-RT

Trimmed (3 SD) z-RT Histogram for above 250 ms & <2s Trials



Repeating z-scoring

Aggregating zRT

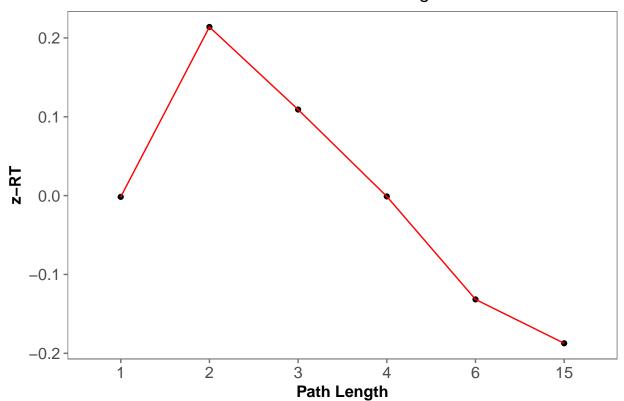
ANOVA

```
z_sem_rt$pathlengthfac = ordered(as.factor(as.character(z_sem_rt$pathlength)),
                            levels = c("1", "2", "3", "4", "6", "15"))
z_sem_rt$subject = as.factor(z_sem_rt$subject)
z_rt_aov = aov(data = z_sem_rt, zRT_trim ~ pathlengthfac +
                     Error(subject/(pathlengthfac)))
summary(z_rt_aov)
##
## Error: subject
             Df
                  Sum Sq
                           Mean Sq F value Pr(>F)
##
## Residuals 9 0.001498 0.0001665
##
## Error: subject:pathlengthfac
                 Df Sum Sq Mean Sq F value
                                             Pr(>F)
## pathlengthfac 5 1.266 0.25316
                                      6.28 0.000169 ***
## Residuals
                45 1.814 0.04031
```

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

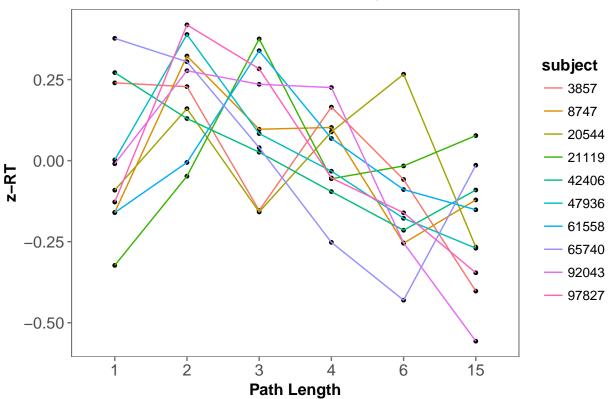
Plotting RTs

z-RT for Relatedness Judgments



Subject-Wise

z-RT for Relatedness Judgments



Regressions

```
## Loading required package: Matrix
## Linear mixed model fit by REML ['lmerMod']
## Formula: rt ~ 1 + (1 | subject) + (1 | trial_index)
     Data: sem
##
##
## REML criterion at convergence: 34877.8
##
## Scaled residuals:
      Min
                1Q Median
                                       Max
## -1.4331 -0.4477 -0.1596 0.2304 18.0486
## Random effects:
  Groups
                Name
                            Variance Std.Dev.
## trial_index (Intercept)
                               707.2 26.59
## subject
                (Intercept)
                            22500.5 150.00
## Residual
                            117846.2 343.29
## Number of obs: 2400, groups: trial_index, 240; subject, 10
## Fixed effects:
##
              Estimate Std. Error t value
```

```
## (Intercept)
                 677.20
                             47.98
                                     14.11
## [1] 0.1645309
## Linear mixed model fit by REML ['lmerMod']
## Formula: rt ~ pathlengthfac + (1 | subject) + (1 | trial_index)
##
     Data: sem
## REML criterion at convergence: 34817.4
##
## Scaled residuals:
      Min
                10 Median
                                3Q
                                       Max
## -1.5644 -0.4463 -0.1554 0.2341 18.0356
## Random effects:
## Groups
                            Variance Std.Dev.
                Name
## trial_index (Intercept)
                               632.7 25.15
                             22503.6 150.01
## subject
                (Intercept)
## Residual
                            117122.7 342.23
## Number of obs: 2400, groups: trial_index, 240; subject, 10
## Fixed effects:
##
                  Estimate Std. Error t value
                               50.468 13.191
## (Intercept)
                   665.739
## pathlengthfac2
                    56.741
                               24.261
                                        2.339
                               24.261
                                        1.578
## pathlengthfac3
                    38.288
## pathlengthfac4
                    17.243
                               24.263
                                        0.711
## pathlengthfac5
                    1.506
                               24.248
                                        0.062
## pathlengthfac6 -45.008
                               24.264 -1.855
##
## Correlation of Fixed Effects:
##
               (Intr) pthln2 pthln3 pthln4 pthln5
## pthlngthfc2 -0.240
## pthlngthfc3 -0.240
                      0.500
## pthlngthfc4 -0.240 0.500
                              0.500
## pthlngthfc5 -0.240 0.500 0.500
## pthlngthfc6 -0.240 0.500 0.500 0.500
## Linear mixed model fit by REML ['lmerMod']
## Formula: rt ~ pathlengthfac + (pathlengthfac | subject)
##
     Data: sem
##
## REML criterion at convergence: 34793.1
##
## Scaled residuals:
##
      Min
                1Q Median
                                3Q
                                       Max
## -1.8827 -0.4447 -0.1548 0.2204 17.8681
## Random effects:
   Groups
                            Variance Std.Dev. Corr
                             34966
                                     186.99
##
   subject
            (Intercept)
##
             pathlengthfac2
                              2082
                                      45.63
                                              -0.08
##
             pathlengthfac3
                             13388
                                     115.71
                                              -0.78 0.67
##
             pathlengthfac4
                              7379
                                      85.90
                                              -0.76 0.71 0.97
##
             pathlengthfac5
                                      75.71
                                              -0.81 0.65 0.96 1.00
                              5731
```

```
pathlengthfac6 2336
                                 48.33 -0.59 0.01 0.57 0.36 0.36
                          115832 340.34
## Residual
## Number of obs: 2400, groups: subject, 10
## Fixed effects:
##
                Estimate Std. Error t value
## (Intercept)
              665.950 61.532 10.823
## pathlengthfac2 56.898
                                    2.028
                            28.060
## pathlengthfac3 38.045
                            43.795 0.869
## pathlengthfac4
                 16.738
                            36.291 0.461
## pathlengthfac5
                 0.925
                            33.945 0.027
## pathlengthfac6 -45.100
                            28.509 -1.582
## Correlation of Fixed Effects:
##
              (Intr) pthln2 pthln3 pthln4 pthln5
## pthlngthfc2 -0.209
## pthlngthfc3 -0.732 0.522
## pthlngthfc4 -0.675 0.557 0.788
## pthlngthfc5 -0.685 0.538 0.760 0.761
## pthlngthfc6 -0.470 0.365 0.489 0.426 0.436
"
```