

Semantic Networks Analysis

Abhilasha Kumar

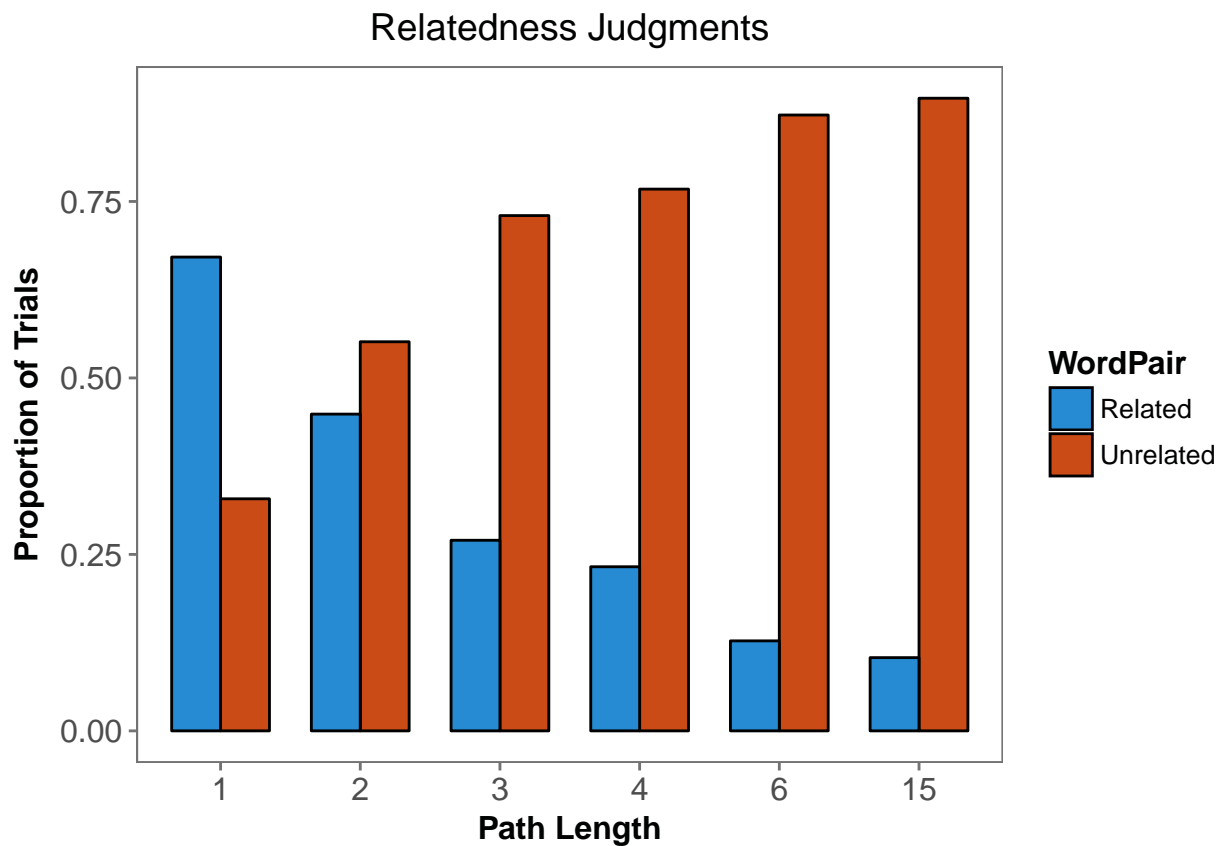
January 16, 2018

Reading the Data

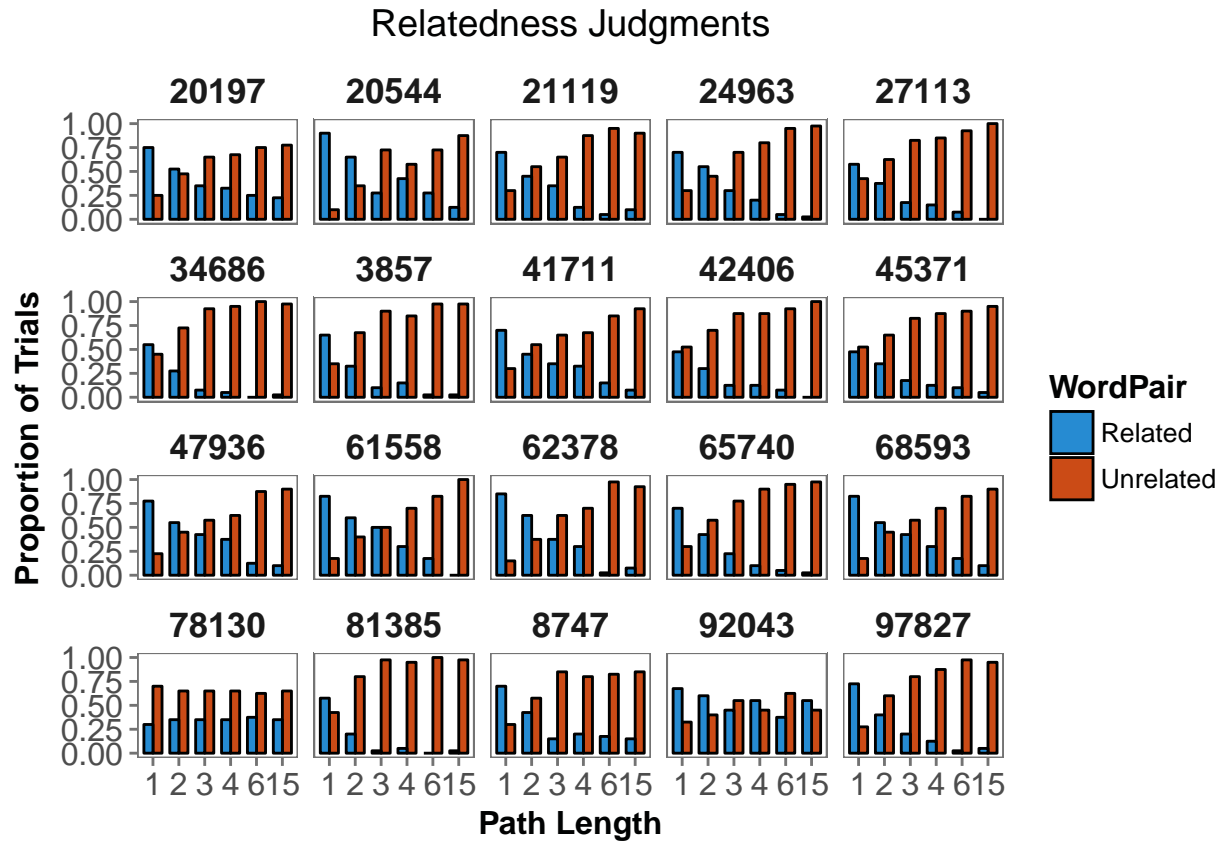
Related-Unrelated Decisions

```
##  
## 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
```

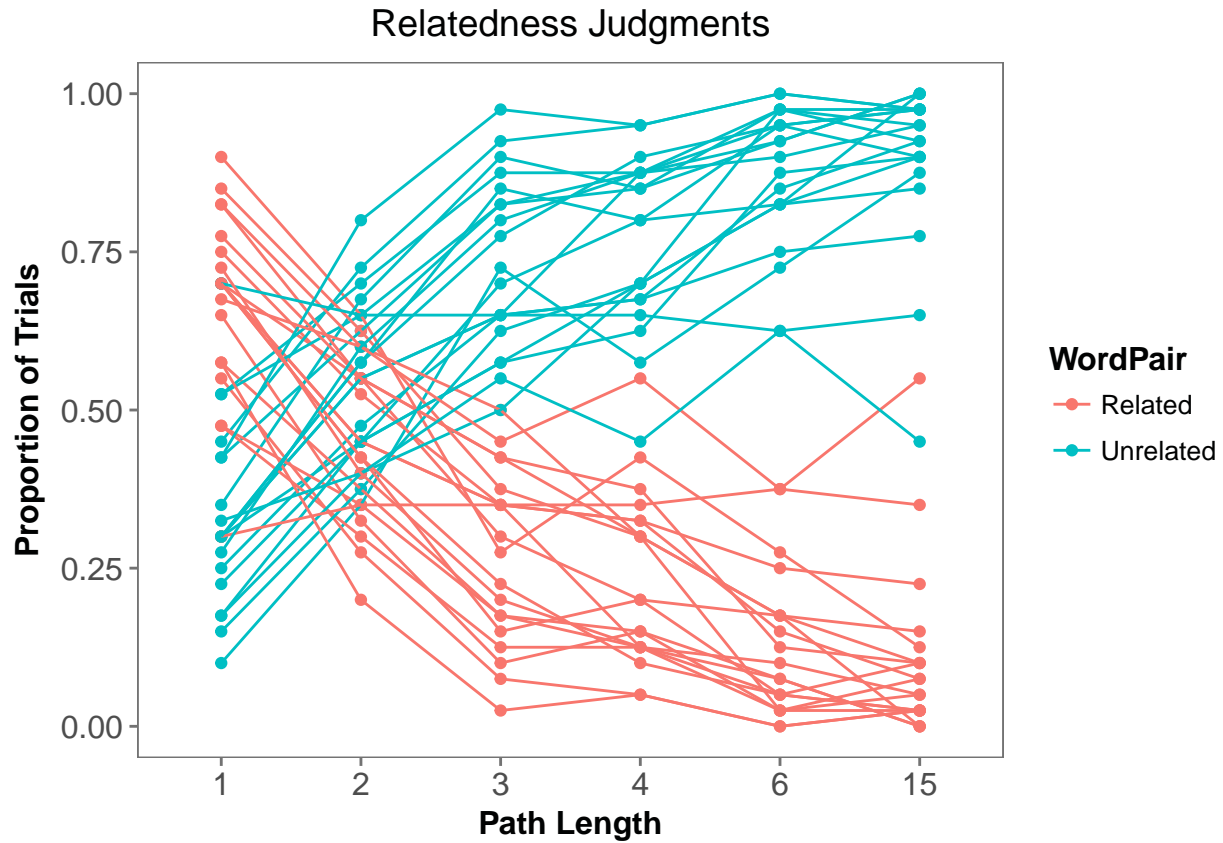
Plotting Proportions



Subject-Wise



Line Plot Subject-Wise



ANOVA

```
relunrel_aov = aov(data = sem_decision, proportion ~ pathlengthfac*Type +
                    Error(subject/(pathlengthfac*Type)))
summary(relunrel_aov)
```

```
##
## Error: subject
##           Df    Sum Sq   Mean Sq F value Pr(>F)
## Residuals 19 4.158e-30 2.188e-31
##
## Error: subject:pathlengthfac
##           Df    Sum Sq   Mean Sq F value Pr(>F)
## pathlengthfac 5 3.330e-30 6.655e-31  1.702  0.142
## Residuals    95 3.716e-29 3.911e-31
##
## Error: subject:Type
##           Df Sum Sq Mean Sq F value Pr(>F)
## Type       1  8.759   8.759   68.57 1e-07 ***
## Residuals 19  2.427   0.128
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
```

```
## Error: subject:pathlengthfac:Type
##               Df Sum Sq Mean Sq F value Pr(>F)
## pathlengthfac:Type  5  9.328  1.8656   109.1 <2e-16 ***
## Residuals          95  1.624  0.0171
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

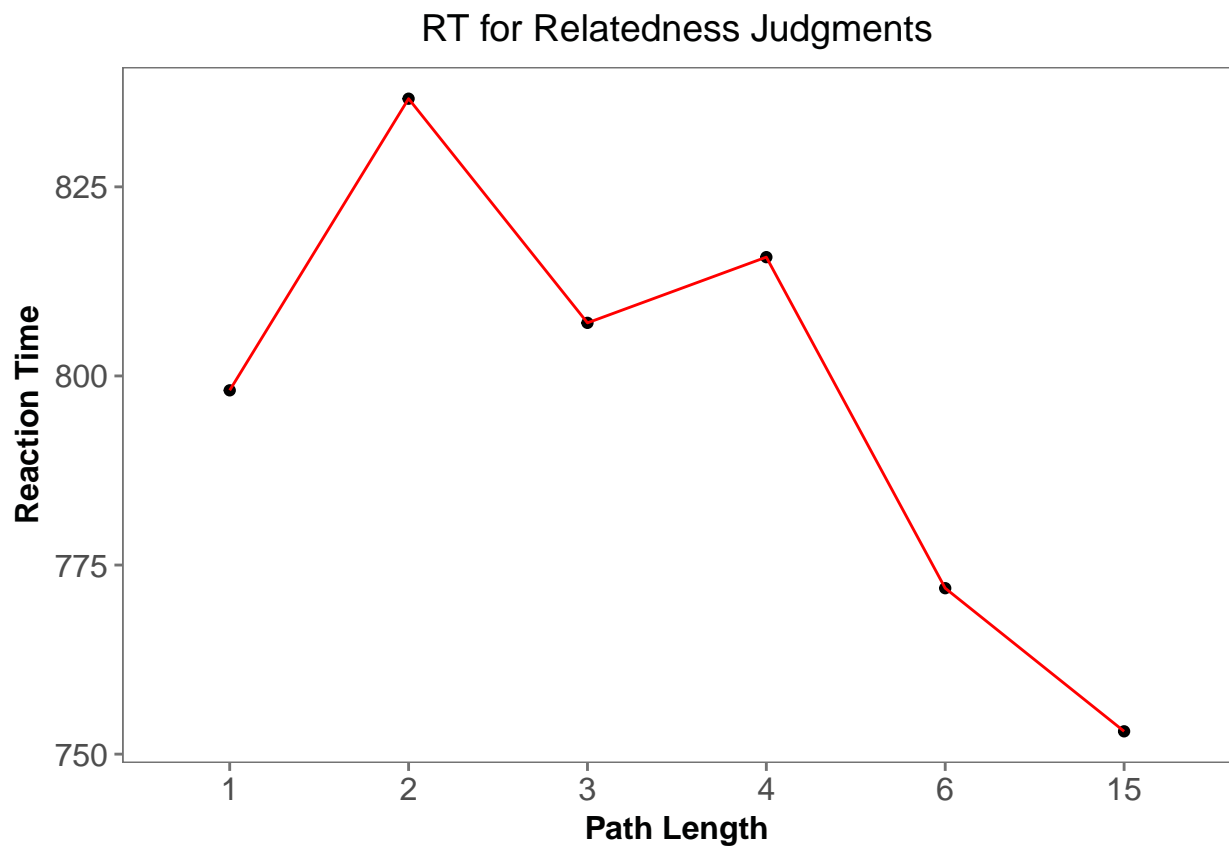
Raw Reaction Time

ANOVA

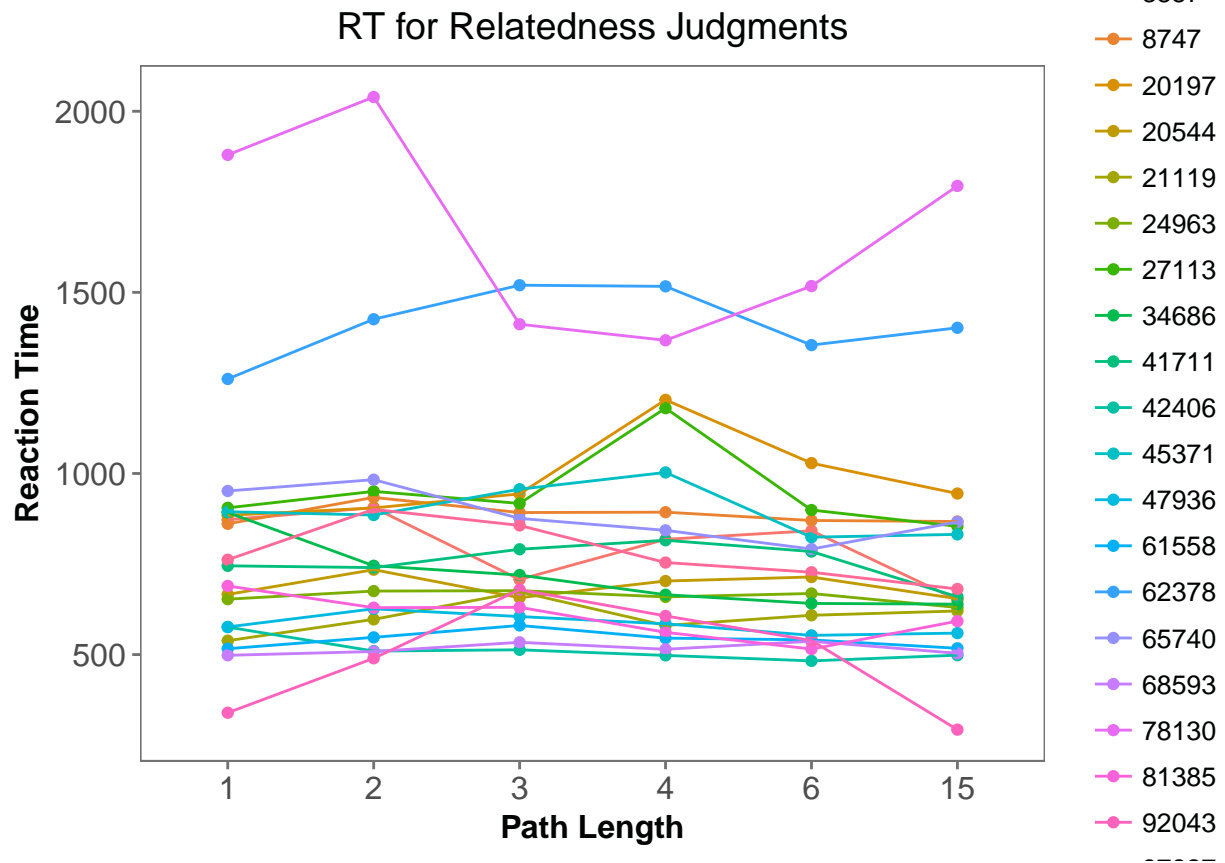
```
sem_rt$pathlengthfac = ordered(as.factor(as.character(sem_rt$pathlength)),
                               levels = c("1", "2", "3", "4", "6", "15"))
sem_rt$subject = as.factor(sem_rt$subject)
rt_aov = aov(data = sem_rt, rt ~ pathlengthfac +
              Error(subject/(pathlengthfac)))
summary(rt_aov)
```

```
##
## Error: subject
##               Df Sum Sq Mean Sq F value Pr(>F)
## Residuals 19 10213162  537535
##
## Error: subject:pathlengthfac
##               Df Sum Sq Mean Sq F value Pr(>F)
## pathlengthfac  5  91732  18346   2.097 0.0724 .
## Residuals      95 831163   8749
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Plotting RTs



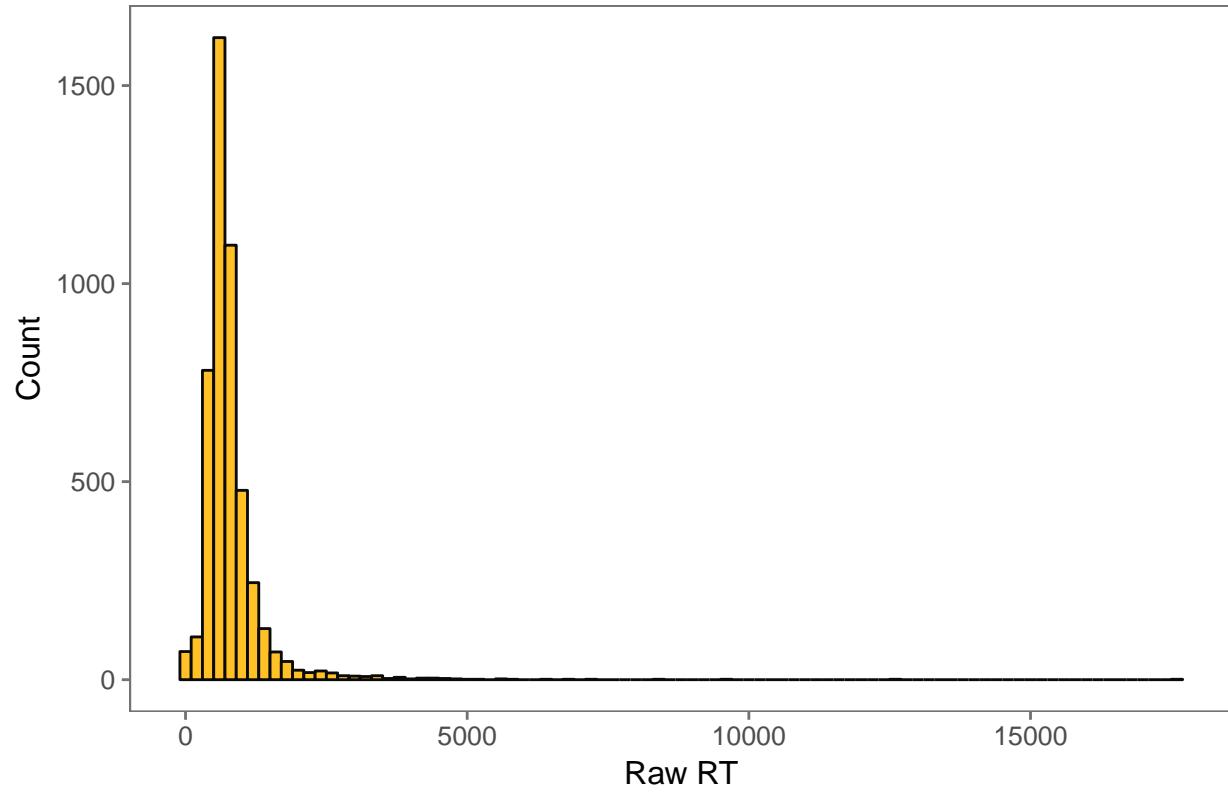
Subject-Wise



z-scored Reaction Time

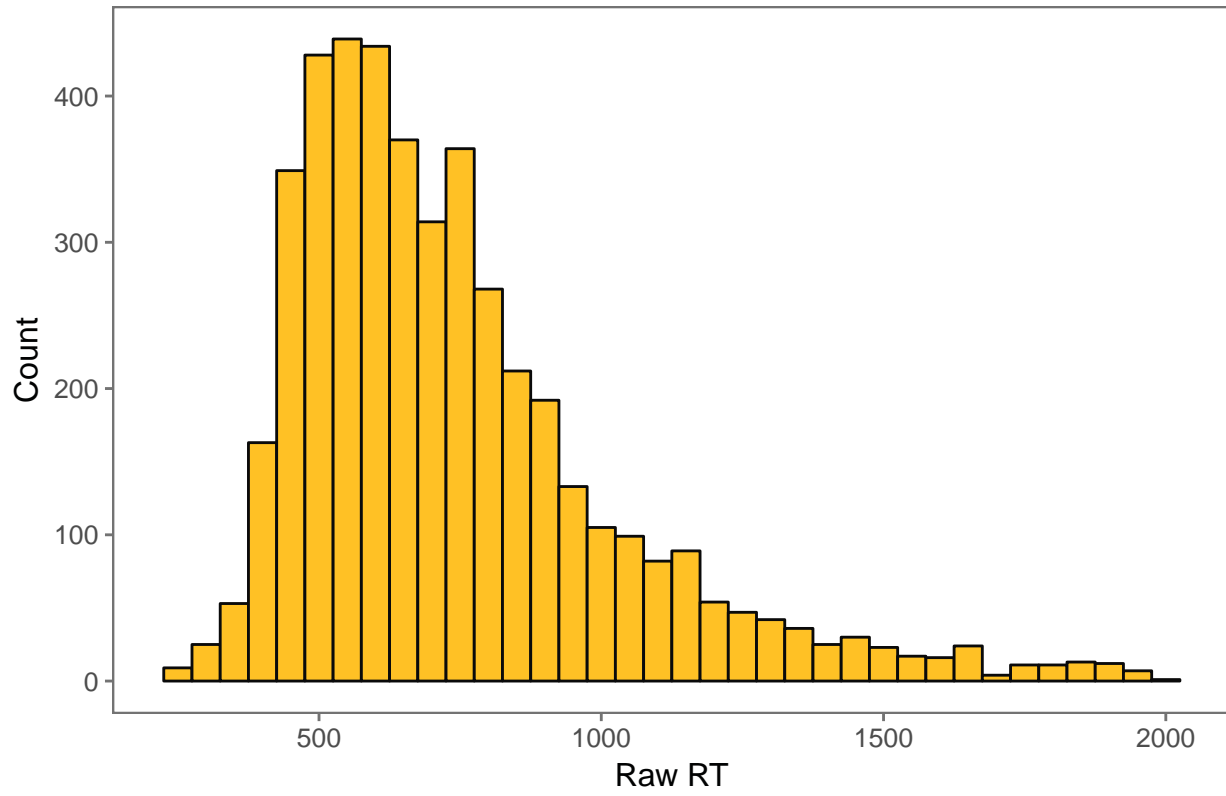
Histogram of RT

Raw RT Histogram for All Trials

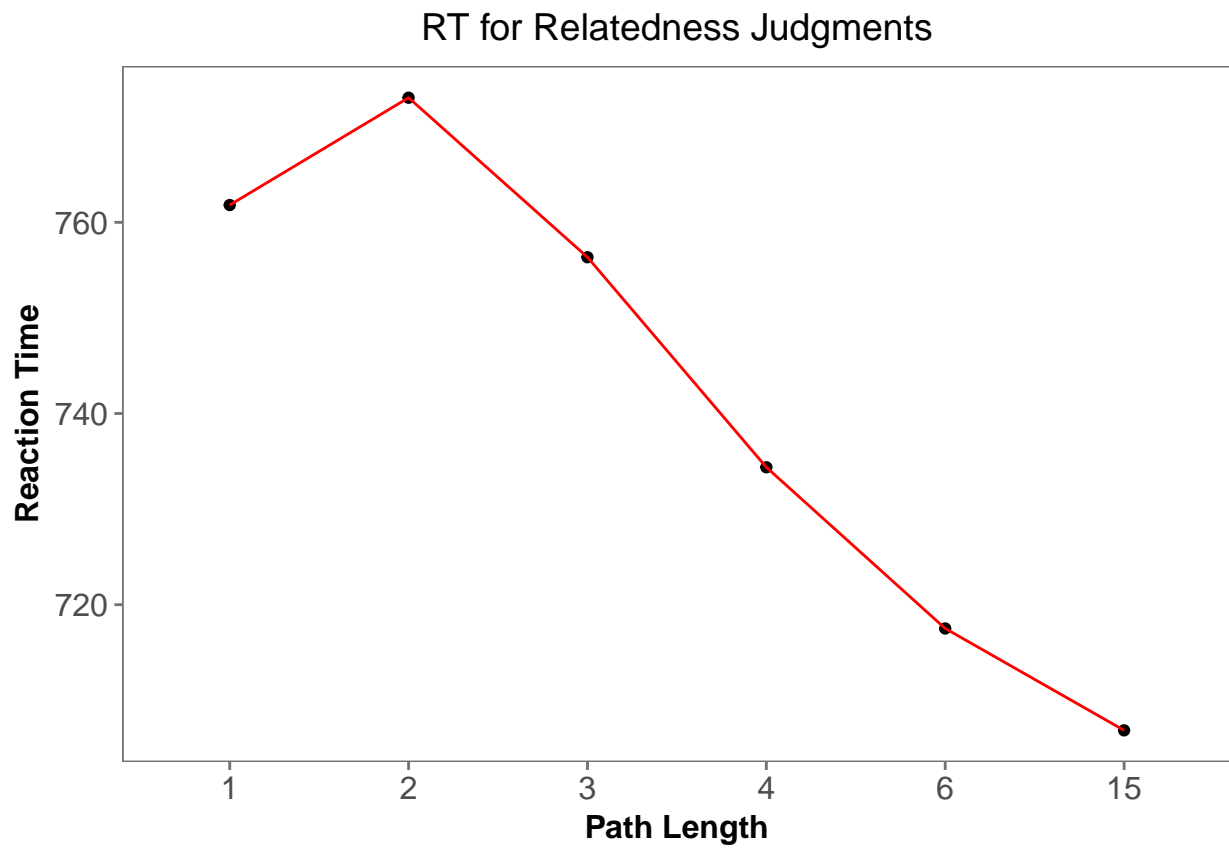


First Trim

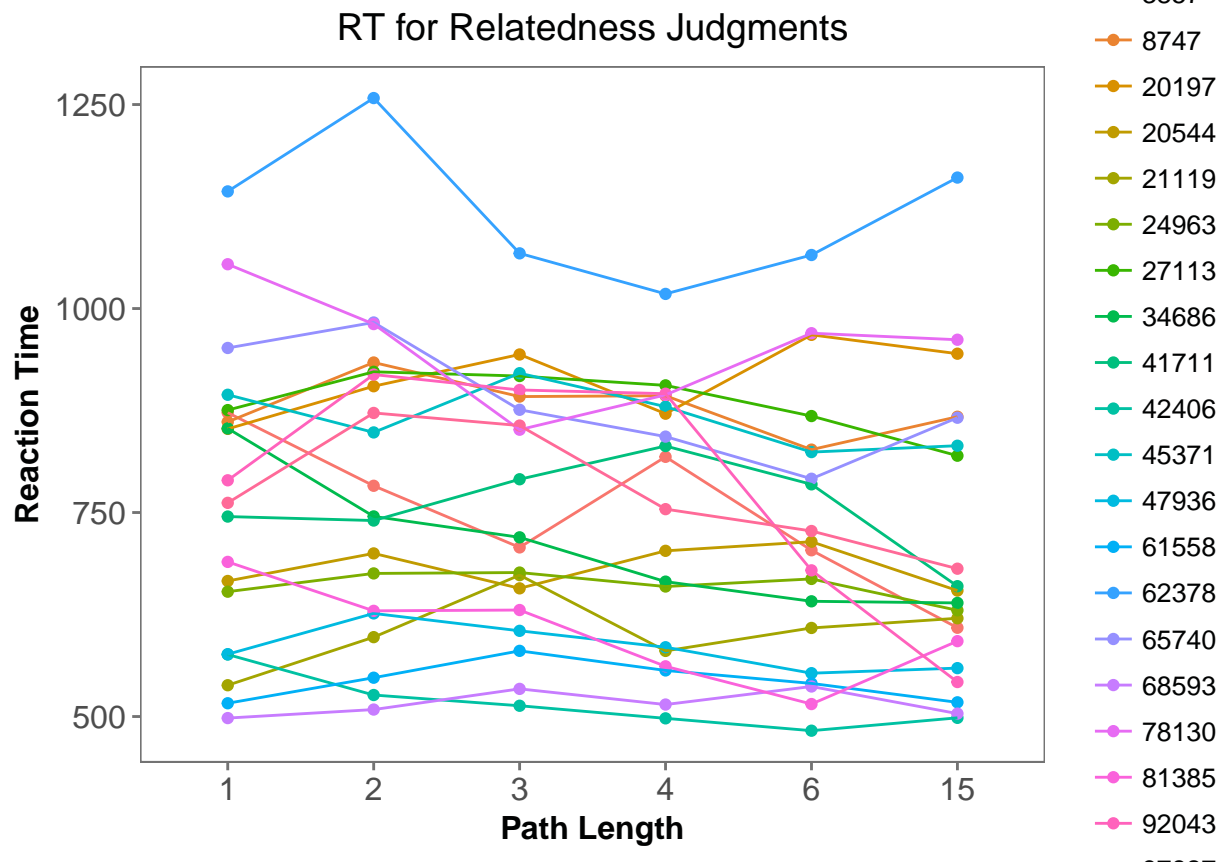
Raw RT Histogram for Trials Above 250 ms and below 2000 ms



Raw RT aggregates After Trimming

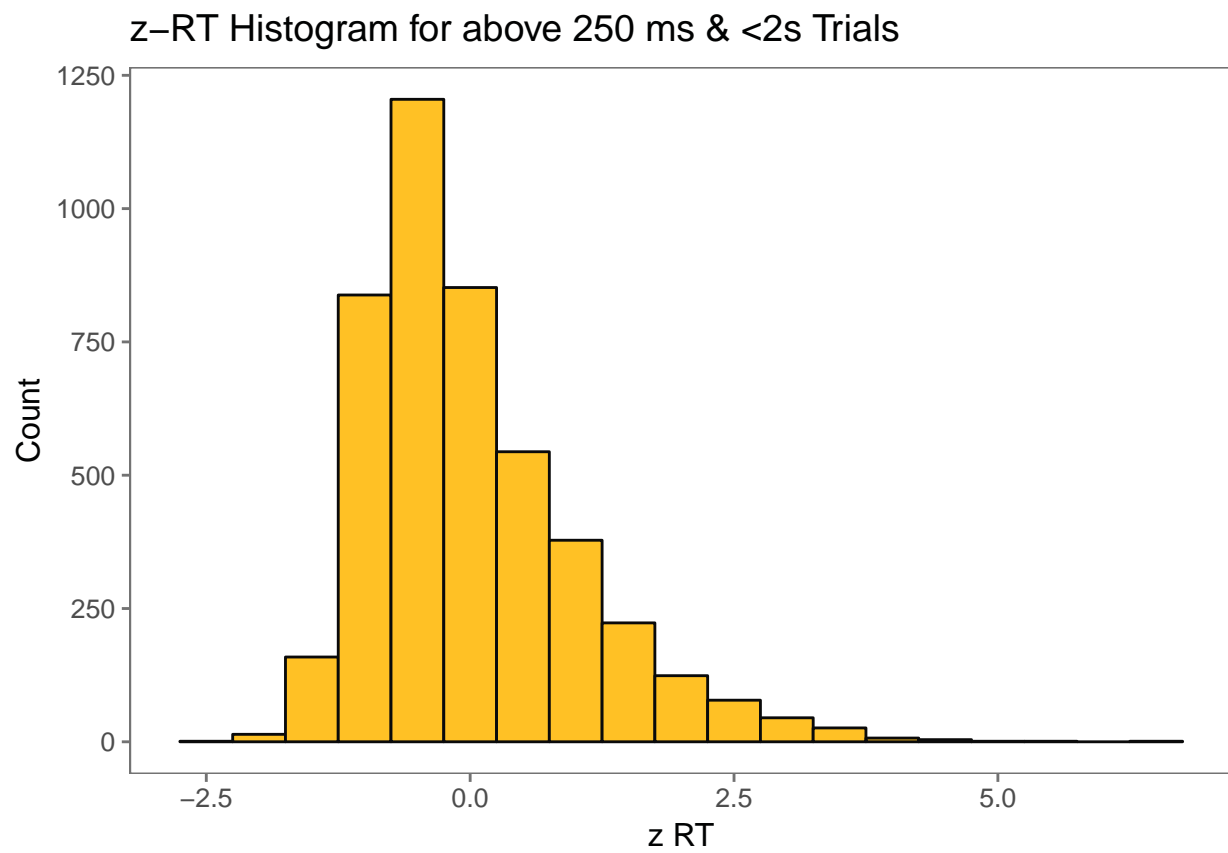


Subject Raw RT again



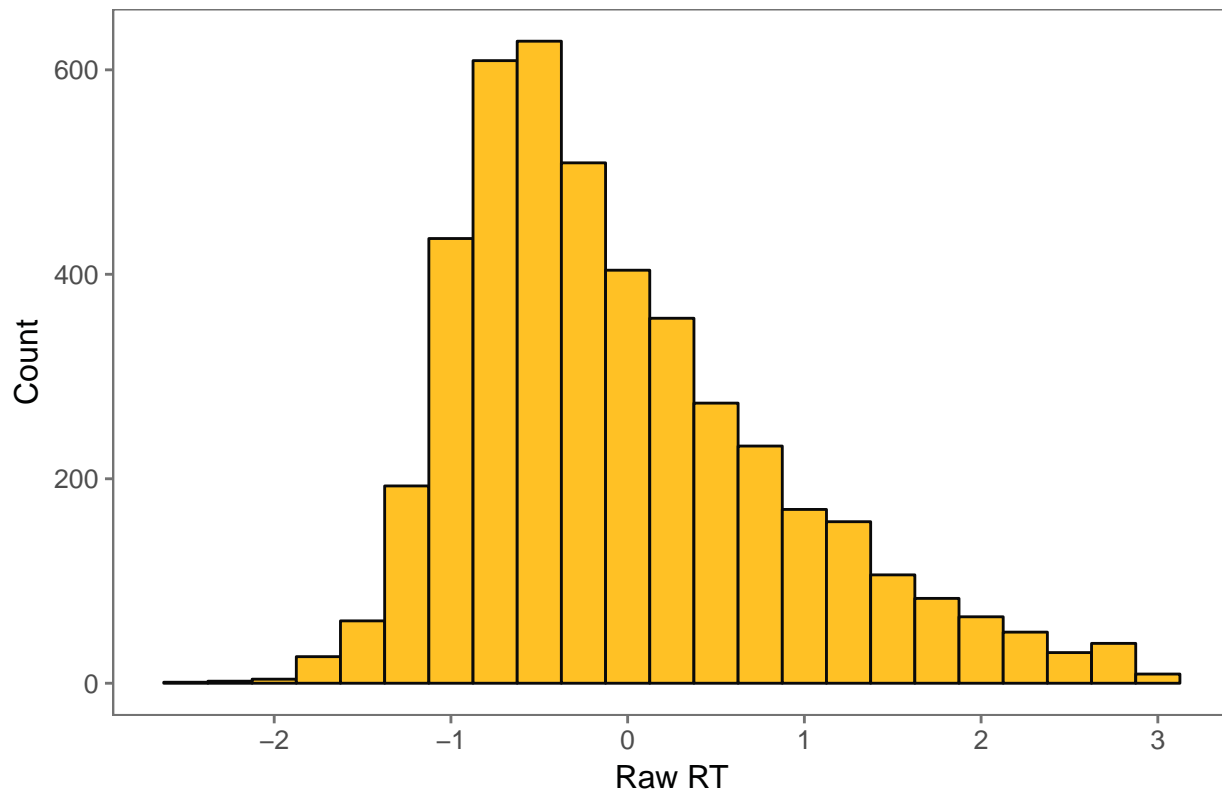
Making the z-scores

z-RT Distribution



Trimming z-RT

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



Repeating z-scoring

Aggregating zRT

ANOVA

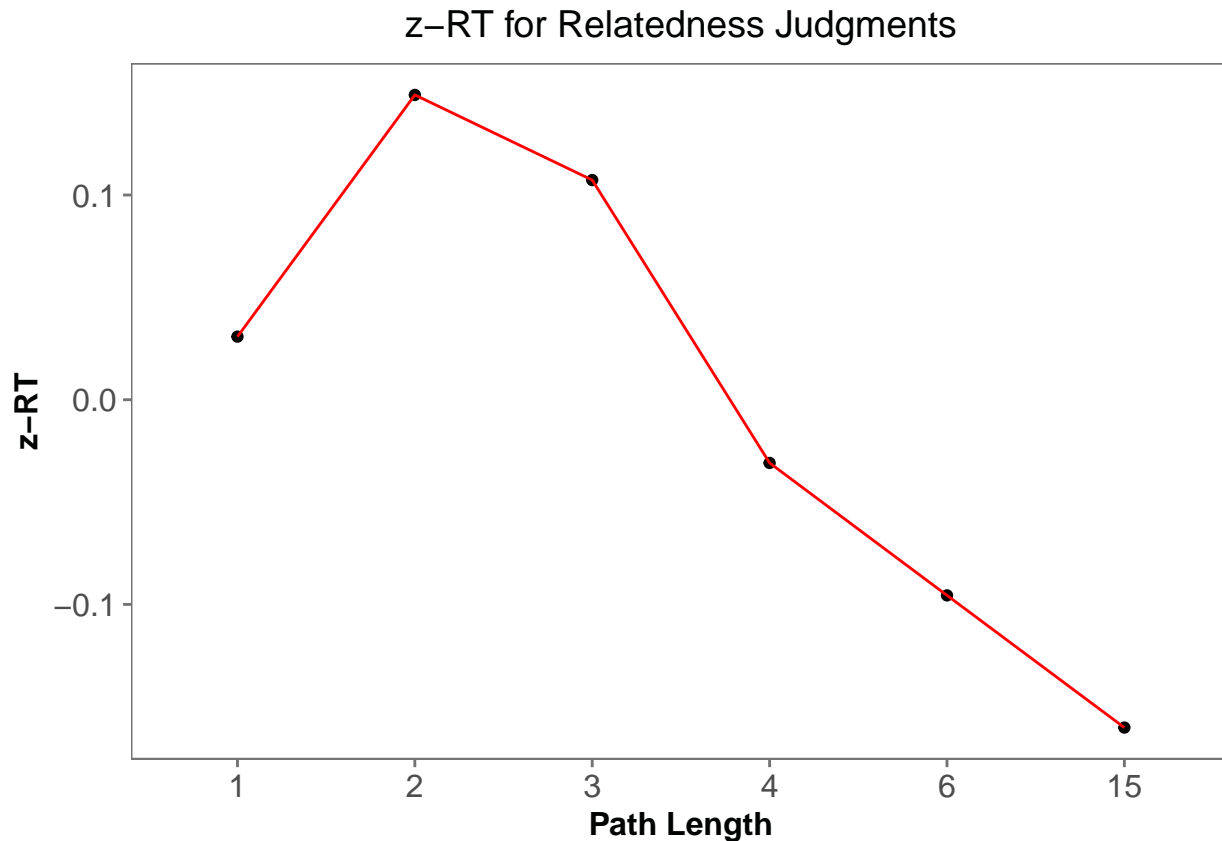
```
z_sem_rt$pathlengthfac = ordered(as.factor(as.character(z_sem_rt$pathlength)),
                                levels = c("1", "2", "3", "4", "6", "15"))
z_sem_rt$subject = as.factor(z_sem_rt$subject)

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 19 0.003357 0.0001767
##
## Error: subject:pathlengthfac
##           Df Sum Sq Mean Sq F value Pr(>F)
## pathlengthfac 5 1.492 0.29848 7.018 1.28e-05 ***
## Residuals    95 4.040 0.04253
```

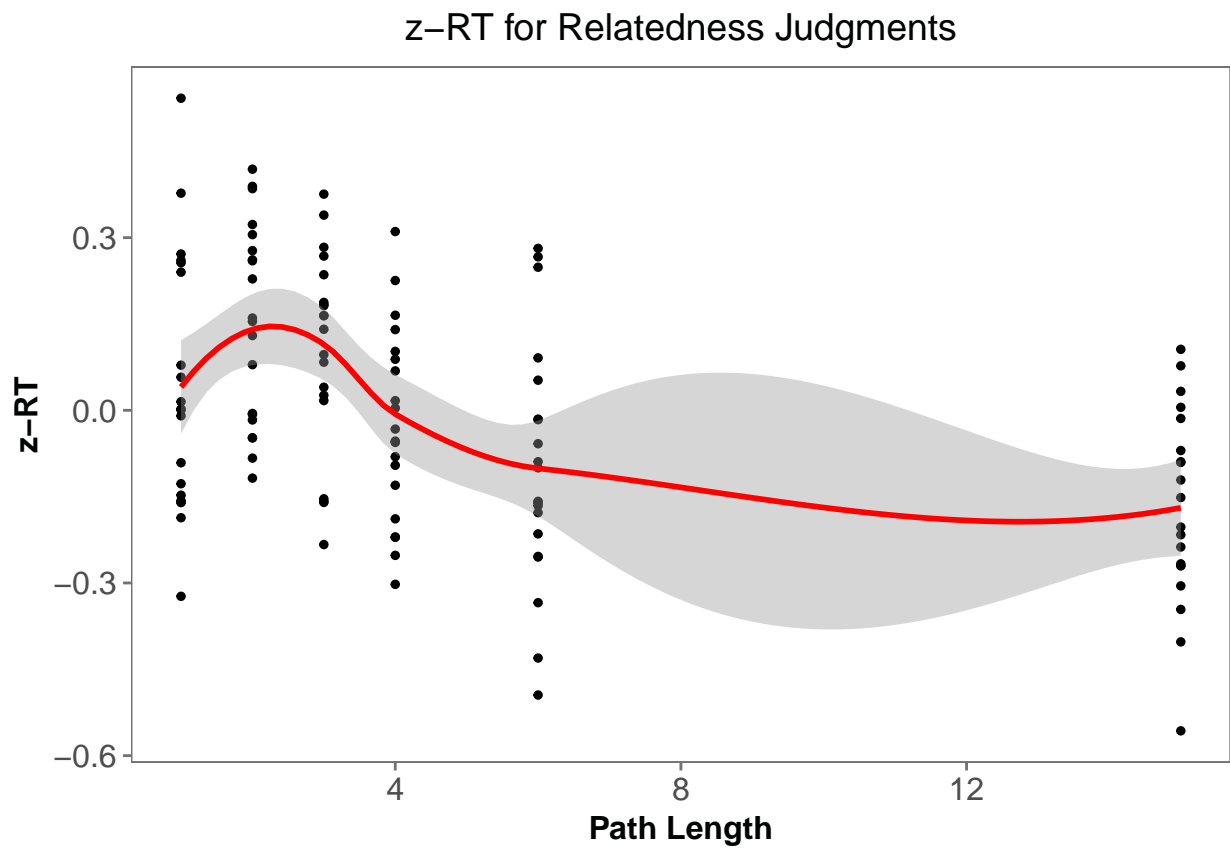
```
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Plotting RTs

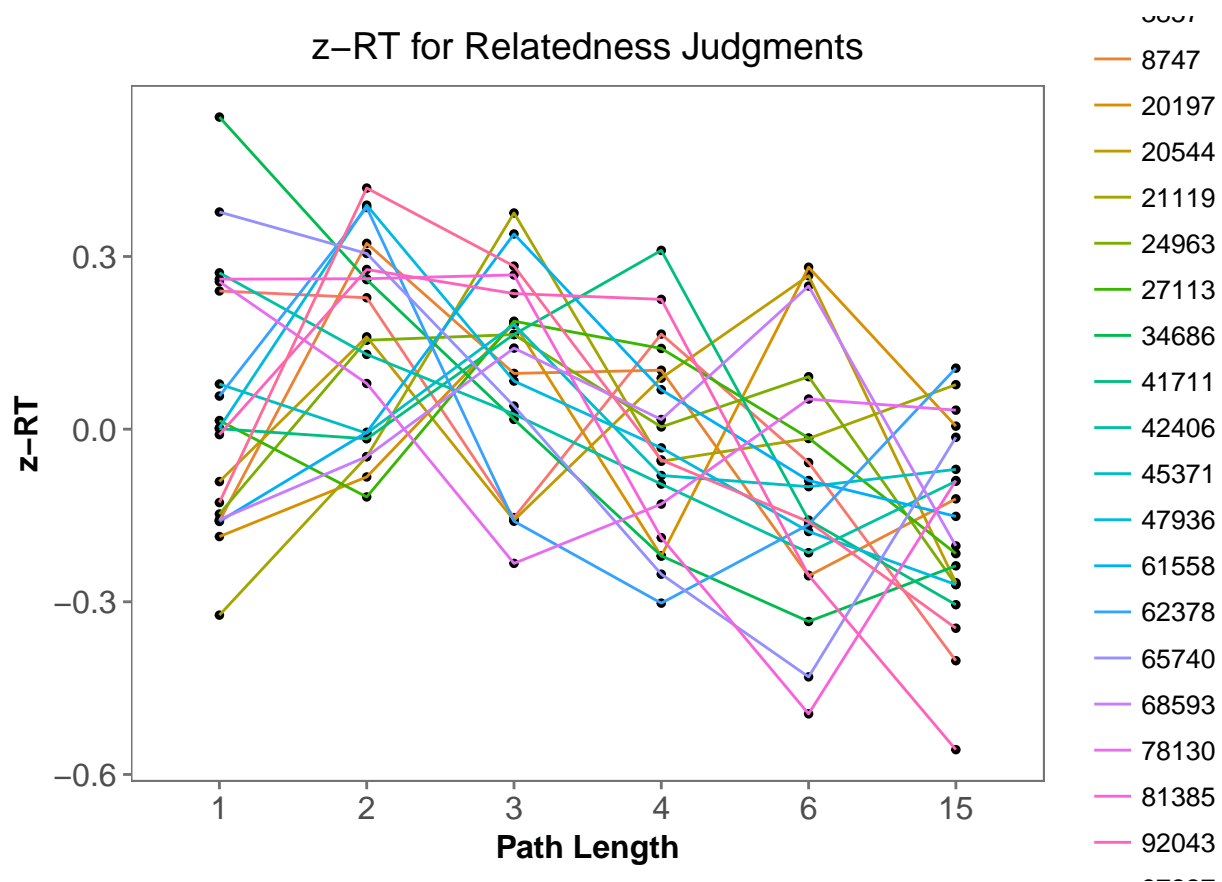


Quadratic Trend

```
z_sem_rt %>%
  ggplot(aes(x = pathlength, y = zRT_trim))+
  geom_point(color = "black", size = 1)+
  geom_smooth(method = "loess", color = "red")+
  #geom_errorbar(aes(ymin=Trials - ci, ymax=Trials + ci),
  #              width=.2, color = "gray26",
  #              position = position_dodge(0.7))+
  theme_few()+
  # scale_x_continuous(breaks = c(1,2,3,4,5,6,10,15,20))+
  xlab("Path Length") + ylab("z-RT") +
  ggtitle("z-RT for Relatedness Judgments") +
  # facet_wrap(~subject)+
  theme(axis.text = element_text(size = rel(1)),
        axis.title = element_text(face = "bold", size = rel(1)),
        legend.title = element_text(face = "bold", size = rel(1)),
        plot.title = element_text(hjust = .5),
        strip.text.x = element_text(face = "bold", size = rel(1.4)))
```



Subject-Wise



Regressions

```
## Loading required package: Matrix
## Linear mixed model fit by REML ['lmerMod']
## Formula: rt ~ 1 + (1 | subject)
## Data: sem
##
## REML criterion at convergence: 74311.4
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.7700 -0.3339 -0.1147  0.1521 28.8328
##
## Random effects:
## Groups Name Variance Std.Dev.
## subject (Intercept) 88318 297.2
## Residual 305062 552.3
## Number of obs: 4800, groups: subject, 20
##
## Fixed effects:
## Estimate Std. Error t value
```

```

## (Intercept)    797.07      66.93    11.91
## [1] 0.2245106
## Warning: NAs introduced by coercion
## Warning: NAs introduced by coercion
## Warning: NAs introduced by coercion
## Warning: NAs introduced by coercion
## Warning: NAs introduced by coercion
## Warning: NAs introduced by coercion
## Warning: NAs introduced by coercion
## Linear mixed model fit by REML ['lmerMod']
## Formula: zRT_trim ~ pathlengthfac + (1 | subject)
##   Data: new_sem_z
##
## REML criterion at convergence: 12570.3
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.5976 -0.7370 -0.2157  0.5595  3.6323
##
## Random effects:
##   Groups   Name      Variance Std.Dev.
##   subject (Intercept) 3.104e-32 1.762e-16
##   Residual              9.851e-01 9.925e-01
## Number of obs: 4445, groups:  subject, 20
##
## Fixed effects:
##              Estimate Std. Error t value
## (Intercept)    0.14891    0.03663   4.065
## pathlengthfac1 -0.11808    0.05169  -2.285
## pathlengthfac3 -0.04160    0.05153  -0.807
## pathlengthfac4 -0.17979    0.05181  -3.470
## pathlengthfac5 -0.24449    0.05163  -4.735
## pathlengthfac6 -0.30908    0.05167  -5.982
##
## Correlation of Fixed Effects:
##              (Intr) pthln1 pthln3 pthln4 pthln5
## pthlngthfc1 -0.709
## pthlngthfc3 -0.711  0.504
## pthlngthfc4 -0.707  0.501  0.503
## pthlngthfc5 -0.709  0.503  0.504  0.502
## pthlngthfc6 -0.709  0.503  0.504  0.501  0.503
## Linear mixed model fit by REML ['lmerMod']
## Formula: rt ~ pathlengthfac + (pathlengthfac | subject)
##   Data: sem
##
## REML criterion at convergence: 74218.1

```



```

##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -3.2746 -0.3265 -0.1081  0.1550 28.4882
##
## Random effects:
##      Groups   Name                Variance Std.Dev. Corr
## subject (Intercept)      103398    321.56
##           pathlengthfac2    1574     39.67   0.82
##           pathlengthfac3   16164    127.14  -0.67 -0.13
##           pathlengthfac4   25041    158.24  -0.46  0.13  0.97
##           pathlengthfac5   10178    100.89  -0.64 -0.10  1.00  0.97
##           pathlengthfac6    1772     42.10   0.09  0.64  0.68  0.85  0.71
## Residual                300580    548.25
## Number of obs: 4800, groups:  subject, 20
##
## Fixed effects:
##              Estimate Std. Error t value
## (Intercept)    798.099      74.469  10.717
## pathlengthfac2   38.555      28.812   1.338
## pathlengthfac3    8.919      39.493   0.226
## pathlengthfac4   17.606      44.760   0.393
## pathlengthfac5  -26.163      35.502  -0.737
## pathlengthfac6 -45.083      28.984  -1.555
##
## Correlation of Fixed Effects:
##              (Intr) pthln2 pthln3 pthln4 pthln5
## pthlngthfc2  0.070
## pthlngthfc3 -0.594  0.301
## pthlngthfc4 -0.461  0.322  0.762
## pthlngthfc5 -0.538  0.348  0.725  0.726
## pthlngthfc6 -0.147  0.514  0.487  0.507  0.511
##
## Linear mixed model fit by REML ['lmerMod']
## Formula: zRT_trim ~ pathlength + (1 | subject)
## Data: new_sem_z
##
## REML criterion at convergence: 12576.2
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.6410 -0.7267 -0.2113  0.5628  3.6608
##
## Random effects:
##      Groups   Name                Variance Std.Dev.
## subject (Intercept) 3.114e-32 1.765e-16
## Residual          9.883e-01 9.941e-01
## Number of obs: 4445, groups:  subject, 20
##
## Fixed effects:
##              Estimate Std. Error t value
## (Intercept)  0.096877   0.022245   4.355
## pathlength  -0.018731   0.003192  -5.869
##

```

```

## Correlation of Fixed Effects:
##      (Intr)
## pathlength -0.742

## Linear mixed model fit by REML ['lmerMod']
## Formula: zRT_trim ~ pathlength + I((pathlength)^2) + (1 | subject)
##      Data: new_sem_z
##
## REML criterion at convergence: 12585.2
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.6778 -0.7286 -0.2136  0.5582  3.6670
##
## Random effects:
##      Groups   Name      Variance Std.Dev.
## subject (Intercept) 3.113e-32 1.764e-16
## Residual          9.878e-01 9.939e-01
## Number of obs: 4445, groups: subject, 20
##
## Fixed effects:
##              Estimate Std. Error t value
## (Intercept)    0.1583255  0.0412449   3.839
## pathlength    -0.0458356  0.0156500  -2.929
## I((pathlength)^2) 0.0016209  0.0009162   1.769
##
## Correlation of Fixed Effects:
##      (Intr) pthlng
## pathlength -0.906
## I((pthl)^2)  0.842 -0.979

```