

Shape Discrimination Analysis

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Imports

Load Data

Data is all of the format `${subject}_Behavioral-2_Discrimination_${date}.csv`.

```
## [1] "Loaded 50 subjects"
```

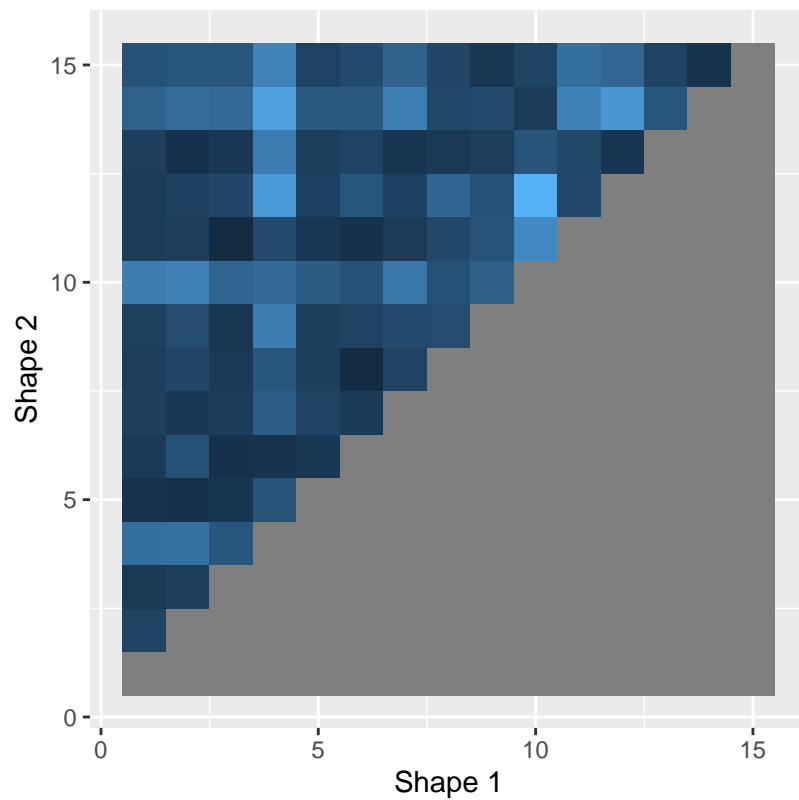
Read existing LOC triu data. This is the shape-by-shape dissimilarity metric, across both modular and lattice subjects.

```
## Warning: The `x` argument of `as_tibble.matrix()` must have unique column names if
```

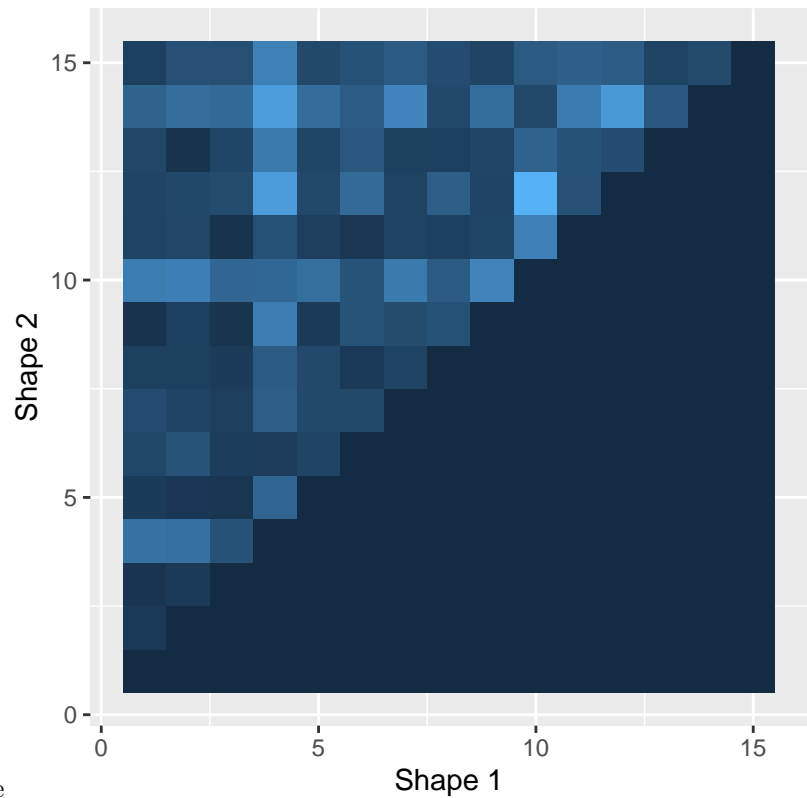
```
## `.name_repair` is omitted as of tibble 2.0.0.
```

```
## i Using compatibility `.name_repair`.
```

LOC RDM

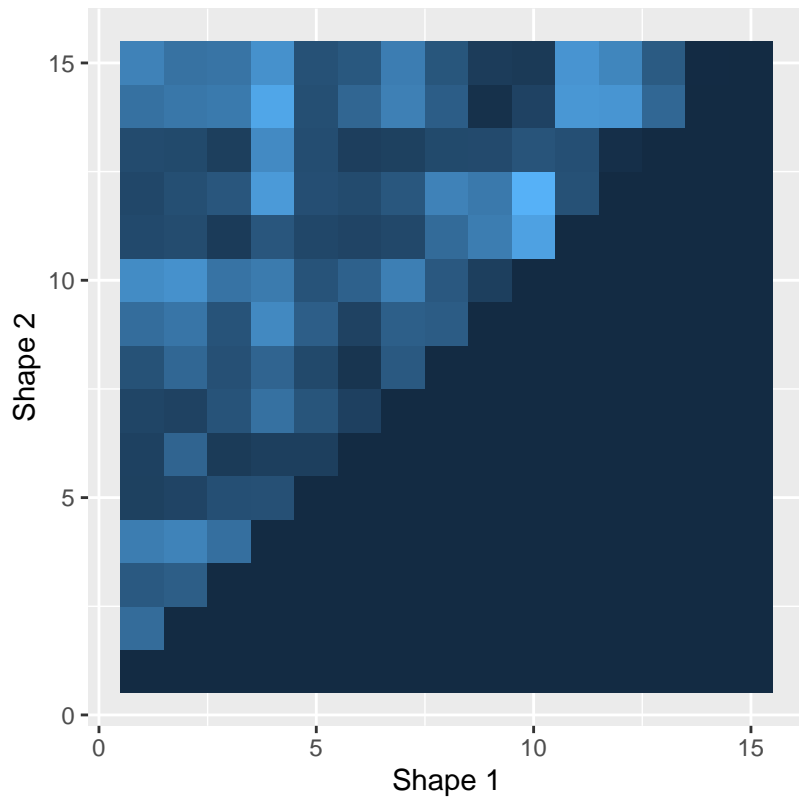


Modular LOC RDM



Also separate for modular and lattice

Lattice LOC RDM



of the results.

And bind the LOC data to the result

Final dataframe:

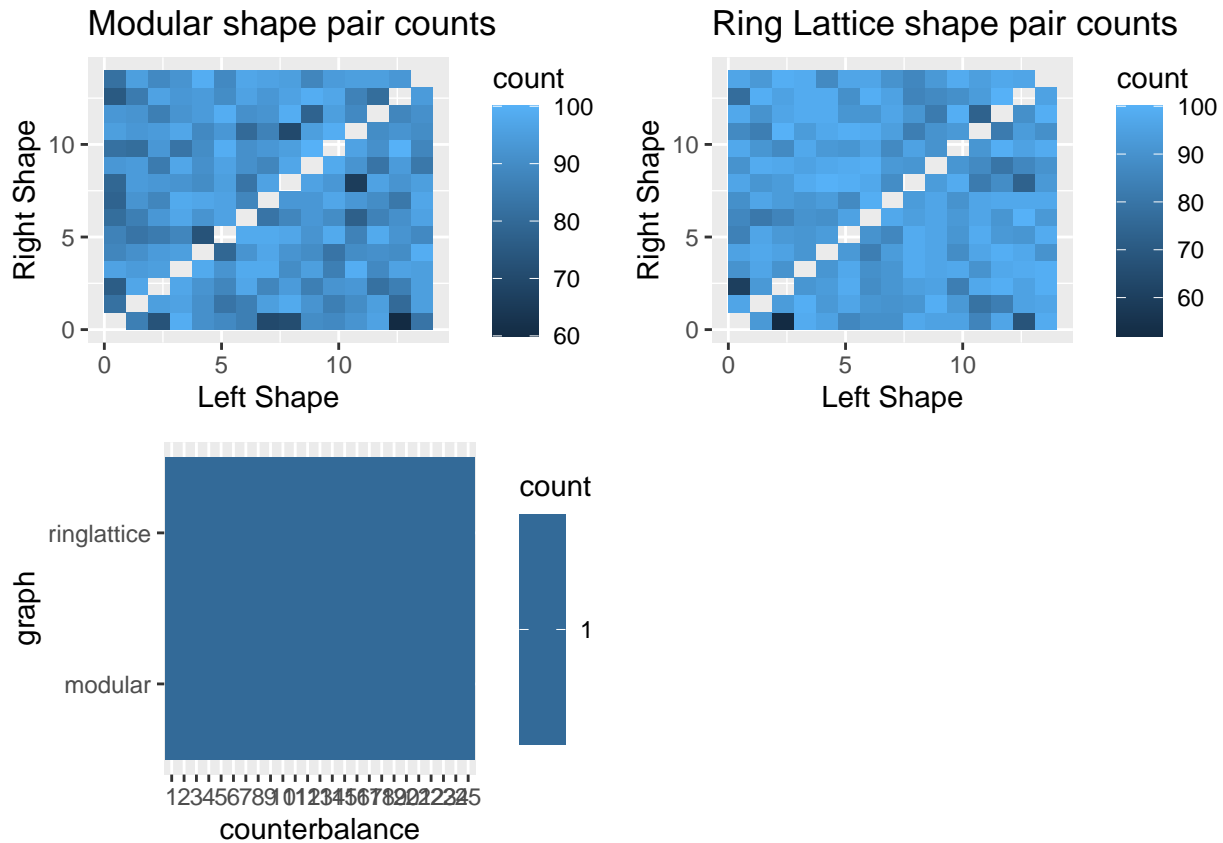
```
## # A tibble: 6 x 44
##   node_left node_~1 shape~2 shape~3 varia~4 varia~5 jitter trial discr~6 discr~7
##   <dbl>    <dbl>    <dbl>    <dbl>    <dbl>    <dbl>    <dbl> <dbl>    <dbl>    <dbl>
## 1         3      12         1         7         0         3 -0.15     3         0         2
## 2        11         9         9         8         3         0 -0.25     4         0         3
## 3         2        14         6         3         4         1 -0.15     6         0         5
## 4         8         2         5         6         4         1  0         7         0         6
## 5        10        12        14         7         1         4  0.1       11        0        10
## 6        10         2        14         6         3         0  0.2       15         0        14
## # ... with 34 more variables: discrimination.thisN <dbl>,
## #   discrimination.thisIndex <dbl>, experiment.time <dbl>, block.type <chr>,
## #   block.name <chr>, block.timings.start <dbl>, trial.correct <lgl>,
## #   trial.response <chr>, trial.timeout <lgl>, trial.timings.stimulus <dbl>,
## #   trial.timings.fixation_one <dbl>, trial.timings.fixation_two <dbl>,
## #   trial.timings.event <dbl>, trial.timings.rt <dbl>, Experiment <chr>,
## #   subject <fct>, graph <fct>, counterbalance <fct>, subset <fct>, ...

## # A tibble: 6 x 44
##   node_left node_~1 shape~2 shape~3 varia~4 varia~5 jitter trial discr~6 discr~7
##   <dbl>    <dbl>    <dbl>    <dbl>    <dbl>    <dbl>    <dbl> <dbl>    <dbl>    <dbl>
## 1         0         0        12        12         4         1  0.05     1         0         0
## 2        12        12         7         7         3         0 -0.05     2         0         1
## 3         3        12         1         7         0         3 -0.15     3         0         2
## 4        11         9         9         8         3         0 -0.25     4         0         3
## 5         0         0        12        12         3         0  0.15     5         0         4
## 6         2        14         6         3         4         1 -0.15     6         0         5
## # ... with 34 more variables: discrimination.thisN <dbl>,
## #   discrimination.thisIndex <dbl>, experiment.time <dbl>, block.type <chr>,
## #   block.name <chr>, block.timings.start <dbl>, trial.correct <lgl>,
## #   trial.response <chr>, trial.timeout <lgl>, trial.timings.stimulus <dbl>,
## #   trial.timings.fixation_one <dbl>, trial.timings.fixation_two <dbl>,
## #   trial.timings.event <dbl>, trial.timings.rt <dbl>, Experiment <chr>,
## #   subject <fct>, graph <fct>, counterbalance <fct>, subset <fct>, ...
```

Data Verification

Are all shape pairs covered?

And make sure that each condition was covered once by condition



Results

RT and LOC Correlation

Is the average $1/RT$ correlated with the LOC dissimilarity across shape pairs?

First plot: - Each point is a shape pair - Plotting mean $1/R$ and LOC Dissimilarity values - We observe that more dissimilar shapes have lower reaction times (higher $1/RT$)

```
cor.test(df.mean$rt.inv.mean, df.mean$z.loc_val.mean)
```

```
##
## Pearson's product-moment correlation
##
## data: df.mean$rt.inv.mean and df.mean$z.loc_val.mean
## t = 3.2476, df = 103, p-value = 0.001572
## alternative hypothesis: true correlation is not equal to 0
## 95 percent confidence interval:
## 0.1201214 0.4690369
## sample estimates:
## cor
## 0.3047699
```

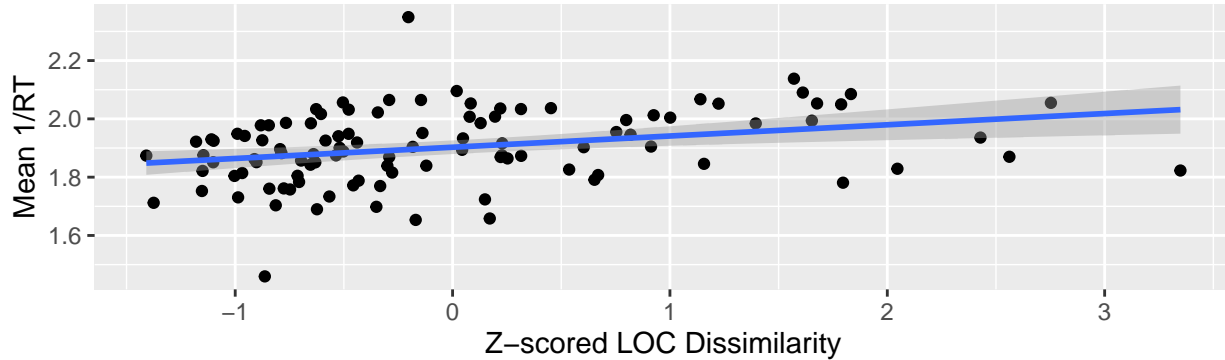
If we compute the correlation on a per-subject basis, nearly all subjects display a positive effect, in most cases very strong.

We find a strong linear relationship between mean of $1/RT$ (across subjects), and LOC dissimilarity, across all shape pairs.

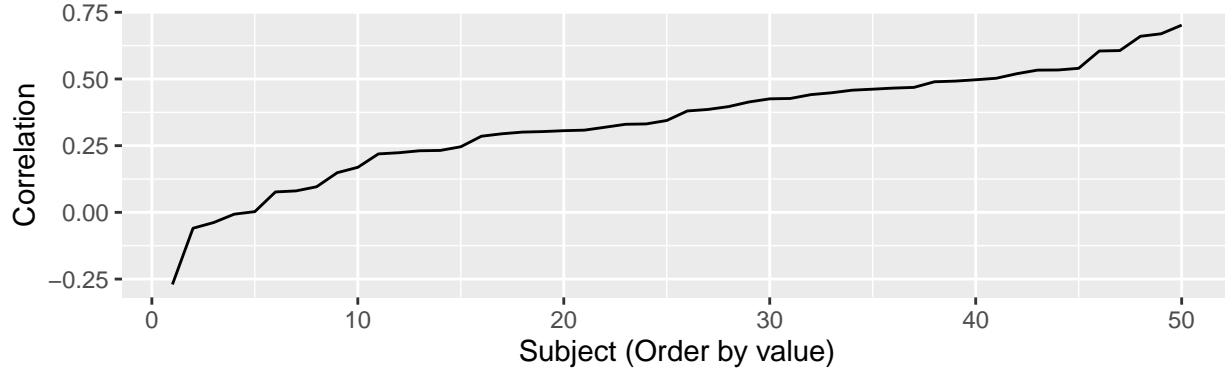
When this correlation is taken on a per-subject basis (for the behavioral data), against the canonical LOC RDM, almost all subjects show individually high correlations.

```
## `geom_smooth()` using formula = 'y ~ x'
```

A) Correlation between 1/RT and LOC, $r=0.3$ $p=0.0016$



B) Correlation between 1/RT and LOC, by subject



```
## Saving 6.5 x 4.5 in image
```

```
ggsave("images/rt_vs_loc.pdf", p1, height=5, width=5)
```

```
## `geom_smooth()` using formula = 'y ~ x'
```

Group LOC Correlation

We can also separate out the correlations by group.

```
df %>%
  group_by(subject, shape_pair_ordered) %>%
  summarise(rt.inv.median = median(rt.inv), graph=first(graph), z.loc_val = first(z.loc_val), .groups =
  group_by(subject) %>%
  summarise(correlation = cor(rt.inv.median, z.loc_val), graph=first(graph), .groups = 'drop') %>%
  group_by(graph) %>%
  summarise(correlation.mean = mean(correlation))
```

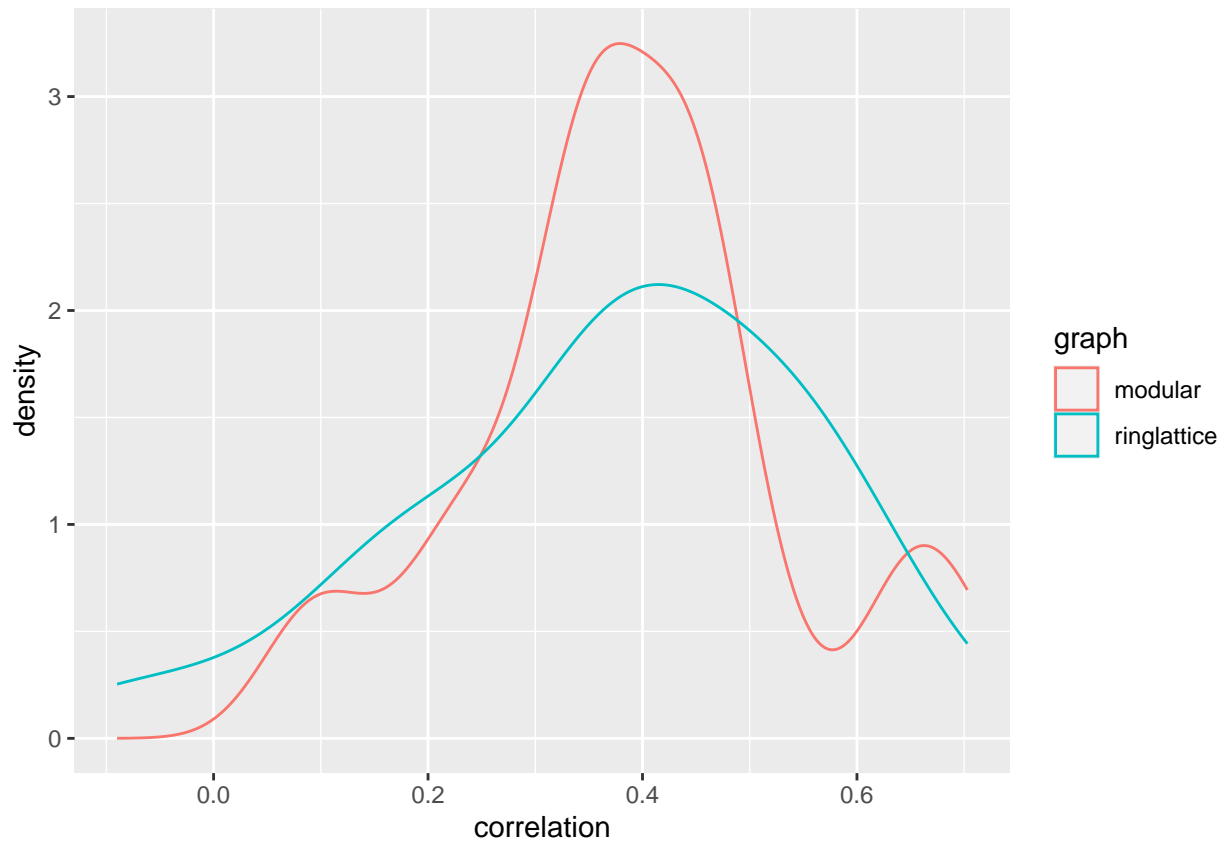
```
## # A tibble: 2 x 2
```

```
##   graph      correlation.mean
##   <fct>          <dbl>
## 1 modular      0.386
## 2 ringlattice  0.368
```

We find roughly similar distributions for correlations between subjects in the modular and lattice behavioral

groups

```
df %>%
  group_by(subject, shape_pair_ordered) %>%
  summarise(rt.inv.median = median(rt.inv), graph=first(graph), z.loc_val = first(z.loc_val), .groups =
  group_by(subject) %>%
  summarise(correlation = cor(rt.inv.median, z.loc_val), graph=first(graph), .groups = 'drop') %>%
  ggplot(aes(x=correlation, color=graph)) + geom_density()
```



Group Differences

```
df.median <- df %>%
  group_by(subject) %>%
  summarise(rt.inv.median = median(rt.inv), graph=first(graph), rt.inv.mean = mean(rt.inv))

df.median %>%
  group_by(graph) %>%
  summarise(rt.inv.median.mean = mean(rt.inv.median), rt.inv.median.sem = sd(rt.inv.median) / sqrt(leng

## # A tibble: 2 x 3
##   graph      rt.inv.median.mean rt.inv.median.sem
##   <fct>          <dbl>          <dbl>
## 1 modular          1.89          0.0583
## 2 ringlattice      1.94          0.0534

x <- df %>%
  group_by(graph) %>%
  summarise(rt.inv.median = median(rt.inv))
```

```
x

## # A tibble: 2 x 2
##   graph      rt.inv.median
##   <fct>          <dbl>
## 1 modular          1.89
## 2 ringlattice      1.97

rt.inv.median.modular <- (df.median %>% filter(graph == "modular"))$rt.inv.median
rt.inv.median.lattice <- (df.median %>% filter(graph == "ringlattice"))$rt.inv.median
t.test(rt.inv.median.modular, rt.inv.median.lattice)

##
## Welch Two Sample t-test
##
## data:  rt.inv.median.modular and rt.inv.median.lattice
## t = -0.67042, df = 47.636, p-value = 0.5058
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
##  -0.2120431  0.1060123
## sample estimates:
## mean of x mean of y
##  1.887699  1.940714
```

Basic Linear Model

First analysis: Purely linear model.

We're averaging across subjects for our behavioral RT, first.

Note that we can't include both `loc_val` and `shape_pair_ordered`, since here the effect for `shape_pair_ordered` already allows a separate fit for each shape, and we end up with collinearity. Since we don't care about `loc_val` here, don't think this is really a problem, though I've included both. `shape_pair_ordered` provides a slightly better fit, but the huge number of parameters result in a higher AIC, though not a 'significant' AIC difference (of <2)

We find graph type to be significant in either case, with decreased reaction times (higher 1/RT) for the ring lattice.

```
df.mean <- df %>%
  group_by(shape_pair_ordered, graph) %>%
  summarize(rt.inv.mean = mean(rt.inv), z.loc_val.mean = mean(z.loc_val))

## `summarise()` has grouped output by 'shape_pair_ordered'. You can override
## using the `.groups` argument.

m1 <- lm(rt.inv.mean ~ graph + z.loc_val.mean, df.mean)
summary(m1)

##
## Call:
## lm(formula = rt.inv.mean ~ graph + z.loc_val.mean, data = df.mean)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -0.51199 -0.07704 -0.00119  0.10080  0.84129
##
## Coefficients:
```

```
##               Estimate Std. Error t value Pr(>|t|)
## (Intercept)      1.86630    0.01475 126.553 < 2e-16 ***
## graphringlattice 0.07419    0.02085   3.558 0.000464 ***
## z.loc_val.mean    0.03835    0.01049   3.655 0.000326 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.1511 on 207 degrees of freedom
## Multiple R-squared:  0.1116, Adjusted R-squared:  0.1031
## F-statistic: 13.01 on 2 and 207 DF,  p-value: 4.771e-06
```

```
AIC(m1)

## [1] -192.768

df.mean <- df %>%
  group_by(shape_pair_ordered, graph) %>%
  summarize(rt.inv.mean = mean(rt.inv), z.loc_val.mean = mean(z.loc_val))

## `summarise()` has grouped output by 'shape_pair_ordered'. You can override
## using the `.groups` argument.

m2 <- lm(rt.inv.mean ~ graph + shape_pair_ordered, df.mean)
summary(m2)
```

```
##
## Call:
## lm(formula = rt.inv.mean ~ graph + shape_pair_ordered, data = df.mean)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -0.36525 -0.05922  0.00000  0.05922  0.36525
##
## Coefficients:
##               Estimate Std. Error t value Pr(>|t|)
## (Intercept)      1.8780181    0.0929362   20.208 < 2e-16 ***
## graphringlattice 0.0741937    0.0180535    4.110 7.92e-05 ***
## shape_pair_ordered0_10 0.0702896    0.1308101    0.537  0.59218
## shape_pair_ordered0_11 -0.1535583    0.1308101   -1.174  0.24312
## shape_pair_ordered0_12 0.0700183    0.1308101    0.535  0.59361
## shape_pair_ordered0_13 -0.1209685    0.1308101   -0.925  0.35723
## shape_pair_ordered0_14 0.1272187    0.1308101    0.973  0.33304
## shape_pair_ordered0_2 -0.4326641    0.1308101   -3.308  0.00129 **
## shape_pair_ordered0_3 -0.0666757    0.1308101   -0.510  0.61133
## shape_pair_ordered0_4 0.0119040    0.1308101    0.091  0.92767
## shape_pair_ordered0_5 -0.2106864    0.1308101   -1.611  0.11029
## shape_pair_ordered0_6 -0.0645646    0.1308101   -0.494  0.62265
## shape_pair_ordered0_7 -0.0666440    0.1308101   -0.509  0.61150
## shape_pair_ordered0_8 -0.0001883    0.1308101   -0.001  0.99885
## shape_pair_ordered0_9 0.0790094    0.1308101    0.604  0.54716
## shape_pair_ordered1_10 -0.0357618    0.1308101   -0.273  0.78510
## shape_pair_ordered1_11 -0.1811704    0.1308101   -1.385  0.16902
## shape_pair_ordered1_12 -0.0941281    0.1308101   -0.720  0.47340
## shape_pair_ordered1_13 0.0848057    0.1308101    0.648  0.51821
## shape_pair_ordered1_14 -0.0402385    0.1308101   -0.308  0.75899
## shape_pair_ordered1_2 -0.0723407    0.1308101   -0.553  0.58144
```


## shape_pair_ordered1_3	0.1367738	0.1308101	1.046	0.29817
## shape_pair_ordered1_4	-0.0387931	0.1308101	-0.297	0.76739
## shape_pair_ordered1_5	0.0859303	0.1308101	0.657	0.51269
## shape_pair_ordered1_6	0.0114275	0.1308101	0.087	0.93055
## shape_pair_ordered1_7	-0.2163550	0.1308101	-1.654	0.10115
## shape_pair_ordered1_8	0.0453769	0.1308101	0.347	0.72937
## shape_pair_ordered1_9	0.1349791	0.1308101	1.032	0.30453
## shape_pair_ordered10_11	-0.0762775	0.1308101	-0.583	0.56108
## shape_pair_ordered10_12	0.1498584	0.1308101	1.146	0.25458
## shape_pair_ordered10_13	-0.1292234	0.1308101	-0.988	0.32551
## shape_pair_ordered10_14	0.1490714	0.1308101	1.140	0.25707
## shape_pair_ordered11_12	-0.1725568	0.1308101	-1.319	0.19002
## shape_pair_ordered11_13	0.0215212	0.1308101	0.165	0.86964
## shape_pair_ordered11_14	0.0305845	0.1308101	0.234	0.81559
## shape_pair_ordered12_13	-0.0040090	0.1308101	-0.031	0.97561
## shape_pair_ordered12_14	0.1159936	0.1308101	0.887	0.37727
## shape_pair_ordered13_14	0.0155912	0.1308101	0.119	0.90536
## shape_pair_ordered2_10	-0.0363804	0.1308101	-0.278	0.78148
## shape_pair_ordered2_11	0.1071230	0.1308101	0.819	0.41470
## shape_pair_ordered2_12	0.0549125	0.1308101	0.420	0.67551
## shape_pair_ordered2_13	0.0980395	0.1308101	0.749	0.45526
## shape_pair_ordered2_14	0.1161485	0.1308101	0.888	0.37663
## shape_pair_ordered2_3	-0.0459731	0.1308101	-0.351	0.72596
## shape_pair_ordered2_4	-0.1109154	0.1308101	-0.848	0.39843
## shape_pair_ordered2_5	0.0052144	0.1308101	0.040	0.96828
## shape_pair_ordered2_6	-0.1084834	0.1308101	-0.829	0.40882
## shape_pair_ordered2_7	-0.1536814	0.1308101	-1.175	0.24274
## shape_pair_ordered2_8	0.0263881	0.1308101	0.202	0.84052
## shape_pair_ordered2_9	0.0354589	0.1308101	0.271	0.78687
## shape_pair_ordered3_10	0.4565251	0.1308101	3.490	0.00071 ***
## shape_pair_ordered3_11	-0.0395316	0.1308101	-0.302	0.76310
## shape_pair_ordered3_12	0.2206175	0.1308101	1.687	0.09469 .
## shape_pair_ordered3_13	0.1410654	0.1308101	1.078	0.28335
## shape_pair_ordered3_14	0.1714035	0.1308101	1.310	0.19297
## shape_pair_ordered3_4	-0.1871759	0.1308101	-1.431	0.15546
## shape_pair_ordered3_5	-0.0633496	0.1308101	-0.484	0.62920
## shape_pair_ordered3_6	-0.0825370	0.1308101	-0.631	0.52945
## shape_pair_ordered3_7	-0.0509304	0.1308101	-0.389	0.69782
## shape_pair_ordered3_8	0.1745343	0.1308101	1.334	0.18503
## shape_pair_ordered3_9	-0.0100859	0.1308101	-0.077	0.93869
## shape_pair_ordered4_10	-0.0531587	0.1308101	-0.406	0.68530
## shape_pair_ordered4_11	0.0278755	0.1308101	0.213	0.83167
## shape_pair_ordered4_12	0.1197721	0.1308101	0.916	0.36199
## shape_pair_ordered4_13	0.1163233	0.1308101	0.889	0.37592
## shape_pair_ordered4_14	-0.0156680	0.1308101	-0.120	0.90489
## shape_pair_ordered4_5	-0.0706438	0.1308101	-0.540	0.59032
## shape_pair_ordered4_6	-0.1266822	0.1308101	-0.968	0.33507
## shape_pair_ordered4_7	-0.2222708	0.1308101	-1.699	0.09227 .
## shape_pair_ordered4_8	0.0984548	0.1308101	0.753	0.45336
## shape_pair_ordered4_9	0.1183908	0.1308101	0.905	0.36753
## shape_pair_ordered5_10	-0.1565470	0.1308101	-1.197	0.23413
## shape_pair_ordered5_11	0.0955970	0.1308101	0.731	0.46654
## shape_pair_ordered5_12	-0.1379655	0.1308101	-1.055	0.29401
## shape_pair_ordered5_13	-0.0419871	0.1308101	-0.321	0.74887

```
## shape_pair_ordered5_14  0.1509095  0.1308101  1.154  0.25129
## shape_pair_ordered5_6   -0.0111558  0.1308101  -0.085  0.93220
## shape_pair_ordered5_7   -0.1965797  0.1308101  -1.503  0.13592
## shape_pair_ordered5_8    0.1413327  0.1308101  1.080  0.28244
## shape_pair_ordered5_9    0.1807396  0.1308101  1.382  0.17003
## shape_pair_ordered6_10  -0.0317031  0.1308101  -0.242  0.80898
## shape_pair_ordered6_11  -0.0458448  0.1308101  -0.350  0.72670
## shape_pair_ordered6_12   0.0337831  0.1308101  0.258  0.79672
## shape_pair_ordered6_13   0.1380786  0.1308101  1.056  0.29361
## shape_pair_ordered6_14  -0.1060428  0.1308101  -0.811  0.41941
## shape_pair_ordered6_7    -0.0354671  0.1308101  -0.271  0.78683
## shape_pair_ordered6_8    -0.0155735  0.1308101  -0.119  0.90546
## shape_pair_ordered6_9     0.0702540  0.1308101  0.537  0.59237
## shape_pair_ordered7_10  -0.0990872  0.1308101  -0.757  0.45047
## shape_pair_ordered7_11   0.0808803  0.1308101  0.618  0.53773
## shape_pair_ordered7_12   0.0656543  0.1308101  0.502  0.61680
## shape_pair_ordered7_13  -0.0431268  0.1308101  -0.330  0.74230
## shape_pair_ordered7_14  -0.1433409  0.1308101  -1.096  0.27570
## shape_pair_ordered7_8    -0.0746202  0.1308101  -0.570  0.56961
## shape_pair_ordered7_9     0.0190746  0.1308101  0.146  0.88435
## shape_pair_ordered8_10   0.0711929  0.1308101  0.544  0.58744
## shape_pair_ordered8_11  -0.0335715  0.1308101  -0.257  0.79796
## shape_pair_ordered8_12  -0.1251884  0.1308101  -0.957  0.34077
## shape_pair_ordered8_13  -0.2517150  0.1308101  -1.924  0.05705
## shape_pair_ordered8_14  -0.1005920  0.1308101  -0.769  0.44364
## shape_pair_ordered8_9    -0.0065262  0.1308101  -0.050  0.96031
## shape_pair_ordered9_10  -0.0804191  0.1308101  -0.615  0.54004
## shape_pair_ordered9_11  -0.0927509  0.1308101  -0.709  0.47988
## shape_pair_ordered9_12  -0.2570267  0.1308101  -1.965  0.05210
## shape_pair_ordered9_13  -0.1505379  0.1308101  -1.151  0.25245
## shape_pair_ordered9_14   0.0336627  0.1308101  0.257  0.79742
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.1308 on 104 degrees of freedom
## Multiple R-squared:  0.6655, Adjusted R-squared:  0.3278
## F-statistic: 1.971 on 105 and 104 DF, p-value: 0.0003073
```

```
AIC(m2)
```

```
## [1] -191.8998
```

Mixed Effects Model

Here we're modeling 1/RT to be predicted by **graph**, LOC values, and random effects for each subject, and each shape pair.

I *think* this is the correct model to be using, but would appreciate a critical eye.

However, it doesn't appear that graph effects are significant here, with presumably a richer dataset allowing us to fit all repetitions of a shape pair per subject.

```
m3 <- lmer(r1.inv ~ graph + z.loc_val + (1 | subject) + (1 | shape_pair_ordered), df)
summary(m3)
```

```
## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
```

```

## Formula: rt.inv ~ graph + z.loc_val + (1 | subject) + (1 | shape_pair_ordered)
## Data: df
##
## REML criterion at convergence: 111249.3
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -1.721  -0.218   0.016   0.214  137.759
##
## Random effects:
## Groups             Name             Variance Std.Dev.
## shape_pair_ordered (Intercept) 0.003454 0.05877
## subject              (Intercept) 0.071356 0.26713
## Residual                                1.050584 1.02498
## Number of obs: 38431, groups: shape_pair_ordered, 105; subject, 50
##
## Fixed effects:
##              Estimate Std. Error      df t value Pr(>|t|)
## (Intercept)   1.871355   0.054249 48.808521  34.496 < 2e-16 ***
## graphrnglattce 0.066073   0.076280 47.704012   0.866  0.391
## z.loc_val      0.033908   0.007827 93.178768   4.332 3.72e-05 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) grphrn
## grphrnglttc -0.703
## z.loc_val    0.001  0.000

```

```

m4 <- lmer(rt.inv ~ graph * z.loc_val + (1 | subject) + (1 | shape_pair_ordered), df)
summary(m4)

```

```

## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula: rt.inv ~ graph * z.loc_val + (1 | subject) + (1 | shape_pair_ordered)
## Data: df
##
## REML criterion at convergence: 111256.5
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -1.722  -0.217   0.016   0.214  137.758
##
## Random effects:
## Groups             Name             Variance Std.Dev.
## shape_pair_ordered (Intercept) 0.003455 0.05878
## subject              (Intercept) 0.071358 0.26713
## Residual                                1.050610 1.02499
## Number of obs: 38431, groups: shape_pair_ordered, 105; subject, 50
##
## Fixed effects:
##              Estimate Std. Error      df t value Pr(>|t|)
## (Intercept)   1.871e+00  5.425e-02 4.881e+01  34.495 < 2e-16
## graphrnglattce 6.607e-02  7.628e-02 4.770e+01   0.866 0.390714
## z.loc_val      3.526e-02  9.434e-03 1.965e+02   3.738 0.000243

```

```
## graphringlattice:z.loc_val -2.704e-03  1.052e-02  3.833e+04  -0.257 0.797102
##
## (Intercept)                ***
## graphringlattice
## z.loc_val                  ***
## graphringlattice:z.loc_val
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) grphrn z.loc_v
## grphrnglttc -0.703
## z.loc_val    0.001  0.000
## grphrngl:._  0.000  0.000 -0.558
```

Figure 5 Exploration

Inter-pattern Distance versus Intra-pattern distance

```
df.full %>%
  filter(match) %>%
  filter(trial.correct) %>%
  group_by(subject, graph) %>%
  summarize(mean.rt.inv = mean(rt.inv)) %>%
  group_by(graph) %>%
  summarize(mean.rt.inv = mean(mean.rt.inv))
```

```
## `summarise()` has grouped output by 'subject'. You can override using the
## `.groups` argument.
```

```
## # A tibble: 2 x 2
##   graph      mean.rt.inv
##   <fct>          <dbl>
## 1 modular          1.98
## 2 ringlattice      2.05
```

```
df.full %>%
  filter(match) %>%
  group_by(graph) %>%
  summarize(mean.correct= mean(trial.correct))
```

```
## # A tibble: 2 x 2
##   graph      mean.correct
##   <fct>          <dbl>
## 1 modular          0.898
## 2 ringlattice      0.911
```

```
df.full %>%
  filter(!match) %>%
  filter(trial.correct) %>%
  group_by(graph) %>%
  summarize(mean.rt.inv = mean(rt.inv))
```

```
## # A tibble: 2 x 2
##   graph      mean.rt.inv
##   <fct>          <dbl>
```

```

## 1 modular          1.86
## 2 ringlattice      1.94

m5 <- lmer(rt.inv ~ graph*z.modular_loc_val + graph:z.lattice_loc_val + (1 | subject) + (1 | shape_pair_
summary(m5)

## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula: rt.inv ~ graph * z.modular_loc_val + graph:z.lattice_loc_val +
## (1 | subject) + (1 | shape_pair_ordered)
## Data: df
##
## REML criterion at convergence: 111269.7
##
## Scaled residuals:
##    Min      1Q  Median      3Q      Max
## -1.732 -0.217  0.016   0.214 137.753
##
## Random effects:
## Groups              Name              Variance Std.Dev.
## shape_pair_ordered (Intercept) 0.003521 0.05934
## subject              (Intercept) 0.071440 0.26728
## Residual              1.050617 1.02500
## Number of obs: 38431, groups: shape_pair_ordered, 105; subject, 50
##
## Fixed effects:
##
##              Estimate Std. Error      df t value
## (Intercept)      1.871e+00  5.429e-02  4.882e+01  34.472
## graphringlattice    6.605e-02  7.632e-02  4.769e+01   0.865
## z.modular_loc_val    2.886e-02  1.307e-02  1.994e+02   2.208
## graphringlattice:z.modular_loc_val -1.186e-02  1.457e-02  3.839e+04  -0.814
## graphmodular:z.lattice_loc_val    9.398e-03  1.301e-02  1.980e+02   0.723
## graphringlattice:z.lattice_loc_val  1.873e-02  1.300e-02  1.976e+02   1.441
##
##              Pr(>|t|)
## (Intercept)      <2e-16 ***
## graphringlattice    0.3911
## z.modular_loc_val    0.0284 *
## graphringlattice:z.modular_loc_val  0.4158
## graphmodular:z.lattice_loc_val    0.4708
## graphringlattice:z.lattice_loc_val  0.1512
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) grphrn z.md__ grphrnglttc:z.m__ grphm:.__
## grphrnglttc      -0.703
## z.mdlr_lc_v        0.001  0.000
## grphrnglttc:z.m__  0.000  0.000 -0.559
## grphmdl:.__        0.000  0.000 -0.684  0.383
## grphrnglttc:z.l__  0.000  0.000 -0.257 -0.382          0.378

df.full.match <-
df.full %>%
  filter(match) %>%
  filter(trial.correct)

```

```

m6 <- lmer(rt.inv ~ graph + (1 | subject) + (1 | shape_pair_ordered), df.full.match)
summary(m6)

## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula: rt.inv ~ graph + (1 | subject) + (1 | shape_pair_ordered)
## Data: df.full.match
##
## REML criterion at convergence: 103607.1
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.048  -0.254   0.004   0.250  127.795
##
## Random effects:
## Groups             Name             Variance Std.Dev.
## subject             (Intercept)  0.067288  0.25940
## shape_pair_ordered (Intercept)  0.003222  0.05676
## Residual                        0.888734  0.94273
## Number of obs: 38003, groups: subject, 50; shape_pair_ordered, 15
##
## Fixed effects:
##              Estimate Std. Error      df t value Pr(>|t|)
## (Intercept)    1.97500    0.05435 54.40298  36.339 <2e-16 ***
## graphrnglattce  0.07689    0.07401 47.97186   1.039  0.304
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr)
## grphrnglttc -0.681

df.full.correct <-
  df.full %>%
  filter(trial.correct)
m7 <- lmer(rt.inv ~ graph*match + (1 | subject) + (1 | shape_pair_ordered), df.full.correct)
summary(m7)

## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula: rt.inv ~ graph * match + (1 | subject) + (1 | shape_pair_ordered)
## Data: df.full.correct
##
## REML criterion at convergence: 215161.8
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -1.830  -0.238   0.008   0.232  143.280
##
## Random effects:
## Groups             Name             Variance Std.Dev.
## shape_pair_ordered (Intercept)  0.004328  0.06579
## subject             (Intercept)  0.067302  0.25943
## Residual                        0.972456  0.98613

```

```

## Number of obs: 76434, groups:  shape_pair_ordered, 120; subject, 50
##
## Fixed effects:
##
##              Estimate Std. Error      df t value Pr(>|t|)
## (Intercept)    1.872e+00  5.277e-02 5.030e+01  35.483 < 2e-16 ***
## graphringlattice    6.656e-02  7.407e-02 4.882e+01   0.899   0.373
## matchTRUE        1.035e-01  2.080e-02 8.200e+01   4.976 3.52e-06 ***
## graphringlattice:matchTRUE 9.329e-03  1.428e-02 7.634e+04   0.653   0.514
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) grphrn mtTRUE
## grphrnglttc -0.702
## matchTRUE   -0.084  0.033
## grphrn:TRUE  0.068 -0.096 -0.346

```