

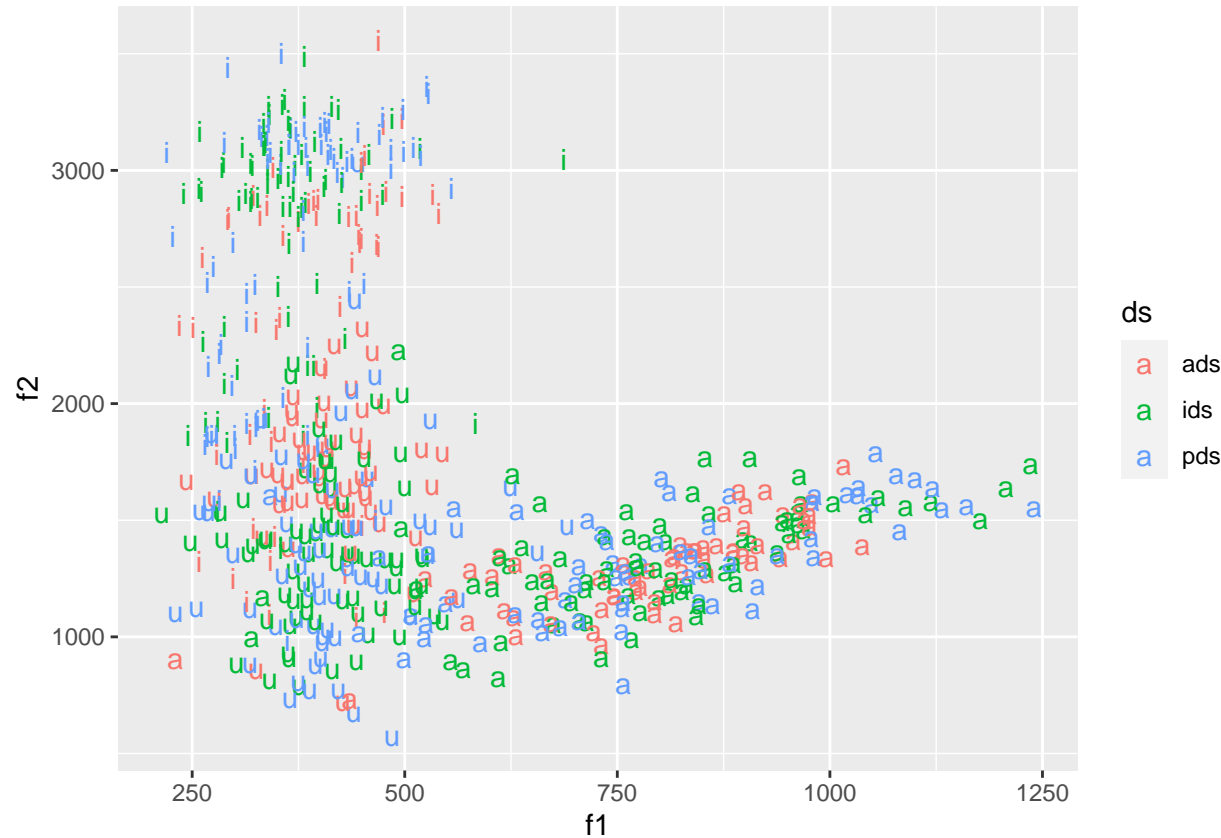
Supplementary materials: Positive Valence Contributes to Hyperarticulation in Maternal Speech to Infants and Puppies

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Initial data cleaning

Note there are no outliers:



```
##
## ball bead boot  box
## 105  209  201   99
```

Next we calculate vowel space for each mom and register.

```
## # A tibble: 3 x 2
##   ds      vss
##   <chr>   <dbl>
## 1 ads    596848.
## 2 ids    886587.
## 3 pds    842918.
```

Additional information on main analyses

Paired contrasts in vowel space size.

```
##
## Paired t-test
##
## data:  vs$vss[vs$ds == "pds"] and vs$vss[vs$ds == "ids"]
## t = -0.4956, df = 9, p-value = 0.6321
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
##  -242997.1  155658.9
## sample estimates:
## mean of the differences
##                -43669.08
```

```
##
## Cohen's d
##
## d estimate: 0.1559337 (negligible)
## 95 percent confidence interval:
##      lower      upper
## -0.7850539  1.0969214
```

```
##
## Paired t-test
##
## data:  vs$vss[vs$ds == "ids"] and vs$vss[vs$ds == "ads"]
## t = 2.4351, df = 9, p-value = 0.03767
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
##   20572.94 558905.61
## sample estimates:
## mean of the differences
##                289739.3
```

```
##
## Cohen's d
##
```

```
## d estimate: -1.121556 (large)
## 95 percent confidence interval:
##      lower      upper
## -2.1322879 -0.1108244

##
## Paired t-test
##
## data: vs$vss[vs$ds == "pds"] and vs$vss[vs$ds == "ads"]
## t = 1.9906, df = 9, p-value = 0.07773
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
##  -33574.0 525714.4
## sample estimates:
## mean of the differences
##      246070.2

##
## Cohen's d
##
## d estimate: -0.9882865 (large)
## 95 percent confidence interval:
##      lower      upper
## -1.983551077  0.006978064
```

Duration and pitch analyses

```
## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula: Vowel.Duration ~ ds * target + (1 + ds | motID)
## Data: dat
##
## REML criterion at convergence: -1202.6
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -3.5120 -0.5243 -0.0814  0.3421  5.3534
##
## Random effects:
## Groups Name Variance Std.Dev. Corr
## motID (Intercept) 0.0007915 0.02813
## dsids 0.0049585 0.07042 0.71
## dspds 0.0016792 0.04098 0.41 0.42
## Residual 0.0067777 0.08233
## Number of obs: 614, groups: motID, 10
##
## Fixed effects:
## Estimate Std. Error df t value Pr(>|t|)
## (Intercept) 0.195420 0.018230 32.341259 10.720 3.54e-12 ***
## dsids 0.071922 0.031004 23.340307 2.320 0.0294 *
## dspds 0.028832 0.025617 43.263609 1.126 0.2666
## targetbead 0.014766 0.019373 253.452590 0.762 0.4467
## targetboot -0.025244 0.019137 254.312247 -1.319 0.1883
```

```

## targetbox          -0.007169   0.022381  84.997033  -0.320   0.7495
## dsids:targetbead   0.018547   0.025530 371.954675   0.726   0.4680
## dspds:targetbead   0.008486   0.026016 382.948878   0.326   0.7445
## dsids:targetboot  -0.038286   0.025470 375.205621  -1.503   0.1336
## dspds:targetboot  -0.012732   0.026305 366.612811  -0.484   0.6287
## dsids:targetbox   -0.014377   0.031213 160.492273  -0.461   0.6457
## dspds:targetbox    0.008059   0.032281 166.800408   0.250   0.8032
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) dsids  dspds  trgtbd trgtbt trgtbx dsds:trgtbd dspds:trgtbd
## dsids          -0.183
## dspds          -0.426  0.457
## targetbead     -0.714  0.405  0.494
## targetboot     -0.723  0.410  0.500  0.677
## targetbox      -0.658  0.365  0.448  0.616  0.624
## dsds:trgtbd    0.528 -0.567 -0.370 -0.746 -0.501 -0.449
## dspds:trgtbd   0.521 -0.300 -0.711 -0.735 -0.494 -0.445  0.552
## dsds:trgtbt    0.529 -0.571 -0.371 -0.496 -0.738 -0.449  0.675      0.365
## dspds:trgtbt   0.512 -0.296 -0.717 -0.480 -0.715 -0.435  0.360      0.686
## dsds:trgtbx    0.446 -0.525 -0.312 -0.418 -0.423 -0.681  0.605      0.308
## dspds:trgtbx   0.436 -0.250 -0.640 -0.409 -0.414 -0.666  0.304      0.603
##              dsds:trgtbt dspds:trgtbt dsds:trgtbx
## dsids
## dspds
## targetbead
## targetboot
## targetbox
## dsds:trgtbd
## dspds:trgtbd
## dsds:trgtbt
## dspds:trgtbt  0.532
## dsds:trgtbx  0.609      0.303
## dspds:trgtbx  0.305      0.613      0.467

## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula: Pitch ~ ds * target + (1 + ds | motID)
## Data: dat
##
## REML criterion at convergence: 6746.6
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.3545 -0.5564 -0.1364  0.4836  8.8010
##
## Random effects:
## Groups   Name                Variance Std.Dev. Corr
## motID    (Intercept)         348.8    18.68
##          dsids              3636.3    60.30  -0.59
##          dspds              1946.0    44.11  -0.44  0.98
## Residual                    4369.5    66.10
## Number of obs: 606, groups:  motID, 10

```

```
##
## Fixed effects:
##      Estimate Std. Error    df t value Pr(>|t|)
## (Intercept)  200.1587    14.7500  42.3778  13.570  <2e-16 ***
## dsids        27.1051    26.6546   24.2858   1.017  0.3192
## dspds        46.6849    22.5136   31.7025   2.074  0.0463 *
## targetbead   -2.0223    16.1501  298.5380  -0.125  0.9004
## targetboot    11.7437    15.9741  294.1606   0.735  0.4628
## targetbox    -3.3188    18.8580  117.0155  -0.176  0.8606
## dsids:targetbead 16.4764    21.4548  503.9741   0.768  0.4429
## dspds:targetbead  0.1623    21.0737  336.0349   0.008  0.9939
## dsids:targetboot 25.0384    21.4710  504.4648   1.166  0.2441
## dspds:targetboot 11.1454    21.1429  288.3700   0.527  0.5985
## dsids:targetbox  10.9551    26.8979  343.5838   0.407  0.6841
## dspds:targetbox -13.5956    25.4359  111.5193  -0.535  0.5941
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##      (Intr) dsids  dspds  trgtbd trgtbt trgtbx dsds:trgtbd dspds:trgtbd
## dsids      -0.660
## dspds      -0.630  0.768
## targetbead -0.762  0.443  0.474
## targetboot -0.772  0.449  0.479  0.701
## targetbox  -0.710  0.423  0.428  0.644  0.651
## dsds:trgtbd  0.595 -0.582 -0.395 -0.772 -0.546 -0.511
## dspds:trgtbd 0.562 -0.349 -0.651 -0.748 -0.517 -0.466  0.596
## dsds:trgtbt  0.597 -0.586 -0.397 -0.541 -0.765 -0.512  0.704      0.419
## dspds:trgtbt 0.554 -0.351 -0.653 -0.505 -0.730 -0.456  0.418      0.690
## dsds:trgtbx  0.536 -0.544 -0.371 -0.485 -0.491 -0.750  0.641      0.387
## dspds:trgtbx 0.485 -0.331 -0.573 -0.441 -0.445 -0.689  0.387      0.602
##      dsds:trgtbt dspds:trgtbt dsds:trgtbx
## dsids
## dspds
## targetbead
## targetboot
## targetbox
## dsds:trgtbd
## dspds:trgtbd
## dsds:trgtbt
## dspds:trgtbt  0.583
## dsds:trgtbx  0.645      0.390
## dspds:trgtbx 0.389      0.605      0.586
```

F1 and F2 analyses

```
## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula: f1 ~ ds * target + (1 + ds | motID)
## Data: dat
##
## REML criterion at convergence: 7352.5
##
```

```

## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -4.9580 -0.5054 -0.0140  0.5005  3.6533
##
## Random effects:
##   Groups   Name      Variance Std.Dev.  Corr
##   motID    (Intercept) 1802     42.46
##           dsids       1336     36.55   -0.69
##           dspds       2056     45.35   -0.75  0.75
##   Residual      11086     105.29
## Number of obs: 611, groups:  motID, 10
##
## Fixed effects:
##              Estimate Std. Error      df t value Pr(>|t|)
## (Intercept)      716.49      25.49    40.48  28.106 < 2e-16 ***
## dsids             16.74      30.00    45.67   0.558   0.580
## dspds             38.90      32.50    49.79   1.197   0.237
## targetbead     -322.74      25.82   458.13 -12.498 < 2e-16 ***
## targetboot     -306.20      25.51   458.03 -12.005 < 2e-16 ***
## targetbox       146.59      31.04   264.54   4.723 3.78e-06 ***
## dsids:targetbead -38.79      33.04   424.12  -1.174   0.241
## dspds:targetbead -55.05      34.15   492.90  -1.612   0.108
## dsids:targetboot -12.69      32.94   417.43  -0.385   0.700
## dspds:targetboot -35.35      34.39   447.67  -1.028   0.305
## dsids:targetbox  -12.15      40.21   148.61  -0.302   0.763
## dspds:targetbox  -37.64      42.66   239.88  -0.882   0.378
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr) dsids  dspds  trgtbd trgtbt trgtbx dsds:trgtbd dspds:trgtbd
## dsids          -0.731
## dspds          -0.736  0.602
## targetbead     -0.707  0.579  0.550
## targetboot     -0.716  0.586  0.557  0.701
## targetbox      -0.666  0.535  0.516  0.652  0.659
## dsds:trgtbd    0.538 -0.758 -0.428 -0.767 -0.534 -0.489
## dspds:trgtbd   0.532 -0.446 -0.749 -0.753 -0.528 -0.489  0.584
## dsds:trgtbt    0.539 -0.762 -0.429 -0.528 -0.759 -0.489  0.683  0.404
## dspds:trgtbt   0.527 -0.445 -0.753 -0.516 -0.738 -0.483  0.402  0.701
## dsds:trgtbx    0.486 -0.694 -0.395 -0.476 -0.482 -0.734  0.613  0.369
## dspds:trgtbx   0.479 -0.404 -0.679 -0.469 -0.474 -0.720  0.364  0.625
##              dsds:trgtbt dspds:trgtbt dsds:trgtbx
## dsids
## dspds
## targetbead
## targetboot
## targetbox
## dsds:trgtbd
## dspds:trgtbd
## dsds:trgtbt
## dspds:trgtbt  0.569
## dsds:trgtbx  0.616      0.369
## dspds:trgtbx  0.364      0.632      0.552

```

```

## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula: f2 ~ ds * target + (1 + ds | motID)
## Data: dat
##
## REML criterion at convergence: 8883.4
##
## Scaled residuals:
## Min 1Q Median 3Q Max
## -4.3137 -0.4833 0.0961 0.6227 3.5400
##
## Random effects:
## Groups Name Variance Std.Dev. Corr
## motID (Intercept) 20427 142.9
## dsids 10077 100.4 -0.54
## dspds 26244 162.0 -0.19 0.93
## Residual 142792 377.9
## Number of obs: 611, groups: motID, 10
##
## Fixed effects:
## Estimate Std. Error df t value Pr(>|t|)
## (Intercept) 1199.73 88.90 39.31 13.495 2.44e-16 ***
## dsids 129.80 101.08 56.79 1.284 0.20434
## dspds 13.88 115.78 53.82 0.120 0.90504
## targetbead 1193.16 91.71 398.32 13.011 < 2e-16 ***
## targetboot 513.56 90.58 398.38 5.670 2.75e-08 ***
## targetbox 212.06 109.18 193.36 1.942 0.05355 .
## dsids:targetbead 255.72 116.74 391.69 2.191 0.02907 *
## dspds:targetbead 312.39 121.74 468.83 2.566 0.01060 *
## dsids:targetboot -462.22 116.27 384.69 -3.975 8.40e-05 ***
## dspds:targetboot -341.26 122.74 433.69 -2.780 0.00567 **
## dsids:targetbox -149.90 137.95 146.23 -1.087 0.27899
## dspds:targetbox 44.91 151.63 224.94 0.296 0.76735
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
## (Intr) dsids dspds trgtbd trgtbt trgtbx dsds:trgtbd dspds:trgtbd
## dsids -0.708
## dspds -0.603 0.641
## targetbead -0.714 0.600 0.539
## targetboot -0.723 0.607 0.546 0.696
## targetbox -0.669 0.549 0.500 0.643 0.650
## dsds:trgtbd 0.542 -0.777 -0.434 -0.768 -0.529 -0.481
## dspds:trgtbd 0.531 -0.474 -0.746 -0.747 -0.518 -0.476 0.591
## dsds:trgtbt 0.544 -0.780 -0.437 -0.524 -0.761 -0.481 0.673 0.407
## dspds:trgtbt 0.525 -0.477 -0.750 -0.505 -0.730 -0.468 0.406 0.697
## dsds:trgtbx 0.494 -0.697 -0.420 -0.475 -0.481 -0.744 0.599 0.386
## dspds:trgtbx 0.469 -0.437 -0.675 -0.451 -0.456 -0.703 0.369 0.619
## dsds:trgtbt dspds:trgtbt dsds:trgtbx
## dsids
## dspds
## targetbead
## targetboot

```

```
## targetbox
## dsds:trgtbd
## dspds:trgtbd
## dsds:trgtbt
## dspds:trgtbt 0.578
## dsds:trgtbx 0.601 0.390
## dspds:trgtbx 0.371 0.627 0.588
## optimizer (nloptwrap) convergence code: 0 (OK)
## boundary (singular) fit: see ?isSingular
```

Variability analyses

We first get standard deviation of f1 and f2, separating by mom and register.

We then fit separate regressions to each.

```
## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula: f1.sd ~ ds * target + (1 | motID)
## Data: sds
##
## REML criterion at convergence: 803.1
##
## Scaled residuals:
## Min 1Q Median 3Q Max
## -2.5964 -0.5157 0.0173 0.4615 3.2291
##
## Random effects:
## Groups Name Variance Std.Dev.
## motID (Intercept) 95.23 9.758
## Residual 1211.56 34.807
## Number of obs: 90, groups: motID, 10
##
## Fixed effects:
## Estimate Std. Error df t value Pr(>|t|)
## (Intercept) 112.08 17.95 77.95 6.245 2.07e-08 ***
## dsids -10.49 24.61 69.22 -0.426 0.671215
## dspds 37.71 24.61 69.22 1.532 0.130086
## targetbead -62.41 20.83 73.75 -2.997 0.003716 **
## targetboot -69.92 20.83 73.75 -3.357 0.001247 **
## targetbox -50.30 23.06 77.33 -2.181 0.032210 *
## dsids:targetbead 21.06 29.12 69.22 0.723 0.472112
## dspds:targetbead -15.93 29.12 69.22 -0.547 0.586190
## dsids:targetboot 19.44 29.12 69.22 0.667 0.506734
## dspds:targetboot -11.49 29.12 69.22 -0.394 0.694478
## dsids:targetbox 111.52 31.77 69.22 3.510 0.000794 ***
## dspds:targetbox 70.58 31.77 69.22 2.221 0.029616 *
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
## (Intr) dsids dspds trgtbd trgtbt trgtbx dsds:trgtbd dspds:trgtbd
## dsids -0.686
## dspds -0.686 0.500
```



```

## targetbead    -0.836  0.591  0.591
## targetboot    -0.836  0.591  0.591  0.721
## targetbox     -0.771  0.534  0.534  0.664  0.664
## dsds:trgtbd    0.580 -0.845 -0.423 -0.699 -0.499 -0.451
## dspds:trgtbd   0.580 -0.423 -0.845 -0.699 -0.499 -0.451  0.500
## dsds:trgtbt    0.580 -0.845 -0.423 -0.499 -0.699 -0.451  0.714      0.357
## dspds:trgtbt   0.580 -0.423 -0.845 -0.499 -0.699 -0.451  0.357      0.714
## dsds:trgtbx    0.531 -0.775 -0.387 -0.458 -0.458 -0.689  0.655      0.327
## dspds:trgtbx   0.531 -0.387 -0.775 -0.458 -0.458 -0.689  0.327      0.655
##              dsds:trgtbt dspds:trgtbt dsds:trgtbx
## dsids
## dspds
## targetbead
## targetboot
## targetbox
## dsds:trgtbd
## dspds:trgtbd
## dsds:trgtbt
## dspds:trgtbt  0.500
## dsds:trgtbx   0.655      0.327
## dspds:trgtbx  0.327      0.655      0.500

## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula: f2.sd ~ ds * target + (1 | motID)
## Data: sds
##
## REML criterion at convergence: 1019.6
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -2.7122 -0.5822 -0.0900  0.4701  3.3161
##
## Random effects:
##  Groups   Name                Variance Std.Dev.
## motID    (Intercept)            0      0.0
## Residual                  20638    143.7
## Number of obs: 90, groups: motID, 10
##
## Fixed effects:
##              Estimate Std. Error      df t value Pr(>|t|)
## (Intercept)      121.25      71.83    78.00   1.688  0.09541 .
## dsids              80.26     101.58    78.00   0.790  0.43189
## dspds              42.89     101.58    78.00   0.422  0.67400
## targetbead       488.77      84.99    78.00   5.751 1.65e-07 ***
## targetboot        65.34      84.99    78.00   0.769  0.44435
## targetbox       -45.07      92.73    78.00  -0.486  0.62831
## dsids:targetbead -396.78     120.20    78.00  -3.301  0.00145 **
## dspds:targetbead -212.95     120.20    78.00  -1.772  0.08035 .
## dsids:targetboot  -29.54     120.20    78.00  -0.246  0.80654
## dspds:targetboot   50.50     120.20    78.00   0.420  0.67553
## dsids:targetbox   31.85     131.14    78.00   0.243  0.80876
## dspds:targetbox   21.80     131.14    78.00   0.166  0.86843
## ---

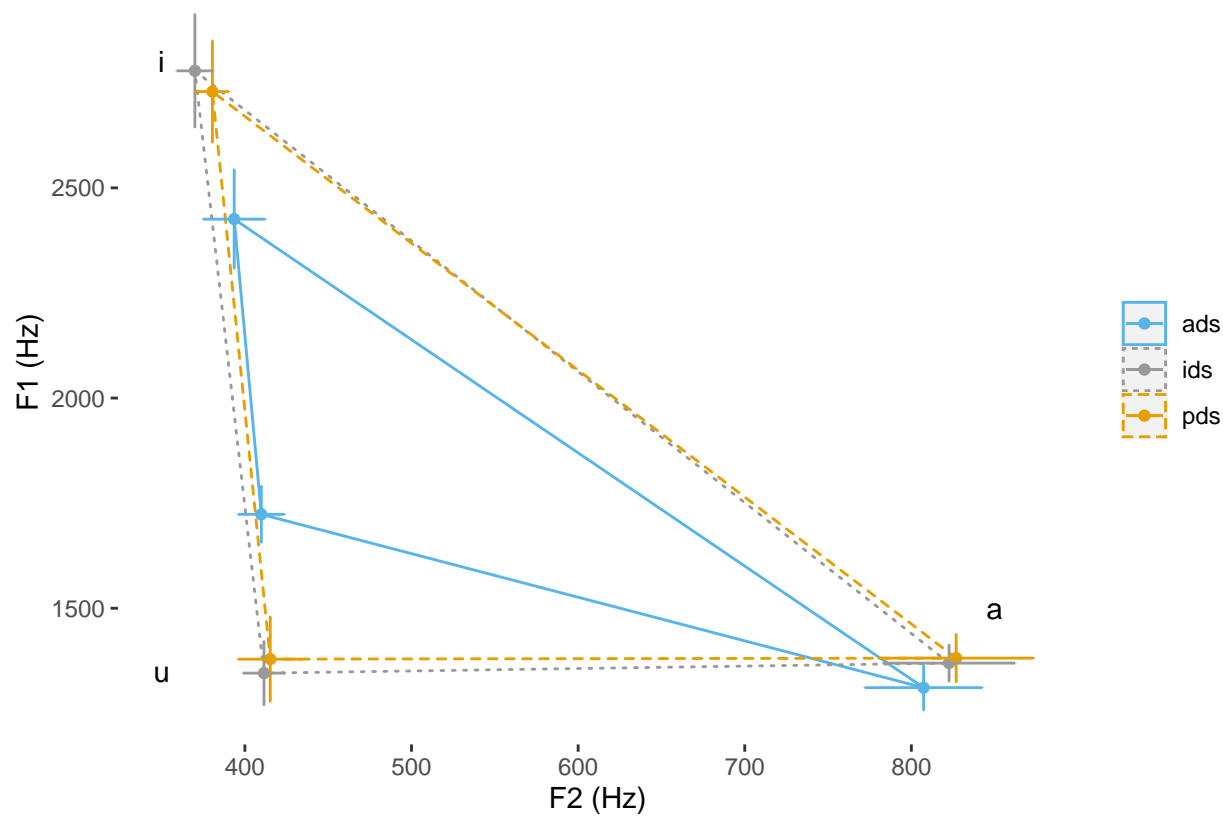
```

```

## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##          (Intr) dsids  dspds  trgtbd trgtbt trgtbx dsds:trgtbd dspds:trgtbd
## dsids          -0.707
## dspds          -0.707  0.500
## targetbead     -0.845  0.598  0.598
## targetboot     -0.845  0.598  0.598  0.714
## targetbox      -0.775  0.548  0.548  0.655  0.655
## dsds:trgtbd     0.598 -0.845 -0.423 -0.707 -0.505 -0.463
## dspds:trgtbd    0.598 -0.423 -0.845 -0.707 -0.505 -0.463  0.500
## dsds:trgtbt     0.598 -0.845 -0.423 -0.505 -0.707 -0.463  0.714      0.357
## dspds:trgtbt    0.598 -0.423 -0.845 -0.505 -0.707 -0.463  0.357      0.714
## dsds:trgtbx     0.548 -0.775 -0.387 -0.463 -0.463 -0.707  0.655      0.327
## dspds:trgtbx    0.548 -0.387 -0.775 -0.463 -0.463 -0.707  0.327      0.655
##          dsds:trgtbt dspds:trgtbt dsds:trgtbx
## dsids
## dspds
## targetbead
## targetboot
## targetbox
## dsds:trgtbd
## dspds:trgtbd
## dsds:trgtbt
## dspds:trgtbt  0.500
## dsds:trgtbx  0.655      0.327
## dspds:trgtbx  0.327      0.655      0.500
## optimizer (nloptwrap) convergence code: 0 (OK)
## boundary (singular) fit: see ?isSingular

```

Figure of the vowel space



pdf
2

pdf
2

