EyeDOC

Simo

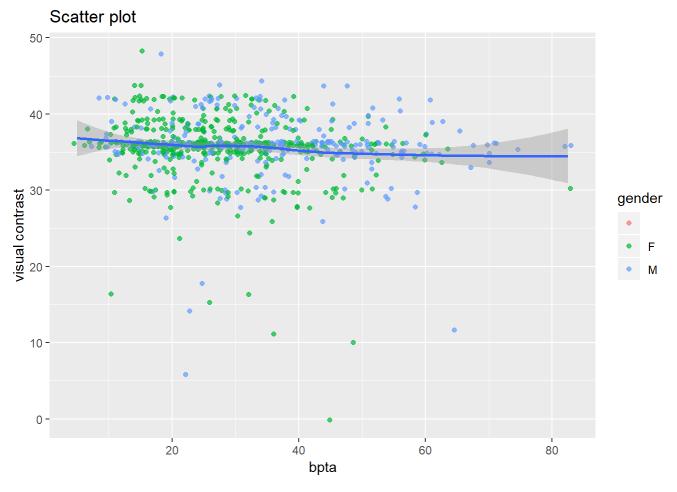
January 31, 2019

Scatter plot

##1 BPTA

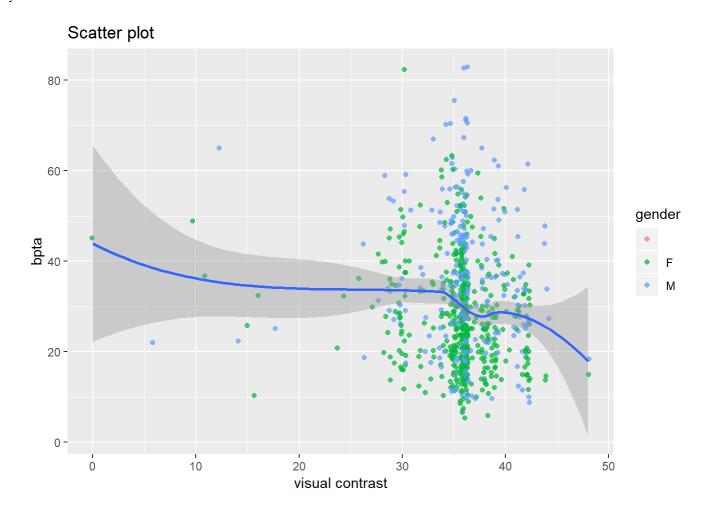
###1.1.1 BPTA & contrast sensitivity

```
\# `geom_smooth()` using method = 'loess' and formula 'y ~ x'
```



###1.1.2

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

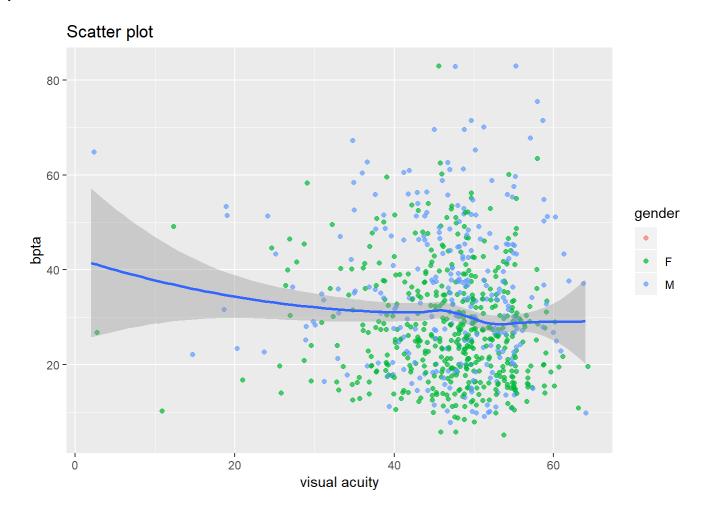


1.2 BPTA & Visual Acuity

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



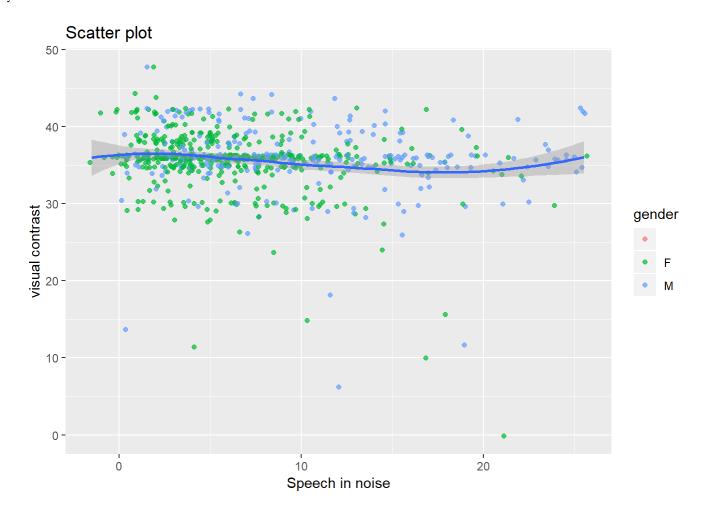
`geom_smooth()` using method = 'loess' and formula 'y \sim x'



2 SNR

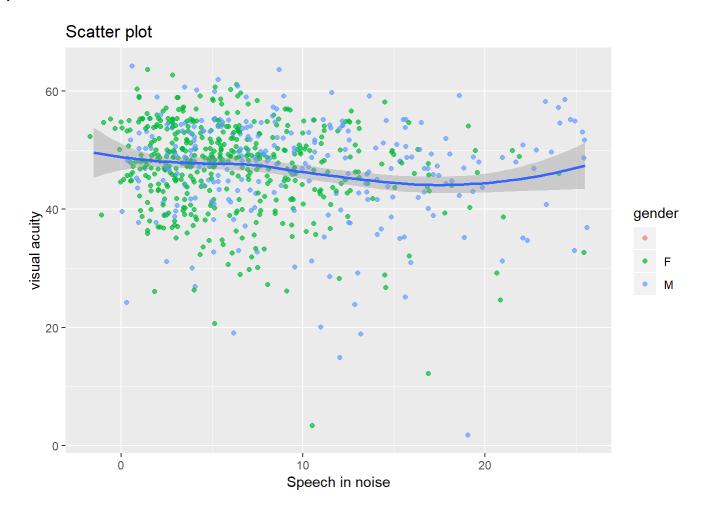
2.1 Signal to noise ratio and Visual contrast sensitivity

```
\# `geom_smooth()` using method = 'loess' and formula 'y ~ x'
```



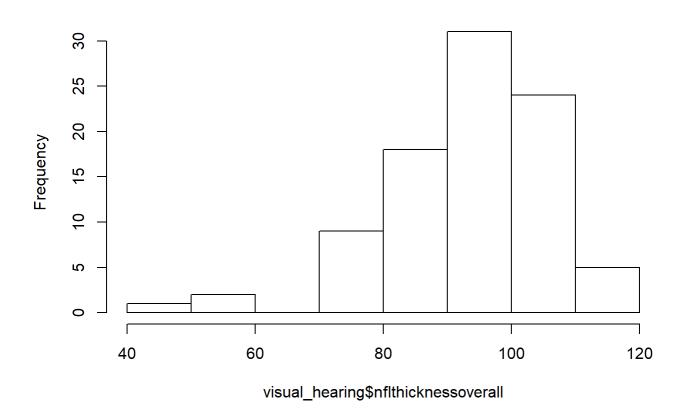
2.2 Signal to noise ratio and Visual acuity

```
## `geom_smooth()` using method = 'loess' and formula 'y \sim x'
```

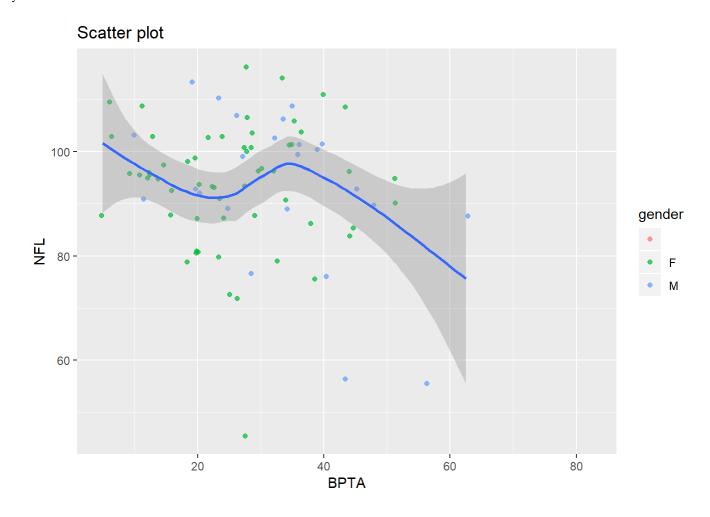


OCT PTA

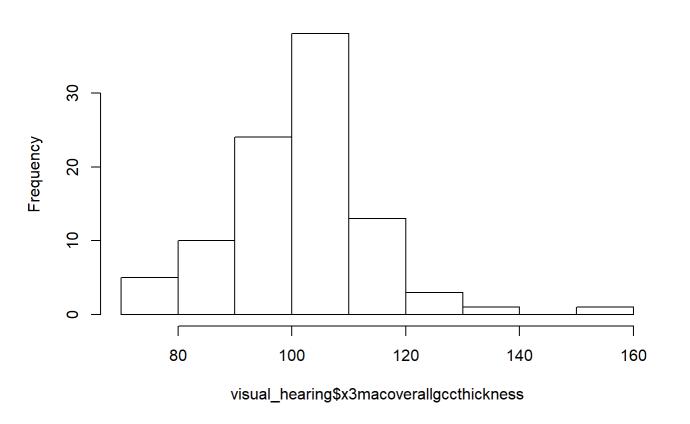
Histogram of visual_hearing\$nflthicknessoverall



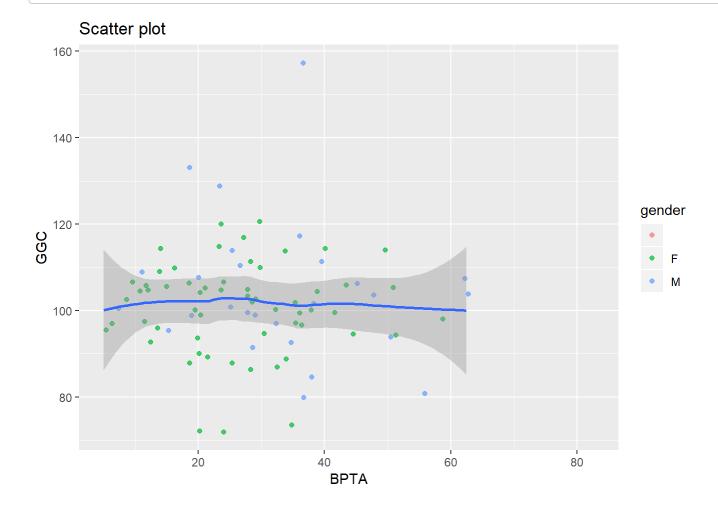
`geom_smooth()` using method = 'loess' and formula 'y \sim x'



Histogram of visual_hearing\$x3macoverallgccthickness

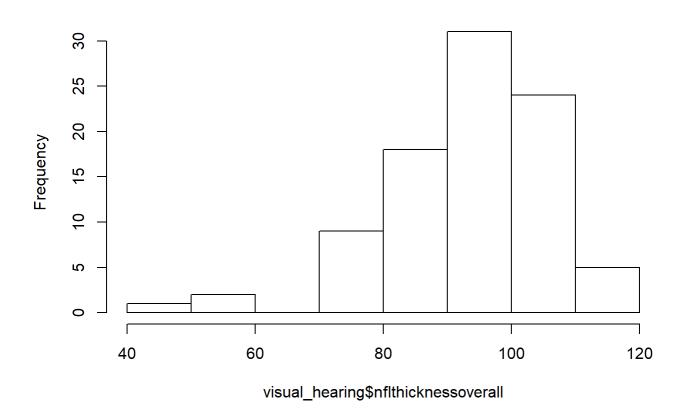


```
\# `geom_smooth()` using method = 'loess' and formula 'y ~ x'
```

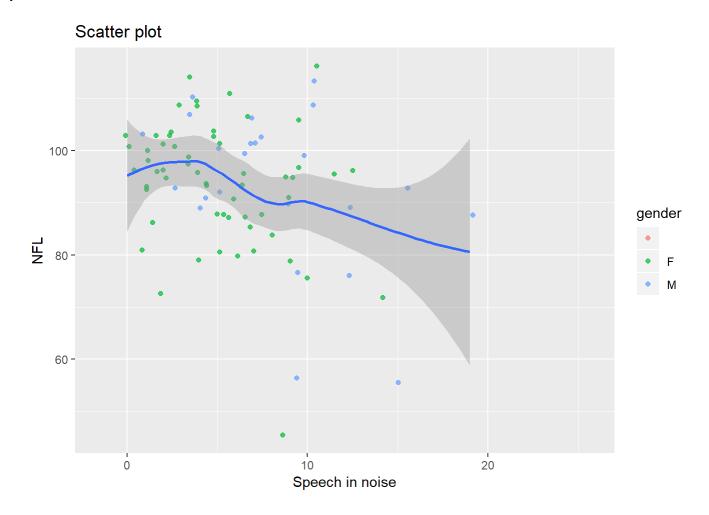


Speech in noise

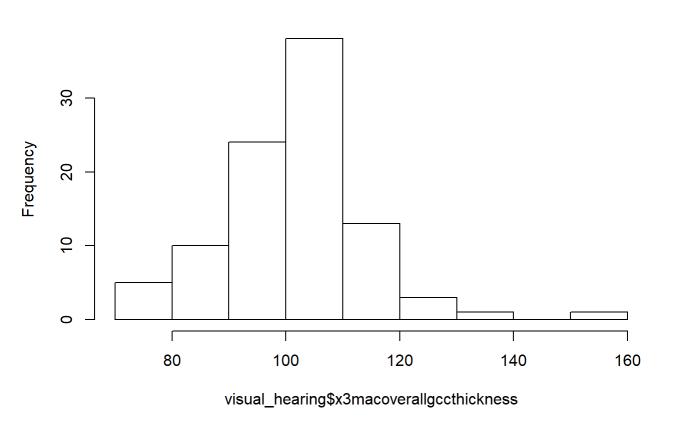
Histogram of visual_hearing\$nflthicknessoverall



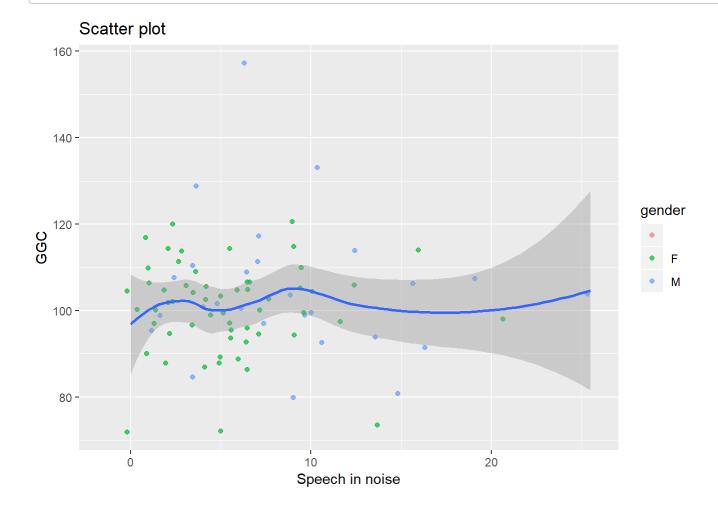
`geom_smooth()` using method = 'loess' and formula 'y \sim x'



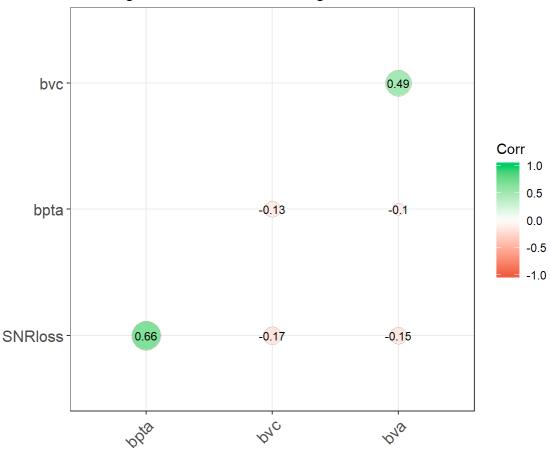
Histogram of visual_hearing\$x3macoverallgccthickness



`geom_smooth()` using method = 'loess' and formula 'y ~ x'



Correlogram of visaul and hearing functions



```
## function (x, y = NULL, use = "everything", method = c("pearson",
       "kendall", "spearman"))
##
## {
##
       na.method <- pmatch(use, c("all.obs", "complete.obs", "pairwise.complete.obs",
           "everything", "na.or.complete"))
##
       if (is.na(na.method))
##
           stop("invalid 'use' argument")
##
       method <- match.arg(method)</pre>
##
##
       if (is.data.frame(y))
##
           y <- as.matrix(y)
       if (is.data.frame(x))
##
           x <- as.matrix(x)
##
##
       if (!is.matrix(x) && is.null(y))
           stop("supply both 'x' and 'y' or a matrix-like 'x'")
##
       if (!(is.numeric(x) || is.logical(x)))
##
           stop("'x' must be numeric")
##
       stopifnot(is.atomic(x))
##
       if (!is.null(y)) {
##
##
           if (!(is.numeric(y) || is.logical(y)))
               stop("'y' must be numeric")
##
           stopifnot(is.atomic(y))
##
##
       Rank <- function(u) {</pre>
##
##
           if (length(u) == 0L)
##
##
           else if (is.matrix(u)) {
```

```
##
                if (nrow(u) > 1L)
                    apply(u, 2L, rank, na.last = "keep")
##
##
                else row(u)
            }
##
            else rank(u, na.last = "keep")
##
##
       if (method == "pearson")
##
            .Call(C_cor, x, y, na.method, FALSE)
##
       else if (na.method %in% c(2L, 5L)) {
##
##
            if (is.null(y)) {
                .Call(C cor, Rank(na.omit(x)), NULL, na.method, method ==
##
##
                     "kendall")
            }
##
##
            else {
                nas <- attr(na.omit(cbind(x, y)), "na.action")</pre>
##
                dropNA <- function(x, nas) {</pre>
##
                    if (length(nas)) {
##
##
                      if (is.matrix(x))
                         x[-nas, , drop = FALSE]
##
##
                       else x[-nas]
##
                    else x
##
                }
##
                .Call(C_cor, Rank(dropNA(x, nas)), Rank(dropNA(y,
##
##
                    nas)), na.method, method == "kendall")
            }
##
##
       else if (na.method != 3L) {
##
           x \leftarrow Rank(x)
##
##
            if (!is.null(y))
                y \leftarrow Rank(y)
##
            .Call(C_cor, x, y, na.method, method == "kendall")
##
       }
##
       else {
##
##
            if (is.null(y)) {
##
                ncy <- ncx <- ncol(x)
                if (ncx == 0)
##
                    stop("'x' is empty")
##
                r <- matrix(0, nrow = ncx, ncol = ncy)
##
##
                for (i in seq_len(ncx)) {
                    for (j in seq_len(i)) {
##
                      x2 < -x[, i]
##
                      y2 <- x[, j]
##
##
                       ok <- complete.cases(x2, y2)
##
                      x2 <- rank(x2[ok])
                       y2 <- rank(y2[ok])
##
##
                       r[i, j] \leftarrow if (any(ok))
                         .Call(C_cor, x2, y2, 1L, method == "kendall")
##
                       else NA
##
##
                     }
##
                r \leftarrow r + t(r) - diag(diag(r))
##
##
                rownames(r) <- colnames(x)
##
                colnames(r) <- colnames(x)</pre>
```

```
##
                r
            }
##
            else {
##
##
                if (length(x) == 0L | length(y) == 0L)
                     stop("both 'x' and 'y' must be non-empty")
##
                matrix_result <- is.matrix(x) || is.matrix(y)</pre>
##
##
                if (!is.matrix(x))
##
                    x \leftarrow matrix(x, ncol = 1L)
                if (!is.matrix(y))
##
                     y <- matrix(y, ncol = 1L)
##
                ncx <- ncol(x)</pre>
##
##
                ncy <- ncol(y)</pre>
                r <- matrix(0, nrow = ncx, ncol = ncy)
##
                for (i in seq_len(ncx)) {
##
                    for (j in seq_len(ncy)) {
\#\#
                       x2 <- x[, i]
##
                       y2 <- y[, j]
##
                       ok <- complete.cases(x2, y2)
##
                       x2 <- rank(x2[ok])
##
                       y2 < - rank(y2[ok])
##
##
                       r[i, j] \leftarrow if (any(ok))
##
                         .Call(C_cor, x2, y2, 1L, method == "kendall")
                       else NA
##
##
                     }
##
                rownames(r) <- colnames(x)</pre>
##
##
                colnames(r) <- colnames(y)</pre>
                if (matrix_result)
##
##
                     r
##
                else drop(r)
            }
##
##
## }
## <bytecode: 0x00000001aebfa18>
## <environment: namespace:stats>
```