

Supplement C

Outcome measures of the *in vivo* experiments

The following code describes the calculations that were performed to compute the absolute and relative changes in anal pressure of the *in vivo* experiments.

MATLAB code

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% Written in MATLAB R2018b
% Edited in MATLAB R2020a

function [s_in] = calcOutcome(s,int_names,fs,tx)
%calcOutcome Calculates desired outcomes from selected windows and data
%   Input/output variables:
%   - s: structure array containing information of all selected files
%   - int_names: interval names of interest
%   - fs: sampling frequency
%   - tx: time of which the inhib pressure slope is calculated, in seconds
%   - s_in: modified structure array containing information of all selected
files

s_in = s; % work with copy of input array
tx(tx==0) = 1; % if tx=0, this value will later cause an error (cannot divide
by 0). So if input tx equals 0, tx will be changed to 1.
slope_fieldname = sprintf('inhib_slope_for_%usec', tx); % fieldname of slope
that includes the amount of seconds it is calculated from

for iRow = 1:numel(s_in) % loop over rows, i.e. selected files
    pressure = s(iRow).pres_adj; % copy values, but with a simpler name

    %% start/end point index definition
    %---(beware this is the index in the pressure vector, not the
    %corresponding time value in seconds) ---%

    %baseline_avg
    start_base = s_in(iRow).baseline(1) * fs;
    end_base = s_in(iRow).baseline(2) * fs;

    %stimulation_avg
    start_stim = round(s_in(iRow).stimulation(1) * fs);

    if isnan(s_in(iRow).inhibition) % in case of absent inhibition
        end_stim = round(s_in(iRow).stimulation(2) * fs); % average
stimulation pressure must be calculated from whole stim interval

    else % in case of present inhibition
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        end_stim = round(s_in(iRow).inhibition(1) * fs); % average
stimulation pressure must be calculated from interval start stim to start
inhib

%inhibition_avg
    start_inhib = round(s_in(iRow).inhibition(1) * fs);
    end_inhib = round(s_in(iRow).inhibition(2) * fs);

%relative pressure drop
    a = max(pressure(start_stim : end_stim)) -
mean(pressure(start_base : end_base));
    b = pressure(end_inhib) - mean(pressure(start_base : end_base));

%pressure_drop_slope
    start_slope = round(start_inhib);
    end_slope = round((s_in(iRow).inhibition(1) + tx) * fs);

%Zoom in on data and set start- and endpoint of onset
    pressure_window = pressure((start_inhib-4*fs):(start_inhib+4*fs));
% select a smaller window of the data for faster analysis of onset, start at
stimulation and end at inhibition + 4 sec's
    pressure_mean = movmean(pressure_window,500); %
create a movingmean vector of the pressure data with a window of 500 samples
    changepts = findchangepts(pressure_mean,'Statistic','linear',
'MaxNumChanges',20,'MinDistance',50); % find 20 changing points in slope and
average with a minimal distance of 50 samples inbetween
    changepts = changepts + (start_inhib-4*fs)-1;
% add the start_simulation to rewrite from window to full data set
    start_onset = dsearchn(changepts,start_inhib); %
find the point in changepts closest to the start_inhib value
    end_onset = dsearchn(changepts,
(find(pressure_window==max(pressure_window))+(start_inhib-4*fs)-1)); % find
the point in changepts closest to the maximum value
    mean_baseline = mean(pressure(start_base:end_base)); %
take the mean pressure of the baseline phase
    end

%% calculation of outcome values
    s_in(iRow).baseline_avg = mean(pressure(start_base : end_base));
    s_in(iRow).stimulation_avg = mean(pressure(start_stim : end_stim));
    s_in(iRow).stimulation_startvalue = pressure(start_stim);
    s_in(iRow).stimulation_peak = max(pressure(start_stim : end_stim)); %
this does include the bias by fluctuating pressure values (even after low pass
filter is applied)

    if isnan(s_in(iRow).inhibition) % in case of absent inhibition
        s_in(iRow).inhibition_avg = NaN;
        s_in(iRow).inhibition_startvalue = NaN;
        s_in(iRow).inhibition_endvalue = NaN;
        s_in(iRow).rel_pressure_drop = NaN;
        s_in(iRow).(slope_fieldname) = NaN;
        s_in(iRow).onset_height = NaN;
        s_in(iRow).onset_increase = NaN;
    else % in case of present inhibition

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        s_in(iRow).inhibition_avg = mean(pressure(start_inhib :
end_inhib));
        s_in(iRow).inhibition_startvalue = pressure(start_inhib);
        s_in(iRow).inhibition_endvalue = pressure(end_inhib);
        %
        s_in(iRow).rel_pressure_drop = (s_in(iRow).inhibition_avg -
s_in(iRow).stimulation_avg) ...
        %
        / (s_in(iRow).stimulation_avg -
s_in(iRow).baseline_avg ) * 100;
        s_in(iRow).rel_pressure_drop = ( b - a ) / a * 100;
        s_in(iRow).(slope_fieldname) = (pressure(end_slope)-
pressure(start_slope)) / tx;

        if end_onset-start_onset > 0          % in case of present onset
            if (pressure(changepts(end_onset))-mean_baseline)-
(pressure(changepts(start_onset))-mean_baseline)> 0
                s_in(iRow).onset_height = (pressure(changepts(end_onset))-
mean_baseline)-(pressure(changepts(start_onset))-mean_baseline); %remove
baseline pressure and calculate the difference between start and end onset
                s_in(iRow).onset_increase =
(pressure(changepts(end_onset))-
mean_baseline)/(pressure(changepts(start_onset))-mean_baseline); % remove
baseline pressure and calculate the increase percentage of the pressure
            else
                s_in(iRow).onset_height = NaN;
                s_in(iRow).onset_increase = NaN;
            end
        else          % in case of absent onset
            s_in(iRow).onset_height = NaN;
            s_in(iRow).onset_increase = NaN;
        end

    end

end

end

```