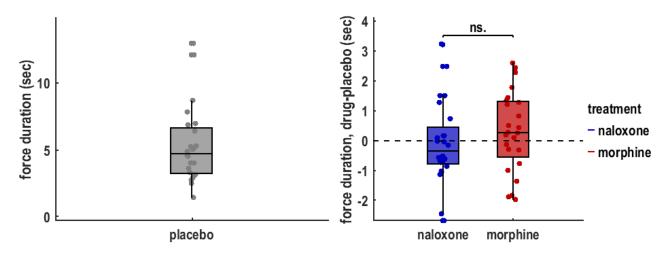
```
%% motiscan opioid 2level gripAccu
% 1. Analysis of major measurements
% define test from dataset
taskname = 'gripAccu';
%% parameters
% define metrics
varname = 'effortDuration';
% ftransform = @ (x) normalize(x, 'zscore');
ftransform = @identity;
fname = @nanmean;
varlegend = 'force duration, drug-placebo (sec)';
% define statistical procedure
statisticalTest = {'opioid contrast'};
% display options
% - plot type
plotType = {'jitter','boxplot'};
%% aggregate data
% - prepare data
nbin = [ ntrt , nsub ];
Y = nan(nbin); % conditional means by treatment
Y2 = nan(nbin); % mean difference to placebo
S = nan(nbin);
% - condition data
for isub = 1:nsub % subject loop
    % select
        tab = data{isub}.(taskname).table;
        selection = (tab.task==taskname);
        tab = tab(selection,:);
    % variables
        y = tab.(varname);
        y = ftransform(y);
        trt = tab.treatment;
        trt = removecats(trt, '0');
        trt = reordercats(trt,treatmentList);
        session = tab.sessionNumber;
    % stats
        [~,subtrt] = ismember(unique(trt),treatmentList);
        ysub = splitapply(fname,y,double(trt));
        sess = splitapply(fname, session, double(trt));
    % store
        Y(subtrt, isub) = ysub;
        S(subtrt,isub) = sess;
end
% - concatenate data
Y2 = Y-Y(2,:); % normalize by the subject mean under placebo
% Y2 = Y - nanmean(Y,1) + nanmean(nanmean(Y)); % normalize by the subject mean
Y = reshape(Y,nsub*ntrt,1);
Y2 = reshape(Y2,nsub*ntrt,1);
T = repmat(nominal(treatmentList'), nsub, 1);
T = reordercats(T, treatmentList);
S = reshape(S,nsub*ntrt,1);
SUB = repmat([1:nsub],ntrt,1);
```

```
Estimate
                          SE
                                    tStat
                                               pValue
            -0.33376
                       0.40571
                                   -0.82267
                                               0.41357
naloxone
           -0.22906
                       0.38373
                                   -0.59693
                                               0.55254
placebo
morphine
           0.071425
                       0.39464
                                   0.18099
                                               0.85692
session
            0.11951
                        0.1613
                                    0.74092
                                               0.46129
```

```
% statistical inference
p = nan(1,numel(statisticalTest));
score = nan(1,numel(statisticalTest));
contrast = [-1 \ 0 \ 1];
for i = 1:numel(statisticalTest)
    try
        contrast = [contrast , zeros(size(contrast,1),ncofactor) ];
        [p(i),score(i),d] = coefTest(stat,contrast)
    end
end
%% display
fig = figure;
clear g;
alpha=0.7;
g(1,1) = gramm('x',T,'y',Y,'color',T,'subset',(T=='placebo'));
g(1,1).set color options('map',vertcat(col{[2]}),'lightness',100);
q(1,1).set order options('x',treatmentList,'color',treatmentList);
g(1,1).geom_jitter('height',0.01);
g(1,1).stat boxplot('width',0.9);
g(1,1).set_names('x','','y','force duration (sec)','color','treatment');
q(1,1).axe property('YLim',[min(Y) max(Y)] + [-0.2 0.2]*mean(Y));
g(1,1).axe property('XLim',[-2 4]);
g(1,2) = gramm('x',T,'y',Y2,'color',T,'subset',(T~='placebo'));
g(1,2).set_color_options('map',vertcat(col{[1 3]}),'lightness',100);
g(1,2).set order options('x',treatmentList([1 3]),'color',treatmentList([1 3]));
g(1,2).geom jitter('height',0.01);
g(1,2).stat boxplot('width',0.9);
g(1,2).set_names('x','','y','force duration, drug-placebo (sec)','color','treatment');
g(1,2).axe_property('YLim',[min(Y2) max(Y2)] + [-0.2 0.2]*mean(Y2));
q(1,2).axe property('XLim',[0 3]);
g.draw;
```

ordering given as values ordering given as values

```
axes(g(1,2).facet_axes_handles);
```



```
l = plot(g(1,2).facet_axes_handles.XLim,[0 0],'--k');
fig.Position = [ 100 100 900 400];
```

