

ANNEXE II

ANALYSE DÉTAILLÉE DE L'EXPÉRIENCE

The analysis of the experiment data is also accessible online: <https://github.com/NormandErwan/HandheldVesadAnalysis/blob/master/Handheld%20VESAD%20Analysis.ipynb>.

0.1 1. Data preparation

Configuration :

```
In [1]: # Imports
import numpy as np
import pandas as pd
from pandas.api.types import CategoricalDtype
import itertools

import seaborn as sns
import matplotlib as mpl
import matplotlib.pyplot as plt
import matplotlib.ticker as ticker
import matplotlib.patches as patches

from scipy import stats
import statsmodels.api as sm
from statsmodels.formula.api import ols
from statsmodels.stats.multicomp import (MultiComparison, pairwise_tukeyhsd)
from statsmodels.stats.multitest import multipletests
from statsmodels.stats.libqsturng import psturng

from ast import literal_eval
from os import listdir
from os.path import join

from IPython.html.services.config import ConfigManager

C:\Users\Erwan\Miniconda\envs\master-thesis\lib\site-packages\IPython\html.py:14: ShimWarning: The
  "`IPython.html.widgets` has moved to `ipywidgets`.", ShimWarning)

In [2]: # Notebook configuration
%matplotlib inline

# Ruler
```

```

ip = get_ipython()
cm = ConfigManager(parent=ip)
cm.update('notebook', {"ruler_column": [80]})

# Style and size of the figures
subplotsize = (5, 4)
legend_fontsize = sns.plotting_context('notebook')['axes.labelsize']
legend_title_fontsize = legend_fontsize + 1
sns.set(context='notebook', style='whitegrid', font_scale=1.25,
        rc={'legend.fontsize': legend_fontsize, 'savefig.dpi': 300,
            'savefig.bbox': 'tight'})

# Render the plots with this language
#language = 'English'
language = 'Français'

```

Data are loaded in the following variables: - The ranks of the participants from the post-questionary: ranks - Trials of the participants: raw_trials - Trials averaged so that each participants ends up with a single observation (Tip 9 of Dragicevic, 2016): trials > Dragicevic, P. (2016). Fair statistical communication in HCI. In Modern Statistical Methods for HCI (pp. 291-330). Springer, Cham.

```

In [3]: trials = pd.read_csv('participant_trials.csv')
        raw_trials = pd.read_csv('participant_trials.csv')
        ranks = pd.read_csv('participant_ranks.csv')

```

Creates independent variables lists (ivs) and dependent variables lists (ranks_dvs, trials_dvs) to make easier to use algorithms and to plot figures.

```

In [4]: # IVs
        if language == 'Français':
            iv_labels = ['IHM', 'Taille du texte', 'Distance', 'Groupe']
        else:
            iv_labels = ['Technique', 'Text Size', 'Distance', 'Ordering']

ivs = ['technique', 'text_size', 'distance', 'ordering']
ivs = pd.DataFrame(columns=ivs, index=['label', 'categorical', 'palette'])

# Shortcuts
technique = ivs['technique']
text_size = ivs['text_size']
distance = ivs['distance']
ordering = ivs['ordering']

# Setup categories and palettes
palette = sns.color_palette('Set2', 8)
for iv_id, iv_label in zip(ivs, iv_labels):
    iv_categories = trials.drop_duplicates(iv_id)\
                        .sort_values([iv_id + '_id'])[iv_id]
    iv_categorical = pd.Categorical(iv_categories, iv_categories, ordered=True)
    ivs[iv_id] = [iv_label, iv_categorical, palette]

```

```

technique['palette'] = [palette[1], palette[0], palette[2]]
text_size['palette'] = sns.light_palette(palette[6], 3)[1:3]
distance['palette'] = sns.light_palette(palette[4], 3)[1:3]
ordering['palette'] = [palette[1], palette[0], palette[2]]

```

In [5]: # Trials DVs

```

if language == 'Français':
    trials_dv_labels = ['Temps de complétion (s)', 'Sélections',
                        'Temps en sélection', 'Distance 3D en sélection',
                        'Distance en sélection', 'Désélections', 'Erreurs',
                        'Disques classés', 'Défilements', 'Temps de défilement',
                        'Distance 3D de défilement', 'Distance de défilement',
                        'Zooms', 'Temps de zoom', 'Distance 3D de zoom',
                        'Distance de zoom', 'Mouvements tête-téléphone',
                        'Distance relative tête-téléphone']
else:
    trials_dv_labels = ['Task Completion Time (s)', 'Selections',
                        'Selection Time', 'Selection Distance',
                        'Selection Distance on Grid', 'Deselections', 'Errors',
                        'Items Classified', 'Pans', 'Pan Time', 'Pan Distance',
                        'Pan Distance on Grid', 'Zooms', 'Zoom Time',
                        'Zoom Distance', 'Zoom Distance on Grid',
                        'Phone-Head Motion', 'Signed Phone-Head Motion']

trials_dvs = trials.loc[:, 'total_time':'signed_head_phone_distance'].columns
trials_dvs = pd.Series(data=trials_dv_labels, index=trials_dvs)

# Shortcuts
participant_id = 'Participant'
total_time = trials_dvs['total_time']

```

In [6]: # Ranks DVs

```

if language == 'Français':
    ranks_dv_labels = ['Facile à comprendre', 'Mentalement facile à utiliser',
                        'Physiquement facile à utiliser', 'Rapidité',
                        'Performance', 'Frustration', 'Préférence']
else:
    ranks_dv_labels = ['Easy to Understand', 'Mentally Easy to Use',
                        'Physically Easy to Use', 'Subjective Speed',
                        'Subjective Performance', 'Frustration', 'Preference']

ranks_dv_scales = [pd.Categorical(list(range(1, 6)), list(
    range(1, 6)), ordered=True)] * len(ranks_dv_labels)

RdYlBu = sns.color_palette('RdYlBu', 5)
ranks_dv_palettes = [sns.color_palette('RdYlBu', 5)] * len(ranks_dv_labels)

ranks_dvs = ranks.loc[:, 'easy_understand':'preference'].columns
ranks_dvs = pd.DataFrame(data=[ranks_dv_labels, ranks_dv_scales,
                                ranks_dv_palettes],
                           columns=ranks_dvs, index=['label', 'scale', 'palette'])

```

```

ranks_dvs.at['scale', 'preference'] = pd.Categorical(
    list(range(1, 4)), list(range(1, 4)), ordered=True)
ranks_dvs.at['palette', 'preference'] = [RdYlBu[4], RdYlBu[2], RdYlBu[0]]

```

Clean the data:

```

In [7]: # Setup columns of trials and ranks
columns = []
for column in trials.columns:
    if (column in ivs.columns):
        columns.append(ivs[column]['label'])
    elif (column in trials_dvs.index):
        columns.append(trials_dvs[column])
    else:
        columns.append(column)
columns[0] = participant_id

trials.columns = columns
ranks.columns = [participant_id, ordering['label'], technique['label']] \
    + ranks_dvs.loc['label', :].tolist()

# Setup translated column values of trials and ranks
for iv_id in ivs:
    iv = ivs[iv_id]
    trials[iv['label']] = trials[iv['label']].astype(iv['categorical'])
    raw_trials[iv_id] = raw_trials[iv_id].astype(iv['categorical'])

for iv_id in ['technique', 'ordering']:
    iv = ivs[iv_id]
    ranks[iv['label']] = ranks[iv['label']].astype(iv['categorical'])

if language == 'Français':
    technique['categorical'].categories = ['Téléphone', 'VESAD tactile', 'VESAD']
    text_size['categorical'].categories = ['Grand', 'Petit']
    distance['categorical'].categories = ['Proche', 'Loin']
    ordering['categorical'].categories = ['Groupe 1', 'Groupe 2', 'Groupe 3']
else:
    ordering['categorical'].categories = ['Group 1', 'Group 2', 'Group 3']

for iv_id in ivs:
    iv = ivs[iv_id]
    trials[iv['label']].cat.categories = iv['categorical'].categories

for iv_id in ['technique', 'ordering']:
    iv = ivs[iv_id]
    ranks[iv['label']].cat.categories = iv['categorical'].categories

In [8]: # Some participants are non valid or don't have complete measures
invapars = [0, 4]
ranks = ranks[~ranks[participant_id].isin(invapars)].reset_index(drop=True)

```

```

trials = trials[~trials[participant_id].isin(invapars)].reset_index(drop=True)
raw_trials = raw_trials[~raw_trials['participant_id'].isin(invapars)
                        ].reset_index(drop=True)

```

```

In [9]: # Some participants have wrong head phone mesures
for dist in ['absolute_head_phone_distance', 'signed_head_phone_distance']:
    trials.loc[trials[trials_dvs[dist]] == 0, trials_dvs[dist]] = np.nan
    raw_trials.loc[raw_trials[dist] == 0, dist] = np.nan

```

```

In [10]: # Eval the arrays in some dvs
def eval_if_str(data):
    return literal_eval(data) if isinstance(data, str) else data

trials['grid_config'] = trials['grid_config'].apply(eval_if_str)

```

```

In [11]: # Average trials to have one observation per participant
trials = trials.groupby([participant_id] + ivs.loc['label', :].tolist(),
                        observed=True).mean().reset_index()
raw_trials = raw_trials.groupby(['participant_id'] + ivs.columns.tolist(),
                                observed=True).mean().reset_index()

```

Utilities:

```

In [12]: # Figure labels in the selected language
labels = pd.Series()

if language == 'Français':
    labels['category'] = 'Catégorie'
    labels['count'] = 'Nombre total'
    labels['distance'] = 'Distance moyenne (m)'
    labels['dv'] = 'VD'
    labels['iv'] = 'VI'
    labels['iv_value'] = 'Valeur VI'
    labels['mean_difference'] = 'Différence des moyennes'
    labels['mean_difference_percentage'] = 'Différence des moyennes (%)'
    labels['mean_rank'] = 'Note moyenne'
    labels['preferences'] = ['Premier', 'Deuxième', 'Troisième']
    labels['p_value'] = 'Valeur p'
    labels['question'] = 'Question'
    labels['rank'] = 'Note'
    labels['time'] = 'Temps moyen (s)'
    labels['t_statistic'] = 'Statistique T'
    labels['votes'] = 'Participants'
else:
    labels['category'] = 'Category'
    labels['count'] = 'Count'
    labels['distance'] = 'Mean Distance (m)'
    labels['dv'] = 'DV'
    labels['iv'] = 'IV'
    labels['iv_value'] = 'IV Value'
    labels['mean_difference'] = 'Mean Difference'

```

```

labels['mean_difference_percentage'] = 'Mean Difference Percentage'
labels['mean_rank'] = 'Mean Rank'
labels['preferences'] = ['First', 'Second', 'Third']
labels['p_value'] = 'p-value'
labels['question'] = 'Question'
labels['rank'] = 'Rank'
labels['time'] = 'Mean Time (s)'
labels['t_statistic'] = 'T statistic'
labels['votes'] = 'Participants'

In [13]: def mean_ci(x, which=95, n_boot=1000):
        """Returns the confidence interval of the mean"""
        x2 = [i for i in x if not np.isnan(i)]
        boots = sns.algorithms.bootstrap(x2, n_boot=n_boot)
        return sns.utils.ci(boots, which=which)

In [14]: def print_mean_ci(x, ci_which=95):
        """Returns a string containing the mean with the CI of x"""
        ci1, ci2 = mean_ci(x, which=ci_which)
        return '{:.2f} [{:.2f}, {:.2f}].format(np.mean(x), ci1, ci2)

In [15]: def exp_mean(x):
        return np.exp(np.mean(x))

In [16]: def print_exp_mean_ci(x, ci_which=95):
        ci1, ci2 = np.exp(mean_ci(x, which=ci_which))
        return '{:.2f} [{:.2f}, {:.2f}].format(exp_mean(x), ci1, ci2)

In [17]: def geometric_mean(x):
        return exp_mean(np.log(x))

In [18]: def print_geo_mean_ci(x, ci_which=95):
        ci1, ci2 = np.exp(mean_ci(np.log(x), which=ci_which))
        return '{:.2f} [{:.2f}, {:.2f}].format(geometric_mean(x), ci1, ci2)

In [19]: def mean_difference(a, b):
        """Returns the mean difference value and percentage between a and b"""
        mean_diff = np.mean(a) - np.mean(b)
        mean_diff_percentage = mean_diff / np.mean(b) * 100
        return (mean_diff, mean_diff_percentage)

In [20]: def p_values_correction(data, alpha=0.05, correction_method='fdr_bh'):
        if correction_method != None:
            reject, p_values_corrected, a1, a2 = \
                multipletests(data[labels['p_value']].tolist(), alpha=alpha,
                             method=correction_method)
            data[labels['p_value']] = p_values_corrected

In [21]: def subplots(nsubplots, ncols_max=2, subplotsize=subplotsize, *plt_args):
        ncols = min(ncols_max, nsubplots)
        nrows = ((nsubplots - 1) // ncols) + 1
        fig, axs = plt.subplots(nrows=nrows, ncols=ncols, \

```

```

figsize=(subplotsize[0]*ncols, subplotsize[1]*nrows),
*plt_args)

if nrows == 1 and ncols == 1:
    axs = [axs]
elif nrows >= 2 and ncols >= 2:
    axs = [ax for ax_row in axs for ax in ax_row]

for ax in axs[::-1][0:len(axs) - nsubplots]:
    fig.delaxes(ax)

return (fig, axs)

In [22]: def fix_legend_fontsize(ax_legend, fontsize=legend_title_fontsize):
plt.setp(ax_legend.get_title(), fontsize=fontsize)

In [23]: def config_legend(ax, iv_id, fontsize=legend_title_fontsize):
legend = ax.legend(title=ivs[iv_id]['label'], frameon=True,
                    loc='center left', bbox_to_anchor=(1, 0.5))
fix_legend_fontsize(legend)
return legend

```

0.2 2. Participant ranks

Some functions for the analysis:

```

In [24]: def get_ranks_count(iv_index, dv_index):
iv, dv = ivs[iv_index], ranks_dvs[dv_index]

ranks_counts_index = pd.MultiIndex.from_product([iv['categorical'],
                                                dv['scale']],
                                                names=[iv['label'],
                                                labels['rank']])

# Zero counts by default and gets the counts
default_counts = pd.Series(0, index=ranks_counts_index)
ranks_counts = ranks.groupby([iv['label'], dv['label']]).size()

# Merge and remove duplicated defaults
ranks_counts = pd.concat([ranks_counts, default_counts])
ranks_counts = ranks_counts[~ranks_counts.index.duplicated(keep='first')]

ranks_counts.sort_index(inplace=True)
ranks_counts.index = ranks_counts_index # Restore the index
return ranks_counts

In [25]: def cumulated_barplot(data, palette, **args):
for row_id, row in data.iloc[::-1].iterrows():
    sns.barplot(y=data.columns, x=row, label=row_id,
                color=palette[row_id-1], orient='h', **args)

In [26]: def plot_ranks_distributions(iv_id, dv_ids):
iv = ivs[iv_id]

```

```

fig, axs = subplots(len(dv_ids))
for dv_id, ax in zip(dv_ids, axs):
    dv = ranks_dvs[dv_id]

    cumulated_ranks_count = get_ranks_count(iv_id, dv_id).unstack(level=0)\
        .cumsum()
    cumulated_barplot(cumulated_ranks_count, palette=dv['palette'], ax=ax)
    ax.set(xlabel=labels['votes'], xlim=(0, cumulated_ranks_count.max()[0]))
    ax.xaxis.set_major_locator(ticker.MultipleLocator(2)) # Fix the axis ticks

    ax_handles, ax_labels = ax.get_legend_handles_labels()
    legend = ax.legend(ax_handles[:-1], ax_labels[:-1], frameon=True,
                       loc='lower center', bbox_to_anchor=(0.5, 1),
                       mode=None, ncol=len(dv['scale']), title=dv['label'],
                       fontsize=legend_fontsize-2)
    fix_legend_fontsize(legend)

fig.tight_layout(h_pad=1) # Add padding to avoid legend and labels overlap
return (fig, axs)

```

```

In [27]: def plot_ranks(iv_id, dv_ids, estimator=np.mean):
    iv = ivs[iv_id]

    fig, axs = subplots(len(dv_ids))
    for dv_id, ax in zip(dv_ids, axs):
        dv = ranks_dvs[dv_id]

        sns.barplot(x=iv['label'], y=dv['label'], palette=iv['palette'],
                    data=ranks, ax=ax, estimator=estimator)
        ax.set(ylim=(0, dv['scale'][-1]))
        ax.yaxis.set_major_locator(ticker.MultipleLocator(1)) # Fix the axis ticks

    return (fig, axs)

```

```

In [28]: def rank_samples(iv_id, dv_id):
    """Returns the list of ranks (DV) for each IV value"""
    samples = []
    iv = ivs[iv_id]
    for iv_value in iv['categorical']:
        dv_label = ranks_dvs[dv_id]['label']
        sample = ranks[ranks[iv['label']] == iv_value][dv_label]\
            .reset_index(drop=True)
        samples.append(sample)
    return samples

```

```

In [29]: def test_ranks(iv_id, dv_ids, **args):
    results = []
    iv = ivs[iv_id]

    for dv_id in dv_ids:

```



```

        dv = ranks_dvs[dv_id]
        samples = rank_samples(iv_id, dv_id)
        H, p = stats.kruskal(*samples)
        results.append([iv['label'], dv['label'], H, p])

results = pd.DataFrame(results, columns=[labels['iv'], labels['dv'],
                                       'Kruskal-Wallis H',
                                       labels['p_value']])

p_values_correction(results, **args)
return results

In [30]: def test_pairwise_ranks(iv_id, dv_ids, **args):
        iv = ivs[iv_id]
        iv_category_ids = range(len(iv['categorical']))

        results = []
        for dv_id in dv_ids:
            dv = ranks_dvs[dv_id]
            samples = rank_samples(iv_id, dv_id)
            sample_pairs = itertools.combinations(iv_category_ids, 2)
            for id1, id2 in sample_pairs:
                U, p = stats.mannwhitneyu(samples[id1], samples[id2])
                mean_diff, mean_diff_per = mean_difference(samples[id1], samples[id2])
                results.append([iv['label'], iv['categorical'][id1],
                              iv['categorical'][id2], dv['label'],
                              mean_diff, mean_diff_per, U, p])

        columns = [labels['iv'], labels['iv_value'] + ' 1',
                   labels['iv_value'] + ' 2', labels['dv'],
                   labels['mean_difference'], labels['mean_difference_percentage'],
                   'Mann-Whitney U', labels['p_value']]
        results = pd.DataFrame(results, columns=columns)
        p_values_correction(results, **args)
        return results

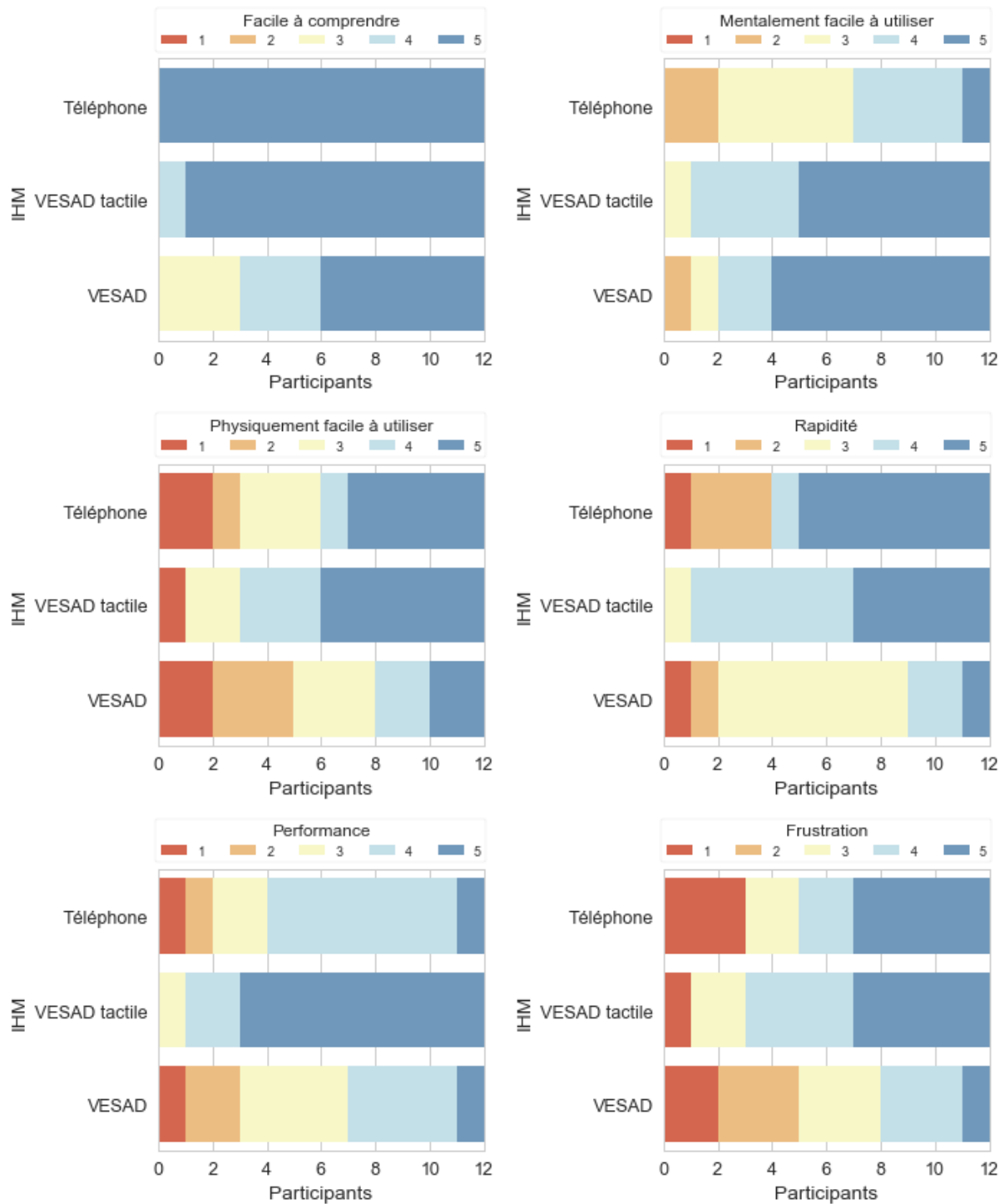
```

We display the rank and preference distributions:

```

In [31]: (fig, axs) = plot_ranks_distributions('technique', ranks_dvs.columns[0:-1])
        fig.savefig('ranks_distributions.png')

```

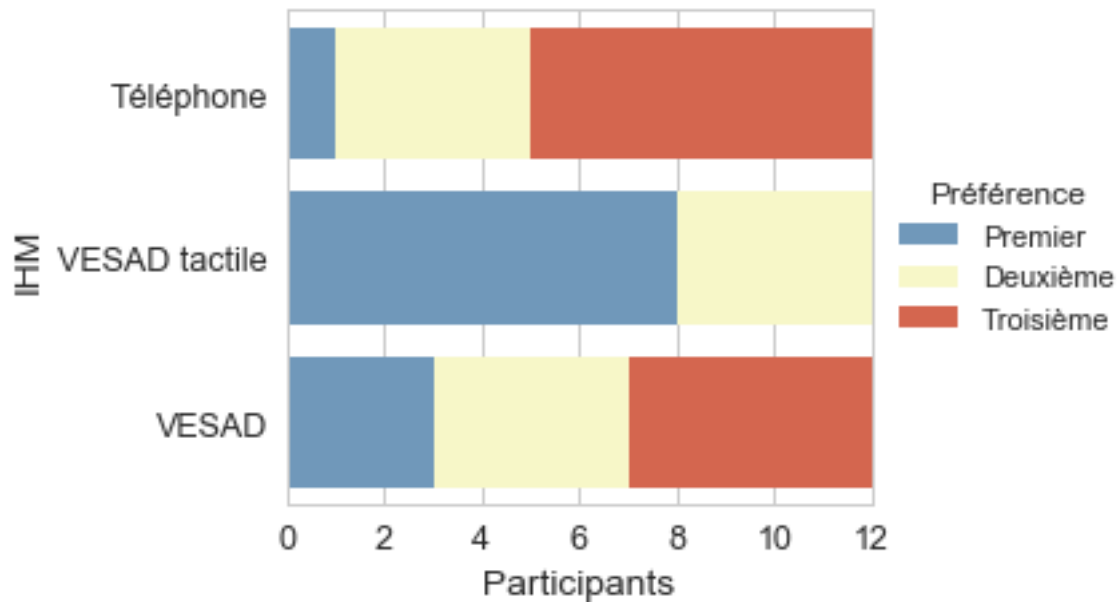


```
In [32]: (fig, axs) = plot_ranks_distributions('technique', ['preference'])
```

```
ax_handles, ax_labels = axs[0].get_legend_handles_labels()
legend = axs[0].legend(ax_handles[::-1], labels['preferences'],
                      loc='center left', bbox_to_anchor=(1, 0.5),
                      title=ranks_dvs['preference']['label'])
```

```
fix_legend_fontsize(legend)

fig.savefig('preferences_distribution.png')
```



We use the Kruskal-Wallis test (Benjamini–Hochberg correction) on each question to check if there is statistical significant differences between the ranks among TECHNIQUE:

```
In [33]: test_ranks('technique', ranks_dvs, correction_method='fdr_bh')
```

```
Out[33]:
```

	VI	VD	Kruskal-Wallis H	Valeur p
0	IHM	Facile à comprendre	11.001420	0.007497
1	IHM	Mentalement facile à utiliser	10.905742	0.007497
2	IHM	Physiquement facile à utiliser	4.148121	0.125674
3	IHM	Rapidité	7.130846	0.039599
4	IHM	Performance	13.784269	0.006303
5	IHM	Frustration	4.500057	0.122962
6	IHM	Préférence	12.638889	0.006303

All the questions, except Physically Easy to Use and Frustration, are statistically significant: Easy to Understand ($p = 0.007$), Mentally Easy to Use ($p = 0.007$), Subjective Speed ($p = 0.04$), Subjective Performance ($p = 0.006$), Preference ($p = 0.006$).

We use then pairwise Mann-Whitney tests (Benjamini–Hochberg correction) for the significant questions above:

```
In [34]: test_pairwise_ranks('technique', ['easy_understand', 'mentally_easy_use',
      'could_go_fast', 'subjective_performance',
      'preference'], correction_method='fdr_bh')
```

```
Out[34]:
```

	VI	Valeur VI 1	Valeur VI 2	VD \
0	IHM	Téléphone	VESAD tactile	Facile à comprendre
1	IHM	Téléphone	VESAD	Facile à comprendre

2	IHM	VESAD tactile	VESAD	Facile à comprendre
3	IHM	Téléphone	VESAD tactile	Mentalement facile à utiliser
4	IHM	Téléphone	VESAD	Mentalement facile à utiliser
5	IHM	VESAD tactile	VESAD	Mentalement facile à utiliser
6	IHM	Téléphone	VESAD tactile	Rapidité
7	IHM	Téléphone	VESAD	Rapidité
8	IHM	VESAD tactile	VESAD	Rapidité
9	IHM	Téléphone	VESAD tactile	Performance
10	IHM	Téléphone	VESAD	Performance
11	IHM	VESAD tactile	VESAD	Performance
12	IHM	Téléphone	VESAD tactile	Préférence
13	IHM	Téléphone	VESAD	Préférence
14	IHM	VESAD tactile	VESAD	Préférence

	Différence des moyennes	Différence des moyennes (%)	Mann-Whitney U \
0	0.083333	1.694915	66.0
1	0.750000	17.647059	36.0
2	0.666667	15.686275	40.5
3	-1.166667	-25.925926	23.0
4	-1.083333	-24.528302	28.5
5	0.083333	1.886792	69.5
6	-0.500000	-11.538462	70.5
7	0.750000	24.324324	48.5
8	1.250000	40.540541	21.0
9	-1.166667	-25.000000	22.5
10	0.333333	10.526316	57.0
11	1.500000	47.368421	17.5
12	1.166667	87.500000	16.0
13	0.333333	15.384615	56.0
14	-0.833333	-38.461538	32.0

	Valeur p
0	0.215732
1	0.008605
2	0.020977
3	0.005031
4	0.010151
5	0.475027
6	0.475027
7	0.126838
8	0.004655
9	0.004655
10	0.215732
11	0.003904
12	0.003904
13	0.215732
14	0.013011

We display the geometric mean and 95% CI of each question:

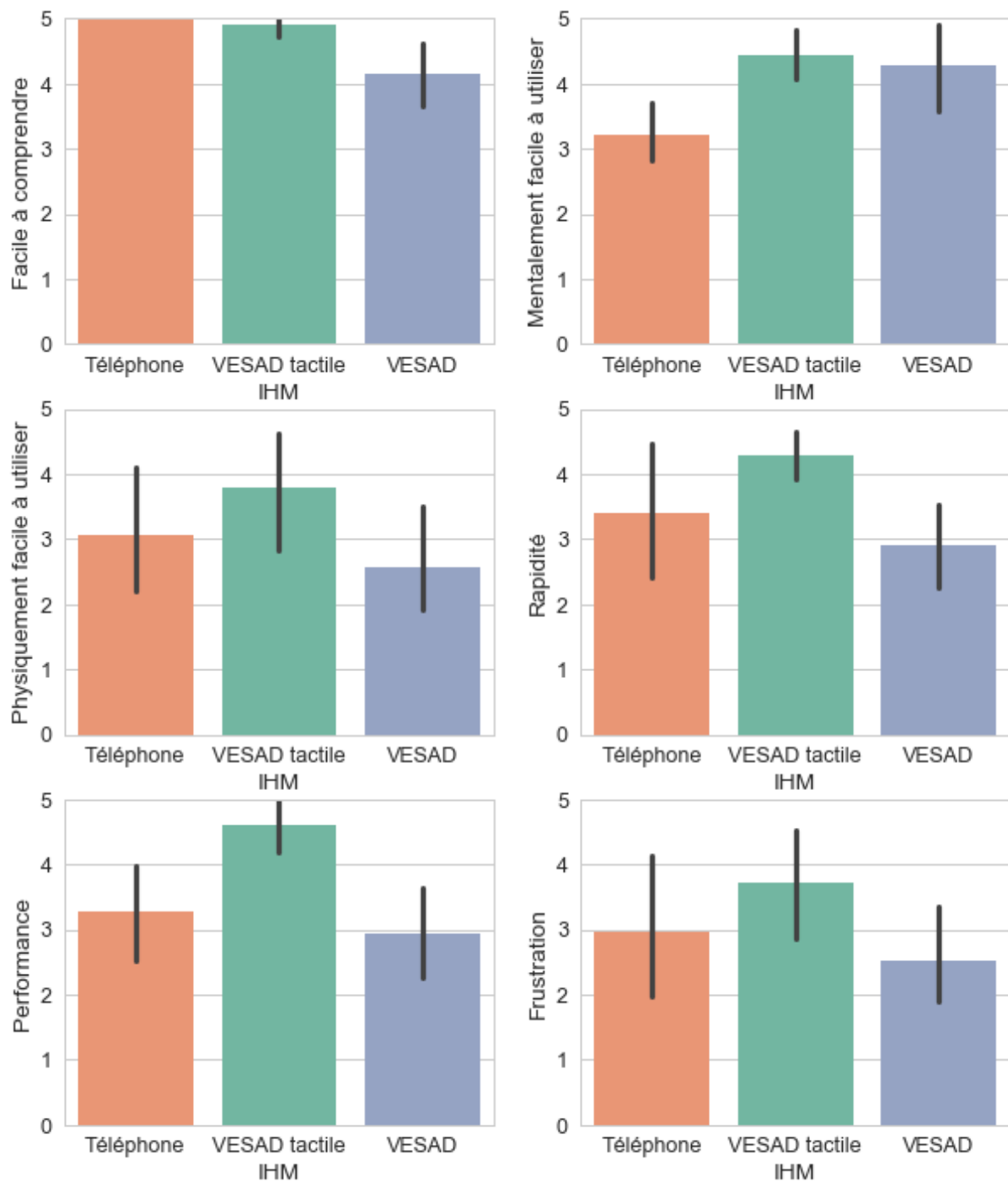
```
In [35]: ranks.groupby([ivs['technique']['label']]).aggregate(print_geo_mean_ci)\
```

```
.loc[:, ranks_dvs.loc['label', :]].transpose()
```

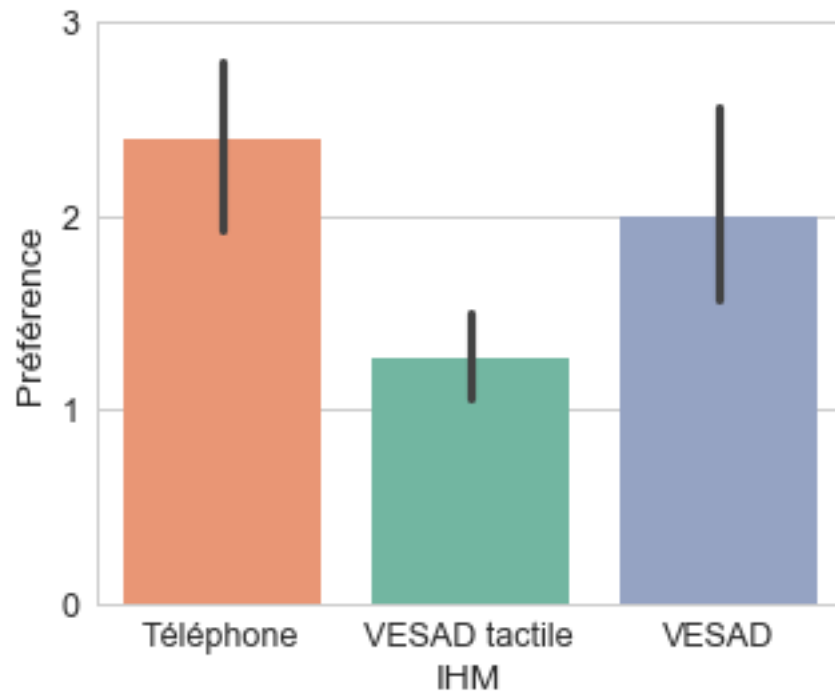
```
Out[35]: IHM          Téléphone          VESAD tactile \
Facile à comprendre      5.00 [5.00, 5.00]  4.91 [4.73, 5.00]
Mentalement facile à utiliser 3.22 [2.77, 3.68]  4.45 [4.03, 4.82]
Physiquement facile à utiliser 3.06 [2.16, 4.16]  3.80 [2.75, 4.62]
Rapidité                  3.41 [2.42, 4.55]  4.29 [3.94, 4.64]
Performance              3.27 [2.51, 3.98]  4.62 [4.16, 5.00]
Frustration              2.96 [2.02, 4.14]  3.73 [2.71, 4.56]
Préférence               2.39 [1.95, 2.80]  1.26 [1.06, 1.50]
```

```
IHM          VESAD
Facile à comprendre      4.16 [3.73, 4.70]
Mentalement facile à utiliser 4.28 [3.60, 4.91]
Physiquement facile à utiliser 2.58 [1.88, 3.42]
Rapidité                2.90 [2.29, 3.51]
Performance            2.94 [2.27, 3.61]
Frustration            2.53 [1.88, 3.31]
Préférence             1.99 [1.55, 2.47]
```

```
In [36]: (fig, axs) = plot_ranks('technique', ranks_dvs.columns[0:-1],
                                estimator=geometric_mean)
fig.savefig('ranks.png')
```



```
In [37]: (fig, axs) = plot_ranks('technique', ['preference'],
                                estimator=geometric_mean)
fig.savefig('preferences.png')
```



Overall significant results are:

- **Easy to Understand:** *PhoneOnly* is significantly better than *MidAirInArOut* ($p = 0.009$), and seems a little better than *PhoneInArOut*.
- **Physically Easy to Use:** There is no significant differences due to TECHNIQUE; they seem scored similar.
- **Mentally Easy to Use:** *PhoneOnly* is statistically and practically worst than *PhoneInArOut* ($p = 0.005$) and *MidAirInArOut* ($p = 0.01$).
- **Subjective Speed:** *PhoneInArOut* is significantly better than *MidAirInArOut* ($p = 0.05$).
- **Subjective Performance:** *PhoneInArOut* is statistically better than *PhoneOnly* ($p = 0.005$) and *MidAirInArOut* ($p = 0.004$).
- **Frustration:** There is no significant differences due to TECHNIQUE; they seem scored similar.
- **Preference:** *PhoneInArOut* is significantly preferred to *PhoneOnly* ($p = 0.004$) and *MidAirInArOut* ($p = 0.01$).

0.3 3. Participant trials

Some functions for the analysis:

```
In [38]: def trial_samples(iv_id, dv_id, data=trials):
          iv, dv = ivs[iv_id], trials_dvs[dv_id]
          return [data[data[iv['label']] == iv_value][dv]\
                  for iv_value in iv['categorical']]

In [39]: def trial_means(iv_ids, dv_ids, data=trials, aggregate=print_mean_ci):
          iv_labels = [ivs.at['label', iv_id] for iv_id in iv_ids]
          dv_labels = [trials_dvs[dv_id] for dv_id in dv_ids]
          return data[iv_labels + dv_labels].groupby(iv_labels)\
                  .aggregate(aggregate)
```

```

In [40]: def melt_trials(value_vars, var_name, value_name, data=trials):
    return pd.melt(data, id_vars=ivs.loc['label', :],
                   value_vars=value_vars, var_name=var_name,
                   value_name=value_name)

In [41]: def test_normality(iv_ids, dv_ids, data=trials, **args):
    results = []

    for dv_id in dv_ids:
        for iv_id in iv_ids:
            iv = ivs[iv_id]
            samples = trial_samples(iv_id, dv_id, data)
            for iv_value, sample in zip(iv['categorical'], samples):
                W, p = stats.shapiro(sample)
                results.append([iv['label'], iv_value, trials_dvs[dv_id], W, p])

    results = pd.DataFrame(results, columns=[labels['iv'], labels['iv_value'],
                                           labels['dv'], 'Shapiro W',
                                           labels['p_value']])

    p_values_correction(results, **args)
    return results

In [42]: def test_equal_variances(iv_ids, dv_ids, data=trials, levene_center='mean',
                                **args):
    results = []

    for dv_id in dv_ids:
        for iv_id in iv_ids:
            samples = trial_samples(iv_id, dv_id, data)
            W, p = stats.levene(*samples)
            results.append([ivs[iv_id]['label'], trials_dvs[dv_id], W, p])

    results = pd.DataFrame(results, columns=[labels['iv'], labels['dv'],
                                           'Levene W', labels['p_value']])

    p_values_correction(results, **args)
    return results

In [43]: def test_pairwise_trials(iv_id, dv_id, data=trials, log_data=False, **args):
    results = []
    iv, dv = ivs[iv_id], trials_dvs[dv_id]
    iv_category_ids = range(len(iv['categorical']))

    samples = trial_samples(iv_id, dv_id, data)
    sample_pairs = itertools.combinations(iv_category_ids, 2)
    for id1, id2 in sample_pairs:
        T, p = stats.ttest_ind(samples[id1], samples[id2])

        mean_diff, mean_diff_per = mean_difference(np.exp(samples[id1]),
                                                    np.exp(samples[id2]))\
            if log_data else mean_difference(samples[id1], samples[id2])

```



```

        results.append([iv['label'], iv['categorical'][id1],
                        iv['categorical'][id2], dv, mean_diff,
                        mean_diff_per, T, p])

columns = [labels['iv'], labels['iv_value'] + ' 1',
           labels['iv_value'] + ' 2', labels['dv'],
           labels['mean_difference'], labels['mean_difference_percentage'],
           labels['t_statistic'], labels['p_value']]
results = pd.DataFrame(results, columns=columns)

p_values_correction(results, **args)
return results

In [44]: def test_non_normal_trials(iv_ids, dv_ids, data=trials, **args):
    results = []

    for dv_id in dv_ids:
        for iv_id in iv_ids:
            samples = trial_samples(iv_id, dv_id, data)
            H, p = stats.kruskal(*samples)
            results.append([ivs[iv_id]['label'], trials_dvs[dv_id], H, p])

    results = pd.DataFrame(results, columns=[labels['iv'], labels['dv'],
                                           'Kruskal-Wallis H',
                                           labels['p_value']])

    p_values_correction(results, **args)
    return results

In [45]: def test_pairwise_non_normal_trials(iv_ids, dv_ids, data=trials, **args):
    results = []

    for dv_id in dv_ids:
        for iv_id in iv_ids:
            iv, dv = ivs[iv_id], trials_dvs[dv_id]
            samples = trial_samples(iv_id, dv_id, data)

            iv_category_ids = range(len(iv['categorical']))
            sample_pairs = itertools.combinations(iv_category_ids, 2)
            for id1, id2 in sample_pairs:
                U, p = stats.mannwhitneyu(samples[id1], samples[id2])
                mean_diff, mean_diff_per = mean_difference(samples[id1],
                                                            samples[id2])
                results.append([iv['label'], iv['categorical'][id1],
                                iv['categorical'][id2], dv, mean_diff,
                                mean_diff_per, U, p])

    columns = [labels['iv'], labels['iv_value'] + ' 1',
               labels['iv_value'] + ' 2', labels['dv'],
               labels['mean_difference'], labels['mean_difference_percentage'],
               'Mann-Whitney U', labels['p_value']]
    results = pd.DataFrame(results, columns=columns)

```

```

    p_values_correction(results, **args)
    return results

In [46]: def plot_trials(iv_ids_list, dv_id, data=trials, kind='bar', **args):
    dv = trials_dvs[dv_id]
    if (len(iv_ids_list) == 0):
        iv_ids_list = [[iv_id] for id_id in ivs.columns]

    fig, axs = subplots(len(iv_ids_list))
    for id_ids, ax in zip(iv_ids_list, axs):

        iv_ids = [ivs[id_id] for id_id in id_ids]
        if (len(iv_ids) == 1):
            if (kind == 'bar'):
                sns.barplot(x=iv_ids[0]['label'], y=dv, data=data,
                           palette=iv_ids[0]['palette'], ax=ax, **args)

            elif (kind == 'box'):
                sns.boxplot(x=iv_ids[0]['label'], y=dv, data=data,
                           palette=iv_ids[0]['palette'], ax=ax, **args)

            elif (kind == 'count'):
                sns.countplot(hue=iv_ids[0]['label'], x=dv, data=data,
                             palette=iv_ids[0]['palette'], ax=ax, **args)
                ax.set(ylabel='Count')
                ax.legend(loc='upper right', title=labels['count'],
                          frameon=True)

        elif (len(ivs) == 2):
            if (kind == 'bar'):
                sns.barplot(x=iv_ids[1]['label'], y=dv, hue=iv_ids[0]['label'],
                           data=data, palette=iv_ids[0]['palette'], ax=ax,
                           **args)
                ax.legend(frameon=True, loc='upper left', bbox_to_anchor=(1, 1))

    return (fig, axs)

```

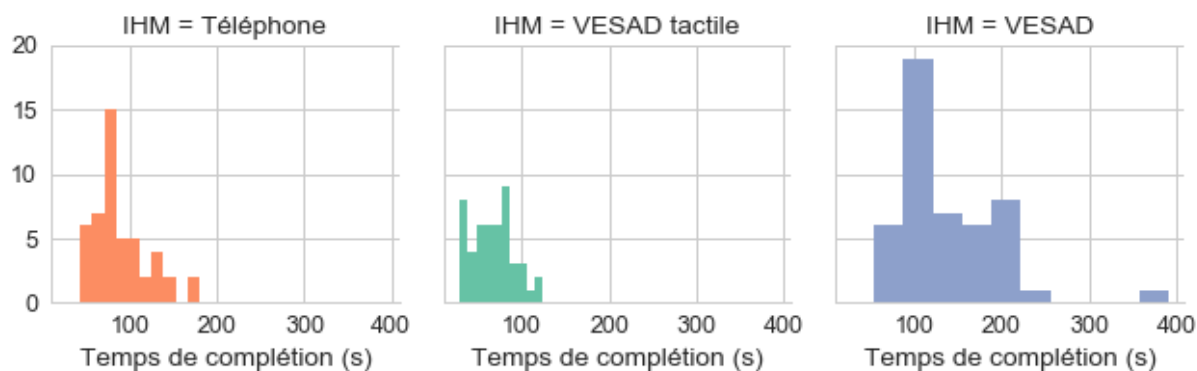
0.3.1 3.1. Task completion time

We first apply a log transform to TCT to approximate a normal distribution.

```

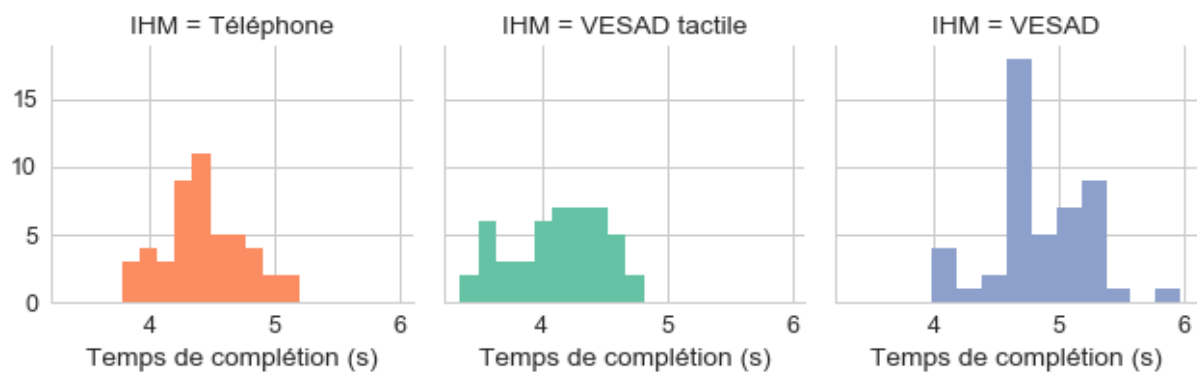
In [47]: g = sns.FacetGrid(trials, col=technique['label'], hue=technique['label'],
                           palette=technique['palette'])
    g = g.map(plt.hist, total_time)
    g.savefig('tct_distributions.png')

```



```
In [48]: trials[total_time] = np.log(trials[total_time])
raw_trials['total_time'] = np.log(raw_trials['total_time'])

In [49]: g = sns.FacetGrid(trials, col=technique['label'], hue=technique['label'],
                           palette=technique['palette'])
g = g.map(plt.hist, total_time)
g.savefig('tct_distributions_log.png')
```



We test the normality of TCT distributions for each TECHNIQUE and their equality of variances, since it's the main factor of interest.

```
In [50]: test_normality(['technique'], ['total_time'])
```

```
Out[50]:
```

	VI	Valeur VI	VD	Shapiro W	Valeur p
0	IHM	Téléphone	Temps de complétion (s)	0.983679	0.735995
1	IHM	VESAD tactile	Temps de complétion (s)	0.971828	0.446912
2	IHM	VESAD	Temps de complétion (s)	0.966510	0.446912

```
In [51]: test_equal_variances(['technique'], ['total_time'])
```

```
Out[51]:
```

	VI	VD	Levene W	Valeur p
0	IHM	Temps de complétion (s)	0.756269	0.471309

We meet all the assumptions of an ANOVA. Trials were done independently, TCT distributions are normal and their variances are equal.

We perform a full factorial ANOVA with the model: $TCT \sim \text{TECHNIQUE} \times \text{TEXT_SIZE} \times \text{DISTANCE} + \text{TECHNIQUE} \times \text{ORDERING}$.

```
In [52]: tct_model = ols('total_time ~ technique * text_size * distance'
                        + '+ technique * ordering', data=raw_trials).fit()
sm.stats.anova_lm(tct_model, typ=2)
```

```
Out[52]:
```

	sum_sq	df	F	PR(>F)
technique	12.328279	2.0	62.210952	1.610604e-19
text_size	0.145175	1.0	1.465169	2.283752e-01
distance	0.029459	1.0	0.297314	5.865353e-01
ordering	2.133256	2.0	10.764832	4.829976e-05
technique:text_size	0.398420	2.0	2.010508	1.381943e-01
technique:distance	0.074854	2.0	0.377730	6.861892e-01
text_size:distance	0.291976	1.0	2.946742	8.850822e-02
technique:ordering	3.086868	4.0	7.788476	1.215733e-05
technique:text_size:distance	0.499563	2.0	2.520896	8.443561e-02
Residual	12.484644	126.0	NaN	NaN

The main significant effect on TCT is TECHNIQUE ($F = 62.2$, $p < 0.0001$), then ORDERING ($F = 10.8$, $p < 0.0001$). There is also an significant interaction effect: TECHNIQUE \times ORDERING ($F = 2.5$, $p < 0.0001$).

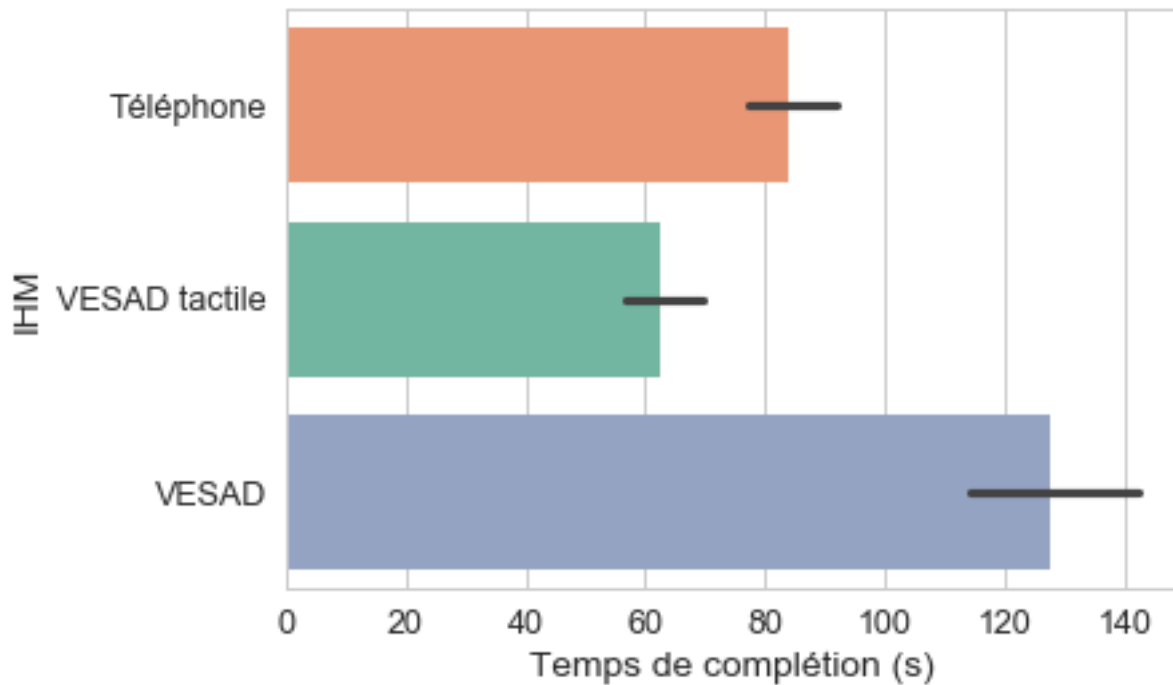
We display mean TCT values with 95% CI for these conditions:

```
In [53]: trial_means(['technique'], ['total_time'], aggregate=print_exp_mean_ci)
```

```
Out[53]:
```

	Temps de complétion (s)
IHM	
Téléphone	84.02 [76.81, 92.76]
VESAD tactile	62.59 [56.06, 69.33]
VESAD	127.71 [114.14, 141.96]

```
In [54]: ax = sns.barplot(x=trials_dvs['total_time'], y=technique['label'],
                        palette=technique['palette'], data=trials, estimator=exp_mean)
ax.get_figure().savefig('tct.png')
```



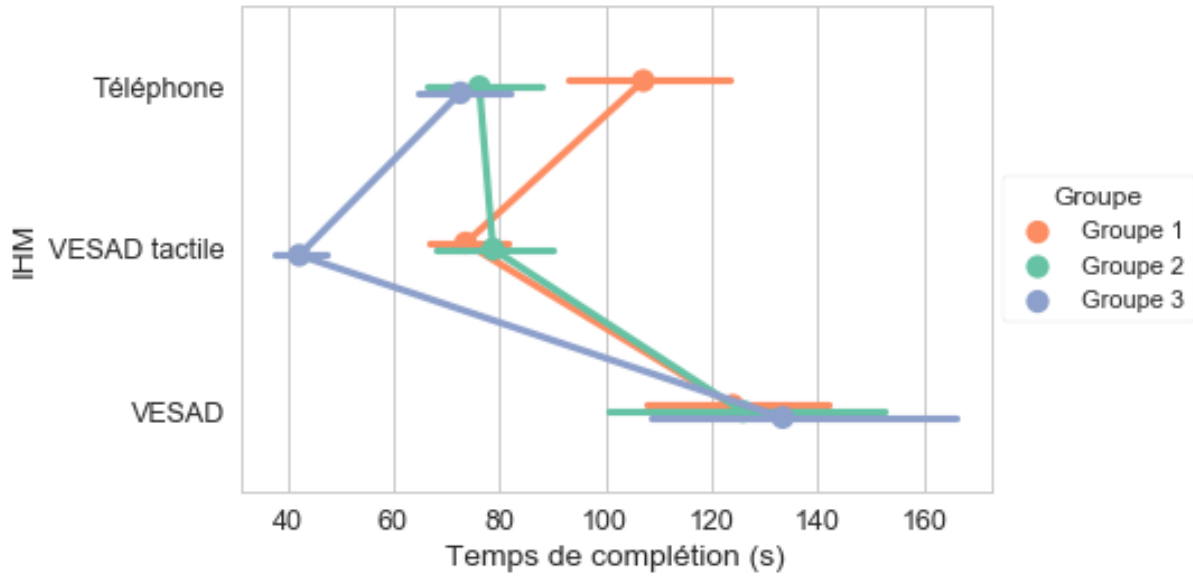
```
In [55]: trial_means(['ordering', 'technique'], ['total_time'],
                    aggregate=print_exp_mean_ci).unstack()
```

```
Out[55]:
```

		Temps de complétion (s)		
IHM		Téléphone	VESAD tactile	\
Groupe				
Groupe 1	107.14	[92.23, 123.88]	73.63	[67.26, 81.12]
Groupe 2	76.22	[66.31, 87.24]	78.79	[68.15, 89.81]
Groupe 3	72.64	[64.96, 82.05]	42.27	[37.99, 47.55]

IHM		VESAD	
Groupe			
Groupe 1	124.00	[108.09, 141.20]	
Groupe 2	125.91	[101.42, 152.20]	
Groupe 3	133.41	[108.95, 169.41]	

```
In [56]: ax = sns.pointplot(x=total_time, y=technique['label'], hue=ordering['label'],
                             palette=ordering['palette'], data=trials, dodge=True,
                             estimator=exp_mean)
config_legend(ax, 'ordering')
ax.get_figure().savefig('tct_ordering.png')
```



It seems that participants who started with *PhoneOnly* were slower with this technique than the other groups. Similarly, participants who finished with *PhoneInArOut* were faster with this technique. It indicates there is a learning curve on the task, but interestingly participants from all groups performed equally with *MidAirInArOut* technique.

We compare the TCT for the three techniques only with pairwise t-tests (Benjamini–Hochberg correction).

```
In [57]: test_pairwise_trials('technique', 'total_time', correction_method='fdr_bh',
                             log_data=True)
```

```
Out[57]:
```

	VI	Valeur VI 1	Valeur VI 2	VD \
0	IHM	Téléphone	VESAD tactile	Temps de complétion (s)
1	IHM	Téléphone	VESAD	Temps de complétion (s)
2	IHM	VESAD tactile	VESAD	Temps de complétion (s)

	Différence des moyennes	Différence des moyennes (%)	Statistique T \
0	21.840362	32.666819	4.090635
1	-49.467731	-35.803106	-5.615875
2	-71.308093	-51.610437	-9.032751

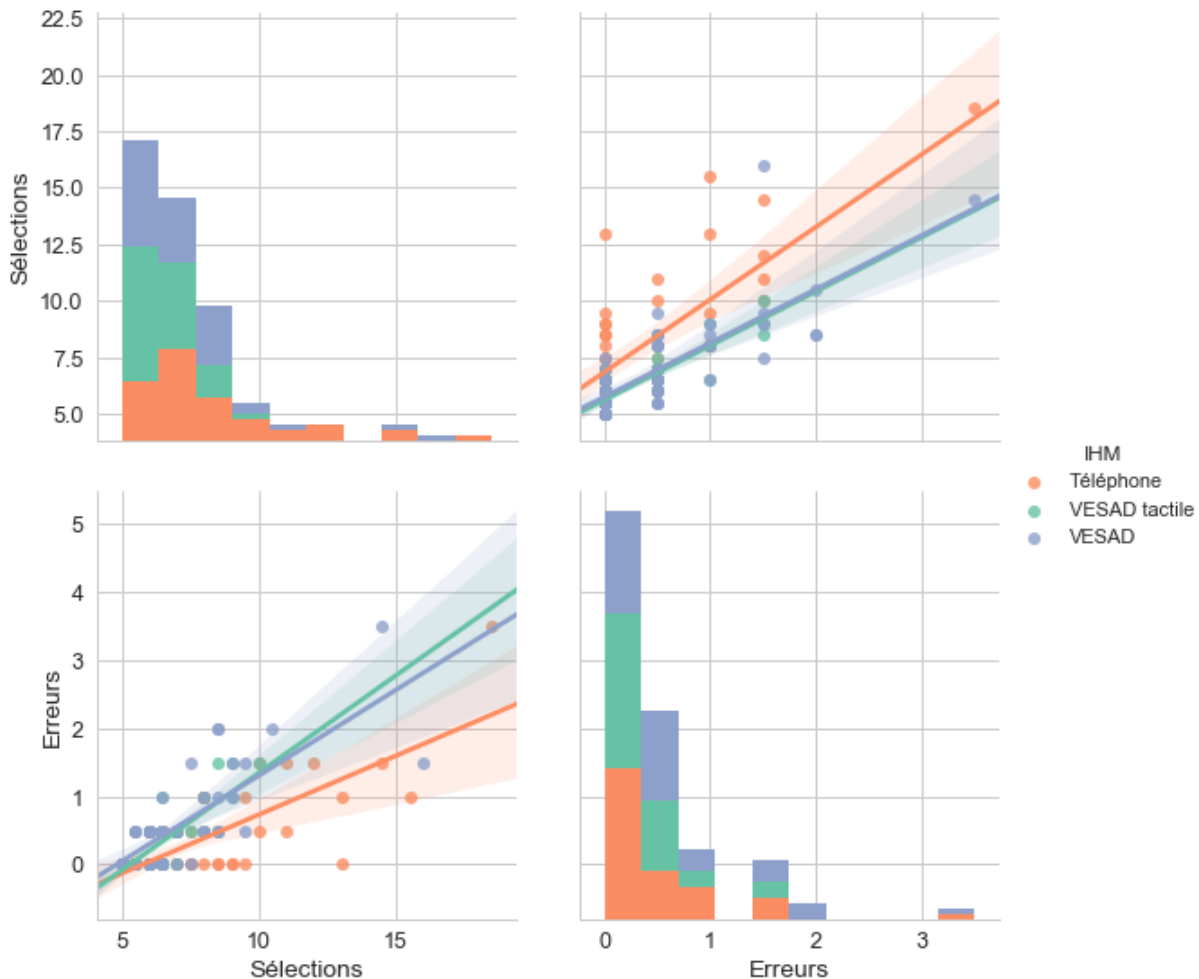
	Valeur p
0	9.077859e-05
1	2.966612e-07
2	6.282833e-14

- *PhoneInArOut* is 71s (+52%) faster than *MidAirInArOut* ($p < 0.0001$).
- *PhoneOnly* is 49s (+36%) faster than *MidAirInArOut* ($p < 0.0001$).
- *PhoneInArOut* is 22s (+33%) faster than *PhoneOnly* ($p < 0.0001$).

0.3.2 3.2. Errors

We visualize first the SELECTIONS and ERRORS distributions:

```
In [58]: g = sns.pairplot(trials, hue=technique['label'], kind='reg',
                        vars=[trials_dvs['selections_count'], trials_dvs['errors']],
                        palette=technique['palette'], size=4);
g.savefig('selections_errors_distributions.png')
```



It seems that a user makes as much errors as she makes selections. The relation is almost the same for each technique, even if users seems to make more selections for the same number of errors with PhoneOnly.

We can't use ANOVA on SELECTIONS and ERRORS variables as their distributions are exponentials. We use instead the Kruskal-Wallis test (Benjamini-Hochberg correction) to check if there is significative differences due to TECHNIQUE, TEXT_SIZE, DISTANCE or ORDERING.

```
In [59]: test_non_normal_trials(['technique', 'text_size', 'distance', 'ordering'],
                                ['selections_count', 'errors'])
```

```
Out[59]:
```

	VI	VD	Kruskal-Wallis H	Valeur p
0	IHM	Sélections	15.125918	0.004155
1	Taille du texte	Sélections	0.003897	0.950224
2	Distance	Sélections	0.052701	0.935347
3	Groupe	Sélections	11.510535	0.010095

4	IHM	Erreurs	5.201541	0.148433
5	Taille du texte	Erreurs	0.103661	0.935347
6	Distance	Erreurs	0.201599	0.935347
7	Groupe	Erreurs	11.153096	0.010095

Only TECHNIQUE ($p = 0.004$) and ORDERING ($p = 0.01$) have a significant effect on SELECTIONS. But, only ORDERING ($p = 0.01$) have a significant effect on ERRORS.

We use then pairwise Mann-Whitney tests (Benjamini–Hochberg correction) for the significant questions above:

```
In [60]: test_pairwise_non_normal_trials(['technique', 'ordering'],
                                         ['selections_count', 'errors'])
```

```
Out[60]:
```

	VI	Valeur VI 1	Valeur VI 2	VD	Différence des moyennes \
0	IHM	Téléphone	VESAD tactile	Sélections	1.864583
1	IHM	Téléphone	VESAD	Sélections	1.041667
2	IHM	VESAD tactile	VESAD	Sélections	-0.822917
3	Groupe	Groupe 1	Groupe 2	Sélections	1.729167
4	Groupe	Groupe 1	Groupe 3	Sélections	1.583333
5	Groupe	Groupe 2	Groupe 3	Sélections	-0.145833
6	IHM	Téléphone	VESAD tactile	Erreurs	0.125000
7	IHM	Téléphone	VESAD	Erreurs	-0.166667
8	IHM	VESAD tactile	VESAD	Erreurs	-0.291667
9	Groupe	Groupe 1	Groupe 2	Erreurs	0.333333
10	Groupe	Groupe 1	Groupe 3	Erreurs	0.343750
11	Groupe	Groupe 2	Groupe 3	Erreurs	0.010417

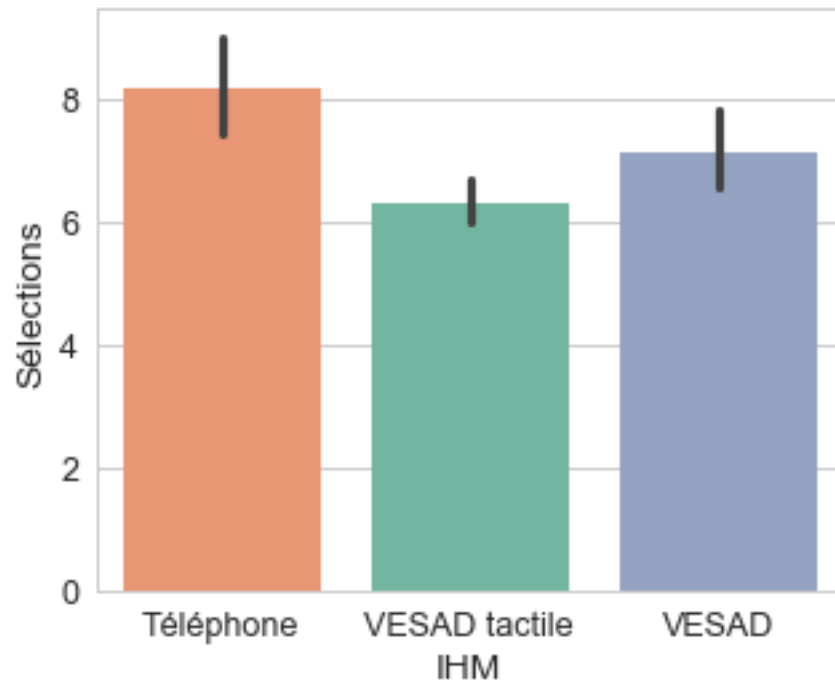
	Différence des moyennes (%)	Mann-Whitney U	Valeur p
0	29.537954	627.0	0.000635
1	14.598540	864.5	0.029356
2	-11.532847	915.0	0.059881
3	26.265823	755.0	0.005196
4	23.529412	755.0	0.005196
5	-2.167183	1136.0	0.454285
6	42.857143	1083.5	0.312007
7	-28.571429	957.0	0.083050
8	-50.000000	873.0	0.026306
9	103.225806	846.5	0.020183
10	110.000000	761.5	0.005196
11	3.333333	1039.0	0.205336

```
In [61]: trial_means(['technique'], ['selections_count', 'errors'])
```

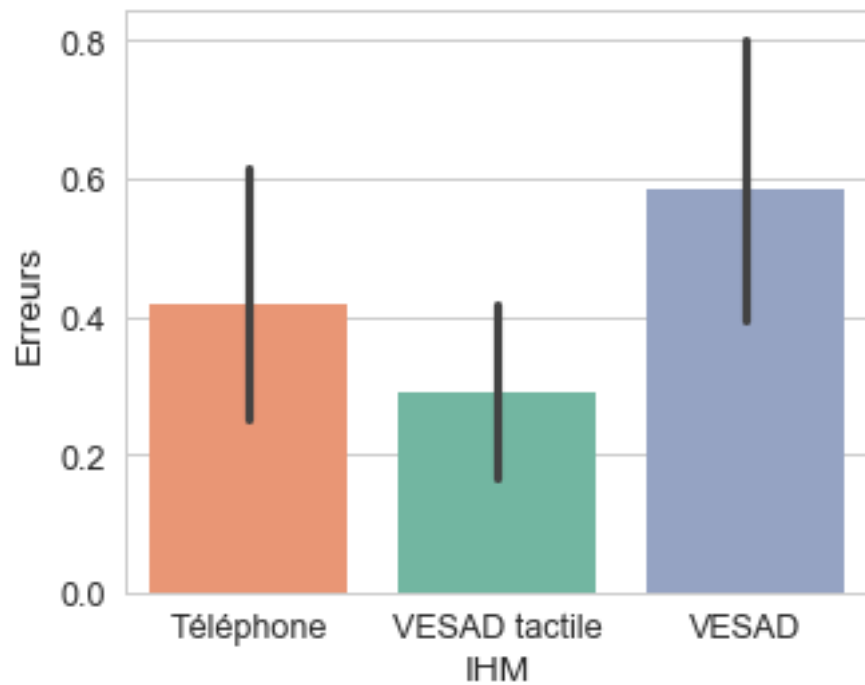
```
Out[61]:
```

	Sélections	Erreurs
IHM		
Téléphone	8.18 [7.42, 8.99]	0.42 [0.24, 0.59]
VESAD tactile	6.31 [5.97, 6.70]	0.29 [0.18, 0.42]
VESAD	7.14 [6.54, 7.86]	0.58 [0.39, 0.80]

```
In [62]: (fig, axs) = plot_trials(['technique'], 'selections_count')
fig.savefig('selections.png')
```

```
In [63]: (fig, axs) = plot_trials([[ 'technique' ]], 'errors')
         fig.savefig('errors.png')
```

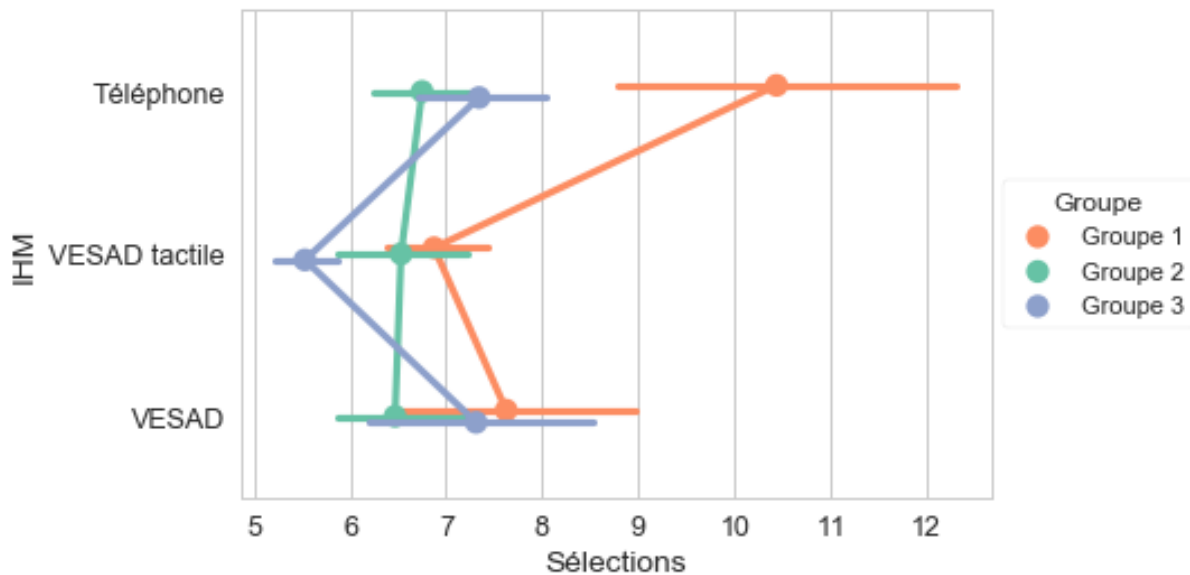


```
In [64]: trial_means([ 'ordering', 'technique' ], [ 'selections_count', 'errors' ])
```

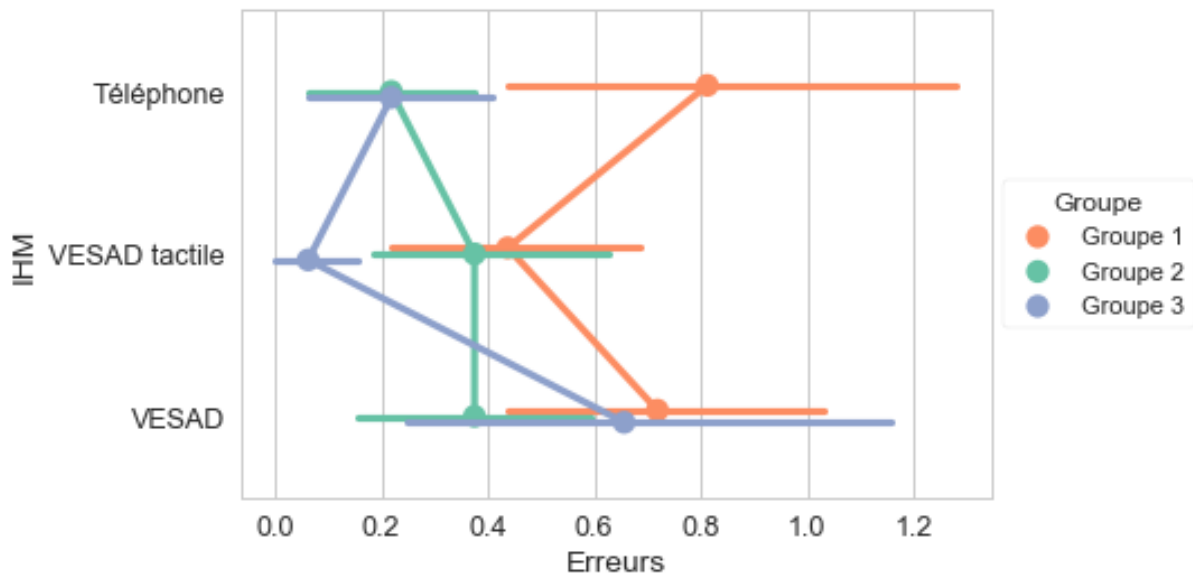
Out [64]:

		Sélections	Erreurs
Groupe	IHM		
Groupe 1	Téléphone	10.44 [8.72, 12.31]	0.81 [0.38, 1.28]
	VESAD tactile	6.88 [6.38, 7.50]	0.44 [0.22, 0.69]
	VESAD	7.62 [6.62, 9.12]	0.72 [0.41, 1.06]
Groupe 2	Téléphone	6.75 [6.25, 7.31]	0.22 [0.06, 0.41]
	VESAD tactile	6.53 [5.87, 7.22]	0.38 [0.16, 0.59]
	VESAD	6.47 [5.91, 7.13]	0.38 [0.19, 0.62]
Groupe 3	Téléphone	7.34 [6.72, 8.06]	0.22 [0.06, 0.38]
	VESAD tactile	5.53 [5.25, 5.84]	0.06 [0.00, 0.16]
	VESAD	7.31 [6.28, 8.69]	0.66 [0.25, 1.13]

```
In [65]: ax = sns.pointplot(x=trials_dvs['selections_count'], y=technique['label'],
                             hue=ordering['label'], palette=ordering['palette'],
                             data=trials, dodge=True)
config_legend(ax, 'ordering')
ax.get_figure().savefig('selections_ordering.png')
```



```
In [66]: ax = sns.pointplot(x=trials_dvs['errors'], y=technique['label'],
                             hue=ordering['label'], palette=ordering['palette'],
                             data=trials, dodge=True)
config_legend(ax, 'ordering')
ax.get_figure().savefig('errors_ordering.png')
```



0.3.3 3.3. Navigation

Variable meanings:

- Selection Time = time spent looking for where to drop an item that had been picked
- Selection Distance = distance travelled by the finger with an item selected
- Head Phone Distance = sum of the distance between the head and the phone

In [67]: *# Data preparation*

```
trial_counts = melt_trials(var_name=labels['category'],
                           value_name=labels['count'],
                           value_vars=[trials_dvs['pan_count'],
                                       trials_dvs['zoom_count']])

trial_times = melt_trials(var_name=labels['category'],
                          value_name=labels['time'],
                          value_vars=[trials_dvs['selections_time'],
                                      trials_dvs['pan_time'],
                                      trials_dvs['zoom_time']])

trial_distances_dvs = [trials_dvs['selections_projected_distance'],
                      trials_dvs['pan_projected_distance'],
                      trials_dvs['zoom_projected_distance'],
                      trials_dvs['absolute_head_phone_distance']]
trial_distances = melt_trials(var_name=labels['category'],
                              value_name=labels['distance'],
                              value_vars=trial_distances_dvs)
```

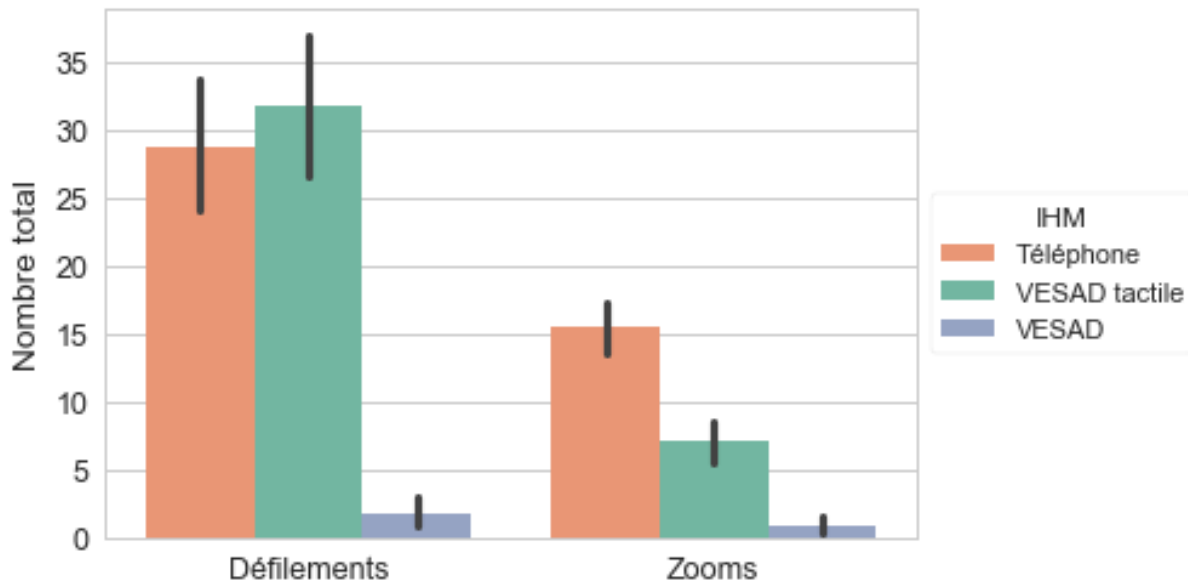
In [68]: trial_means(['technique'], ['pan_count', 'zoom_count'])

Out[68]:

	Défilements	Zooms
IHM		

Téléphone	28.70 [24.34, 33.28]	15.53 [13.62, 17.53]
VESAD tactile	31.81 [26.62, 37.60]	7.14 [5.72, 8.81]
VESAD	1.83 [0.81, 3.08]	0.83 [0.27, 1.53]

```
In [69]: ax = sns.barplot(x=labels['category'], y=labels['count'],
                        hue=technique['label'], palette=technique['palette'],
                        data=trial_counts)
config_legend(ax, 'technique')
ax.set(xlabel='')
ax.get_figure().savefig('navigation_count.png')
```



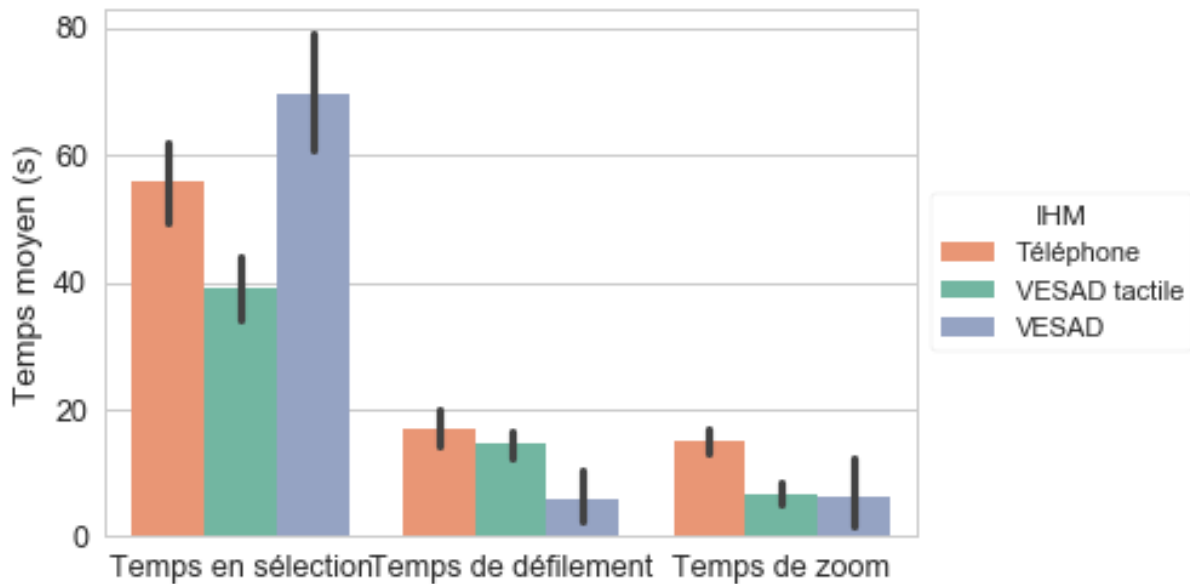
```
In [70]: trial_means(['technique'], ['selections_time', 'pan_time', 'zoom_time'])
```

```
Out[70]:
```

	Temps en sélection	Temps de défilement \
IHM		
Téléphone	55.74 [50.20, 61.79]	17.12 [14.48, 19.97]
VESAD tactile	38.98 [34.32, 43.80]	14.53 [12.51, 16.50]
VESAD	69.68 [60.35, 79.78]	5.97 [2.15, 10.07]

	Temps de zoom
IHM	
Téléphone	15.18 [13.31, 17.18]
VESAD tactile	6.70 [5.11, 8.62]
VESAD	6.24 [1.67, 11.87]

```
In [71]: ax = sns.barplot(x=labels['category'], y=labels['time'], data=trial_times,
                        hue=technique['label'], palette=technique['palette'])
config_legend(ax, 'technique')
ax.set(xlabel='')
ax.get_figure().savefig('navigation_time.png')
```



```
In [72]: trial_means(['technique'], ['selections_projected_distance',
                                     'pan_projected_distance',
                                     'zoom_projected_distance',
                                     'absolute_head_phone_distance'])
```

```
Out[72]:
```

	Distance en sélection	Distance de défilement	Distance de zoom \
IHM			
Téléphone	4.87 [4.09, 5.68]	1.50 [1.20, 1.82]	2.11 [1.75, 2.55]
VESAD tactile	2.83 [2.40, 3.28]	1.09 [0.91, 1.29]	0.69 [0.51, 0.89]
VESAD	8.72 [7.19, 10.53]	0.47 [0.18, 0.86]	0.62 [0.15, 1.19]

Mouvements tête-téléphone

IHM	
Téléphone	3.18 [2.34, 4.12]
VESAD tactile	1.57 [1.27, 1.91]
VESAD	6.12 [4.93, 7.43]

```
In [73]: g = sns.factorplot(x=technique['label'], y=labels['distance'],
                             col=labels['category'], data=trial_distances,
                             palette=technique['palette'], kind='bar', col_wrap=2)
```

```
g.set_titles('{col_name}') # Replace subplot titles
```

```
for ax in g.axes:
    ax.title.set_position([0.5, -0.12])
```

```
g.set_axis_labels('') # Custom legend
```

```
g.set_xticklabels([])
```

```
legend_handles = [patches.Patch(color=color, label=value)\
                    for value, color in zip(technique['categorical'],
                                             technique['palette'])]
```

```

legend = plt.legend(handles=legend_handles, loc='center left',
                    bbox_to_anchor=(1, 1.1), title=technique['label'])
fix_legend_fontsize(legend)

g.savefig('navigation_distance.png')

```

