Introduction

This project archives the results for the following, in-submission publication:

Inducing musical-interval learning by combining task practice with periods of stimulus exposure alone

Abstract

A key component of musical expertise is the ability to discriminate between and identify musical intervals, such as a perfect 4th and a major 3rd. This ability is necessary for playing by ear, performance self-correction, and the analysis of music. However, even musicians can struggle with 'ear training' and mastery is typically associated with years of musical experience. Here we report a training regimen that yields learning on musical-interval discrimination and generalization to musical-interval identification across four training sessions. Critically, the regimen combines periods of practice on musical-interval discrimination and periods of stimulus exposure without practice within each session. Continuous practice yields no improvement at all. The Practice+Exposure regimen generates improvement in both musicians and non-musicians, providing a practical means for the rapid acquisition of an aural skill in music that has long been viewed to take many years to learn.

Installation

Once you have donwloaded this project, to setup the development environment will need to install Anaconda Python 2.7. Once you're finished with the installation, run the following commands on the command line (replacing [project directory] with the location of the folder containing the downloaded project).

```
cd [project directory]
conda env create -f environment.yml
source activate apmusic 2017 12 11
```

In windows these command can be run on the command line that is installed with Anaconda. You will also need to replace the last line as follows:

```
activate apmusic_2017_12_11
```

This setup will take some time to run on your machine as various python and R software packages are installed.

Usage

Once installed, to run the anlayses or re-create a figure you will need to have an open prgoramming environment ready. If you haven't closed the window you opened during "Installation", you can use that, otherwise, re-open the command line and run the following

```
cd [project directory]
source activate apmusic_2017_12_11
```

Again, in Windows these command can be run on the command line that is installed with Anaconda and you will need to replace the last line with activate apmusic_2017_12_11 (no source).

Each figure panel from the paper generally has at least two coresponding files. They are either python or R source files, and they re-create the figure pdf and run the corresponding analyses, respectively. These files are located under the plots and analyses subfolders. There are some figures with auxillary analyses (e.g. figure 2A has a secondary analysis by foil instead of regimen), which are included as seperate scripts. All scripts associated with a figure include that figure name as a prefix of the filename.

To run any of these files, call either Rscript or python (as appropriate for the file type) from the same command line you used above.

For example, to re-run the analyses corresponding to Figure 1A you can call the following from the command line.

```
python analyses/fig1A.py
```

And to re-create Figure 1A's pdf:

Rscript plots/fig1A.R

Yet again, Windows will differ: use \setminus instead of / in your directory names.

Recreating the preprocessed data

The scripts for each figure and analysis load some preprocessed data. All preprocessed data files are listed in preprocessing/files.txt. If you want to recreate one of these files you can do so by finding the script with the same name prefix as the file.

For example, the preprocessed data file preprocessing/data/discrim_2017-12-31.csv can be recreated (with a new date suffix) by calling

Rscript preprocessing/discrim.R

All preprocessed files are stored under the preprocessing/data subfolder, while the data subfolder contains the original data, as collected during the experiment.

WARNING: some of the preprocessing steps can be time consuming to run.

To start all analyses over from scratch, delete all of the preprocessing data files under the preprocessing/data directory. Then, rerun the preprocessing scripts in the following order

- 1. Aggregate discrimination data preprocessing/discrim.R
- 2. Aggregate classification data preprocessing/class.R
- 3. Multi-level model fitting preprocessing/multi_level_sample.py
- 4. Multi-level model fitting with musical experience preprocessing/music_multi_level_sample.py
- 5. Each figure's preprocessing scripts preprocessing/fig1A.py, preprocessing/fig1B.py etc...

Steps 2 and 3 will be the most time consuming (e.g. ~ 30 minutes on a 2015 MacBook Air).

If you re-process any data you will have to update the file name listed in preprocessing/files.txt so that any scripts relying on this data will use the newly created files.

Editing the scripts

To edit the R files you can use R Studio and to edit python you can use Spyder. Alternatively you can use a decent text editor to edit both types of files:

- Emacs
- Vim
- Sublime

Regimen Abbreviations

The training regimens have been given abbreviations: AP (Practice+Exposure), LA (All-Practice), SA (Practice+Silence), and P (Exposure+Silence).

Comparison to Published Figures and Statistics

The resulting pdfs from the analysis were imported into Adobe Illustrator and the aesthetics modified for improved legibility and consistency. Thus the published figures will differ somewhat from those produced by these scripts: however, the actual data and reported statistics as represented in the final paper should remain *unchanged*. Small variations in model fits and the resulting statistics may occur due to the MCMC sampling if any of the model samples are re-processed.