

# pygamma-agreement: Gamma $\gamma$ measure for inter/intra-annotator agreement in Python

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#### **Software**

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## Introduction

In the last decades, it became easier to collect large audio recordings in naturalistic conditions and large corpora of text from the Internet. This broadens the scope of questions that can be addressed regarding speech and language.

Scientists need to challenge their hypotheses and quantify the observed phenomenons on speech and language; that is why researchers add different layers of annotations on top of speech and text data. Some type of human intervention is used to reliably describe events contained in the corpus's content (ex: Wikipedia articles, conversations, child babbling, animal vocalizations, or even just environmental sounds). These events can either be tagged at a particular point in time, or over a stretch of time. It is also commonplace to provide a categorical annotation or - in the case of speech - even precise transcriptions (Serratrice, 2000) for these events. Depending on the difficulty of the annotation task and the eventual expertise of the annotators, the annotations they produce can include a certain degree of interpretation. A common strategy when building annotated corpora is to have small parts of a corpus annotated by several annotators, to be able quantify their consensus on that reduced subset of the corpus. If that consensus is deemed robust (i.e., agreement is high), we infer that the annotation task is well defined, less prone to interpretation, and that annotations that cover the rest of the corpus are reliable (Gwet, 2012). An objective measure of the agreement (and subsequent disagreement) between annotators is thus desirable.

## Statement of Need

The Gamma  $(\gamma)$  Inter-Annotator Agreement Measure was proposed by (Mathet et al., 2015) as a way to quantify inter-rater agreement for sequences of annotations. The  $\gamma$ -agreement measure solves a number of the shortcomings of other pre-existing measures. This quantification will have to satisfy some constraints: segmentation, unitizing, categorization, weighted categorization and the support for any number of annotators. They should also provide a chance-corrected value. Other measures, such as the  $\kappa$  (Carletta, 1996) or Krippendorff's  $\alpha$ 's (Krippendorff, 2011), have existed for some time to deal with these constraints, but never could address all of them at once. A detailed comparison between metrics is available in (Mathet et al., 2015).

To solve all of these constraints at once, the  $\gamma$ -agreement works in three steps: 1) a disorder (i.e. a cost) is computed for each potential alignments between the different annotators' units, using an annotation-dependent dissimilarity (akin to a distance between units). This disorder models the disagreement between annotators. 2) Using a convex optimization algorithm, a global alignment with the lowest total disorder is found. 3) By sampling random and synthetic



annotations from the original annotation and computing their own disorders, the original annotation's disorder value is chance-corrected to provide the final  $\gamma$ -agreement measure. The authors of (Mathet et al., 2015) provided a Java freeware GUI implementation along with their paper. This implementation has already been used by some researchers (Da San Martino et al., 2019) to compute an inter-rater agreement measure on their annotations.

However, linguist and automated speech researchers today use analysis pipelines that are either Python or shell scripts. To this day, no open-source implementation allows for the  $\gamma$ agreement to be computed in a programmatical way, and researchers that are already proficient in Python and willing to automate their work might be hindered by the graphical nature of the original Java implementation. Moreover, the original  $\gamma$ -agreement algorithm has several parameters that are determinant in its computation and cannot be configured as of now. For this reason, we thought it would greatly benefit the speech and linguistic scientific community if we could provide them with a fully open-source Python implementation of the original algorithm. In that end, we made sure that our implementation had several key features. It is comparatively as fast as the original implementation, taking about 10s to compute a highconfidence  $\gamma$ -agreement measure on a middle-range processor. The pygamma-agreement package is modular and users can easily extend one of the modules without the burden of rebuilding an optimized code base from scratch (Ex: Users can easily add a new dissimilarity measure). Our code allows for fine-grained constructions of an annotation Continuum and an advanced configurability of the  $\gamma$ -agreement's different parameters. We also made sure it eases the interpretability of the input (i.e. the annotation continuum) and the output (I.e the alignments) data structures, as both can be visualized in a Jupyter Notebook (see Figure 1). Finally, we support most of the commonly used data formats and analysis pipelines in the speech and linguistic fields.

# The pygamma-agreement Package

The pygamma-agreement package provides users with two ways to compute the  $\gamma$ -agreement for a corpus of annotations. The first one is to use the package's Python API.

```
import pygamma_agreement as pa
continuum = pa.Continuum.from_csv("data/PaulAlexSuzann.csv")
dissimilarity = pa.CombinedCategoricalDissimilarity(categories=list(continuum.categamma_results = continuum.compute_gamma(dissimilarity, precision_level=0.02)
print(f"Gamma is {gamma_results.gamma}")
```

The most important primitives from our API (the Continuum Figure 1 and Alignment Figure 2 classes) can be displayed using the matplotlib.pyplot backend if the user is working in a Jupyter notebook.

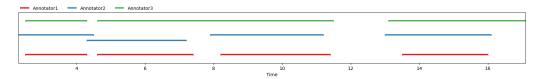
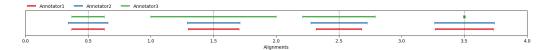


Figure 1: Displaying a Continuum in a jupyter notebook. This is a temporally accurate representation of the annotated data.





**Figure 2:** Displaying an Alignment in a jupyter notebook. This is a visual and schematic representation of the alignment computed between annotations of the original continuum (the order of units is respected but the annotations are scaled for visual clarity).

The second one is a command-line application that can be invoked directly from the shell, for those who prefer to use shell scripts for corpus processing:

pygamma-agreement corpus/\*.csv --confidence\_level 0.02 --output\_csv results.csv

We support a variety of commonly used annotation formats among speech researchers and linguists: RTTM, ELAN, TextGrid, CSV and pyannote.core.Annotation objects.

Computing the  $\gamma$ -agreement requires both array manipulation and the solving of multiple optimization problem formulated as Mixed-Integer Programming (MIP) problems. We thus used the *de facto* standard for all of our basic array operations, NumPy (Harris et al., 2020). Since some parts of the algorithm are fairly demanding, we made sure that these parts were heavily optimized using numba (Lam et al., 2015). We used cvxpy's (Diamond & Boyd, 2016) MIP-solving framework to solve the optimization problem. For time-based annotations, we rely on primitives from pyannote.core (Bredin et al., 2020). We made sure that it is robustly tested using the widely-adopted pytest testing framework. We also made sure that pygamma-agreement's outputs matched both the theoretical values from the original paper and those of the Java freeware. We set-up an automated Travis CI to use these tests to ensure our package's quality. Most of our package's code is type-hinted and has descriptive docstrings, both of which can be leveraged by IDEs to ease the use of our API.

We provide a user documentation as well as an example Jupyter notebook in our package's repository. Additionally, we have used and tested pygamma-agreement in conjunction with the development of our own custom-built annotation platform, Seshat (Titeux et al., 2020). In Table , we present two use cases for our implementation of the  $\gamma$ -agreement measure on two corpora. These two corpora, ranging from medical interviews to child recording, allowed us to evaluate the performance of the  $\gamma$ -agreement on a wide panel of annotation types.

Corpus	Annotation	# Classes	Mean of $\gamma$
Clinical Interviews	Turn-Takings	3	0.64
	Utterances	1	0.61
Child Recordings	Speech Activity	1	0.46
Child Recordings	Child/Adult-directed speech	2	0.27

Table :  $\gamma$  Inter-rater agreement for clinical interviews (16 samples) and child-centered day-long recordings (20 samples).

We have uploaded our package to the Pypi repository, thus, pygamma-agreement can be installed using pip.

#### **Future Work**

Our implementation of the  $\gamma$ -agreement opens the path for a number of potential extensions:



- An obvious improvement is to add support for the " $\gamma$ -cat" metric, a complement measure (Mathet, 2017) for the  $\gamma$ -agreement.
- The  $\gamma$ -agreement's theoretical framework allows for some useful improvements such as:
  - the implementation of new dissimilarities, such as a sequence-based dissimilarity (based on the Levenshtein distance), an ordinal dissimilarity (for ordered sets of categories) and a scalar dissimilarity.
  - for categorical annotations, the support for an undefined set of categories, with annotators using different sets of categories. This would be solved using an adapted implementation of the Hungarian Algorithm. This could be useful in unconstrained diarization annotation tasks.
  - more hypothetically, the possibility to have so-called "soft alignments," where a unit has weighted alignments with other units in its continuum.

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