

¹ Peekbank: An open, large-scale repository for developmental eye-tracking data of children's
² word recognition

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25 **Open Practices Statement.** All code for reproducing the paper is available at
26 <https://github.com/langcog/peekbank-paper>. Raw and standardized datasets are available
27 on the Peekbank OSF repository (<https://osf.io/pr6wu/>) and can be accessed using the
28 peekbankr R package (<https://github.com/langcog/peekbankr>).

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37

Abstract

38 The ability to rapidly recognize words and link them to referents is central to children's
39 early language development. This ability, often called word recognition in the developmental
40 literature, is typically studied in the looking-while-listening paradigm, which measures
41 infants' fixation on a target object (vs. a distractor) after hearing a target label. We present
42 a large-scale, open database of infant and toddler eye-tracking data from
43 looking-while-listening tasks. The goal of this effort is to address theoretical and
44 methodological challenges in measuring vocabulary development. We first present how we
45 created the database, its features and structure, and associated tools for processing and
46 accessing infant eye-tracking datasets. Using these tools, we then work through two
47 illustrative examples to show how researchers can use Peekbank to interrogate theoretical
48 and methodological questions about children's developing word recognition ability.

49 *Keywords:* word recognition; eye-tracking; vocabulary development;
50 looking-while-listening; visual world paradigm; lexical processing

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53 word recognition

54 Across their first years of life, children learn words at an accelerating pace (Frank,
55 Braginsky, Yurovsky, & Marchman, 2021). While many children will only produce their first
56 word at around one year of age, most children show signs of understanding many common
57 nouns (e.g., *mommy*) and phrases (e.g., *Let's go bye-bye!*) much earlier in development
58 (Bergelson & Swingley, 2012, 2013; Tincoff & Jusczyk, 1999). Although early word
59 understanding is a critical element of first language learning, the processes involved are less
60 directly apparent in children's behaviors and are less accessible to observation than
61 developments in speech production (Fernald, Zangl, Portillo, & Marchman, 2008;
62 Hirsh-Pasek, Cauley, Golinkoff, & Gordon, 1987). To understand a spoken word, children
63 must process the incoming auditory signal and link that signal to relevant meanings – a
64 process often referred to as word recognition. One of the primary means of measuring word
65 recognition in young infants is using eye-tracking techniques that gauge where children look
66 in response to linguistic stimuli (Fernald, Zangl, Portillo, & Marchman, 2008). The logic of
67 these methods is that if, upon hearing a word, a child preferentially looks at a target
68 stimulus rather than a distractor, the child is able to recognize the word and activate its
69 meaning during real-time language processing. Measuring early word recognition offers
70 insight into children's early word representations: children's speed of response (i.e., moving
71 their eyes; turning their heads) to the unfolding speech signal can reveal children's level of
72 comprehension (Bergelson, 2020; Fernald, Pinto, Swingley, Weinberg, & McRoberts, 1998).
73 Word recognition skills are also thought to build a foundation for children's subsequent
74 language development. Past research has found that early word recognition efficiency is
75 predictive of later linguistic and general cognitive outcomes (Bleses, Makransky, Dale, Højlen,
76 & Ari, 2016; Marchman et al., 2018).

77 While word recognition is a central part of children's language development, mapping

78 the trajectory of word recognition skills has remained elusive. Studies investigating children's
79 word recognition are typically limited in scope to experiments in individual labs involving
80 small samples tested on a handful of items. The limitations of single datasets makes it
81 difficult to understand developmental changes in children's word knowledge at a broad scale.

82 One way to overcome this challenge is to compile existing datasets into a large-scale
83 database in order to expand the scope of research questions that can be asked about the
84 development of word recognition abilities. This strategy capitalizes on the fact that the
85 looking-while-listening paradigm is widely used, and vast amounts of data have been
86 collected across labs on infants' word recognition over the past 35 years (Golinkoff, Ma, Song,
87 & Hirsh-Pasek, 2013). Such datasets have largely remained isolated from one another, but
88 once combined, they have the potential to offer general insights into lexical development.

89 Similar efforts to collect other measures of language development have borne fruit in recent
90 years. For example, WordBank aggregated data from the MacArthur-Bates Communicative
91 Development Inventory, a parent-report measure of child vocabulary, to deliver new insights
92 into cross-linguistic patterns and variability in vocabulary development (Frank, Braginsky,
93 Yurovsky, & Marchman, 2017, 2021). In this paper, we introduce *Peekbank*, an open
94 database of infant and toddler eye-tracking data aimed at facilitating the study of
95 developmental changes in children's word recognition.

96 Measuring Word Recognition: The Looking-While-Listening Paradigm

97 Word recognition is traditionally studied in the looking-while-listening paradigm
98 (Fernald, Zangl, Portillo, & Marchman, 2008; alternatively referred to as the intermodal
99 preferential looking procedure, Hirsh-Pasek, Cauley, Golinkoff, & Gordon, 1987). In these
100 studies, infants listen to a sentence prompting a specific referent (e.g., *Look at the dog!*)
101 while viewing two images on the screen (e.g., an image of a dog – the target image – and an
102 image of a bird – the distractor image). Infants' word recognition is evaluated by how

103 quickly and accurately they fixate on the target image after hearing its label. Past research
104 has used this basic method to study a wide range of questions in language development. For
105 example, the looking-while-listening paradigm has been used to investigate early noun
106 knowledge, phonological representations of words, prediction during language processing, and
107 individual differences in language development (Bergelson & Swingley, 2012; Golinkoff, Ma,
108 Song, & Hirsh-Pasek, 2013; Lew-Williams & Fernald, 2007; Marchman et al., 2018; Swingley
109 & Aslin, 2002).

110 While this research has been fruitful in advancing understanding of early word
111 knowledge, fundamental questions remain. One central question is how to accurately capture
112 developmental change in the speed and accuracy of word recognition. There is ample
113 evidence demonstrating that infants become faster and more accurate in word recognition
114 over the first few years of life (e.g., Fernald, Pinto, Swingley, Weinberg, & McRoberts, 1998).
115 However, precisely measuring developmental increases in the speed and accuracy of word
116 recognition remains challenging due to the difficulty of distinguishing developmental changes
117 in word recognition skill from changes in knowledge of specific words. This problem is
118 particularly thorny in studies with young children, since the number of items that can be
119 tested within a single session is limited and items must be selected in an age-appropriate
120 manner (Peter et al., 2019). More broadly, key differences in the design choices (e.g., how
121 distractor items are selected) and analytic decisions (e.g., how the analysis window is defined)
122 between studies can obscure developmental change if not appropriately taken into account.

123 One approach to addressing these challenges is to conduct meta-analyses aggregating
124 effects across studies while testing for heterogeneity due to researcher choices (Bergmann et
125 al., 2018; Lewis et al., 2016). However, meta-analyses typically lack the granularity to
126 estimate participant-level and item-level variation or to model behavior beyond
127 coarse-grained effect size estimates. An alternative way to approach this challenge is to
128 aggregate trial-level data from smaller studies measuring word recognition with a wide range

129 of items and design choices into a large-scale dataset that can be analyzed using a unified
130 modeling approach. A sufficiently large dataset would allow researchers to estimate
131 developmental change in word recognition speed and accuracy while generalizing across
132 changes related to specific words or the design features of particular studies.

133 A related open theoretical question is understanding changes in children's word
134 recognition at the level of individual items. Looking-while-listening studies have been limited
135 in their ability to assess the development of specific words. One limitation is that studies
136 typically test only a small number of trials for each item, reducing power to precisely measure
137 the development of word-specific accuracy (DeBolt, Rhemtulla, & Oakes, 2020). A second
138 limitation is that target stimuli are often yoked with a narrow set of distractor stimuli (i.e., a
139 child sees a target with only one or two distractor stimuli over the course of an experiment),
140 leaving ambiguous whether accurate looking to a particular target word can be attributed to
141 children's recognition of the target word or their knowledge about the distractor.
142 Aggregating across many looking-while-listening studies has the potential to meet these
143 challenges by increasing the number of observations for specific items at different ages and by
144 increasing the size of the inventory of distractor stimuli that co-occur with each target.

145 Replicability and Reproducibility

146 A core challenge facing psychology in general, and the study of infant development in
147 particular, are threats to the replicability and reproducibility of core empirical results (Frank
148 et al., 2017; Nosek et al., 2022). In infant research, many studies are not adequately powered
149 to detect the main effects of interest (Bergmann et al., 2018). This issue is compounded by
150 low reliability in infant measures, often due to limits on the number of trials that can be
151 collected from an individual infant in an experimental session (Byers-Heinlein, Bergmann, &
152 Savalei, 2021). One hurdle to improving power in infant research is that it can be difficult to
153 develop a priori estimates of effect sizes and how specific design decisions (e.g., the number

154 of test trials) will impact power and reliability. Large-scale databases of infant behavior can
155 aid researchers in their decision-making by allowing them to directly test how different
156 design decisions affect power and reliability. For example, if a researcher is interested in
157 understanding how the number of test trials could impact the power and reliability of their
158 looking-while-listening design, a large-scale infant eye-tracking database would allow them to
159 simulate possible outcomes across a range of test trials, providing the basis for data-driven
160 design decisions.

161 In addition to threats to replicability, the field of infant development also faces
162 concerns about analytic reproducibility – the ability for researchers to arrive at the same
163 analytic conclusion reported in the original research article, given the same dataset. A recent
164 estimate based on studies published in a prominent cognitive science journal suggests that
165 analyses can remain difficult to reproduce, even when data are made available to other
166 research teams (Hardwicke et al., 2018). Aggregating data in centralized databases can aid
167 in improving reproducibility in several ways. First, building a large-scale database requires
168 defining a standardized data specification. Recent examples include the **brain imaging**
169 **data structure** (BIDS), an effort to specify a unified data format for neuroimaging
170 experiments (Gorgolewski et al., 2016), and the data formats associated with **ChildProject**,
171 for managing long-form at-home language recordings (Gautheron, Rochat, & Cristia, 2021).
172 Defining a data standard – in this case, for infant eye-tracking experiments – supports
173 reproducibility by guaranteeing that critical information will be available in openly shared
174 data and by making it easier for different research teams to understand the data structure.
175 Second, open databases make it easy for researchers to generate open and reproducible
176 analytic pipelines, both for individual studies and for analyses aggregating across datasets.
177 Creating open analytic pipelines across many datasets also serves a pedagogical purpose,
178 providing teaching examples illustrating how to implement analytic techniques used in
179 influential studies and how to conduct reproducible analyses with infant eye-tracking data.

180 Peekbank: An open database of developmental eye-tracking studies.

181 What all of these open challenges share is that they are difficult to address at the scale
182 of a single research lab or in a single study. To address this challenge, we developed
183 *Peekbank*, a flexible and reproducible interface to an open database of developmental
184 eye-tracking studies. The Peekbank project (a) collects a large set of eye-tracking datasets
185 on children’s word recognition, (b) introduces a data format and processing tools for
186 standardizing eye-tracking data across heterogeneous data sources, and (c) provides an
187 interface for accessing and analyzing the database. In the current paper, we introduce the
188 key components of the project and give an overview of the existing database. We then
189 provide two worked examples of how researchers can use Peekbank. In the first, we examine
190 a classic result in the word recognition literature, and in the second we aggregate data across
191 studies to investigate developmental trends in the recognition of individual words.

192 Design and Technical Approach**193 Database Framework**

194 One of the main challenges in compiling a large-scale eye-tracking database is the lack
195 of a shared data format: both labs and individual experiments can record their results in a
196 wide range of formats. For example, different experiments encode trial-level and
197 participant-level information in many different ways. Therefore, we have developed a
198 common tabular format to support analyses of all studies simultaneously.

199 As illustrated in Figure 1, the Peekbank framework consists of four main components:
200 (1) a set of tools to *convert* eye-tracking datasets into a unified format, (2) a relational
201 database populated with data in this unified format, (3) a set of tools to *retrieve* data from
202 this database, and (4) a web app (using the Shiny framework) for visualizing the data. These

203 components are supported by three packages. The `peekds` package (for the R language, R
204 Core Team, 2021) helps researchers convert existing datasets to use the standardized format
205 of the database. The `peekbank` module (Python) creates a database with the relational
206 schema and populates it with the standardized datasets produced by `peekds`. The database
207 is served through MySQL, an industry standard relational database server, which may be
208 accessed by a variety of programming languages, and can be hosted on one machine and
209 accessed by many others over the Internet. As is common in relational databases, records of
210 similar types (e.g., participants, trials, experiments, coded looks at each timepoint) are
211 grouped into tables, and records of various types are linked through numeric identifiers. The
212 `peekbankr` package (R) provides an application programming interface, or API, that offers
213 high-level abstractions for accessing the tabular data stored in Peekbank. Most users will
214 access data through this final package, in which case the details of data formatting,
215 processing, and the specifics of connecting to the database are abstracted away from the user.

216 Database Schema

217 The Peekbank database contains two major types of data: (1) metadata regarding
218 experiments, participants, and trials, and (2) time course looking data, detailing where a
219 child is looking on the screen at a given point in time (Fig. 2).

220 **Metadata.** Metadata can be separated into four parts: (1) participant-level
221 information (e.g., demographics), (2) experiment-level information (e.g., the type of eye
222 tracker used to collect the data), (3) session information (e.g. a participant's age for a
223 specific experimental session), and (4) trial information (e.g., which images or videos were
224 presented onscreen, and paired with which audio).

225 *Participant Information.*

226 All information about individual participants in Peekbank is completely de-identified

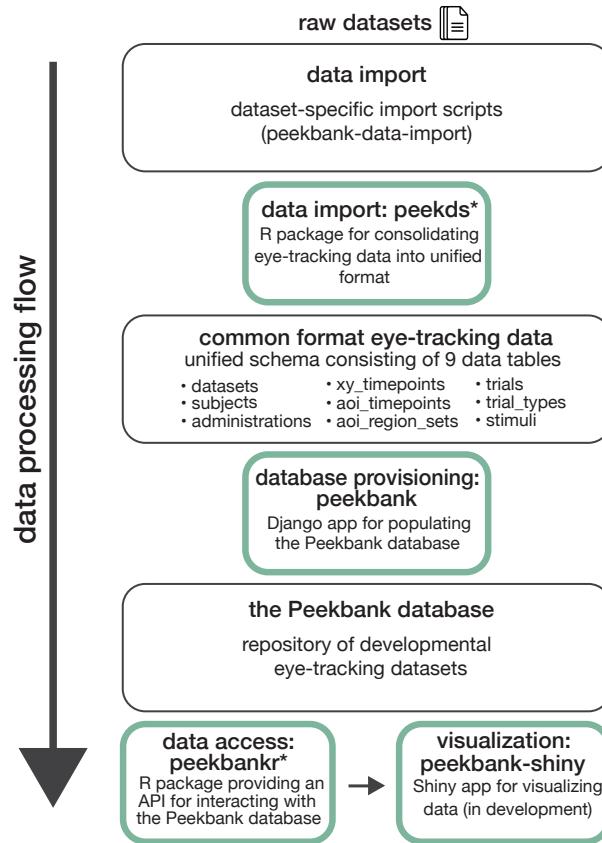


Figure 1. Overview of the Peekbank data ecosystem. Peekbank tools are highlighted in green. * indicates R packages introduced in this work.

under United States law, containing none of the key identifiers listed under the “Safe Harbor” standard for data de-identification. All participant-level linkages are made using anonymous participant identifiers.

Invariant information about individuals who participate in one or more studies (e.g., a participant’s first language) is recorded in the **subjects** table, while the **administrations** table contains information about each individual session in a given study (see Session Information, below). This division allows Peekbank to gracefully handle longitudinal designs: a single participant can complete multiple sessions and thus be associated with multiple administrations.

Participant-level data includes all participants who have experiment data. In general,

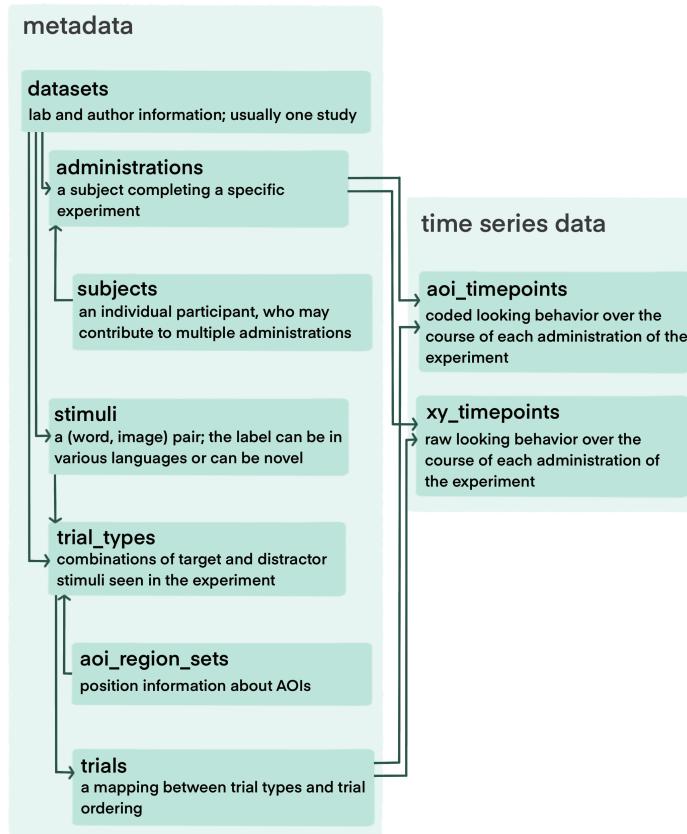


Figure 2. The Peekbank schema. Each darker rectangle represents a table in the relational database.

²³⁷ we include as many participants as possible in the database and leave it to end-users to
²³⁸ apply the appropriate exclusion criteria for their analysis.

²³⁹ ***Experiment Information.***

²⁴⁰ The **datasets** table includes information about the lab conducting the study and the
²⁴¹ relevant publications to cite regarding the data. In most cases, a dataset corresponds to a
²⁴² single study.

²⁴³ Information about the experimental design is split across the **trial_types** and
²⁴⁴ **stimuli** tables. The **trial_types** table encodes information about each trial *in the design*
²⁴⁵ *of the experiment*,¹ including the target stimulus and location (left vs. right), the distractor

¹ We note that the term *trial* is ambiguous and could be used to refer to both a particular combination of

246 stimulus and location, and the point of disambiguation for that trial. If a dataset used
247 automatic eye-tracking rather than manual coding, each trial type is additionally linked to a
248 set of area of interest (x, y) coordinates, encoded in the `aoi_region_sets` table. The
249 `trial_types` table links trial types to the `aoi_region_sets` table and the `trials` table.
250 Each trial_type record links to two records in the `stimuli` table, identified by the
251 `distractor_id` and the `target_id` fields.

252 Each record in the `stimuli` table is a (word, image) pair. In most experiments, there
253 is a one-to-one mapping between images and labels (e.g., each time an image of a dog
254 appears it is referred to as *dog*). For studies in which there are multiple potential labels per
255 image (e.g., *dog* and *chien* are both used to refer to an image of a dog), images can have
256 multiple rows in the `stimuli` table with unique labels. This structure is useful for studies on
257 synonymy or using multiple languages. It is also possible for an image to be associated with
258 a row with no label, if the image appears solely as a distractor (and thus its label is
259 ambiguous). For studies in which the same label refers to multiple images (e.g., the word *dog*
260 refers to an image of a dalmatian and a poodle), the same label can have multiple rows in
261 the `stimuli` table with unique images.

262 ***Session Information.***

263 The `administrations` table includes information about the participant or experiment
264 that may change between sessions of the same study, even for the same participant. This
265 includes the age of the participant, the coding method (eye-tracking vs. hand-coding), and
266 the properties of the monitor that was used.

267 ***Trial Information.***

268 The `trials` table includes information about a specific participant completing a

stimuli seen by many participants and a participant seeing that particular combination at a particular point in the experiment. We track the former in the `trial_types` table and the latter in the `trials` table.

269 specific instance of a trial type. This table links each record in the time course looking data
270 (described below) to the trial type and specifies the order of the trials seen by a specific
271 participant.

272 **Time course data.** Raw looking data is a series of looks to areas of interest (AOIs),
273 such as looks to the left or right of the screen, or to (x, y) coordinates on the experiment
274 screen, linked to points in time. For data generated by eye-trackers, we typically have (x, y)
275 coordinates at each time point, which we encode in the `xy_timepoints` table. These looks
276 are also recoded into AOIs according to the AOI coordinates in the `aoi_region_sets` table
277 using the `add_aois()` function in `peekds`, and encoded in the `aoi_timepoints` table. For
278 hand-coded data, we typically have a series of AOIs (i.e., looks to the left vs. right of the
279 screen), but lack information about exact gaze positions on-screen; in these cases the AOIs
280 are recoded into the categories in the Peekbank schema (target, distractor, other, and
281 missing) and encoded in the `aoi_timepoints` table; however, these datasets do not have any
282 corresponding data in the `xy_timepoints` table.

283 Typically, timepoints in the `xy_timepoints` table and `aoi_timepoints` table need to
284 be regularized to center each trial’s time around the point of disambiguation – such that 0 is
285 the time of target word onset in the trial (i.e., the beginning of *dog* in *Can you find the*
286 *dog?*). We re-centered timing information to the onset of the target label to facilitate
287 comparison of target label processing across all datasets.² If time values run throughout the
288 experiment rather than resetting to zero at the beginning of each trial, `rezero_times()` is
289 used to reset the time at each trial. After this, each trial’s times are centered around the
290 point of disambiguation using `normalize_times()`. When these steps are complete, the
291 time course is ready for resampling.

² While information preceding the onset of the target label in some datasets such as co-articulation cues (Mahr, McMillan, Saffran, Ellis Weismer, & Edwards, 2015) or adjectives (Fernald, Marchman, & Weisleder, 2013) can in principle disambiguate the target referent, we use a standardized point of disambiguation based on the onset of the label for the target referent. Onset times for other potentially disambiguating information (such as adjectives) can typically be recovered from the raw data provided on OSF.

To facilitate time course analysis and visualization across datasets, time course data must be resampled to a uniform sampling rate (i.e., such that every trial in every dataset has observations at the same time points). All data in the database is resampled to 40 Hz (observations every 25 ms), which represents a compromise between retaining fine-grained timing information from datasets with dense sampling rates (maximum sampling rate among current datasets: 500 Hz) while minimizing the possibility of introducing artifacts via resampling for datasets with lower sampling rates (minimum sampling rate for current datasets: 30 Hz). Further, 25 ms is a mathematically convenient interval for ensuring consistent resampling; we found that using 33.333 ms (30 Hz) as our interval simply introduced a large number of technical complexities. The resampling operation is accomplished using the `resample_times()` function. During the resampling process, we interpolate using constant interpolation, selecting for each interpolated timepoint the location for the earlier-observed time point in the original data for both `aoi_timepoints` and `xy_timepoints` data. Compared to linear interpolation (see e.g., Wass, Smith, & Johnson, 2013) – which fills segments of missing or unobserved time points by interpolating between the observed locations of timepoints at the beginning and end of the interpolated segment –, constant interpolation has the advantage that it is more conservative, in the sense that it does not introduce new look locations beyond those measured in the original data. One possible application of our new dataset is investigating the consequences of other interpolation functions for data analysis.

312 Processing, Validation, and Ingestion

313 Although Peekbank provides a common data format, the crux issue of populating the
314 database is the conversion of existing datasets to this format. Each dataset is imported via a
315 custom import script, which documents the process of conversion. Often various decisions
316 must be made in this import process (for example, how to characterize a particular trial type

317 within the options available in the Peekbank schema); these scripts provide a reproducible
318 record of this decision-making process. Our data import repository (available on GitHub at
319 <https://github.com/langcog/peekbank-data-import>) contains all of these scripts, links to
320 internal documentation on data import, and a set of generic import templates for different
321 formats.

322 Many of the specific operations involved in importing a dataset can be abstracted
323 across datasets. The `peekds` package offers a library of these functions. Once the data have
324 been extracted in a tabular form, the package also offers a validation function that checks
325 whether all tables have the required fields and data types expected by the database. In an
326 effort to double check the data quality and to make sure that no errors are made in the
327 importing script, we also typically perform a visual check of the import process, creating a
328 time course plot to replicate the results in the paper that first presented each dataset. Once
329 this plot has been created and checked for consistency and all tables pass our validation
330 functions, the processed dataset is ready for reprocessing into the database using the
331 `peekbank` library. This library applies additional data checks, and adds the data to the
332 MySQL database using the Django web framework.

333 To date, the import process has been carried out by the Peekbank team using data
334 offered by other research teams. There is no technical obstacle to data contributors also
335 providing an import script to facilitate contribution, though in practice creating these scripts
336 requires familiarity with both R scripting and the specific Peekbank schema; writing a first
337 import script can be somewhat time-consuming.

338 Current Data Sources

339 The database currently includes 20 looking-while-listening datasets comprising $N=1594$
340 total participants (Table 1). The current data represents a convenience sample of datasets

Table 1
Overview of the datasets in the current database.

Study Citation	Dataset name	N	Mean age (mos.)	Age range (mos.)	Method	Language
Adams et al., 2018	adams_marchman_2018	69	17.1	13–20	manual coding	English
Byers-Heinlein et al., 2017	byers-heinlein_2017	48	20.1	19–21	eye-tracking	English, French
Casillas et al., 2017	casillas_tseltal_2015	23	31.3	9–48	manual coding	Tseltal
Fernald et al., 2013	fmw_2013	80	20.0	17–26	manual coding	English
Frank et al., 2016	frank_tablet_2016	69	35.5	12–60	eye-tracking	English
Garrison et al., 2020	garrison_bergelson_2020	35	14.5	12–18	eye-tracking	English
Hurtado et al., 2007	xsectional_2007	49	23.8	15–37	manual coding	Spanish
Hurtado et al., 2008	hurtado_2008	76	21.0	17–27	manual coding	Spanish
Mahr et al., 2015	mahr_coartic	29	20.8	18–24	eye-tracking	English
Perry et al., 2017	perry_cowpig	45	20.5	19–22	manual coding	English
Pomper & Saffran, 2016	pomper_saffran_2016	60	44.3	41–47	manual coding	English
Pomper & Saffran, 2019	pomper_salientme	44	40.1	38–43	manual coding	English
Potter & Lew-Williams, unpub.	potter_canine	36	23.8	21–27	manual coding	English
Potter et al., 2019	potter_remix	44	22.6	18–29	manual coding	Spanish, English
Ronfard et al., 2021	ronfard_2021	40	20.0	18–24	manual coding	English
Swingley & Aslin, 2002	swingley_aslin_2002	50	15.1	14–16	manual coding	English
Weisleder & Fernald, 2013	weisleder_stl	29	21.6	18–27	manual coding	Spanish
Yurovsky & Frank, 2017	attword_processed	288	25.5	13–59	eye-tracking	English
Yurovsky et al., 2013	reflook_socword	435	33.6	12–70	eye-tracking	English
Yurovsky et al., unpub.	reflook_v4	45	34.2	11–60	eye-tracking	English

341 that were (a) datasets collected by or available to Peekbank team members, (b) made
 342 available to Peekbank after informal inquiry or (c) datasets that were openly available. Most
 343 datasets (14 out of 20 total) consist of data from monolingual native English speakers. They
 344 span a wide age spectrum with participants ranging from 9 to 70 months of age, and are
 345 balanced in terms of gender (47.30% female; 50.40% male; 2.30% unreported). The datasets
 346 vary across a number of design-related dimensions, and include studies using manually coded
 347 video recordings and automated eye-tracking methods (e.g., Tobii, EyeLink) to measure gaze
 348 behavior. All studies tested familiar items, but the database also includes 5 datasets that
 349 tested novel pseudo-words in addition to familiar words. Users interested in a subset of the
 350 data (e.g., only trials testing familiar words) can filter out unwanted trials using columns
 351 available in the schema (e.g., using the column `stimulus_novelty` in the `stimuli` table).

352 Versioning and Reproducibility

353 The content of Peekbank will change as we add additional datasets and revise previous
 354 ones. To facilitate reproducibility of analyses, we use a versioning system by which

355 successive releases are assigned a name reflecting the year and version, e.g., 2022.1. By
356 default, users will interact with the most recent version of the database available, though the
357 `peekbankr` API allows researchers to run analyses against any previous version of the
358 database. For users with intensive use-cases, each version of the database may be
359 downloaded as a compressed .sql file and installed on a local MySQL server.

360 Peekbank allows for fully reproducible analyses using our source data, but the goal is
361 not to reproduce precisely the analyses – or even the datasets – in the publications whose
362 data we archive. Because of our emphasis on a standardized data importing and formatting
363 pipeline, there may be minor discrepancies in the time course data that we archive compared
364 with those reported in original publications. Further, we archive all of the data that are
365 provided to us – including participants that might have been excluded in the original studies,
366 if these data are available – rather than attempting to reproduce specific exclusion criteria.
367 We hope that Peekbank can be used as a basis for comparing different exclusion and filtering
368 criteria – as such, an inclusive policy regarding importing all available data helps us provide
369 a broad base of data for investigating these decisions.

370 **Interfacing with Peekbank**

371 **Peekbankr**

372 The `peekbankr` API offers a way for users to access data from the database and
373 flexibly analyze it in R. The majority of API calls simply allow users to download tables (or
374 subsets of tables) from the database. In particular, the package offers the following functions:

- 375 • `connect_to_peekbank()` opens a connection with the Peekbank database to allow
376 tables to be downloaded with the following functions
377 • `get_datasets()` gives each dataset name and its citation information

- 378 • `get_subjects()` gives information about persistent participant identifiers (e.g., native
379 languages, sex)
- 380 • `get_administrations()` gives information about specific experimental
381 administrations (e.g., participant age, monitor size, gaze coding method)
- 382 • `get_stimuli()` gives information about word–image pairings that appeared in
383 experiments
- 384 • `get_trial_types()` gives information about pairings of stimuli that appeared in the
385 experiment (e.g., point of disambiguation, target and distractor stimuli, condition,
386 language)
- 387 • `get_trials()` gives the trial orderings for each administration, linking trial types to
388 the trial IDs used in time course data
- 389 • `get_aoi_region_sets()` gives coordinate regions for each area of interest (AOI)
390 linked to trial type IDs
- 391 • `get_xy_timepoints()` gives time course data for each participant’s looking behavior
392 in each trial, as (x, y) coordinates on the experiment monitor
- 393 • `get_aoi_timepoints()` gives time course data for each participant’s looking behavior
394 in each trial, coded into areas of interest

395 Once users have downloaded tables, they can be merged using `join` commands via their

396 linked IDs. A set of standard merges are shown below in the “Peekbank in Action” section;
397 these allow the common use-case of examining time course data and metadata jointly.

398 Because of the size of the XY and AOI data tables, downloading data across multiple

399 studies can be time-consuming. Many of the most common analyses of the Peekbank data
400 require downloading the `aoi_timepoints` table, thus we have put substantial work into
401 optimizing transfer times. In particular, `connect_to_peekbank` offers a data compression
402 option, and `get_aoi_timepoints` by default downloads time courses via a compressed
403 (run-length encoded) representation, which is then uncompressed on the client side. More

404 information about these options (including how to modify them) can be found in the
405 package documentation.

406 **Shiny App**

407 One goal of the Peekbank project is to allow a wide range of users to easily explore and
408 learn from the database. We therefore have created an interactive web application –
409 `peekbank-shiny` – that allows users to quickly and easily create informative visualizations
410 of individual datasets and aggregated data (<https://peekbank-shiny.com/>).

411 `peekbank-shiny` is built using Shiny, a software package for creating web apps for data
412 exploration with R, as well as the `peekbankr` package. All code for the Shiny app is publicly
413 available (<https://github.com/langcog/peekbank-shiny>). The Shiny app allows users to
414 create commonly used visualizations of looking-while-listening data, based on data from the
415 Peekbank database. Specifically, users can visualize:

- 416 1. the *time course of looking data* in a profile plot depicting infant target looking across
417 trial time
- 418 2. *overall accuracy*, defined as the proportion target looking within a specified analysis
419 window
- 420 3. *reaction times* in response to a target label, defined as how quickly participants shift
421 fixation to the target image on trials in which they were fixating on the distractor
422 image at onset of the target label
- 423 4. an *onset-contingent plot*, which shows the time course of participant looking as a
424 function of their look location at the onset of the target label

425 Users are given various customization options for each of these visualizations, e.g.,
426 choosing which datasets to include in the plots, controlling the age range of participants,
427 splitting the visualizations by age bins, and controlling the analysis window for time course

analyses. Plots are then updated in real time to reflect users' customization choices. A screenshot of the app is shown in Figure 3. The Shiny app thus allows users to quickly inspect basic properties of Peekbanks datasets and create reproducible visualizations without incurring any of the technical overhead required to access the database through R.

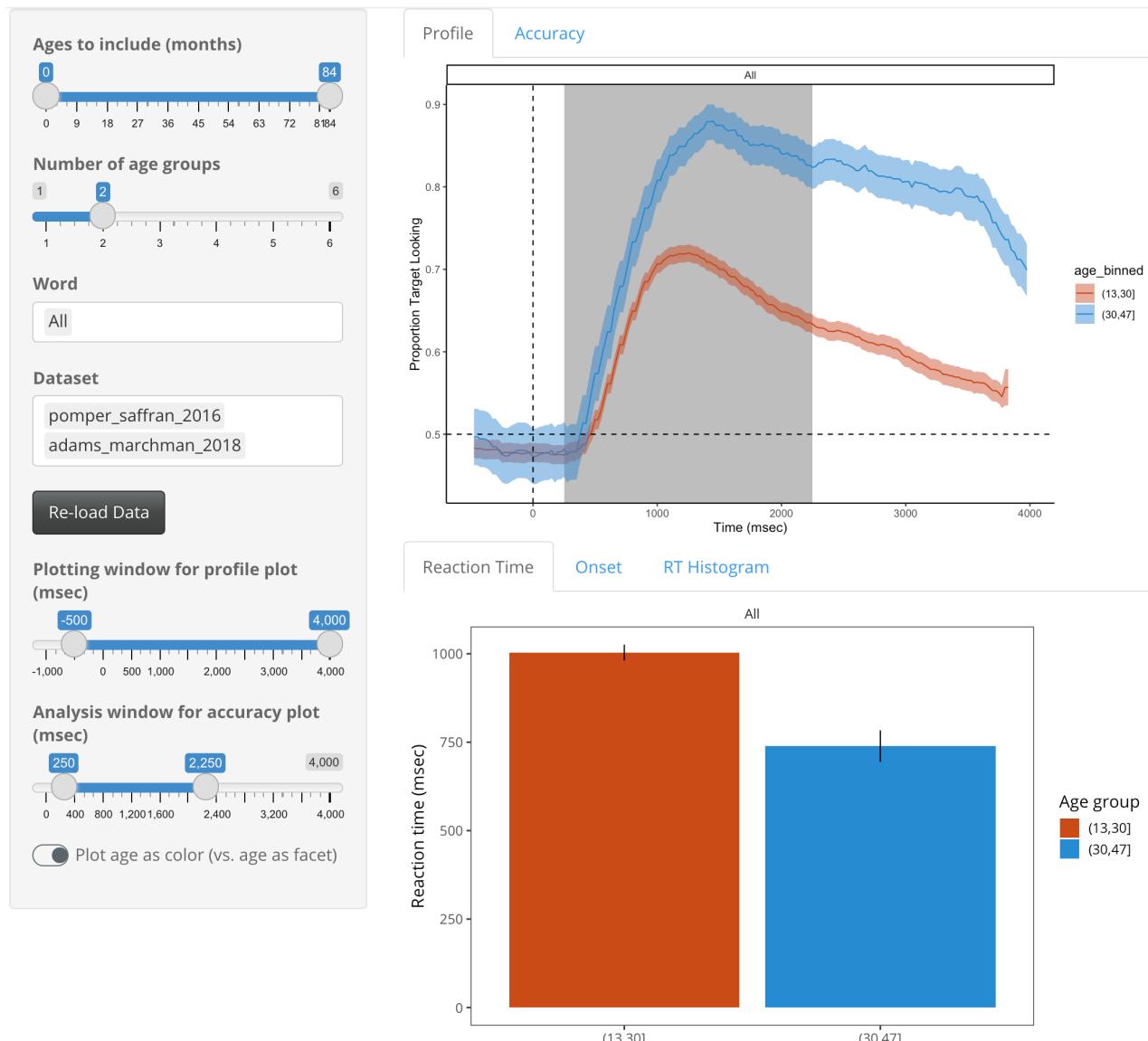


Figure 3. Screenshot of the Peekbank Shiny app, which shows a variety of standard analysis plots as a function of user-selected datasets, words, age ranges, and analysis windows. Shown here are mean reaction time and proportion target looking over time by age group for two selected datasets.

432 OSF site

433 In addition to the Peekbank database proper, all data is openly available on the
434 Peekbank OSF webpage (<https://osf.io/pr6wu/>). The OSF site also includes the original raw
435 data (both time series data and metadata, such as trial lists and participant logs) that was
436 obtained for each study and subsequently processed into the standardized Peekbank format.
437 Where available, the OSF page also includes additional information about the stimuli used in
438 each dataset, including in some instances the original stimulus sets (e.g., image and audio
439 files).

440 Peekbank in Action

441 In the following section, we provide examples of how users can access and analyze the
442 data in Peekbank. First, we provide an overview of some general properties of the datasets
443 in the database. We then demonstrate two potential use-cases for Peekbank data. In each
444 case, we provide sample code to demonstrate the ease of doing simple analyses using the
445 database. Our first example shows how we can investigate the findings of a classic study.
446 This type of investigation can be a very useful exercise for teaching students about best
447 practices for data analysis (e.g., Hardwicke et al., 2018) and also provides an easy way to
448 explore looking-while-listening time course data in a standardized format. Our second
449 example shows an exploration of developmental changes in the recognition of particular
450 words. Besides its theoretical interest (which we will explore more fully in subsequent work),
451 this type of analysis could in principle be used for optimizing the stimuli for new
452 experiments, especially as the Peekbank dataset grows and gains coverage over a greater
453 number of items. All analyses are conducted using R [Version 4.1.1; R Core Team (2021)]³

³ We, furthermore, used the R-packages *dplyr* [Version 1.0.7; Wickham, François, Henry, and Müller (2021)], *forcats* [Version 0.5.1; Wickham (2021a)], *ggplot2* [Version 3.3.5; Wickham (2016)], *ggthemes* [Version 4.2.4; Arnold (2021)], *here* [Version 1.0.1; Müller (2020)], *papaja* [Version 0.1.0.9997; Aust and Barth (2020)], *peekbankr* [Version 0.1.1.9002; Braginsky, MacDonald, and Frank (2021)], *purrr* [Version 0.3.4; Henry and

454 **General Descriptives**

Study Citation	Unique Items	Prop. Target	95% CI
Adams et al., 2018	8	0.65	[0.63, 0.67]
Byers-Heinlein et al., 2017	6	0.55	[0.52, 0.58]
Casillas et al., 2017	30	0.59	[0.54, 0.63]
Fernald et al., 2013	12	0.65	[0.63, 0.67]
Frank et al., 2016	24	0.64	[0.6, 0.68]
Garrison et al., 2020	87	0.60	[0.56, 0.64]
Hurtado et al., 2007	8	0.59	[0.55, 0.63]
Hurtado et al., 2008	12	0.61	[0.59, 0.63]
Mahr et al., 2015	10	0.71	[0.68, 0.74]
Perry et al., 2017	12	0.61	[0.58, 0.63]
Pomper & Saffran, 2016	40	0.77	[0.75, 0.8]
Pomper & Saffran, 2019	16	0.74	[0.72, 0.75]
Potter & Lew-Williams, unpub.	16	0.65	[0.61, 0.68]
Potter et al., 2019	8	0.63	[0.58, 0.67]
Ronfard et al., 2021	8	0.69	[0.65, 0.73]
Swingley & Aslin, 2002	22	0.57	[0.55, 0.59]
Weisleder & Fernald, 2013	12	0.63	[0.6, 0.66]
Yurovsky & Frank, 2017	6	0.63	[0.62, 0.65]
Yurovsky et al., 2013	6	0.61	[0.6, 0.63]
Yurovsky et al., unpub.	10	0.61	[0.57, 0.65]

Table 2

Average proportion target looking in each dataset.

455 One of the values of the uniform data format we use in Peekbank is the ease of
 456 providing cross-dataset descriptions that can give an overview of some of the general
 457 patterns found in our data. A first broad question is about the degree of accuracy in word
 458 recognition found across studies. In general, participants demonstrated robust, above-chance
 459 word recognition in each dataset (chance=0.5). Table 2 shows the average proportion of
 460 target looking within a standard critical window of 367-2000ms after the onset of the label
 461 for each dataset (Swingley & Aslin, 2002). Proportion target looking was generally higher for
 462 familiar words ($M = 0.66$, 95% CI = [0.65, 0.67], $n = 1543$) than for novel words learned
 463 during the experiment ($M = 0.59$, 95% CI = [0.58, 0.61], $n = 822$).

464 A second question of interest is about the variability across items (i.e., target labels)

Wickham (2020)], *readr* [Version 2.0.1; Wickham and Hester (2021)], *stringr* [Version 1.4.0; Wickham (2019)], *tibble* [Version 3.1.4; Müller and Wickham (2021)], *tidyR* [Version 1.1.3; Wickham (2021b)], *tidyverse* [Version 1.3.1; Wickham et al. (2019)], *tinylabels* (Barth, 2021), *viridis* [Version 0.6.1; Garnier et al. (2021a); Garnier et al. (2021b)], *viridisLite* [Version 0.4.0; Garnier et al. (2021b)], and *xtable* [Version 1.8.4; Dahl, Scott, Roosen, Magnusson, and Swinton (2019)].

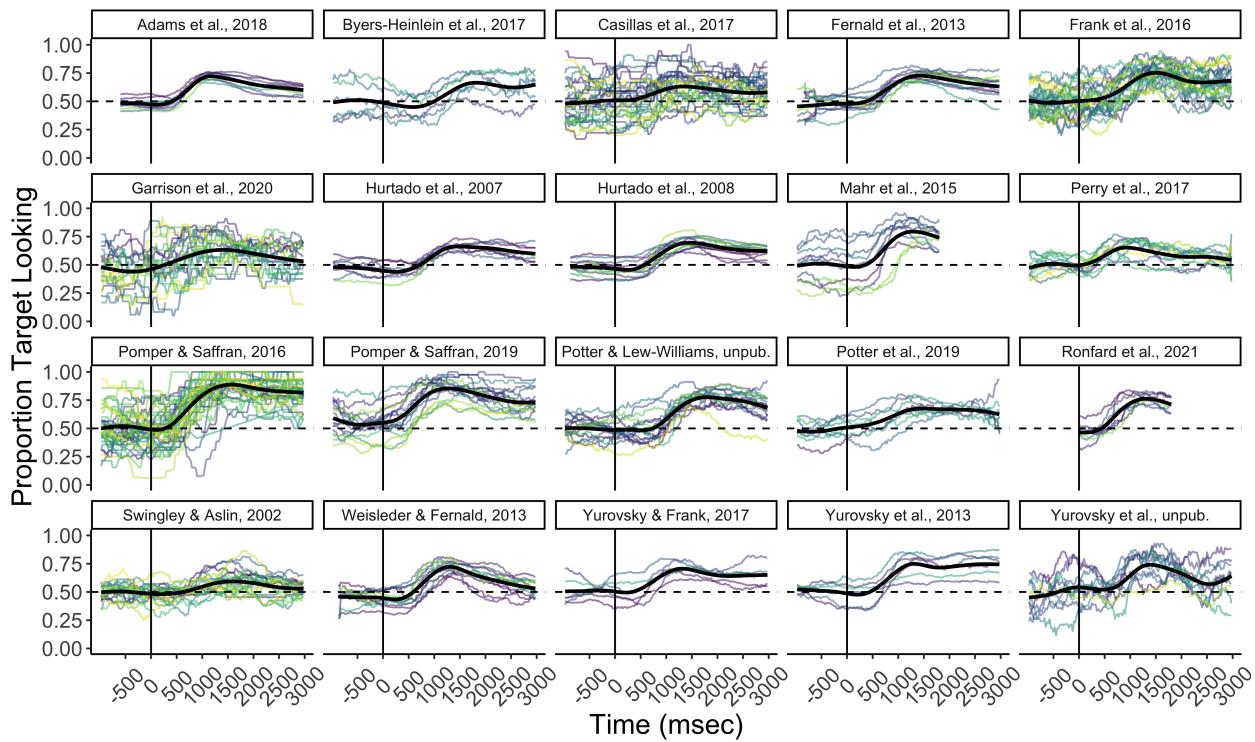


Figure 4. Item-level variability in proportion target looking within each dataset (chance=0.5). Time is centered on the onset of the target label (vertical line). Colored lines represent specific target labels. Black lines represent smoothed average fits based on a general additive model using cubic splines.

465 within specific studies. Some studies use a smaller set of items (e.g., 8 nouns, Adams et al.,
 466 2018) while others use dozens of different items (e.g., Garrison, Baudet, Breitfeld, Aberman,
 467 & Bergelson, 2020). Figure 4 gives an overview of the variability in proportion looking to the
 468 target item for individual words in each dataset. Although all datasets show a gradual rise in
 469 average proportion target looking over chance performance, the number of unique target
 470 labels and their associated accuracy vary widely across datasets.

471 Investigating prior findings: Swingley and Aslin (2002)

472 Swingley and Aslin (2002) investigated the specificity of 14-16-month-olds' word
 473 representations using the looking-while-listening paradigm, asking whether recognition would
 474 be slower and less accurate for mispronunciations, e.g. *opal* (mispronunciation) instead of

475 *apple* (correct pronunciation).⁴ In this short vignette, we show how easily the data in
 476 Peekbank can be used to visualize this result. Our goal here is not to provide a precise
 477 analytical reproduction of the analyses reported in the original paper, but rather to
 478 demonstrate the use of the Peekbank framework to analyze datasets of this type. In
 479 particular, because Peekbank uses a uniform data import standard, it is likely that there will
 480 be minor numerical discrepancies between analyses on Peekbank data and analyses that use
 481 another processing pipeline.

```
library(peekbankr)
aoi_timepoints <- get_aoi_timepoints(dataset_name = "swingley_aslin_2002")
administrations <- get_administrations(dataset_name = "swingley_aslin_2002")
trial_types <- get_trial_types(dataset_name = "swingley_aslin_2002")
trials <- get_trials(dataset_name = "swingley_aslin_2002")
```

482 We begin by retrieving the relevant tables from the database, `aoi_timepoints`,
 483 `administrations`, `trial_types`, and `trials`. As discussed above, each of these can be
 484 downloaded using a simple API call through `peekbankr`, which returns dataframes that
 485 include ID fields. These ID fields allow for easy joining of the data into a single dataframe
 486 containing all of the information necessary for the analysis.

```
swingley_data <- aoi_timepoints |>
  left_join(administrations) |>
  left_join(trials) |>
  left_join(trial_types) |>
  filter(condition != "filler") |>
  mutate(condition = if_else(condition == "cp", "Correct", "Mispronounced"))
```

487 As the code above shows, once the data are joined, condition information for each
 488 timepoint is present and so we can easily filter out filler trials and set up the conditions for
 489 further analysis.

⁴ The original paper investigated both close (e.g., *opple*, /apl/) and distant (e.g., *opal*, /opl/) mispronunciations. For simplicity, here we combine both mispronunciation conditions since the close vs. distant mispronunciation manipulation showed no effect in the original paper.

```

accuracies <- swingley_data |>
  group_by(condition, t_norm, administration_id) |>
  summarize(correct = sum(aoi == "target") /
    sum(aoi %in% c("target", "distractor"))) |>
  group_by(condition, t_norm) |>
  summarize(mean_correct = mean(correct),
            ci = 1.96 * sd(correct) / sqrt(n())))

```

490 The final step in our analysis is to create a summary dataframe using `dplyr`
 491 commands. We first group the data by timestep, participant, and condition and compute the
 492 proportion looking at the correct image. We then summarize again, averaging across
 493 participants, computing both means and 95% confidence intervals (via the approximation of
 494 1.96 times the standard error of the mean). The resulting dataframe can be used for
 495 visualization of the time course of looking.

496 Figure 5 shows the average time course of looking for the two conditions, as produced
 497 by the code above. Looks after the correctly pronounced noun appeared both faster
 498 (deviating from chance earlier) and more accurate (showing a higher asymptote). Overall,
 499 this example demonstrates the ability to produce this visualization in just a few lines of code.

500 Item analyses

501 A second use-case for Peekbank is to examine item-level variation in word recognition.
 502 Individual datasets rarely have enough statistical power to show reliable developmental
 503 differences within items. To illustrate the power of aggregating data across multiple datasets,
 504 we select the four words with the most data available across studies and ages (apple, book,
 505 dog, and frog) and show average recognition trajectories.

506 Our first step is to collect and join the data from the relevant tables including
 507 timepoint data, trial and stimulus data, and administration data (for participant ages). We
 508 join these into a single dataframe for easy manipulation; this dataframe is a common

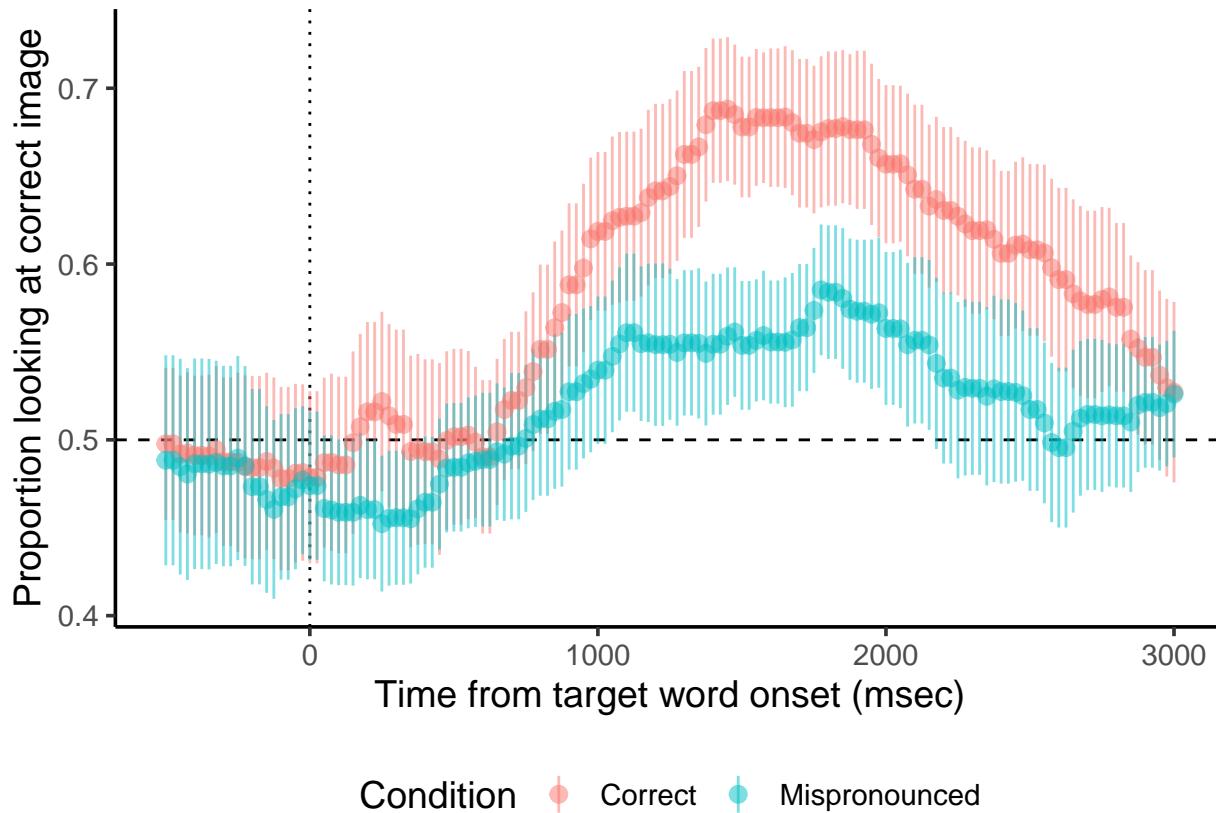


Figure 5. Proportion looking at the correct referent by time from the point of disambiguation (the onset of the target noun) in Swingley & Aslin (2002). Colors show the two pronunciation conditions; points give means and ranges show 95% confidence intervals. The dotted line shows the point of disambiguation and the dashed line shows chance performance.

509 starting point for analyses of item-level data.

```
all_aoi_timepoints <- get_aoi_timepoints()
all_stimuli <- get_stimuli()
all_administrations <- get_administrations()
all_trial_types <- get_trial_types()
all_trials <- get_trials()

aoi_data_joined <- all_aoi_timepoints |>
  right_join(all_administrations) |>
  right_join(all_trials) |>
```

```

right_join(all_trial_types) |>
  mutate(stimulus_id = target_id) |>
  right_join(all_stimuli) |>
  select(administration_id, english_stimulus_label, age, t_norm, aoi)

```

510 Next we select a set of four target words (chosen based on having more than 100
 511 children contributing data for each word across several one-year age groups). We create age
 512 groups, aggregate, and compute timepoint-by-timepoint confidence intervals using the z
 513 approximation.

```

target_words <- c("book", "dog", "frog", "apple")

target_word_data <- aoi_data_joined |>
  filter(english_stimulus_label %in% target_words) |>
  mutate(age_group = cut(age, breaks = seq(12, 48, 12))) |>
  filter(!is.na(age_group)) |>
  group_by(t_norm, administration_id, age_group, english_stimulus_label) |>
  summarise(correct = sum(aoi == "target") /
    sum(aoi %in% c("target", "distractor"))) |>
  group_by(t_norm, age_group, english_stimulus_label) |>
  summarise(ci = 1.96 * sd(correct, na.rm=TRUE) / sqrt(length(correct)),
            correct = mean(correct, na.rm=TRUE),
            n = n())

```

514 Finally, we plot the data as time courses split by age. Our plotting code is shown below
 515 (with styling commands removed for clarity). Figure 6 shows the resulting plot, with time
 516 courses for each of three (rather coarse) age bins. Although some baseline effects are visible
 517 across items, we still see clear and consistent increases in looking to the target, with the

518 increase appearing earlier and in many cases asymptoting at a higher level for older children.

519 This simple averaging approach is a proof-of-concept to demonstrate some of the
 520 potential of the Peekbank dataset. An eye-movement trajectory on an individual trial reflects
 521 myriad factors, including the age and ability of the child, the target and distractor stimuli on
 522 that trial, the position of the trial within the experiment, and the general parameters of the
 523 experiment (for example, stimulus timing, eye-tracker type and calibration, etc.). Although
 524 we often neglect these statistically in the analysis of individual experiments – for example,
 525 averaging across items and trial orders – they may lead to imprecision when we average
 526 across multiple studies in Peekbank. For example, studies with older children may use more
 527 difficult items or faster trial timing, leading to the impression that children’s abilities overall
 528 increase more slowly than they do in fact. Even in our example in Figure 6, we see hints of
 529 this confounding – for example, the low baseline looks to *apple* may be an artifact of an
 530 attractive distractor being paired with this item in one or two studies. In future work, we
 531 hope to introduce model-based analytic methods that use mixed effects regression to factor
 532 out study-level and individual-level variance in order to recover developmental effects more
 533 appropriately (see e.g., Zettersten et al., 2021 for a prototype of such an analysis).

```
ggplot(target_word_data,
       aes(x = t_norm, y = correct, col = age_group)) +
  geom_line() +
  geom_linerange(aes(ymin = correct - ci, ymax = correct + ci),
                 alpha = .2) +
  facet_wrap(~english_stimulus_label)
```

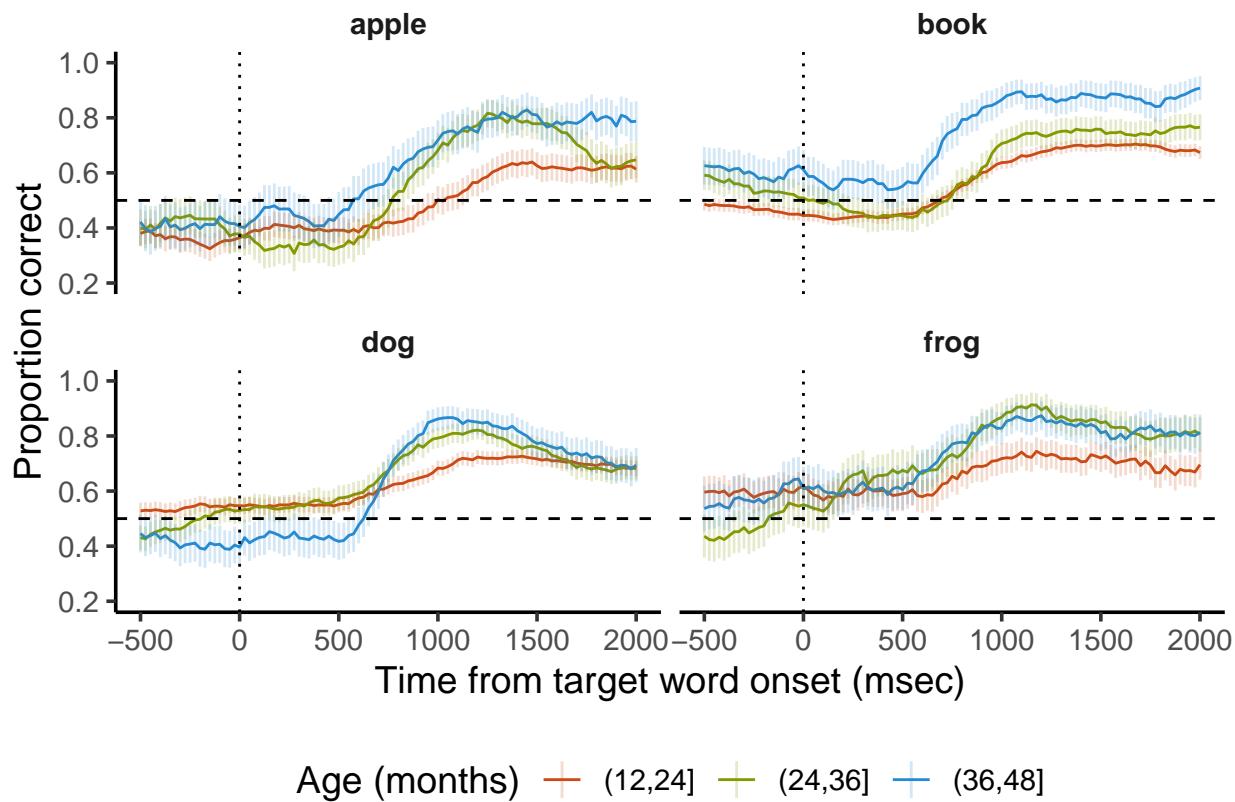


Figure 6. Time course plot for four well-represented target items in the Peekbank dataset, split by three age groups. Each line represents children’s average looking to the target image after the onset of the target label (dashed vertical line). Error bars represent 95% CIs.

534

Discussion

535 Theoretical progress in understanding child development requires rich datasets, but
 536 collecting child data is expensive, difficult, and time-intensive. Recent years have seen a
 537 growing effort to build open source tools and pool research efforts to meet the challenge of
 538 building a cumulative developmental science (Bergmann et al., 2018; Frank, Braginsky,
 539 Yurovsky, & Marchman, 2017; Sanchez et al., 2019; The ManyBabies Consortium, 2020).
 540 The Peekbank project expands on these efforts by building an infrastructure for aggregating
 541 eye-tracking data across studies, with a specific focus on the looking-while-listening
 542 paradigm. This paper presents an overview of the structure of the database, shows how users
 543 can access the database, and demonstrates how it can be used both to investigate prior
 544 experiments and to synthesize data across studies.

545 The current database has a number of limitations, particularly in the number and
546 diversity of datasets it contains. With 20 datasets currently available in the database,
547 idiosyncrasies of particular designs and condition manipulations still have a substantial
548 influence on the results of particular analyses, as discussed above in our item analysis
549 example. Expanding the set of distinct datasets will allow us to increase the number of
550 datasets that contain specific items, leading to more robust generalizations across the many
551 sources of variation that are confounded within studies (e.g., item set, participant age range,
552 and specific experimental parameters). A critical next step will be the development of
553 analytic models that take this structure into account in making generalizations across
554 datasets.

555 A second limitation stems from the fact that the database represents a convenience
556 sample of data readily available to the Peekbank team, which leads the database to be
557 relatively homogeneous in a number of key respects. First, the datasets primarily come from
558 labs that share similar theoretical perspectives and implement the looking-while-listening
559 method in similar ways. The current database is also limited by the relatively homogeneous
560 background of its participants, both with respect to language (almost entirely monolingual
561 native English speakers) and cultural background (Henrich, Heine, & Norenzayan, 2010;
562 Muthukrishna et al., 2020). Increasing the diversity of lab sources, participant backgrounds,
563 and languages will expand the scope of the generalizations we can form about child word
564 recognition, while also providing new opportunities for describing cross-lab, cross-cultural,
565 and cross-linguistic variation.

566 Towards the goal of expanding our database, we invite researchers to contribute their
567 data. On the Peekbank website we provide technical documentation for potential
568 contributors. Although we anticipate being involved in most new data imports, as discussed
569 above, our import process is transparently documented and the repository contains examples
570 for most commonly-used eye-trackers.

571 Contributing data to an open repository also can raise questions about participant

572 privacy. Potential contributors should consult with their local institutional review boards for

573 guidance on any challenges, but we do not foresee obstacles because of the de-identified

574 nature of the data. Under United States regulation, all data contributed to Peekbank are

575 considered de-identified and hence not considered “human subjects data”; hence, institutional

576 review boards should not regulate this contribution process. Under the European Union’s

577 Generalized Data Protection Regulation (GDPR), labs may need to take special care to

578 provide a separate set of participant identifiers that can never be re-linked to their own

579 internal records.

580 While the current database is focused on studies of word recognition, the tools and

581 infrastructure developed in the project can in principle be used to accommodate any

582 eye-tracking paradigm, opening up new avenues for insights into cognitive development.

583 Gaze behavior has been at the core of many key advances in our understanding of infant

584 cognition (Aslin, 2007; Baillargeon, Spelke, & Wasserman, 1985; Bergelson & Swingley, 2012;

585 Fantz, 1963; Liu, Ullman, Tenenbaum, & Spelke, 2017; Quinn, Eimas, & Rosenkrantz, 1993).

586 Aggregating large datasets of infant looking behavior in a single, openly-accessible format

587 promises to bring a fuller picture of infant cognitive development into view.

588

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