NEBULA101: an open dataset for the study of language aptitude in behaviour, brain structure and function

Alessandra Rampinini* [a,e], Irene Balboni* [a-d-e], Olga Kepinska[b,c], Raphael Berthele [d,e], Narly Golestani [a,b,c,e]

[a] Department of Psychology, Faculty of Psychology and Education Science, University of Geneva, Geneva, Switzerland

[b] Vienna Cognitive Science Hub, University of Vienna, Vienna, Austria.

[c] Department of Behavioural and Cognitive Biology, Faculty of Life Sciences, University of Vienna, Vienna, Austria.

[d] Institute of Multilingualism, University of Fribourg, Fribourg, Switzerland.

[e] National Centre for Competence in Research Evolving Language, Switzerland.

Abstract

This paper introduces the "NEBULA101 - Neuro-behavioural Understanding of Language Aptitude" dataset, which comprises behavioural and brain imaging data from 101 healthy adults to examine individual differences in language and cognition. Human language, a multifaceted behaviour, varies significantly among individuals, at different processing levels. Recent advances in cognitive science have embraced an integrated approach, combining behavioural and brain studies to explore these differences comprehensively. The NEBULA101 dataset offers brain structural, diffusion-weighted, task-based and resting-state MRI data, alongside extensive linguistic and non-linguistic behavioural measures to explore the complex interaction of language and cognition in a highly multilingual sample. By sharing this multimodal dataset, we hope to promote research on the neuroscience of language, cognition and multilingualism, enabling the field to deepen its understanding of the multivariate panorama of individual differences and ultimately contributing to open science.

Background & Summary

Individual differences in language. This paper describes the "NEBULA101 - Neuro-behavioural Understanding of Language Aptitude" dataset. The dataset collects behavioural and brain imaging data of 101 healthy adults for the study of individual differences in language and cognition.

Human language is a complex behaviour, and crucial to understanding its workings is the fact that individuals differ in the way they manifest this and other cognitive skills[1]. The science of individual differences has expanded in recent years, thanks to a more integrated, less modular take on cognition: by combining the study of behaviour and the brain in a deep phenotyping approach, mindful of individual differences, researchers can gain a more comprehensive understanding of complex cognitive functions, language included[2].

Language aptitude. One aspect of language shown to display large individual differences is language

aptitude. Language aptitude was originally proposed to explain why some people display remarkable

abilities when learning additional languages [3–5]. We use the term *additional* language for any language that

is not the (or one of the) individual's *first* language(s), i.e., language(s) to which they were exposed from

birth. The term "additional" thus includes *second languages* (e.g. in the context of migration), *foreign*

languages (e.g. in classroom learning), or *third/additional languages* of multilinguals who use more than

two.

Initially, researchers viewed language aptitude as a stable trait, comprising phonetic coding,

grammatical sensitivity, inductive learning, and rote learning abilities. These skills were seen as

componential, in the climate of the first cognitive revolution, where the brain-mind-behaviour interaction

was seen as the execution of algorithms operating within cognitive modules[6]: in this view, phonetic coding

was at the core of production and perception of speech sounds, grammatical sensitivity underlay the capacity

to identify structure in language (i.e. morphosyntax), inductive learning supported the generalisation of

language rules from the input, and rote learning skills the construction of vocabulary via routinary use.

Research has since suggested a more parsimonious structure, combining grammatical sensitivity and

inductive learning into a global language analytic ability[7].

Nowadays, with advances in neuroscience and experimental psychology methods, the language

aptitude construct overall still describes a set of skills operating *across* the hierarchy of all language

components, from lower to higher levels of complexity. What has changed is the way we conceptualise

language itself, which has also changed the way we view language aptitude: in the second wave of the

cognitive revolution, thanks to usage-based linguistics and neural-network psychology, language is seen as

a complex adaptive system, rather than a set of rigid structures and fixed operations[8], and one of the

interacting components of human cognition, rather than an isolated function. Language aptitude is part of

this dynamic: it can vary with age[9–11] or with multilingual experience (likely increasing meta-linguistic

awareness[12–14], with patterns yet to be clarified[15,16]), and ultimately, with cognition more generally. In this

context, we view the brain as a network of interrelated functions giving rise to complex behaviours, including

language: to this end, this dataset comprises extensive general and domain-specific cognition tests, together

with brain measures. The underlying concept of an integrated mind, arising from an integrated brain, is at

the core of our methodological choice and of our first exploration of these data with graph theoretical

methods[17].

Mindful that any quantitative analysis will always require some degree of operationalisation (i.e. to

derive scores from tests, we need to identify components of scientific constructs that such tests might tap

into), our position is that this view of language as a dynamic system branching out and connecting to more

general mechanisms holds promise for better understanding language itself and the human cognitive system

more generally.

The importance of research on the multilingual brain. The views expressed above are pivotal to the study of
multilingualism and its behavioural and brain dynamics. In today's globalised society multilingualism and
multicompetence (i.e. using and knowing multiple languages) are becoming normalised[18.] There is, however,
a theoretical issue with defining a sociolinguistic construct such as that of any "-lingualism", due to its
multidimensionality[19], without incurring in stereotypes (e.g. the definition of *native-speakerism*) or
perpetrating problematic practices, such as that of selecting certain profiles[20] or matching "the unmatchable"
based on intrinsically tricky parameters (e.g. the number of languages[21]). Nonetheless, we need to better
define these concepts, as they constitute important characteristics of our present-day society. Knowing and
using multiple languages demands a fundamental cognitive (re)organisation[18], with several psycho-
neurobiological correlates that are somewhat hard to reconcile in a comprehensive view[22]. Therefore, we
must seek to imagine language competence and use in newer, more naturalistic, and multidimensional ways
to better understand their influence on the brains (and lives) of language users.

To this aim, we believe that multimodal datasets with rich phenotypical information, such as the one
presented here, are a step forward in this direction. Language aptitude, viewed as embedded in (and
interacting with) cognition might be one of the engines driving multilingualism (in its many forms), and
understanding its underpinnings might ultimately influence the way we view, use, and even teach
language(s). In this context, Switzerland holds a special status as a country with four official languages and
as a destination for expatriates from all over the world, including users of Minority, Indigenous, Non-
standard, and Dialect (MIND) varieties. These languages have often been disregarded not only by
researchers but by users themselves, who might not even recognise their multilingual status when knowing
an additional MIND language[20]. However, these *do* appear in the multicompetent panorama of many of our
participants, when asked explicitly (see Figure 2).

These issues have recently been tackled in work calling for a more diverse view of cognitive science
in general[23] and neurolinguistics[24] in particular, and underlining the contribution *non-English* (Romance[25] ,
non-Indo-European[26] as well as MIND [20,21]) languages to the field. The ongoing discourse on language
policies and teaching[27–29], as well as the thriving field of instructed language learning research, especially in
the Swiss context[30], are just a part of the puzzle. What is still lacking in the psycho- and neurolinguistics of
multilingualism is a naturalistic perspective rooted in the brain itself, also likely due to the intrinsic difficulty
of quantifying language use in a nonparametric and dynamic way. Humans, however, possess a skill,
language, that varies at all levels, challenging us to consider the remarkable plasticity of advanced human
abilities by harnessing language diversity as a tool for advancing cognitive science[31]. Thus, while leveraging
French knowledge and fluency of our participants - a condition necessary for performance comparability -
this dataset also captures and documents their linguistic diversity and multicompetence. This comprehensive
documentation has the potential to facilitate investigations into how such diversity relates to both the
phenotypical (behavioural) and endophenotypical (brain structural and functional) characteristics of
individuals.

Aptitude for language(s) and individual differences in other domains of cognition. The construct of language
aptitude was developed in the domain of foreign language learning, as explained, but the idea that aptitude
only manifests in foreign languages has now been surpassed, since individual differences can be observed
in first language skills too, even if it is harder to pinpoint them and isolate them from experiential factors[32].
Moreover, such individual differences might co-exist more globally with individual differences in other
domains of cognition[17,33], giving rise to "neurocognitive profiles" of language aptitude involving the
mnemonic domain, fluid reasoning, auditory abilities, and even musicality[34]. It is therefore relevant to ask
ourselves whether language, in this modernised and dynamic view, is part of a positive manifold [35,36]
originating from the beneficial interactions between cognitive processes, as we proposed in a recent analysis
of part of these data[17]. Both the stable and malleable (or *plastic*) features of the human cognitive system are
fundamental to such interactions.

Stability and malleability, predisposition and experience. This dataset presents cross-sectional data.
Nonetheless, to understand the nature of the language (aptitude) construct, it is important to consider it as
the result of a complex interaction between stable traits and malleable states, the first ascribed to genetic
predisposition and the latter to environmental influences (what we generally define as "experience")[37–39].
While we provide a *snapshot* of individual profiles at a given moment in time, in order to formulate relevant
questions on (and interpretations of) these data, it is important to remember that the observed measures arise
from both stable traits and from malleable skills, or states[40]. Cognitive activity[41–44], including language
learning and use[45–48], and brain *plasticity* are intrinsically related. Changes in regional activity, network
connectivity, or morphology, arising from underlying molecular, neurobiochemical and other changes[41],
subtend this state of malleability in brain function and structure[49], both during development and in skill
learning. However, brain functional and structural architecture is also highly polygenic (i.e. controlled by
the complex interaction of multiple genes, which in turn are expressed in multiple variants across
individuals), and in addition, different cortical loci and tissue features (thickness, surface area) are affected
to different degrees by genetics[50]. Further, genetic factors also likely influence the degree to which
neuroplasticity manifests[51].

The open science of language aptitude. Language aptitude has come a long way since its study was confined
to the foreign language classroom, and research restricted to military and government access[52]. The construct
now encompasses some of the most promising avenues for research on the human cognitive system more
generally: the roles of predisposition and experience, the nature of neuroplasticity and the integrated and
multivariate organisation of cognitive domains. Because of its relevance for studying questions on language
and cognition more generally (as well as investigating the role of the environment, e.g. the experience of
multilingualism), this development in the conceptualisation of language aptitude is of particular interest to
the ever-growing world of Findable, Accessible, Interoperable, and Reusable (FAIR) neuroscience data[53],
which has faced a substantial growth in the last 15 years[54], and possibly as many challenges[55,56]. FAIR
principles are accelerating our comprehension of the human brain[57]. Concurrently, the existence of

standardised protocols such as the Brain Imaging Data Structure (BIDS)[58] supports this recent strive towards

data and code accessibility, ensuring that data is organized and its analysis reproducible, enabling more

efficient and effective use of shared datasets and streamlining data preparation, thus allowing scientists to

focus more on discovery.

Mindful that the panorama of shared neuroimaging data is vast and that datasets might be scattered

around repositories of the commercial and institutional type, focusing on either a specific population

(clinical, paediatric) or modality[59], we searched for openly available data similar to ours (MRI and

behavioural data from healthy adult participants focusing on language) on OpenNeuro[60], one of the most

recent, easy to access and growing neuroimaging databases. At the time of writing, a search on

http://openneuro.org for BIDS-valid MRI raw/derivative datasets with more than 50 healthy adults, including

the keyword 'language' and accompanied by a published or pre-registered data descriptor, yields the

following entries (Table 1), with two of the datasets (ds004215, ds000243) being listed but having no

relationship with language (and thus not being included in the below table).

OpenNeuro Accession number	Dataset name and reference	Modality	Number of participants	Complete dataset (all participants-measures)	Associated tasks/measures	Main topic
ds003481[61]	A Dataset to Study Pragmatic Language and Its Underlying Cognitive Processes[62]	sMRI, fMRI, behaviour	145	No	• Comprehension of metaphorical phrases and proverbs • Recognition and emotional categorisation of speech acts • Lexical-semantic processing	Language pragmatics
ds004765[63]	Relationship between resting state functional connectivity and reading-related behavioural measures in 69 adults[64]	sMRI, fMRI (resting state), DWI	69	Yes	• Word/nonword reading • Spelling • Lexical decision • Spoonerisms • Rapid automatised naming • Non-word repetition • Vocabulary	Reading
ds002382[65]	Age-related differences in auditory cortex activity during spoken word recognition[66]*	sMRI, fMRI	61	Yes	• Word listening • Word repetition	Word processing
ds004285[67]	Listening task*	sMRI, fMRI	78	Yes	• Word repetition	Word processing
ds004073[68]	Comparing language lateralisation using fMRI and fTCD[69]	fMRI, functional transcranial doppler sonography (fTCD)	51	No (minimally missing information)	• Word generation • Sentence generation • Phonological decision • Word comprehension • Sentence comprehension • Syntactic decision	Language lateralisation
ds001747[70]	Exploring the Resting State Neural Activity of Monolinguals and Late and Early Bilinguals[71]	sMRI, fMRI (resting state)	92	Yes	• L1 and L2 (where applicable) proficiency • Language background	Bilingualism
ds001796[72]	Bilingualism and the brain**	sMRI, fMRI (task-based), fMRI (resting state), DWI	64	Yes	• Flanker task	Bilingualism
ds002345[73]	Narratives: fMRI data for evaluating models of naturalistic	sMRI, fMRI	345	No	• Naturalistic story listening	Language perception

	language comprehension[74]					
ds003643[75]	Le Petit Prince: A multilingual fMRI corpus using ecological stimuli[76]	sMRI, fMRI	112	Yes	• Naturalistic story listening	Language perception

Table 1. Available neuroimaging datasets on language with more than 50 healthy participants, validated in BIDS. (*)Possibly
including or revising ds002382 but no information on dataset version is provided. (**) No data descriptor available.

A similar, broad search on Google Scholar for data descriptors associated with openly available
language datasets ("open AND mri dataset AND language") yielded more datasets: the MOUS (Mother Of
Unification Studies)[77], a dataset comprising sMRI, DWI, resting state and task-based fMRI and MEG in 204
participants assigned to either sentence reading or listening; the Alice dataset[78], where 75 participants
listened to the same chapter of Alice in Wonderland during fMRI or during EEG. A third dataset presents a
large quantity of data collected in fewer participants with state-of-the-art technology: this is the high-
resolution, 7T fMRI Forrest Gump database[79], during which participants watched the 'Forrest Gump' movie.
This data can be used to study naturalistic language processing, even though the study did not primarily
focus on language. Then, the search yielded the LanA dataset, a probabilistic language atlas derived from
brain data in more than 800 individuals[80]. Finally, the search also yielded two speech production datasets of
vocal tract MRI[81,82].

At the behavioural level there is more heterogeneity in the type and quantity of shared data[59]. The
origins of such heterogeneity have been discussed for quite some time. In a Nature Neuroscience perspective
article from ten years ago, the challenges linked to standardisation and accessibility of behavioural data were
already discussed[83]. Importantly, behaviour was defined as a complex, highly dimensional, dynamic, and
interconnected phenomenon without distinct separable scales, and this was discussed as being one of the
leading causes for lack of standardisation and scarce FAIR compliance. The authors insisted on its
foundational and unifying nature, and called for improved standards: "Behaviour (…) is the principal
function of the brain. (…) Copious, quantitative and open behavioural data has the potential (…) to solidify
the foundations of other [disciplines], including neuroscience" [83] (p.1455). However, even in the cited
datasets, which represent timely and relevant steps towards accessibility of neuroscience data, the
phenotypical (behavioural) information being shared beyond demographics is still relatively less prominent
than the endophenotypical (neural) data: oftentimes, only data from the behavioural tasks being performed
during fMRI are available. Moreover, even when behaviour was tested outside the scanner, if the original
project did not focus on *both* phenotypical and endophenotypical data and on a specific topic (such as, in the
cited examples, pragmatics[84] or reading[64]), the accompanying behavioural information is scarce. Two
notable, recent exceptions in the field of individual differences in language are represented by a behavioural
dataset including 33 measures from 112 adult Dutch speakers[85], and by recent work by Berthele and
colleagues in children[86], both representing an important milestone for shared behavioural data.

Given the lack of standardisation in the way we administer behavioural tasks outside the scanner,
compared to fMRI task delivery, it seems daunting to force the structure of behavioural paradigms and log
files produced by a plethora of software and online platforms to accommodate information within the

structure required by the BIDS standard, a difficulty that we encountered in our work, as well. Some
initiatives, such as the Behaverse project[85], are proposing data structures which, they claim, can
accommodate phenotypical information better than what BIDS can do. However, it is important to note that
any individual differences dataset including both phenotypical and endophenotypical information will have
the added strength of multimodality, and to date, BIDS is the only data structure protocol that can
accommodate *both* in a relatively straightforward way, even if it requires extra (and sometimes *post-hoc*)
work to prepare the materials intended to be shared. Finally, we must note that when sharing mixed raw and
derivative datasets, behavioural data can be easily included as derivatives, these having a more liberal
structure, lifting from the end-user the load of reprocessing and calculating basic scores starting from item-
level data. This is the route we chose for this dataset, including raw (minimally processed) phenotypic and
behavioural data with varying underlying structures, coming from questionnaires and tasks respectively,
crucially accompanied by their derivate scores.

In sum, the NEBULA101 dataset aims to promote the study of individual differences in language
to better understand a multivariate cognitive system, via the sharing of a truly multimodal dataset in an
adequately sized participant sample[87]. We provide sMRI, DWI, task-based and resting state fMRI in 101
individuals, together with broad phenotypic and behavioural data on linguistic but also on non-linguistic,
domain general and domain specific tasks, including cognitive and perceptuomotor tasks. This includes
measures of all language aptitude components from phonetics to syntax, measures of reading and reading
mediators (e.g. phonological awareness), domain-general cognitive skills, numerical processing, musicality
and musical experience and rich multilingual language experience measures. By providing these data to the
public domain, we hope to contribute new discoveries to the over-arching construct of language (aptitude),
embracing the components of individual behavioural and neural phenotypes as widely as possible.

Data types. Behaviour is the phenotype that can be related to, or unify, genetics, neural architecture, neural
activity, body structure, physical limitations, and environmental factors[83]. Given the importance of
behavioural data in exploring individual differences at the neural level, this dataset includes 28 scores derived
from 8 questionnaires, and 74 behavioural measures derived from 21 tasks. Functional neuroimaging data
provide information about the neural correlates of cognitive processes, allowing to elucidate how the brain
supports specific behaviours and skills. Here, we provide resting-state and task-based functional imaging
(fMRI) data, the latter obtained during a language localiser[88]. Finally, NEBULA101 also includes anatomical
T1-weighted and diffusion-weighted (DWI) imaging of the brain, which will allow to study brain anatomy
and white-matter structural connectivity, to shed light on the brain structural correlates of linguistic
behaviours, aptitudes and experiences.
An overview of all measures provided in the NEBULA101 dataset is provided in Table 2. In the
Supplementary Information file, Table S1 contains more details specific to the version of the tests used in
this dataset, such as any adaptations, modality of administration and derivate scores.

Test	Modality	Construct	Derived score	Reference
Language Experience and Proficiency Questionnaire (LEAP-Q)	Questionnaire	Multilingual language experience.	Entropy	Marian et al., 2007[64]
Code Switching questionnaire	Questionnaire	Code switching habits.	Voluntary switching Involuntary switching	Rodriguez-Fornells et al., 2012[89]
Motivational Factors Questionnaire (MFQ)	Questionnaire	Motivation and attitude towards foreign languages (FL).	Ideal foreign language self Instrumentality International contact Foreign language interest Foreign language anxiety Foreign language confidence Milieu Usage willingness of foreign languages	Ryan, 2008[90] Thompson & Lee, 2018[91]
Adult Reading History Questionnaire (AHRQ)	Questionnaire	Reading history.	Reading history score	Lefly & Pennington, 2000[92]
Internal Representations Questionnaire (IRQ)	Questionnaire	Modes of internal reasoning.	Manipulation Factor Orthographic Factor Verbal Factor Visual Factor	Roebuck & Lupyan, 2020[93]
Music Use and Background Questionnaire (MUSEBAQ)	Questionnaire	Music training, capacity, preferences, and motivations.	Index of music training Index of music listening Index of musical instrument playing Cognitive and Emotional Regulation Social Connection Engaged Production Dance Physical Exercise	Chin et al., 2018[94]
Barratt's Simplified Measure of Socioeconomic Status (BSMSS)	Questionnaire	Socioeconomic status.	Barratt's Simplified Measure of Socioeconomic Status	Barratt, 2006[95] Rakesh & Whittle, 2021[96]
Edinburgh Handedness Inventory (EHI)	Questionnaire	Handedness.	Handedness score	Oldfield, 1971[97] Nedjar et al., 1989[98]
Artgram	Behavioural task	Language analytic abilities / Morphosyntax.	Accuracy RT	Developed in-house.
Modern Language Aptitude Test 5 (MLAT5)	Behavioural task	Rote learning.	Accuracy RT	Stansfield, 1989[99]
Farsi uvular Production Task	Behavioural task	Foreign sound production.	Accuracy	Golestani & Pallier, 2007[100]
Hindi Dental Retroflex Contrast	Behavioural task	Phonological categorisation/discrimination.	Accuracy	Golestani et al., 2002[101]
Brocanto	Behavioural task	Language analytic abilities / Pattern recognition.	Accuracy RT (For each block 1-3 and overall)	Kepinska et al., 2017[102] Opitz and Friederici, 2003[103]

Raven's Advanced Progressive Matrices (APM)	Behavioural task	Non-verbal intelligence.	Accuracy RT	Raven, 1998[104]
Corsi block	Behavioural task	Visuospatial memory	Forward span Backward span Total span	Corsi, 1972[105] Arce & McMullen, 2021[106]
Digit Span	Behavioural task	Auditory working memory	Forward span Backward span Total span	Wechsler, 1955[107] Ryan et al., 2014[108] Conway et al., 2005[109]
Revised Tempo Test	Behavioural task	Arithmetic abilities	Accuracy RT (Additions, subtractions, overall)	Bellon et al., 2022[110]
Advanced Measures of Music Audiation (AMMA)	Behavioural task	Music audiation, musicality, musical aptitude.	Tonal score Rhythm score Total score	Gordon, 1989[111]
Attention Network Test - Interaction (ANT-I)	Behavioural task	Attention networks: executive control, alerting, orienting.	Alerting gain Orienting gain Reorienting gain Inhibition gain	Callejas et al., 2005[112]
California Verbal Learning Task (CVLT)	Behavioural task	Verbal working memory	Accuracy immediate recall Accuracy immediate and short-term recall Accuracy long-term recognition RT long term recognition	Deweer et al., 2008[113]
Finger tapping Test	Behavioural task	Fine motor skills	Average finger tapping score dominant hand overall Average finger tapping score nondominant hand overall Average finger tapping score of dominant/nondominant overall (Calculated over blocks)	Strauss et al., 2006[114] Ashendorf et al., 2015[115]
Purdue Pegboard Test	Behavioural task	Fine motor skills	Accuracy, dominant hand Accuracy, nondominant hand Accuracy, simultaneous hands Accuracy, assembly task Average accuracy, dominant-nondominant-simultaneous hands (Calculated over blocks)	Tiffin & Asher, 1948[116]
Rapid Automatised Naming (RAN)	Behavioural task	Naming automatisation	Accuracy RT	Frederickson et al., 1997[117]
Phoneme suppression	Behavioural task	Phonological awareness	Accuracy RT	Rutten et al., 2019[118]
Text Reading	Behavioural task	Reading skills	Accuracy RT	"Le Pollueur", Gola-Asmussen et al., 2010[119] "L'Alouette", Lefavrais, 1967[120]
Word and Pseudoword Reading	Behavioural task	Reading skills	Accuracy RT	EVALEC: Sprenger-Charolles et al., 2005[121]

			(Per stimulus type: regular words, irregular words and pseudowords; overall)	ECLA16+ : Gola-Asmussen et al., 2010[119]
Spelling task	Behavioural task	Spelling skills	Accuracy (regular words, irregular words and pseudowords; overall)	Gola-Asmussen et al., 2011[119]
Spoonerisms	Behavioural task	Phonological awareness	Accuracy RT	Szenkovitz & Ramus, 2005[122]
Non-word repetition	Behavioural task	Phonological working memory	Accuracy of repetition Span, i.e. maximum number of repeated words	Majerus, & Van der Linden 2004[123]
Magnetization Prepared - RApid Gradient Echo (MPRAGE) imaging	sMRI	Brain structural anatomy	n/a	Work-in-progress compressed sensing sequence © Siemens Healthineers
Diffusion-weighted Imaging (DWI)	dMRI	Fibre tractography	n/a	Fedeli et al., 2021[124]
Language network brain functional localiser	fMRI	Functional activation for language	n/a	Malik-Moraleda et al., 2022[88]
Resting-state brain function	fMRI	Resting-state functional activation	n/a	Developed in-house
Field maps	MRI	Intensity of the magnetic field across space	n/a	Developed in-house

Table 2. Overview of all tasks and modalities.

Methods

Participants

According to the Organisation for Economic Cooperation and Development (OECD), approximately 40%

of Swiss individuals aged 25 to 34 possess upper secondary or post-secondary non-tertiary education, around

50% have attained tertiary education, and about 10% have below-secondary education[125,126]. Additionally,

Switzerland is recognized for its linguistic diversity, as we have extensively discussed in the above

paragraphs. The Federal Population Census 2022 Structural Survey[127] indicates that French is spoken by

22.8% of the population. In the survey, 70.1% and 23.4% of the population declared that they speak other

national (German, Italian, and/or Romansch) or non-national languages (participants could declare more

than one language, therefore there is a degree of overlap in these data). Considering that any study examining

the interaction between diverse cognitive domains and language skills will always be somewhat culture-

dependent[128], we chose our target participants with the aim to test at least 100 healthy, relatively multilingual

individuals having French as their first or dominant language, which is the primary language of the Canton

where the data collection took place.

We recruited 104 adult participants who matched these criteria from the Geneva area, the

surrounding French-speaking cantons of Switzerland and neighbouring France through flyers and online

advertisements. Participants provided their consent for disclosing their medical history and filled in an online

screening survey. To safeguard coherence and avoid confounding factors within the sample, prospective

participants were excluded *a priori* if they had musical or simultaneous interpreting professional

qualifications (known to interact with language); vision defects that could not be corrected; body implants

incompatible with MRI or known claustrophobia; neurological or psychiatric conditions; traumatic head

injuries with loss of consciousness; ongoing pregnancies; and past or present serious illnesses that required

invasive and/or continued medical treatment (such as cancer, chronic and/or autoimmune diseases).

Participants with MRI-conditional implants were evaluated on an individual basis upon providing further

documentation. Participants with diagnosed developmental dyslexia, as well as those who reported knowing

more than 10 languages with a self-reported proficiency equal to or higher than 4 out of 10 in reading,

speaking and listening, are not included in this dataset. Figure 1 reports the number of declared languages

for the final sample.

FIGURE 1 HERE

Once eligibility was established, all participants provided a signed informed consent to all

subsequent experimental procedures, including anonymised data reuse for open science. One participant

subsequently withdrew their consent to data sharing and two were not able to undergo brain imaging due to

claustrophobia. Therefore, the final sample includes 101 individuals (M_{age}= 23.35 years, SD = 4.08, 68F,

$M_{education}$= 15.34 years, SD = 2.35). At study completion, all participants received monetary compensation,

an image of their brain and a simplified report on their performance on the behavioural tests. Language

background and social status show that our participants were quite representative of the well educated and

economically stable Swiss society. This information is shown in Figure 2, with social status represented as

a cumulative measure derived from the education and job category of the participant, as well as those of their

family of origin and of their partner, if present, calculated with Barratt's Simplified Measure of Social Status

(BSMSS)[95].

FIGURE 2 HERE

Data Collection

Data collection. All interactions and documents provided to participants were in French. Data were collected

by six individuals who were either first-language French users, or who had learned French as their

second/additional language at an advanced level. All procedures were approved by the Geneva Cantonal

Ethical Commission (CCER Protocol N. 2021-01004) prior to the beginning of the study. All participants

signed a waiver for anonymised data release in the public domain, and no identifying information was

retained, to the best of our knowledge.

Data were collected in two online and two in-person sessions, hereafter named Session 1, 2, 3 and

4, which always occurred on different days but in the same order (1-4) between July 2022 and June 2023.

Session 1 was unsupervised, and required participants to fill out online questionnaires. Session 2 involved

online behavioural data collection, and was supervised by an experimenter. During Session 3 we collected

more behavioural data, this time in person. Finally, in Session 4 we collected neuroimaging (MRI) data.

Behavioural testing. Using established and published tests in cognitive psychology is crucial for addressing

the reproducibility crisis and curbing the proliferation of one-off tests, for ensuring that findings are accurate

and that they can be compared and replicated by other researchers[129]. Therefore, where possible, we chose

existing questionnaires and behavioural tasks, with the exception of an explicit morphosyntax learning test,

which we developed and piloted ourselves due to the lack of such a test in the field (see section "Pilot study").

All instructions and tests were delivered in French. Where the French version of a test was not available, it

was *adapted* and checked by at least one first-language French user in the team. Questionnaire instructions

(Session 1) were presented in written form. Behavioural task instructions (Sessions 2 and 3) were read by a

commercial natural reader (https://www.naturalreaders.com/commercial/read), using the voice "Renee

France" (if not otherwise specified), at speed -1. fMRI task instructions were given by the experimenter via

a microphone. All technical alterations and task adaptations were related to 1) language of delivery, or 2)

adapting a paper & pencil test for computer-based delivery. All the above information is thoroughly

documented for each test in the Supplementary Information file, Table S1. An overview of the data collection

structure is given in Figure 3 and an extensive theoretical explanation of the tests included in sessions 1-3

can be found in our recent exploratory work on the behavioural correlates of language aptitude, which

included this sample[17].

FIGURE 3 HERE

Pilot study. Before data collection, an explicit grammar learning task called ArtGram was developed and

pilot-tested. ArtGram, designed for adults, extends the PLAB4 task used for language aptitude testing in

children and adolescents, for which we could not find an equivalent in adults[130,131]. The test involves learning

an artificial, declensional language lexicon with inflected sample sentences, followed by a self-paced

multiple-choice, speeded translation task with novel sentences, as described in Table 2 and more extensively

in Rampinini et al., (2024)[17]. The pilot study aimed to assess: (1) the task's feasibility, (2) the reliability of

the online platform, (3) redundancy with an implicit grammar learning task (Brocanto). Twenty first-

language French speakers (11F, M_{age} = 27.65, SD = 8.9) without language or reading disorders participated

via video conference. Results showed an unsignificant correlation of $r(18)$ = .44 between the two grammar

tasks.

Session 1: questionnaires. Participants initially filled out a series of questionnaires online using Qualtrics

XM©. The questionnaire sequence was fixed: first came the Edinburgh Handedness Test and the BSMSS

questionnaire, completed upon recruitment. Then came the demographics, MFQ, IRQ, AHRQ, MUSEBAQ,

Code Switching and LEAPQ (see Table 2). When available from the questionnaire manual, automatic scoring

was implemented in Qualtrics via the "Formula Field" and resulted in one column per score in the derivative

questionnaire dataset, for each participant.

Session 2: online behavioural testing. During session 2, supervised behavioural data collection was

conducted via Zoom©. Participants were guided through a demonstration video on how to share both their

sound and entire screen, use wired headphones for reliable online measurements[132,133], and ensure their

microphone was functioning properly. Tasks were delivered through the Gorilla web interface[134]. Headphone

and microphone tests from the Gorilla open materials section were mandatory before starting the task

sequence. Researchers supervised the session, intervening only if technical issues arose. The system

prevented participants from logging into the session from mobile phones or tablets, and only allowed the use

of Mozilla Firefox© or Google Chrome©.

Participants navigated through tasks autonomously in a predetermined order among 15 possible

pseudo-randomizations, one of which was automatically assigned by the system at the start of the testing

sequence. Before each task, participants received written instructions in French on screen, and could not

proceed until the natural reader finished delivering the same instructions orally. Fixed-length breaks (3 or 5

minutes) were included after the most intensive tasks to optimize concentration and compliance. Participants

could end the break early if ready to continue, with a 60-second timer appearing in the last minute of the

break if they had not yet proceeded to the next task.

Session 3: in-person behavioural testing. In this session, participants were tested in person for tasks that

required closer supervision due to their length (e.g. the ANT-I), or that required manual measurements of

reaction times (RT), (e.g. the literacy and literacy mediator tests). Testing was conducted at the Human Neuroscience Platform of "Campus Biotech" in Geneva, in a dedicated, sound-protected room, using the same laptop, mouse, and headset (headphone with microphone) for all participants (see Figure 3). Tasks were organized and delivered via the Gorilla interface using the previously described pseudo-randomization strategy. The same microphone check was included to ensure the safe recording of tasks requiring vocal responses for later assessment.

An experimenter closely supervised the session through a connected screen, mouse, and keyboard while facing the participant, intervening only when necessary due to task requirements or technical issues. For tasks requiring a vocal response, the experimenter manually recorded accuracy and/or RT (assessed via a chronometer) in the session booklet. To prevent data loss, these tasks were also audio-recorded, and the responses were later verified for accuracy by a team member with French as their first language. Task events not requiring manual measurement or verification were recorded directly in Gorilla. During this session, participants also performed three extra tasks that were part of another project[17].

Session 4: Magnetic Resonance Imaging. Following behavioural testing, participants were invited to a brain imaging session on a Siemens 3T Magnetom-Prisma scanner equipped with a 64-channel head coil, again at Campus Biotech. Prior to scanning, they filled in and signed an MRI safety questionnaire to again verify their eligibility for the procedure. When required, participants were fitted with MRI-compatible goggles to correct their vision. The language localiser task was administered in Matlab r2021b with Psychtoolbox3, through a computer connected to a screen in the back of the scanner room, that participants could see through a mirror placed on top of the RF head coil. During the whole session, we could observe their eyes through an eye-tracking camera, to check that they were awake. During the imaging session, the field map, resting state sequence and the language localiser were administered in random order, but the resting state sequence was always preceded by the T1-weighted anatomical scan to avoid spurious activations due to carrying out an active task just before. After a short break outside the scanner, participants were repositioned and underwent the DWI sequence and its corresponding field map acquisition. During this session, participants also performed three fMRI tasks and one anatomical scan that were part of different projects. An overview of the imaging session parameters is provided in Table S1 of the Supplementary Information file.

fMRI language localiser. Participants were instructed to keep their eyes open and look at a black fixation cross on a white background while listening to intact and degraded snippets of the story 'Alice in Wonderland', in their first and second most dominant language of choice (L1, L2)[88,135]. This localiser can be used to inspect individual differences in language activation during quasi-naturalistic listening, and their relationship with behavioural measures of language and cognition[136]. The original localiser paradigm is publicly available from the authors' webpage. As described in Table S1 in the Supplementary Information file, and the Data Records section, we modified this paradigm to include a degraded L2 condition.

Resting-state fMRI sequence. Brain connectivity at rest can be linked to individual differences in
language[137,138] , reading[139,140], and other domains of cognition such as executive function[141,142]. To collect
information on resting-state brain activity, we asked participants to lie down with their eyes open, instructing
them to relax their body and mind as best as they could, while projecting a white fixation cross on a black
background.

BIDS conversion. This dataset conforms to BIDS v1.9.0 and was validated using the command line version
of `bids-validator v1.14.14` (http://bids-standard.github.io/bids-validator/). A LINUX Terminal
print of the output is provided in the Supplementary Information file (Figure S1). The dataset has been
annotated using the Neurobagel annotation tool (https://neurobagel.org/) for enhanced findability[143], the
annotations have been saved in `/neurobagel` at root level, and all JSON sidecars have been validated
with the online version of JSONLint. The BIDS data format conversion occurred in several steps, part of
which could be planned before data collection (such as naming of the MRI sequences, folder structure, and
participant codes), while others were performed *post hoc* to adapt data generated by environments not
optimised for BIDS. In general, the procedure aimed at having a BIDS-coherent data structure and file
names, (re)organising the content of tabular files, adding sidecar files to accompany data and customising
code to work in a BIDS folder structure. These steps were performed on all data collected during the same
testing session, and the NEBULA101 data were subsequently imported into the `/nebula101` data space.
Nonetheless, for clarity, we provide the specific heuristic for the construction of this dataset after DICOM
to NiFTI conversion in `/code/heudiconv/heuristic.py`. Considering this procedure, all code
described is specific to the NEBULA101 dataset but is not meant to be rerun, and is given with paths relative
to the dataset BIDS root folder, but might reference to folders outside this structure, for example to source
data. Outside of the code performing data import or behavioural data cleaning, no further reference is
generally made to external/unavailable files. We describe the steps in Table 3.

Step	Description	Language and environment	Custom code	Code availability	Code location
1	DICOM to NifTI conversion: conversion was handled in heudiconv via dcm2niix.	Python heudiconv	No	No	n/a
2	File structure heuristic: we set up logging, defined constants, and included functions to create keys for file paths and categorise imaging sequences into different modalities based on their metadata.	Python heudiconv	Yes	Yes	`code/heudiconv/heuristic.py`
3	Modifying language localiser log files for BIDS compatibility.	Python	Yes	Yes	`/code/create_nebula/ nebula101_bidsify_mri_logs_alic eloc.py`
4	Importing NEBULA101 data from main BIDS data directory, including behaviour and phenotype data	Python	Yes	Yes	`/code/create_nebula/create_nebu la.py` `/code/create_nebula/make_beh.py` `/code/create_nebula/make_phenot ype.py`
5	Creation of JSON sidecar files for derivative scores.	Python	Yes	Yes	`/code/create_nebula/create_nebu la.py` `/code/create_nebula/create_scor e_id_json.py.`

6	Modifying field map sidecar "intendedFor" field for BIDS compatibility (to remove references to extra scans) and correcting label naming errors in field maps (fid- to acq-).	Python	Yes	Yes	`/code/create_nebula/create_nebula.py`
7	Importing stimuli from fMRI language localiser to store in BIDS dataset and adjusting their folder structure.	Python	Yes	Yes	`/code/create_nebula/create_nebula.py`
9	Defacing T1w MPRAGE scans.	Python PyDeface	Yes*	Yes	`code/1_anat/0_pydeface/nebula101_loop_pydeface.py` * The (minimal) customisation involved creating a script to run the code on the NEBULA101 sample
10	Creation of dataset-level sidecar files	Python	Yes	Yes	`/code/create_nebula/create_nebula.py`

418   *Table 3. List of the steps taken to make the dataset BIDS-compliant.*

Data Records

The dataset is publicly accessible and downloadable from OpenNeuro under a Creative Commons CC0 1.0

(Universal Public Domain Dedication) license with accession number ds005613 at the address

doi:10.18112/openneuro.ds005613.v1.0.1.

In this section we further describe the data structure and its contents. The folder `/nebula101` (Figure 4)

constitutes the root level of the dataset. It contains 101 participant folders with raw imaging data denoted by

the code `sub-pp` followed by three digits, the folders `/phenotype`, `/code`, `/derivatives`,

`/stimuli`, `/neurobagel`, and the mandatory files required by BIDS (README, participant and

dataset description files). The `/phenotype` folder contains tabular data from questionnaires, while

`/neurobagel` contains subject-level annotations (harmonized phenotypic properties and imaging

metadata) that can be encoded in a knowledge graph (see Technical Validation). As concerns the subject

folders, inconsistent numbering is due to non-included participants (see Participants section), the participant

who denied consent to share their data, and the two participants who could not undergo brain imaging). We

describe the other contents of `/nebula101` in detail here below.

FIGURE 4 HERE

The `/code` folder in Figure 5 contains subfolders with 1) fMRI paradigm data for the language localiser; 2)

code to import, modify or generate the BIDS-compliant files as described in Table 3; 3) the BIDS conversion

heuristic; 4) the preprocessing information and code folder; 5) the validation materials folder.

FIGURE 5 HERE

The `/derivatives` folder in Figure 6 contains the preprocessed behavioural data and their sidecar files,

as well as the results of all validation pipelines.

FIGURE 6 HERE

• `cumulative_farsi_rater*` = derivative scores from the Farsi task ratings and their sidecar files.

• `nebula_101_all_questionnaire_scores*` = scores from questionnaires and their sidecar
file.

• `nebula_101_all_task_scores*` = scores from tasks and their sidecar file.

• `nebula_101_leapq_annotation_iso_glottolog*` = mapping of language names to ISO an
Glottolog and its sidecar file.

• `nebula_101_leapq_data*` = LEAPQ data and their sidecar file.

• `nebula_101_leapq_langname_order*` = LEAPQ languages in order of acquisition and their
sidecar file.

The `/stimuli` folder contains a subfolder where the language localiser materials are stored and can be
referenced by the BIDS validator.

The participant folders, named `sub-pp*/ses-01/` contain `/anat, /dwi, /fmap, /func` and `/beh`
folders. These host the raw imaging and behavioural data of each participant, the latter having been
minimally preprocessed to remove metadata and information unrelated to the scoring, as described in
Technical Validation. An example with `sub-pp001/ses-01/` is shown in Figure 7, Figure 8, Figure 9,
Figure 10 and Figure 11.

FIGURE 7 HERE

FIGURE 8 HERE

FIGURE 9 HERE

FIGURE 10 HERE

FIGURE 11 HERE

The `/code/AliceLocalizer-4conditions/` folder contains our own version of the materials
provided in the public domain by the creators of the task[88,135], which we modified to suit our needs, as
described in the fMRI language localiser description (see: Data collection). This version of the code can read
the stimuli from the structure of the BIDS folder. It will still require manual intervention to create the
`events.tsv` files from the Matlab log files, as Matlab currently provides a different tabular structure. We
remind the reader that we altered the structure of the log files to create simplified BIDS events with no

redundant or unneeded information in
`/code/create_nebula/bidsify_mri_logs_aliceloc.py`.

The `/code/preprocessing` folder includes numerically ordered subfolders containing code to
reprocess the anatomical, field map, functional and diffusion data (see Technical Validation), and the
reference code for the raw-to-derivative conversion of behavioural data. Any steps contained in these
subfolders are meant to be run sequentially to recreate the materials needed for technical validation, or if the
user decides to process our raw brain data using the same pipelines.

• `0_beh` contains behavioural data preprocessing code.
• `1_anat` contains the brain extraction code.
• `2_fmap` contains code for field map preparation.
• `3_func` contains code for minimal fMRI preprocessing, to get the displacement parameters used
in the technical validation.
• `4_dwi` contains the preprocessing code for the diffusion data necessary to obtain technical
validation reports.

The `/code/validation` folder contains the following subfolders (see Technical Validation for details
on the operations performed):

• `/anat`: segmentation and sample homogeneity plotting code for the anatomical scans (auto-
generated by Matlab): `/cat12/cat_stat_homogeneity.m` and
`/cat12/cat_stat_homogeneity_autoplot.m`
• `/beh` contains the following items:
• The code to run reliability analysis on behavioural data:
`calculate_cronbach_alpha.py`
• Three additional Python notebooks for generating the correlation data presented here and
additional tables and matrices contained in the Supplementary Information file (Table S2, Table
S3 and Figure S2):
• `correlate_matrix.py`
• `calculate_descriptives.py`
• `farsi_inter_rater.py`
• `/data_checks`: various Python scripts to create data lists of behavioural and imaging data,
and missing data heatmaps, as explained in Technical Validation.
• `/dwi`: individual participant pdf QC reports, the code for generating them, and the group report
folder.
• `/func`: two Python scripts, one for importing displacement `.rms` data from FSL, and the
other for violin plots of average absolute and relative displacement during fMRI (resting state
and task based).

•   `1_copy_motion_params.py`

•   `2_mean_abs_rel_disp_violin.py`

For each of the described folders in `/code/validation/`, there is a mirror
`/derivatives/validation/` folder where the results of the validation pipelines are stored, and
specifically:

•   `/derivatives/validation/anat/cat12/` contains the single subject CAT12 reports in
PDF form.

•   `/derivatives/validation/beh/` contains the item-level reliability data used to calculate
Cronbach alpha and all correlation measures.

•   `/derivatives/validation/data_checks/` contains data presence checks generated by
the respective code for brain imaging, phenotype and behaviour.

•   `/derivatives/validation/dwi/fsl/` contains the reports from subject- and group-level
quality assessment of DWI data.

•   `/derivatives/validation/func/fsl/aliceloc/` contains contains the reports from
subject- and group-level quality assessment of task fMRI data.

•   `/derivatives/validation/func/fsl/rest/` contains contains the reports from
subject- and group-level quality assessment of resting state fMRI data.

All Python notebooks and executables are heavily commented for user-friendliness.

Technical Validation

540 In compliance with the BIDS indications for mixed raw and derivative datasets, to improve user experience
541 and reduce redundancy, we chose to include pre-processed behavioural data as derivative tables. We also
542 provide quality control (QC) reports for all imaging modalities, together with the code for reproducing them.
543 Below we specify the details of technical validation for each data type.

Questionnaires

Questionnaire source data were retrieved from Qualtrics XM© and preprocessed in Python. Each
questionnaire was cleaned with its own script and then merged with the others via another script. The overall
process aimed at data cleaning, as scoring was generally performed via the Qualtrics graphical user interface.

1)   Log file data collection: for each questionnaire, from the source tabular file containing all participants,
we eliminated irrelevant metadata. In rare cases, when duplicates were identified we eliminated them at
source level by keeping the first non-empty instance of the questionnaire, blind to the contents, to avoid
selection bias.

2)   Data cleaning and handling of missing data: in all questionnaires, string responses were recoded as
numbers. Missing data (NaN) was handled as follows: as a general rule, within-questionnaire, all
responses were forced (i.e. progression was halted if a response was missing or on-screen warnings
appeared to highlight unfilled responses). When information was indeed unavailable for a participant,

we set up options to input specific strings, signalling to us that the information was missing. When missing information was nonetheless identified in a questionnaire with *optional* responses, but without the string signalling unavailability, we evaluated on a case-by-case basis whether it was possible to obtain it from the participant. In such rare cases, the questionnaire was retaken by the participant in session 4, assisted by an experimenter (to avoid data alteration), and only missing responses were filled in. In even rarer cases, when an *entirely* missing questionnaire was identified, participants were invited to complete it through an individual link to that specific questionnaire. Any remaining entries with unavailable information were converted to NaN during data preprocessing. With the strategies we put in place, all questionnaires have been completed by all participants in this dataset, except for one monolingual participant not having taken the code-switching questionnaire due to only speaking one language.

3) LEAPQ cleaning: the LEAPQ, due to its length and structure, required additional, *ad-hoc* handling. In Qualtrics, participants reported the number of languages they knew, including extinct ones and dialects, in order of dominance, that is, how much they used each of them over the others, and in order of acquisition, that is, the temporal order in which they learned their languages. They then provided general information for each language in order of dominance (e.g., names, exposure) and specific information (e.g., learning modality, use context). The resulting data file was restructured by language number and question type, filling gaps with NaN to create a 'staircase-like' structure where participants (rows) and questions (columns) are listed by language number, in ascending order. The resulting file is available in `/derivatives/nebula101_leapq_data.tsv` with a corresponding sidecar file explaining the column contents. The list of languages in order of acquisition was extracted into a separate file to avoid confusion, as it was not always the case that the dominance and acquisition lists coincided. Language order information can be accessed at `/derivatives/nebula_101_leapq_langname_order.tsv`.

4) Raw language names as entered by the participants were conformed to ISO 639-3 and Glottolog codes[144] for standardised reference. The conversion map can be found in `/derivatives/nebula_101_leapq_annotation_iso_glottolog.tsv` and its sidecar file.

5) Multilingual language experience entropy. The LEAP-Q is a widely used instrument for assessing multilingual language experience, encompassing contexts of use, learning, use choices, history with the language, and native-likeness. Given that composite measures of multilingualism derived from the LEAP-Q would be beneficial for quantitative analyses, we calculated four different continuous 'multilingualism scores' for each participant, to reflect their multilingual experience cumulatively[64]. Three scores were based on their self-reported proficiency in speaking, reading, and comprehension across all reported languages, while the fourth score was based on current exposure to all reported languages. Following previous research[17,145,146], each participant's multilingualism score combined proficiency or exposure across their different languages using Shannon's entropy equation[147] within the R entropy package[148].

Item-level questionnaire data are stored in `/nebula101/phenotype/` as individual tabular files, each containing data for one questionnaire, where each subject is represented in a single row as a `participant_id`

595 and their responses in the subsequent columns. Each phenotypic tabular file is accompanied by a JSON sidecar
596 file describing each column:

- 597 • `ahrq.tsv` and `ahrq.json` contain data for the Adult Reading History Questionnaire.
- 598 • `bsmss.tsv` and `bsmss.json` contain data for the Barratt Simplified Measure of Social Status.
- 599 • `code_swt.tsv` and `code_swt.json` contain data for the Code Switching Questionnaire.
- 600 • `handedness.tsv` and `handedness.json` contain data for the 10-item French version of the
- 601 Edinburgh Handedness Inventory.
- 602 • `irq.tsv` and `irq.json` contain data for the Internal Reasoning Questionnaire.
- 603 • `mfq.tsv` and `mfq.json` contain data for the Motivation Factors Questionnaire.
- 604 • `musebaq.tsv` and `musebaq.json` contain data for the Music Experience Use and Engagement
- 605 Questionnaire.

606 Given the higher level of processing that the LEAP-Q data went through, its tabular files (responses, language
607 annotations and language order information) reside in `/derivatives`, together with the comprehensive
608 derivate score file of all the questionnaires, `nebula_101_all_questionnaire_scores.tsv`.

Behavioural tasks

611 Behavioural tasks were pre-processed in Python for data cleaning, derivate score calculation and BIDS
612 conversion. The process happened in steps:

1) Log file collection: data were downloaded from Gorilla. Given that participants entered 1 out of 15 possible randomised task sequences, automatically assigned to people by Gorilla, the empty log files from the 14 unused randomisations for each participant had to be deleted. Microphone and headphone check files were identified and renamed.

2) Log file cleaning: irrelevant metadata were eliminated. Task log files were renamed to reflect the actual task (as Gorilla provides them in encrypted form) and to contain BIDS-compliant strings by adding the relevant labels, then converted to tab-separated values. Within each task's log file, column names were renamed to more easily interpretable strings where necessary. Audio files from voice-recorded tasks were renamed as well, for easier reference.

3) Derivate score calculation and tabular file merging: for each task, one or more derivate scores were calculated, based on the task's protocol. See Table 2. and Table S1 for the references and scores that were selected.

625 The behavioural data preprocessing code is provided for reference in
626 `/nebula101/code/preprocessing/0_beh`, and specifically:

a. `2DataCleaning_gorilla.py`: this script cleans the source logfiles (not provided in this dataset) from metadata, as explained.

b. `3Scoring_gorilla.py`: this script scores the files that were generated by Gorilla and cleaned, based on an automated pipeline by using the 'correct' and 'incorrect' columns and/or reaction time information (depending on the task's specific scoring protocol).

c. `4Scoring_manual.py`: this script scores logfiles that were created from tasks that
required human intervention (for example, all tasks requiring manual RT measurements of
voice responses via a chronometer, and/or live pencil scoring by the experimenter, such as
the Text or Word and Pseudoword reading tasks, and/or the subject to write down answers
like in the Spelling task). Here, Gorilla was used to run the task within the pseudo-
randomisation pipeline but scoring necessitated manual intervention.
d. `5Scores-merging.py`: this script merged the scores into a tabular file, from which
they were later imported to the `/nebula101` data space via the script
`create_nebula.py`.
e. `7farsi_raters_assessment.py`: The script calculates cumulative and average
scores for the Farsi uvular sound production task, based on assessments made by two
independent raters. It produces the derivative file provided in
`/nebula101/derivatives/`.

These actions were performed in subsequent steps by Python scripts acting on source data. Users will not be
required to rerun these scripts since we provide cleaned tabular files containing the minimally processed raw
accuracy and RT scores rather than the source data, as well as, crucially, the derivative scores calculated and
ready for use.

For tasks whose data were recorded automatically in Gorilla (when a simple correct/incorrect input was
sufficient, and RT could be measured by key press or mouse click), the raw data of each subject will contain
their `participant_id`, as minimally mandated by BIDS, accuracy and RT columns for each trial, if these
are sufficient to derivate a score provided in `/derivatives` via the code provided in
`/code/preprocessing/0_beh/`. For the manually scored tasks, as described in steps c) and e) above,
single subject raw data resulting from the digitisation of paper-and-pencil materials are provided: such might
include block-level data (e.g. each iteration and type of recall trials in the case of the CVLT) or condition-
level data (e.g. word types in the case of the Word and Pseudoword reading task, story types in the case of
the Text reading task), whose structure will be inherent and specific to the task itself. In the case of the
Spelling, Nonword repetition and Spoonerisms tests, scoring was digitised at task level (as per protocol). We
therefore included these only in the derivative table. Despite any differences in structure, all behavioural
files conform to the minimal BIDS requirements for their data type and are accompanied by extensive JSON
descriptions. All materials used to process source data can be made available upon request, as well as source
data that can be anonymised.

A comprehensive file containing all derivate scores for all behavioural tasks, obtained with steps b) and c)
lives in `/derivatives` and is called `nebula_101_all_task_scores.tsv`.

In this location the Farsi task derivate scores obtained with step e) can also be found: these are called
`cumulative_farsi_rater1_filtered.tsv` and
`cumulative_farsi_rater1_filtered.tsv`. All derivate files are accompanied by JSON
sidecars.

Behavioural data: correlations and reliability. Behavioural scores, whether resulting from tasks or questionnaires (here we generally refer to them together as "behavioural data", and we use "scores" or "measures" when referring specifically to their outcomes), derive from item-level data. To validate these measures, we first explored them via correlations, and then ran reliability analyses in the form of internal consistency coefficients, in a conceptually similar approach to our recent exploratory analysis of behavioural data with a larger cohort[17]. Intercorrelation of the scores obtained from behavioural data (tasks and questionnaires) was minimal, and reliability was acceptable to good in most cases (please refer to the Supplementary Information file for com. The code for calculating Pearson's correlations on the z-scored data as well as Cronbach Alpha, along with item-level reliability data and summary output (figures, tables) are available in `/code/validation/beh/` and in the Supplementary Information (Figures S2 and S3, Tables S2, S3, S4, S5).

To better understand the correlation patterns and reliability of these data, some considerations must be made about correlations and about reliability measured by internal consistency. Regarding correlations, on the one hand, high intercorrelations can be informative about shared variance between tests, but on the other hand, having an excessively intercorrelated dataset can lead to problems linked to multicollinearity[149–152]. On a conceptual level, the separability of latent variables facilitates the interpretation of the overall construct. Our choices, when planning this data collection and selecting the tests, aimed at isolating the best measures to represent each variable that we wanted to test in the context of this exploratory project on language aptitude. All things considered, a relative degree of independence of our measures was not only expected, but desirable. In the Supplementary Information (Table S2), we report all Pearson pairwise linear correlations having a coefficient of at least $|r(100)| > .5$ and significant at $p < .05$. In `/code/validation/beh/` code is provided to generate Table S2 and a matrix (Supplementary Figure S2) to visualise these data (`correlate_matrix.py`). Keeping in mind that correlations are just a preliminary way to explore data, most significant correlations were between metrics within the same test, and between these and their cumulative score, when present. As concerns correlations across tasks (marked with * and ** for positive and negative relationships respectively, in Table S2), the most evident patterns emerging are the correlations between reading measures and those typically related to or predictive of reading skill or deficit (all sub-scores from the RAN, text reading, word and pseudoword reading, spelling, spoonerisms, phoneme suppression and non-word repetition); between the Brocanto and CVLT long-term recognition score; between AMMA (total and tonal scores) and the musical training score from the MUSEBAQ (time, intensity and level of practice reached). No other measures were highly and significantly correlated.

Regarding reliability, the main advantage of internal consistency measured via Cronbach alpha is that, as long as covariance is inspected and there are minimal to no *nonignorable nonresponses* (MNAR data, i.e. missing not at random), it can be applied quite flexibly, unlike other methods[153]. Minimal reliability requirements are a matter of debate, as the α value can be affected by the number of items and improved by the independence of the construct these tap into[154–156]. This being said, it is harder to obtain a reliable

measurement in tasks that measure *spans* (e.g. Corsi blocks, Digit span) and tasks where participants stop

after reaching a maximum of answers with high variability (e.g. Revised tempo test): these will generate a

MNAR pattern, where each row ends with a certain number of NaN because the span has been reached, and

thus will be less suited to internal consistency measures, providing hardly interpretable coefficients[157]. This

can explain the somewhat unreliable Revised tempo test RT, reflected in a pattern where negative or null

covariances outweigh positive ones in a MNAR fashion (Figure 12),

(`/code/validation/beh/cronbach_alpha_results.tsv`).

FIGURE 12 HERE

Rapid automatised naming accuracy (α = .21) as well as accuracy on the first run of Brocanto (α = .25) also

had low reliability. We ascribe the former to the difficulty of the task for the participant, and therefore

reaction times may provide more reliable measures than accuracy: RAN measures naming latency[158], whose

motor component may show more consistent RTs than accuracy (and this indeed shows in our data, as the

two had α coefficients of .93 and .21, respectively). The lower reliability of the first run of Brocanto

is conceptually interesting if compared to the subsequent runs and to the RT data: it shows that upon learning

an artificial language completely inductively, responses are more variable at the beginning of the learning

process, where people tend to guess more frequently, while the time to make a grammaticality judgment is

overall always consistent. We provide the correlation matrices for accuracy and RT across the 3 blocks of

Brocanto in Figure 13. In a few cases, data were recorded at the item level but digitised at the task level

(spoonerisms, non-word repetition and phoneme suppression): item-level data are available upon request.

To facilitate readers, in the Supplementary Information file we show the internal consistency values for tasks

where α > .5, and tasks with α > .6 are additionally highlighted in bold (Table S3). Complete reliability data

are available in `/code/validation/beh/` and can be regenerated with

`calculate_cronbach_alpha.py`. Descriptive statistics and plots of the test metrics are available in

the Supplementary Information file: Table S4, S5; Figure S4, S5, S6).

FIGURE 13 HERE

The Farsi uvular sound production task was rated by two first-language Farsi speakers who heard

each recorded utterance from participants and gave it a 'native-like production score': therefore, any internal

consistency metric would reflect the way the raters scored the task, more than the way the participant

performed it (even though these are clearly related). Thus, for this task, we chose to calculate inter-rater

reliability between the first and second rater, in a procedure identical to a previous study where this task was

used[159]. For the ANT-I task, given its structure (each trial reflecting more than one possible condition from

which a score can be derived), we chose to measure the maximal split-half coefficient to assess reliability[160]

(the Supplementary Information file contains the code for this procedure).

Brain imaging

Anatomical imaging anonymisation, QC and brain extraction. Facial features in anatomical MRI scans

violate the principle of anonymity in open data. Therefore, T1-weighted MPRAGE scans were defaced with

PyDeface (https://pypi.org/project/pydeface/) and their quality was assessed in the Computational Anatomy

Toolbox for SPM (hereon, CAT12)[161]. Specifically, we used the CAT12 segmentation and sample

homogeneity toolboxes, providing easily interpretable quality measures at the participant and group level.

The weighted overall image quality (IQR) and the quartic mean Z-score are the two key indicators of image

quality. IQR combines noise and spatial resolution measurements before pre-processing, while the mean

quartic Z-score assesses the homogeneity of data after pre-processing, with deviations increasing variance

and reducing statistical power. The *product* of IQR and quartic mean Z-scores combines these quality

measures, with a low number indicating high quality. For each participant, we provide a PDF with the CAT12

report, as well as information on sample homogeneity, in `/code/validation/anat/cat12/`. The

CAT12 toolbox can be easily run via a graphical user interface (GUI) in Matlab and requires no custom code.

We provide the group distribution of the described measures in Figure 14.

FIGURE 14 HERE

To prepare T1 MPRAGE anatomical scans to be fed to FSL[162] (see *Functional neuroimaging* section), we

provide code for running an improved skull-stripping as a loop in the NEBULA101 dataset in

`/code/preprocessing/1_anat/1_optiBET/nebula101_run_optiBET.sh`. This code will

process T1-weighted MRI images stored in a BIDS directory to perform improved brain extraction with

optiBET[136] via the following actions:

1) Iterating through all participant directories in the BIDS directory.

2) For each participant, creating an output directory in the derivatives folder if it doesn't exist.

3) Copying the T1-weighted images (`sub-pp*_ses-01_T1w.nii.gz`) to the

corresponding output directory.

4) Checking if each T1-weighted image has already been processed.

5) If an image has not been processed, running the `nebula101_optiBET.sh` master script

on the image, which performs actual brain extraction.

The script referenced in step 5 is the master optiBET script, which needs to be downloaded from

https://montilab.psych.ucla.edu/fmri-wiki/optibet/ and stored in the location where the loop script is stored.

Field maps. We acquired field maps to accompany the DWI and fMRI sequences. Field map preparation

consists of a few steps that are partly scanner-specific, and in the case of Siemens Prisma, consisted in

recomposing the phase and magnitude images. We provide field map preprocessing code in

`/code/preprocessing/2_fmap/fsl/1_nebula101_run_fmap_prep_all.sh`

The script will perform the following steps:

1) Transformation Calculation: calculating the transformation matrix from the anatomical T1 image to
the magnitude image and applying the calculated transformation to the T1 brain mask to align it
with the magnitude image (`FLIRT`).
2) Magnitude Image Masking: using the transformed brain mask to mask the magnitude image,
creating an untrimmed brain image (`fslmaths`).
3) Brain Mask Trimming (`fslmaths`):
a. Binarising the untrimmed mask.
b. Smoothing the mask with an 8 mm kernel.
c. Thresholding the smoothed mask at 75%.
d. Binarising the thresholded mask.
e. Trimming the original brain image using the final binary mask.
f. Removing intermediate untrimmed files.
4) Preparing the final field map (`fsl_prepare_fieldmap`): preparing the final field map using
the processed magnitude and phase difference images, with a specified echo spacing of 2.46 ms.
795

Diffusion-weighted imaging. To validate our diffusion data, we run the `QUAD` (participant) and `SQUAD`
(group) programs within the `FSL-FDT`[162,163] suite as part of the diffusion pre-processing pipeline. The DWI
data and prepared field map of each participant were fed to `EDDY` to correct for susceptibility, eddy currents,
inter- and intra-volume participant movement and signal dropout, using the prepared field map via the `--`
`field` flag. Finally, we ran `eddyQC`[164] for quality assessment. We provide a pdf with QC statistics for each
NEBULA101 participant and for the group, as well as the code to reproduce this procedure, in
`/code/validation/dwi/fsl/dwi_qc_quad_squad.py`. The code is configured to
automatically grab the acquisition parameters and the correct number of shells from the DWI sidecars to
generate the `acqparam.txt` and `index.txt` files required by `EDDY,` and will raise flags if the
expected numbers do not match with the data. An `eddy_quad_qc_paths.txt` file has also been set up
to this aim in the same location, which the user can alter for their needs, should they decide to preprocess
DWI data (which we provide in raw form) with the same pipeline
(`/code/preprocessing/dwi/fsl/1_dwi_prep.py`). Figure 15 reports the signal- and contrast-
to-noise ratios from the group QC metrics generated using `SQUAD`. The complete PDF report is available in
`code/validation/dwi/fsl/squad/group_qc.pdf`.
811

812 FIGURE 15 HERE
813

Functional imaging. As concerns fMRI, we provide QC metrics obtained from preprocessing of the whole
dataset in FSL, and the code to reproduce the procedure on the NEBULA101 sample.
Figure 16 and Figure 17 report violin plots of the absolute and relative displacement by run for the language
localiser and for the resting state fMRI sequence, respectively. Absolute displacement reflects the movement
with respect to the reference volume, while relative displacement reports volume-to-volume information.

Both can impact SNR but while absolute displacement can be counteracted, relative displacement is usually

more harmful, and both must be checked when assessing the quality of fMRI data[165]. Displacement

information and plots (Figure 16 and Figure 17) can be obtained by running

`code/validation/func/fsl/1_copy_motion_params.py` and

`/code/validation/func/fsl/2_mean_abs_rel_disp_violin_py`. In principle, if the data

are not being reprocessed, `1_copy_motion_params` does not need to be rerun as we provide `.rms`

output from FSL, containing absolute and relative displacement data for each participant, in

`/code/validation/func/fsl/aliceloc/` and `/rest`.

FIGURE 16 HERE

FIGURE 17 HERE

A folder containing the code for performing the basic preprocessing steps of fMRI for the language localiser

and resting state sequences has been set up in

`/code/preprocessing/3_func/fsl/1_preproc1/1_loop_fsl_preproc1.py`. This code

performs the first step of fMRI preprocessing with FSL FEAT, consisting of:

1) Motion correction

2) 4D mean intensity normalization

3) Spatial smoothing (5 mm FWHM)

4) B0 unwarping (includes `BBR reg`)

5) Registration (`BBR, FNIRT`)

This code will run using `/code/preprocessing/3_func/fsl/1_preproc1/design_pp.fsf`,

a design file required by FSL which has been set up to contain information on the preprocessing loop, should

the user wish to preprocess our data with the same pipeline.

Overall validation of data structure. This dataset is mostly complete, with minimal missing information.

However, to provide the user with information that is traceable without necessarily having this data

descriptor at hand, we have created a series of scripts within the `/code/data_checks/` folder aimed at

providing report logs on the presence of behavioural and brain imaging data. Such logs are saved to

`/derivatives/validation/data_checks/`.

• `nebula101_bids_report_beh.py`: this script plots behavioural data presence checks on the raw

and derivate dataset and compiles the findings to a report log called

`nebula101_beh_report_[timestamp].txt`. It will also create:

• A tabular file `nebula101_datalist_[timestamp].tsv` listing data as 1 and 0 if

present or absent, for easy reference.

• A heatmap with the above information, called
`data_presence_heatmap_[timestamp].png`.
• A report specific to the LEAPQ data, to have missing and present information at a glance, called
`nebula101_leapq_data_report_[timestamp].txt`
• `nebula101_bids_report_mri.py`: this script plots imaging data presence checks and compiles
the findings to a report log called `nebula101_mri_report_[timestamp].txt`. This can be
used for easy referencing to participants having undergone different conditions (e.g. different field map,
see: Usage Notes).
864

Usage Notes

866 *Behavioural data.* This dataset is mostly complete, but a few technical malfunctions during testing have
867 caused some minimal missing data, as described in Table 4. This information and a heatmap for visualising
868 missing data at a glance can easily be regenerated using the scripts in
869 `/code/validation/data_checks/` called `nebula101_bids_report_beh.py` and
870 `nebula101_missing_data_log.py`.

Participant ID	Columns with missing data at task level
sub-pp031	phon_suppr_rt_manual
sub-pp050	brocanto_corr1, brocanto_incorr1, brocanto_rt1, cvlt_long_corr, cvlt_long_incorr, cvlt_long_rt, brocanto
sub-pp116	cvlt_long_rt
sub-pp118	apm_corr, apm_incorr, apm_rt
sub-pp155	non_word_rep_span, non_word_rep_acc

Table 4 Missing behavioural data.

Imaging data. Here we provide some details to facilitate easier navigation of the imaging data. This
information can be generated by
`/code/validation/data_checks/nebula101_bids_report_mri.py`. JSON sidecar files
have already been adjusted to report the below nuances:
• `sub-pp161`: for this participant, the `beta` field map must be used for resting state fMRI
processing, and the `delta` field map must be used for DWI processing.
• `sub-pp001, sub-pp010, sub-pp013, sub-pp023, sub-pp043, sub-pp053`:
for these participants, the `gamma` field map must be used for resting state fMRI processing.
• `sub-pp032`: the voxel dimensions of `/sub-pp032/ses-01/anat/sub-pp032_ses-`
`01_rec-defaced_T1w.nii.gz` slightly differ in the y and z directions (pixdim1 = 1.000000,
pixdim2 = 1.039060, pixdim3 = 1.039060).
When running the BIDS validator, this information is reported as *warnings* and saved in the log, unless the
`--ignoreWarnings` flag is raised. Warnings to not impede BIDS validation and the dataset will pass it
every time. When processing the data, this information will be read automatically by software that can work

primarily in BIDS (such as fMRIprep[166] in the case of functional imaging). When using software that lacks

this functionality, such as FSL, we recommend processing these participants manually. The `.bidsignore`

file contains instructions for skipping the validation of `/neurobagel` as it contains extra files.

Code Availability

892   All code that can be rerun on these data is provided within the `/code` folder, as described in the Technical

893   Validation section. Raw-to-derivative code is provided for all raw data. Given that we do not provide *source*

894   behavioural data, source-to-raw preprocessing code for tasks and questionnaires is not included, but can be

895   shared upon request.

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Author Contributions

901   CRediT (Contributor Roles Taxonomy) is provided, as follows: Conceptualization: AR IB OK RB NG;

902   Investigation: AR IB; Methodology and Formal Analysis: AR IB OK; Data Curation: AR IB; Technical

903   Validation: AR; Writing - Original Draft AR; Writing - Review & Editing: AR IB OK RB NG;

904   Visualization AR; Resources: RB NG; Supervision: RB NG; Funding acquisition: RB NG.

905   AR and IB (*) contributed equally to this work.

Competing Interests

908   The authors declare no competing interests.

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Figure legends

Figure 1. Frequency distribution of the number of reported languages in the final sample (N = 101).

Figure 2. Participants' language background up until their third language (left, languages coded by colour)
and Socioeconomic status (bottom right) measured via Barratt's simplified measure of social status[95].

Figure 3. Data acquisition and processing structure (session duration is approximate for behavioural data
acquisition due to individual variability in task completion times). [Illustration created with BioRender.com]

Figure 4. Root dataset folder.

Figure 5. Code folder.

Figure 6. Derivatives folder.

Figure 7. Defaced T1 MPRAGE data and their sidecar file.

Figure 8. DWI data: beta values, vectors, and sequence data, with its sidecar file.

Figure 9. Field maps: phase and magnitude images with their sidecars. This participant has an extra field
map (gamma), as explained in the Usage Notes.

Figure 10. Functional MRI data with their sidecar files and event files. Rest refers to resting state fMRI,
aliceloc refers to the language localiser.

Figure 11. Raw behavioural data files with their sidecar files.

Figure 12 Revised tempo test correlation patterns for subtractions (left) and sums (right) RT data. The
different number of items across the 2 figures is due to the fact that 1) participants reached a different number
of completed operations across the two conditions and 2) no one reached the end of either condition (60
operations). Knowing that for this task, there is data missing for a reason, this can be interpreted as a MNAR
pattern.

Figure 13 Top row: correlation matrix for the RT data of Brocanto across 3 blocks. Bottom row: accuracy
data for Brocanto across 3 blocks. Plots show that overall, RTs were more reliable than was accuracy.

Figure 14. CAT12 output. Boxplot showing the weighted IQR by mean absolute z-score scaled by a factor
of 4 (i.e. quartic) to emphasize outliers. As the plot shows, all but 1 participant in this distribution lie in the
good to optimal range of the quality measure. Of note, even when outliers are detected, these are not
necessarily data to discard on an absolute basis, as the score is calculated relatively to the specific sample
being analysed.

Figure 15. Signal-to-noise and contrast-to-noise ratios from the QC-ed DWI data.

Figure 16. Absolute (ABS) and relative (REL) displacement (mm) across runs of the language localiser.

Figure 17. Absolute (ABS) and relative (REL) displacement (mm) during the single resting state sequence
run.

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1 **2**

Questionnaires
Remote tests
3h45'

3

In-person tests
2h45'

sound-proof environment

esperimenter laptop

00:00:00

130cm~

data

testing
laptop

Spelling Sheet

Purdue Pegboard

21 behavioural tasks - 8 questionnaires - 102 metrics

MINIMAL
PREPROCESSING

TECHNICAL
VALIDATION

ANONYMISATION

T1 Structural Imaging
N = 101

4

Brain
Imaging
50'

Resting State Functional imaging
N = 101

Task-based Functional Imaging
N = 101
First & Second Language Localiser

Diffusion-weighted Imaging
N = 101

BIDS
BRAIN IMAGING DATA STRUCTURE

behaviour

brain

Public domain release
(further data exploration)

NEBULA 101
NEURO-BEHAVIOURAL UNDERSTANDING
OF LANGUAGE APTITUDE

/nebula101/

code	sub-pp011	sub-pp031	sub-pp052	sub-pp088	sub-pp114	sub-pp135	sub-pp162
derivatives	sub-pp012	sub-pp032	sub-pp053	sub-pp091	sub-pp115	sub-pp137	sub-pp163
neurobagel	sub-pp013	sub-pp033	sub-pp054	sub-pp092	sub-pp116	sub-pp145	sub-pp164
phenotype	sub-pp018	sub-pp035	sub-pp067	sub-pp093	sub-pp117	sub-pp147	sub-pp166
stimuli	sub-pp019	sub-pp036	sub-pp068	sub-pp094	sub-pp118	sub-pp149	sub-pp168
sub-pp001	sub-pp020	sub-pp038	sub-pp069	sub-pp095	sub-pp119	sub-pp150	sub-pp169
sub-pp003	sub-pp021	sub-pp039	sub-pp072	sub-pp099	sub-pp120	sub-pp151	sub-pp170
sub-pp004	sub-pp022	sub-pp042	sub-pp073	sub-pp105	sub-pp124	sub-pp152	sub-pp171
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sub-pp007	sub-pp025	sub-pp045	sub-pp077	sub-pp108	sub-pp127	sub-pp158	participants.tsv
sub-pp008	sub-pp026	sub-pp046	sub-pp078	sub-pp110	sub-pp128	sub-pp159	README
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- AliceLocalizer-4conditions
- create_nebula
- heudiconv
- preprocessing
- validation

/nebula101/derivatives/

📁 validation   ☐ nebula_101_all_task_scores.tsv

☐ cumulative_farsi_rater1_filtered.json   ☐ nebula_101_leapq_annotation_iso_glottolog.json

☐ cumulative_farsi_rater1_filtered.tsv   ☐ nebula_101_leapq_annotation_iso_glottolog.tsv

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sub-pp001_ses-01_task-aliceloc_run-001_bold.json   sub-pp001_ses-01_task-aliceloc_run-002_events.tsv
sub-pp001_ses-01_task-aliceloc_run-001_bold.nii.gz   sub-pp001_ses-01_task-aliceloc_run-003_bold.json
sub-pp001_ses-01_task-aliceloc_run-001_events.json   sub-pp001_ses-01_task-aliceloc_run-003_bold.nii.gz
sub-pp001_ses-01_task-aliceloc_run-001_events.tsv   sub-pp001_ses-01_task-aliceloc_run-003_events.json
sub-pp001_ses-01_task-aliceloc_run-002_bold.json   sub-pp001_ses-01_task-aliceloc_run-003_events.tsv
sub-pp001_ses-01_task-aliceloc_run-002_bold.nii.gz   sub-pp001_ses-01_task-rest_run-001_bold.json
sub-pp001_ses-01_task-aliceloc_run-002_events.json   sub-pp001_ses-01_task-rest_run-001_bold.nii.gz

/nebula101/sub-pp001/ses-01/beh/

sub-pp001_ses-01_task-amma_beh.json
sub-pp001_ses-01_task-amma_beh.tsv
sub-pp001_ses-01_task-ant_beh.json
sub-pp001_ses-01_task-ant_beh.tsv
sub-pp001_ses-01_task-apm_beh.json
sub-pp001_ses-01_task-apm_beh.tsv
sub-pp001_ses-01_task-artgram_beh.json
sub-pp001_ses-01_task-artgram_beh.tsv
sub-pp001_ses-01_task-brocanto_beh.json
sub-pp001_ses-01_task-brocanto_beh.tsv
sub-pp001_ses-01_task-corsiback_beh.json
sub-pp001_ses-01_task-corsiback_beh.tsv
sub-pp001_ses-01_task-corsifor_beh.json
sub-pp001_ses-01_task-corsifor_beh.tsv
sub-pp001_ses-01_task-cvlt_beh.json

sub-pp001_ses-01_task-cvlt_beh.tsv
sub-pp001_ses-01_task-cvltlong_beh.json
sub-pp001_ses-01_task-cvltlong_beh.tsv
sub-pp001_ses-01_task-digitback_beh.json
sub-pp001_ses-01_task-digitback_beh.tsv
sub-pp001_ses-01_task-digitfor_beh.json
sub-pp001_ses-01_task-digitfor_beh.tsv
sub-pp001_ses-01_task-farsi_beh.json
sub-pp001_ses-01_task-farsi_beh.tsv
sub-pp001_ses-01_task-fingertapping_beh.json
sub-pp001_ses-01_task-fingertapping_beh.tsv
sub-pp001_ses-01_task-hindi_beh.json
sub-pp001_ses-01_task-hindi_beh.tsv
sub-pp001_ses-01_task-mlat5_beh.json
sub-pp001_ses-01_task-mlat5_beh.tsv

sub-pp001_ses-01_task-purdue_beh.json
sub-pp001_ses-01_task-purdue_beh.tsv
sub-pp001_ses-01_task-ran_beh.json
sub-pp001_ses-01_task-ran_beh.tsv
sub-pp001_ses-01_task-rttsub_beh.json
sub-pp001_ses-01_task-rttsub_beh.tsv
sub-pp001_ses-01_task-rttsum_beh.json
sub-pp001_ses-01_task-rttsum_beh.tsv
sub-pp001_ses-01_task-spelling_beh.json
sub-pp001_ses-01_task-spelling_beh.tsv
sub-pp001_ses-01_task-textreading_beh.json
sub-pp001_ses-01_task-textreading_beh.tsv
sub-pp001_ses-01_task-wordreading_beh.json
sub-pp001_ses-01_task-wordreading_beh.tsv

Boxplot: Weighted IQR x Mean absolute Z-score
Common filename: ./mprage/sub-pp*