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Impact of analytic decisions on test-retest reliability of individual and group estimates in functional magnetic resonance imaging: a multiverse analysis using the monetary incentive delay task

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Abstract

Empirical studies reporting low test-retest reliability of individual blood oxygen-level dependent (BOLD) signal estimates in functional magnetic resonance imaging (fMRI) data have resurrected interest among cognitive neuroscientists in methods that may improve reliability in fMRI. Over the last decade, several individual studies have reported that modeling decisions, such as smoothing, motion correction and contrast selection, may improve estimates of test-retest reliability of BOLD signal estimates. However, it remains an empirical question whether certain analytic decisions *consistently* improve individual and group level reliability estimates in an fMRI task across multiple large, independent samples. This study used three independent samples (*N*s: 60, 81, 119) that collected the same task (Monetary Incentive Delay task) across two runs and two sessions to evaluate the effects of analytic decisions on the individual (intraclass correlation coefficient [ICC(3,1)]) and group (Jaccard/Spearman *rho*) reliability estimates of BOLD activity of task fMRI data. The analytic decisions in this study vary across four categories: smoothing kernel (five options), motion correction (four options), task parameterizing (three options) and task contrasts (four options), totaling 240 different pipeline permutations. Across all 240 pipelines, the median ICC estimates are consistently low, with a maximum median ICC estimate of .43 - .55 across the three samples. The analytic decisions with the greatest impact on the median ICC and group similarity estimates are the *Implicit Baseline* contrast, Cue Model parameterization and a larger smoothing kernel. Using an *Implicit Baseline* in a contrast condition meaningfully increased group similarity and ICC estimates as compared to using the *Neutral* cue. This effect was largest for the Cue Model parameterization; however, improvements in reliability came at the cost of interpretability. This study illustrates that estimates of reliability in the MID task are consistently low and variable at small samples, and a higher test-retest reliability may not always improve interpretability of the estimated BOLD signal.

Keywords: Test-rest reliability, Intraclass Correlation, Jaccard Similarity, Functional Magnetic Resonance Imaging, Monetary Incentive Delay task, Individual Differences

Introduction

Reliability in functional magnetic resonance imaging (fMRI) is essential to individual differences research as well as for the development of clinical biomarkers. Unfortunately, numerous studies have demonstrated that reliability of individual estimates in fMRI is low (Elliott et al., 2020; Noble et al., 2019) and the reliability of group estimates in statistical maps is sensitive to varying analytical decisions made by researchers (Botvinik-Nezer et al., 2020)[1]. Poor reliability can hamper validity in cognitive neuroscience research, reducing the ability to uncover brain-behavior effects (Hedge et al., 2018; Nikolaidis et al., 2022) and the ability to detect differences in distinct brain states and individual traits (Gell et al., 2023; Kragel et al., 2021). It remains to be seen whether certain analytic decisions *consistently* reduce individual and/or group reliability estimates of blood oxygen-level dependent (BOLD) activity across measurement occasions in univariate task fMRI analyses.

FMRI analysis involves a range of analytic decisions (Caballero-Gaudes & Reynolds, 2017; Soares et al., 2016) that can result in a vast number of statistical brain maps across which BOLD activity can vary subtly or substantially (Bowring et al., 2022; Carp, 2012; Li et al., 2021). Simple decisions, such as using different MNI template brains, can greatly affect the agreement between parameter estimates between two preprocessing pipelines (Li et al., 2021). Furthermore, the approach used to model a task design can also alter interpretations (Botvinik-Nezer et al., 2020). As a result of numerous arbitrary choices, preprocessing and task modeling decisions can significantly impact the reliability of voxel/region of interest (ROI) estimates (Dubois & Adolphs, 2016).

Different metrics of reliability provide quantitative indices of the consistency (or similarity) of estimates of BOLD activity in specific brain regions (or voxels) during fMRI task activation across repeated measurement occasions (Bennett & Miller, 2013). Researchers can quantify the consistency of two repeated measures in terms of estimated effects (continuous)

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[1] Reliability of parameter estimates at the individual level and thresholded activation maps at the group level have previously been distinguished as "reliability" and "reproducibility" of BOLD activity, respectively (Bennett & Miller, 2013; Plichta et al., 2012; Zuo et al., 2014). We elect to refer to individual and group estimates as distinct forms of reliability and use 'reproducibility' to refer to a broader set of concepts describing various aspects of the ability to reproduce or generalize a research finding (e.g. Goodman et al. [2016]).

and/or the presence/absence of a significant effect (binary). In terms of the continuous effects,

reliability is an estimate of the consistency of the numerical representation of a measure (e.g.,

BOLD activity in the supplementary motor area during a finger tapping task [Witt et al., 2008])

of a mental process (e.g., index finger movement) across repeated measurement occasions within

an *individual* (e.g., task fMRI contrasts across two or more sessions, which can be hours, days or

weeks). This form of reliability is usually calculated using an intraclass correlation (ICC) at the

whole brain (i.e., voxel-wise) and/or ROI level. In terms of binary estimates of an effect,

reliability is an estimate of an experimental task's (e.g., finger tapping task [Witt et al., 2008])

ability to evoke statistically significant activation (above a pre-specified threshold) in the same

regions for *groups* of subjects for a specific condition (e.g., finger movement versus rest) across

measurement occasions (e.g., task fMRI contrasts across two or more scanning sessions). Binary

estimates of reliability are often calculated using Dice (Rombouts et al., 1998) or Jaccard's

similarity coefficients (Maitra, 2010). Together, these two forms of reliability reflect the

consistency (or agreement) in either the magnitude or the binary statistical significance of an

experimental effect occurring during task fMRI.

Traditionally, empirical studies have referred to the "robustness" of above-threshold

activation signals in group fMRI analyses as an implicit indicator of reliability of an fMRI task.

While a useful heuristic, Fröhner et al. (2019) argued that robustness across measurement

occasions only represents reliability of *group* (overall average) BOLD activity and does not

accurately represent *individual* variability in BOLD activity. In addition, thresholding is a

nonlinear operation that can result in substantial variability (Cohen & DuBois, 1999). When

quantifying reliability of BOLD activity in the brain, researchers often report an ICC or a

similarity coefficient for task fMRI (Bennett & Miller, 2013; Fröhner et al., 2019). The lack of

standardization makes it challenging to precisely quantify reliability, relative to individual

differences, and assess the impact of different fMRI analysis decisions on continuous and binary

estimates of reliability.

To date, several studies have examined the impact of analytic decisions, such as spatial

smoothing, motion correction and contrast modeling, on individual estimates of reliability of task

fMRI. Caceres et al. (2009, $n = 10$) found that an optimal smoothing kernel size of 8-10 FWHM

(full-width half-maximum) on a 1.5T scanner with 3.75mm voxels improved reliability. Results

regarding the impact of motion correction on reliability are mixed, with Gorgolewski et al.

(2013, $n = 11$) reporting a positive effect on reliability while Plichta et al. (2012, $n = 25$)

reporting no effect during a reward task and a negative effect during a faces and N-back task on

reliability. However, in a large, young sample, Kennedy et al. (2022, $n = 5,979 - 6,593$) reported

that excluding high motion subjects modestly improved reliability. Finally, Han et al. (2022, $n =$

$29 - 120$) and Kennedy et al. (2022, $n = 5,979 - 6,593$) reported that using an implicit baseline

for different tasks (e.g., rest phase during the task) rather than a neutral cue increased reliability

across measurement occasions. Some, but not all, of these findings are consistent with a previous

review of the fMRI reliability literature (Bennett & Miller, 2013), which suggests that motion,

spatial smoothing and task signal likely impacts reliability in task fMRI. However, differences in

modeling decisions across these studies leaves an important question unanswered: Are there

certain analytic decisions that *consistently* improve reliability (e.g., ICC) of neural activity for an

fMRI task across samples?

The ICC is a statistic adopted from behavioral research to estimate reliability of observed

scores across measurement occasions (Bartko, 1966; Fisher, 1934; Shrout & Fleiss, 1979;

Spearman, 1904). In the context of multi-session data, there are several ways to estimate an ICC,

but for typical univariate fMRI studies, two specific types (ICC[2,1] and ICC[3,1]) are

recommended (For a discussion, see Noble et al., 2021). As described elsewhere (Bennett &

Miller, 2013; Fisher, 1934), the ICC is similar to the product moment correlation. Unlike the

product moment correlation, which estimates separate means and variances between distinct

classes (e.g., age and height), the ICC estimates the mean and variances within a single class

(e.g., measure). For two or more variables from a single class, test-retest reliability estimates the

consistency (or agreement) of the observed scores across the measurement occasions. Using the

correlation coefficient as an example, if there are no differences in subjects' scores across two

measurement occasions, the correlation coefficient would be 1.0. However, if the measure is

affected by systematic and/or unsystematic error across measurement occasions, this would

impact the covariance between observed scores across subjects and decrease the linear

association between measures across the two occasions. Unlike the product moment correlation,

however, the ICC factors out measurement bias which reflects the reproducibility of observed

scores across measurement occasions (Liu et al., 2016). While the correlation between two

occasions ($\mathbf{A} = [1, 3, 6, 9, 12]$ & $\mathbf{B} = 3x\mathbf{A} = [3, 9, 18, 27, 36]$) may be perfect ($r_{AB} = 1.0$), the

consistency in observed scores between the two measurement occasions would be lower

(ICC[3,1] = .60). In fMRI, the reliability of the BOLD signal may be impacted by biological

(e.g., differences in BOLD across brain region), analytic (e.g., task design and analytic

decisions), and participant-level factors (e.g., practice effects, motion, habituation and/or

development). These fluctuations, whether typical or atypical, may contribute to observed

differences and the reduced consistency in scores across measurement occasions, leading to

decreased estimates of reliability.

As discussed in prior work on fMRI reliability (Bennett & Miller, 2010, 2013; Caceres et

al., 2009; Chen et al., 2017; Herting et al., 2017; Noble et al., 2021), the ICC decomposes the

total variance of the data across all subjects and sessions into two key parts: *Between-subject* and

Within-subject variance (for statistical formulas and discussion of ICC, see Liljequist et al.,

[2019] and flowchart in McGraw & Wong [1996, p. 40]). The ICC estimate can be altered by

increasing the differences in BOLD activity between subjects (e.g., subjects differ more in

BOLD activity in index finger movements) and/or ensure that BOLD activity within subjects is

more similar across scans (e.g., BOLD activity in response to finger movements versus rest for

Subject A is consistent across Session 1 and Session 2). Some have argued that the low *between-*

subject variability may be a reason for low reliability of behavioral responses in experimental

tasks that are commonly used in fMRI (Hedge et al., 2018). However, there is little empirical

research on whether the culprit in the reportedly low reliability of fMRI signal across

measurement occasions is a *decreased between-subject* and/or an *increased within-subject*

variability. It also remains an open question whether certain analytic decisions differentially

impact the between/within subject variance and consistently improve reliability across different

samples with the same task. As it relates to prediction and global signal-to-noise ratio, evidence

from Churchill et al. (2015; $n = 25$) suggest that there are likely to be optimal preprocessing

pipelines; however, the degree to which these differ across datasets and individuals is currently

unknown.

The current study uses a multiverse (Steegen et al., 2016) of analytic alternatives to

simultaneously evaluate the effects of analytic decisions on the continuous and binary reliability

estimates of neural activity in task fMRI in three samples. The three samples administered with

the comparable Monetary Incentive Delay (MID) task during fMRI across two runs and two

sessions. The purpose of multiple samples with the same task design is to evaluate the

consistency in findings across studies that vary in their sample populations and task design as

little evidence exists on the *consistency* of reliability estimates for the same task across

independent samples. **Aim 1** evaluates the effects of analytic decisions including task model

smoothing, motion correction, parameterization (i.e., modeling) and task contrasts on the impacts

on reliability, calculated using ICC(3,1) for individual [continuous] beta estimates and Jaccard's

similarity coefficient using significance thresholded group [binary] estimates ($p < .001$,

uncorrected) and Spearman correlation group [continuous] estimates. The decisions are noted in

Table 1. **Aim 1 Hypothesis** is that the highest produced ICC and similarity

coefficient/correlation is for the model decisions indicated by **blue** for A-D decisions in Table 1.

This, in part, is because the analytic strategy includes 1) motion correction techniques that limit

the number of noisy (high motion) subjects and reduce the number of degrees of freedom that are

lost due to censoring, 2) an optimal smoothing for the size of voxels, and 3) the highest

activation contrast from a task modeling phase that is relatively efficient. We hypothesize this to

be more so the case for the older (e.g., AHRB/MLS) than younger samples (e.g., ABCD) due to

changes occurring as a result of development (Herting et al., 2017; Noble et al., 2021). Due to

the lack of information regarding how the between-subject variance (BS) and within-subject

variance (WS) is impacted by analytic choices in task fMRI analyses, **Aim 2** evaluates the

change in BS and WS components. Due to the poor reliability of individual estimates in task

fMRI (Elliott et al., 2020), reported evidence of high between-subject variability in BOLD

activity (Turner et al., 2018), and limited evidence on changes in BS and WS variance

components in the MID task, we do not have a specific **Aim 2 Hypothesis**. Finally, seeing as the

ICC is, in some ways, similar to a moment product correlation (Bennett & Miller, 2010) which

stabilizes at larger sample sizes (Grady et al., 2020; Marek et al., 2022; Schönbrodt & Perugini,

2013), **Aim 3** evaluates at what sample the ICC stabilizes using the most optimal pipeline (e.g.,

highest median ICC) used in Aim 2. Stability of Jaccard coefficient group maps is not considered

in Aim 3 as these estimates are sensitive to significance thresholding. Using the evidence from

prior work on correlations (Grady et al., 2020; Schönbrodt & Perugini, 2013), the **Aim 3**

Hypothesis is that the ICC will stabilize a sample size between 150 to 500.

Table 1. Proposed Analytic Permutations: 360 Total
Modeling Combinations for MID task

First-level Pipeline Decisions	Options
A. Smoothing (FWHM)	

1. 1.5x voxel	ON / OFF
2. 2x voxel	ON / OFF
3. 2.5x voxel	ON / OFF
4. 3x voxel	ON / OFF
5. 3.5x voxel	ON / OFF
B. Motion Correction	
1. None	ON / OFF
2. Regress: Translation/Rotation (x,y,z) + Derivative (x,y,z)	ON / OFF
3. Regress: Regress: Translation/Rotation (x,y,z) + Derivative (x,y,z) + First 8 aCompCor Components	ON / OFF
4. Regress: Translation/Rotation (x,y,z) + Derivative (x,y,z) + First 8 aCompCor Components + Censor High Motion Volumes (FD $\geq$ .9)	ON / OFF
[#]5. Regress: Translation/Rotation (x,y,z) + Derivative (x,y,z) + First 8 aCompCor Components, Exclude mean FD $\geq$ .9	ON / OFF
[#]6. Regress: Translation/Rotation (x,y,z) + Derivative (x,y,z) + First 8 aCompCor Components + Censor High Motion Volumes, Exclude mean FD $\geq$ .9	ON / OFF
C. Task Modeling	
1. MID: Cue Onset, Cue Duration only	ON / OFF
2. MID: Cue Onset, Cue + Fixation Duration	ON / OFF
3. MID: Fixation onset, Fixation Duration	ON / OFF
D. Task Contrasts	
1. MID: Big Win > Neutral	ON / OFF
2. MID: Big Win > Implicit	ON / OFF
3. MID: Small Win > Neutral	ON / OFF

4. MID: Small Win > Implicit	ON / OFF

Blue text: Model hypothesized to produce the highest test-retest reliability; aCompCor: Anatomical Component Based Noise Correction; MID: Monetary Incentive Delay task; FD: Framewise displacement.
[#]Due to the lack of low motion subjects (zero mean FD <.90 in 2/3 samples), this decision was not included in the Stage 2 analyses, resulting in 240 analytic models.

Methods

To answer the questions proposed in Aim 1 and Aim 2, this study will require multiple samples and tasks to obtain a comprehensive view of how analytic decisions impact group and individual reliability metrics (Aim 1) and how BS and WS is impacted (Aim 2) across multiple samples and similar MID task. We use three samples with subjects that have at least two repeated sessions of data. To answer the question about the sample at which ICC stabilizes (Aim 3), we use the repeated session data from a large consortium sample.

The studies were selected based on two criteria. First, the goal is to derive group and individual estimates of reliability using sample sizes that are larger than the reported median sample size in fMRI research. The median reported sample size in fMRI is <30 subjects (Poldrack et al., 2017; Szucs & Ioannidis, 2017). From the review of task fMRI reliability by Bennet and Miller (2010), the median sample for individual (continuous) reliability is 10 subjects (mean = 10.5 [range = 1 to 26]) and for group (binary) reliability is 9.5 subjects (mean = 11.2 [range = 4 to 45]). A recent review and analysis of task fMRI reliability suggests sample sizes are increasing but remain lower than the median sample size in task fMRI, whereby the median sample size for individual reliability in the meta-analysis are 18 subjects (mean = 26.4 [range = 5 to 467]) and the analyses are 45 & 20 subjects (Elliott et al., 2020). Second, the goal is to limit the interaction between reliability estimates and unknown features of the data, such as the mental processes, to get a sense of how the analytic pipeline impacts reliability estimates *consistently* across a similar task design. Thus, the three samples described below exceed N > 50 and use a nearly identical task that is known to evoke a strong BOLD response in specific brain regions to achieve these two goals.

Participants[2]

Adolescent Brain Cognitive Development (ABCD) Study

The ABCD Study® is a longitudinal national study that was designed to study the change

in behavioral and biological measurements across development (Volkow et al., 2018). The focus

here is on the 4.0 brain imaging data that is released by the ABCD-BIDS Community Collection

(ABCC; Feczko et al. [2021]). As of February 2024, the ABCC data contains year 1

(approximately 11,000, participants Aged 9-10) and year 2 (approximately 7,000 participants,

Age 11-13) fMRI data. For Aims 1 and 2, we use a subsample of ABCD participants at the

University of Michigan site (site = 13) with maximum clean data available as this would be

sufficient to test the hypotheses and limit site and scanner effects. For Aim 3, we use a

subsample of N = 2,000 of the maximum clean data available from the ABCC sample and use an

adaptive design to answer at which N ICC stabilizes. To reduce the use of unnecessary

computational resources, the analyses are first performed in N = 525. If the difference between

average ICC estimate for interval N_i & N_{i-1} is > .15, the sample will be extended to $N = 1000$,

adding $N = 500$, until the plotted estimates are stable. As described elsewhere (Casey et al.,

2018), the study collected fMRI data during the Stopsignal, Emotional N-back and MID tasks.

Reliability of consortium-derived region of interest level data for year 1 and year 2 has been

reported elsewhere (Kennedy et al., 2022). We expand on these findings by evaluating how

consistent these results are across studies and which analytic decisions impact estimates of

reliability. Here, we use the raw BOLD timeseries from the MID task as this is consistent with

the two other studies described below.

Michigan Longitudinal Study (MLS)

The MLS is a longitudinal study focused on the change in behavioral and biological

measurements across development. As described elsewhere (Martz et al., 2016; Zucker et al.,

2000), the MLS includes the Neuropsychological Risk cohort. The MLS Neuropsychological

Risk cohort contains year 1 (approximately 159 participants, Age 18-24) and year 2

(approximately 150 participants, Age 20-26) fMRI data. The study collected fMRI data during

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[2] For the Stage 1 submission, the data for the different studies was not fully accessed, inspected, preprocessed or analyzed. Thus, the sample size approximations. The final N for each sample is expected to deviate from the approximated values because of complete data availability and quality control exclusions.

the affective word and MID tasks. Here, we use the raw BOLD data from the MID task as it is

consistent with the ABCD study and Adolescent Risk Behavior Study (described below).

Adolescent Risk Behavior (AHRB) Study

The AHRB study is a longitudinal study focused on the change in behavioral and

biological measurements across development. The AHRB study contains year 1 (approximately

108 participants, Age 17-20) and year 2 (approximately 66 participants, Age 19-22). The study

collected fMRI data during the Emotional Faces and MID tasks. Here, we use the raw BOLD

data from the MID task as it is consistent with the MLS and AHRB study.

FMRI Task, Data, Preprocessing

FMRI Tasks

Across the ABCD, AHRB and MLS studies, reward processing was measured using

comparable versions of the MID task. The MID task (Knutson et al., 2000) is used to model

BOLD signatures of the anticipation and receipt of monetary gains or losses. The MID task and

their nuanced differences across the ABCD, AHRB and MLS studies are described in

supplemental **Section 1.2**. The focus of the present work is on the anticipatory phase of the task.

MRI Acquisition Details

The acquisition details for the AHRB, ABCD and MLS datasets are summarized in

supplemental **Section 1.3 Table S2**.

Data Quality Control and Preprocessing

First, quantitative metrics reported from MRIQC version 23.1.0 (Esteban et al., 2023) for

the structural and BOLD data are evaluated to assess data quality and potentially problematic

subjects. Second, behavioral data were inspected to confirm that participants have the behavioral

data for each run and that participants performed at the targeted probe hit rate (e.g., at or near

60% overall probe hit rate, see supplemental **Section 1.2**). Then, structural and functional MRI

preprocessing is performed using fMRIPrep v23.1.4 (Esteban et al., 2022; RRID:SCR_016216),

which is based on Nipype 1.8.3 (Esteban, Markiewicz, Burns, et al., 2022; RRID:SCR_002502)

and the results are inspected to confirm no subjects' preprocessing steps failed.

Preprocessing between the ABCD, AHRB and MLS are held constant except for two
differences. First, the MLS datasets did not collect fieldmaps and the repetition time for MLS
(2000ms) is slower than the repetition time (800ms) in ABCD/AHRB. Therefore, fMRIPrep's
fieldmap-less distortion correction (SyN-SDC) is used to estimate and correct for fieldmap
distortions in MLS and slice-timing correction is applied *only* on the MLS data. For the ABCD
and AHRB data, fieldmap-less distortion correction is used *only* when a subject does not have
the necessary fieldmaps. Outside of these two exceptions, the preprocessing of the BIDS data
were preprocessed using identical pipelines. The complete preprocessing details are included in
supplemental **Section 1.4**

Analyses

This project is focused on the effects of analytic decisions on estimates of reliability
across (run/session) measurement occasions in task fMRI. As a reminder, reliability is the
estimate of how similar two measures (in this case, voxels for a given contrast from a fMRI 3D
volume) are in terms of estimated effects (continuous) and/or the presence/absence of a
significant effect (binary). We distinguish individual and group estimates in **Figure 1** and
describe the calculations below. For the continuous estimates of reliability described below, the
analyses will be performed separately on task voxels that exceed and do not exceed an *a priori*
specified threshold applied on the NeuroVault (Gorgolewski et al., 2015) meta-analysis
collection that comprises the anticipatory win phase across 15 whole brain maps for the MID
task (Wilson et al., 2018; Collection: 4258, Image ID: 68843). The *suprathreshold* task-positive
voxels are those that exceed the threshold ($z > 3.1$) and the *subthreshold* task voxels are those
that do not exceed the threshold ($z < 3.1$) in the map. We acknowledge that the threshold of $z = $
3.1 is arbitrary (uncorrected, *p*-value = .001) and that the voxels that fall below and above this
threshold may not be significantly different (Gelman & Stern, 2006). However, to constrain the
problem space this is a researcher's decision that is made in these analyses (Gelman & Loken,
2014; Simmons et al., 2011).

Figure 1. Diagram of (**A**) Continuous (individual), (**B**/**C**) binary/continuous (group) and (**D**) random subsampling of Estimates of Reliability across Measurement Occasions in 3D volumes of fMRI data.

Group = group average of activation; Sub = Subject; ICC = Intraclass Correlation; Supra- and Sub-threshold mask is > 3.1 of NeuroVault Vault Image ID #68843 (Collection #4258)

Descriptive Statistics

The mean, standard deviation, count and frequencies are reported for demographic variables from the ABCD, AHRB and MLS datasets. For ABCD, AHRB and MLS, participants self-reported on Age, Sex and Race/Ethnicity. ABCD: Sex is reported as sex at birth (Male, Female, Other, or Not Reported); Race/Ethnicity is reported on a 5-item scale: White, Black, Hispanic, Asian, Other. AHRB: Sex is reported as sex at birth (Male or Female); Race/Ethnicity is available on a 4-item scale: White, Non-Hispanic, Black, Non-Hispanic, Hispanic/Latinx, Other. MLS: Sex is reported as Sex at Birth; Race is available on an 8-item scale: Caucasian,

African American, Native American, Asian American, Filipino or Pacific Islander, Bi-Racial,

Hispanic-Caucasian, and Other.

Behavioral data from the MID task, such as the mean and distribution of probe hit rate

and mean response times (RT) across subjects, will be reported as supplemental information. The

task design is programmed to achieve a probe hit rate of approximately 60% for each subject. It

should be noted that the RT for the probe is not consistently collected across the ABCD, AHRB,

and MLS datasets.

Impact of Analytic Decisions on Reliability in fMRI Data

First-, second- and group-level analyses are performed using Python 3.9.7 and Nilearn

0.9.2 (Abraham et al., 2014). Details about these three analytic steps are described below and the

code is provided on Github. As listed in **Table 1** and described next, the analytic decisions will

be limited to the first-level analysis.

Analytic Decisions: For reasons described in the introduction, the focus of analytic

decisions in this paper will be on **four** categories: Smoothing, Motion Correction, Task Contrast

and Task Parametrization. As reported in empirical studies and meta-analyses of task fMRI

reliability (Bennett & Miller, 2010; Caceres et al., 2009), one way to improve reliability of fMRI

data is by increasing the signal-to-noise ratio in the BOLD data through different smoothing

kernels (Caceres et al., 2009), reducing motion effects in the fMRI data (Gorgolewski et al.,

2013; Kennedy et al., 2022) and using task designs/contrasts that evoke increased neural activity

(Han et al., 2022; Kennedy et al., 2022). These analytic decisions are described in greater detail

in supplemental **Section 1.1**.

Within-run Analysis: A general linear model (GLM) is fit using Nilearn (e.g.,

FirstLevelModel) to estimate the response to task-relevant conditions in the BOLD timeseries for

each participant/voxel. The BOLD timeseries are masked and spatially smoothed using specified

full-width half-maximum (FWHM) Gaussian kernel options (see 'Smoothing' in **Table 1**) and

the timeseries are prewhitened using an 'ar1' noise model. A GLM is fit (using *FirstLevelModel*)

for a design matrix that includes the 15 task-relevant regressors (see task details in supplemental

Section 1.2) and a set of nuisance regressors. Depending on the decision criteria (see 'Motion

Correction' in **Table 1**), nuisance regressors may include, for example, **A**) estimated translation

and rotation (+ derivatives) of head motion or **A** + first eight aCompCor noise components and

the corresponding cosine regressors for high pass filtering (with a cutoff of 128 seconds) that are

calculated by fMRIPrep (see preprocessing of functional data). Task regressors are convolved

with the SPM hemodynamic response function (HRF). The resulting beta estimates from the

GLM, for each individual subject and run, are used to compute four contrasts for the MID task

(see 'Task Contrasts' in **Table 1**).

366   *Within-session Analysis*: Per subject, each study collected two runs for each of two

367   sessions. For each of the four contrast types, the beta and variances estimates from the two MID

368   runs for each subject are averaged using Nilearn's precision-weighted fixed effects model (i.e.,

369   *compute_fixed_effects*).

370   *Group-level Analysis (within-session)*: The MID task weighted fixed effects contrast files

371   are used in a group-level mixed effect model (i.e., Nilearn's *SecondLevelModel*) to average the

372   within-subject estimates across subjects. These group maps are used as measures of the average

373   activation patterns during the MID task in each of the studies across each of the four contrast

374   types within each session.

375   The resulting individual and group maps from the four contrasts are used in calculating

two different estimates of reliability (described in detail below). First, the resulting *within-run*

analysis maps (i.e., for each run) are used for the continuous estimate of reliability *within* each

session (i.e., reliability across runs). Then, the resulting *within-session analysis* maps, computed

from the weighted fixed effects model, are used in the continuous estimate of reliability *between*

the two sessions. Due to the temporal difference within and between sessions, the reliability

within sessions would be hypothesized to be greater than between sessions. The resulting group-

level analysis maps are used in the binary estimate of reliability *between* sessions.

Estimate of Reliability for Continuous Outcomes: Intraclass Correlation

384   Reliability for continuous outcomes at the individual level is estimated using ICC. The

ICC is an estimate of between-subject and within-subject variance that summarizes how similar

the signal intensities are for a given voxel from a 3D volume across sessions. As described in

Liljequist et al. (2019), there are several versions of the ICC, which vary in whether the subjects

and sessions are considered to be fixed (e.g., ICC[1]), subjects are considered to be random and

sessions are considered to be fixed (e.g., consistency, estimated via ICC[3,1]) or the subjects and

sessions are considered to be random (e.g., agreement, estimated via ICC[2,1]). In the case of

these analyses, we assume that subjects are random but do not assume that sessions are random

for two reasons. First, in the case of reliability of runs within a session, the runs are administered

in a fixed manner and the state of the participant cannot be assumed to be random for each.

Second, in the case of reliability across sessions, during the follow-up session subjects have

experienced the MRI environment and the task design in the scanner. In this case, again, it is

difficult to assume that sessions are in fact random as the practice and session effects may be

present. Thus, we estimate the consistency (ICC[3,1]) of the signal intensity for a given voxel

across measurement occasions.

Several packages exist to calculate ICC and Jaccard/Dice coefficients. For example,

_ICC_rep_anova_ & _Similarity_ in Python (Gorgolewski et al., 2011), _fmreli_ in MATLAB (Fröhner

et al., 2019) and _3dICC_ in AFNI (Chen et al., 2017). However, these packages are either a)

limited to a specific ICC calculation (e.g., ICC[3,1]), b) not easy to integrate into reproducible

python code (e.g., _fmreli_), c) do not include similarity calculations (e.g., _3dICC_), or do not return

information about between-subject, within-subject and between-measure variance components.

Thus, to have the flexibility to estimate ICC(1), ICC(2,1) and ICC(3,1), Dice and Jaccard

similarity coefficients and Spearman correlations simultaneously, we wrote and released an

open-source Python package with reliability and similarity functions that works on 3D NifTi

fMRI images.

The _PyReliMRI_ v2.1.0 (Demidenko, Mumford & Poldrack, 2024) Python package is used

to calculate continuous estimates of reliability. _PyReliMRI_ implements a voxel-wise ICC

calculation (e.g., _voxelwise_icc_) for 3D NIfTI images between runs and/or between sessions (see

the ICC example in study flowchart, **Figure 1A**). The function takes in a list of lists (e.g., list of

session 1 and list of session 2) of ordered paths to the preprocessed data [in MNI space] for

session 1 (or run 1) and session 2 (or run 2) subjects, and a binary [MNI space] brain mask. The

package is flexible to take in more than 2 sessions (or runs). An ICC type option (e.g., 'icc_1',

'icc_2' or 'icc_3') indicates the type of ICC estimate that is calculated across the voxels within

the masked 3D volume. The function returns a dictionary with five separate 3D volumes

containing the voxel-wise (1) ICC estimate, (2) lower bound ICC, (3) upper bound ICC, (4)

Between-subject variance (BS) and (5) Within-subject variance (WS) and, in case of ICC(2,1),

(5) Between-measure variance, or the measurement additive bias. Like the ICC & 95%

confidence calculation in the _pingouin_ package (Vallat, 2018), the ICC confidence interval in

PyReliMRI is calculated using the *f*-statistic (Bonett, 2002) to reduce the computation time

compared to using bootstrapped estimates.

$$ICC(3,1) = \frac{MSBS - MSError}{MSBS + MSError} = \frac{\sigma_r^2}{\sigma_r^2 + \sigma_v^2} \qquad \text{Equation 1}$$

Aim 1a: evaluated the effect of analytic decisions (see **Table 1**; **Figure 1A**) on the

ICC(3,1) (equation 1 for two measurement occasions) for individual [continuous] estimates of

voxel activity across the ABCD, AHRB and MLS studies. The parameters in Equation 1 are:

MSBS is the Mean Squared Between-subject Error and *MSError* is the Mean Squared Error. As

described in Liljequist et al. (2019), the differences in the numerator is the between-subject

variance (σ_r^2) and the denominator is the sum of the between-subject variance (σ_r^2) and the

within-subject variance (or noise, [σ_v^2]). For each study, *voxelwise_icc* within the *brain_icc.py*

script is used to estimate the voxel-wise ICC(3,1) for between run and between session reliability

across the 360 model permutations. First, voxel-wise average and standard deviation from the

resulting ICCs for the 360 model permutations are reported in two 3D volumes. Second, the

range and distribution of median ICCs across each study (three) and analytic decision category

(four) are plotted across suprathreshold task-positive and subthreshold ICCs using Rainclouds

(Allen et al., 2019) and the median and standard deviation are reported in a table. Third, to

visualize the ordered median ICCs across the 360 model permutations for suprathreshold task-

positive and subthreshold ICCs, specification curve analyses are used (Simonsohn et al., 2020).

Specifically, results across the 360 model permutations are reported using a specification curve

to represent the range of estimated effects across the variable permutations. This consists of two

panels: Panel A represents the *ordered* median ICC coefficients and the associated 95%

confidence interval (across samples) colored based on no significance (gray), negative (red) or

positive (blue) significance from the Null (Null here is 0) and Panel B represents the analytic

decisions from each of the four categories (see **Table 1)** that produced the median ICC estimates.

The median ICC estimates from the 360 models are reported separately for suprathreshold task-

positive and subthreshold activation (the specification curve for all ICC estimates for

suprathreshold task-positive and subthreshold activation are provided as supplemental

information). Finally, to evaluate the effect of the analytic decisions on the median ICC,

hierarchical linear modeling (HLM) is performed as implemented in the *lmer()* function from the

lme4 R package (Bates et al., 2020). HLM is used to regress the median ICC on the [four]

analytic decisions as fixed effects with a random intercept model is fit (Matuschek et al.,

2017)for samples across the suprathreshold task-positive and subthreshold maps. Multiple

comparisons corrections are applied using the Tukey adjustment as implemented in the *emmeans*

package (Lenth et al., 2023). For these HLM models, the interpretation focuses on the

significant, non-zero effect of an independent variable (e.g., smoothing) on the dependent

variable (e.g., median ICC) while the remaining independent variables are assumed to be zero.

Aim 2: evaluated the change in between- and within-subject variance across the analytic

model permutations. Similar to Aim 1 (**Figure 1A**), *voxelwise_icc* within the *brain_icc.py* script

is used to estimate the BS and WS across the 360 model permutations. The range and

distribution of median BS and WS across each study and analytic decision category are plotted

across suprathreshold task-positive and subthreshold BS/WS using Rainclouds. Then, two

separate specification curve analyses report the *ordered* median BS and WS coefficients in one

panel and the analytic decisions that produced the BS and WS estimates in a second panel

separately for suprathreshold task-positive and subthreshold activation. Finally, like Aim 1, two

HLMs are used to regress the median BS and median WS on the [four] analytic decisions as

fixed effects with a random intercept only for sample across the suprathreshold task-positive and

subthreshold maps. Multiple comparisons corrections are applied using the Tukey adjustment.

Like Aim 1, the interpretation focuses on the significant, non-zero effect of an independent

variable (e.g., smoothing) on the dependent variable (e.g., median BS or median WS) while the

remaining independent variables are assumed to be zero.

Aim 3: evaluated the sample size at which the ICC stabilizes (**Figure 1D**). The chosen

pipeline is based on the highest median ICC across the studies for the suprathreshold task-

positive mask from Aim 1a and is rerun for the ABCD sample. Based on this pipeline, the first-

level analysis steps are repeated for N = 525 from the N = 2000 subsample for only the ABCD

data. Then, *voxelwise_icc* within the *brain_icc.py* script is used to derive estimates of the median

ICC, BS and WS for the between runs (e.g., measurement occasions) reliability across randomly

sampled subjects for 25 to 525 subjects in intervals of 50. Similar to the methods in Liu et al.

(2023), 100 iterations are performed at each N (with replacement) and the median ICC, the

associated BS and WS estimates are retained from *voxelwise_icc*. The average and 95%

confidence interval for the estimates across the 100 iterations is plotted for each interval of *N*

with the y-axis representing the median ICC and x-axis representing *N*. The plotted values will

be used to infer change and stability in the estimated median ICCs and variance components

across the sample size. If stability is not achieved by *N* = 500, the sample is extended to *N* =

1,000 and the analyses are repeated.

Estimate of Reliability: Jaccard Coefficient for Binary & Spearman Correlation for Continuous

Outcomes

The estimate of reliability for group analyses is estimated using the Jaccard Similarity for

binary and Spearman correlation for continuous outcomes. The estimates are used to evaluate

how the MID task evokes BOLD activation above a pre-specified threshold (*p* < .001) in the

same voxels for *groups* of subjects across measurement occasions (run/session) in the ABCD,

AHRB and MLS studies.

The *PyReliMRI* package is used. *PyReliMRI* calculates the similarity between two 3D

volumes using a Jaccard's coefficient which, in short, is the intersection divided by the union

between two binary images (see **Figure 1B**) or the Spearman correlation, which is ranked

correlation between two continuous variables (see **Figure 1C**). The Jaccard coefficient ranges

from 0 to 1, whereby higher values reflect greater similarity between two images. Like the

product-moment correlation, the Spearman correlation ranges from -1 to 1, whereby values >0

indicate a positive association between images and values <0 indicate a negative association

between images. The function (i.e., *image_similarity*) takes in the paths for MNI *image file1* and

image file2, a specified MNI mask and integer (i.e., z-stat/t-stat) at which to threshold the image.

The images are masked (if a mask is provided), thresholded at the specified integer (if a

threshold is provided) and the resulting images are binarized per user's input (i.e., if threshold =

0, the resulting similarity = 1). Based on the specified similarity metric, the resulting estimates

are similarity (e.g., Dice/Jaccard) or correlation coefficient (e.g., Spearman) between the two 3D

NIfTI images. For similarity between 2+ NIfTI images, *pairwise_similarity* is used. Similar to

image_similarity, *pairwise_similaity* takes in paths for an MNI mask, a threshold integer for the

3D volumes and the similarity type. Unlike *image_similarity*, *pairwise_similarity* allows for a

list (2+) of paths pointing to 3D volumes and creates pairwise-combinations across the image

paths between which to estimate similarity. The function returns the similarity coefficient in a

dataframe with the resulting similarity (or correlation coefficient) and the image label (e.g.,

basename of the provided path for given volume).

$$J(A, B) \ = \ \frac{|A \cap B|}{|A \cup B|} \qquad \text{Equation 2}$$

$$Spearman\ Correlation_{A,B} \ = \ \frac{6 \Sigma d_i^2}{n(n^2 - 1)} \qquad \text{Equation 3}$$

Aim 1b: evaluated the effect of analytic decisions (see Table 1) in the Jaccard's similarity

coefficient (Equation 2; **Figure 1B**) and Spearman correlation (Equation 3; **Figure 1C**) using the

group binary & continuous estimates. In Equation 2, J(A, B) is the similarity coefficient between

A (session 1) and B (session 2). This is derived from intersection, |A ∩ B|, which represents the

elements that are common to both A and B divided by the union, |A ∪ B|, or the elements that are

both in A and/or B. In Equation 3, the Spearman Rank Coefficient, as implemented in Scipy stats

using *spearmanr* (Virtanen et al., 2020)*,* is ranked correlation between unthresholded images A

and B, whereby Σd^2 is the sum of squared differences between ranked values in session A and B,

normalized by (n * (n² - 1)).

Since the Jaccard similarity coefficient is sensitive to thresholding and sample size

(Bennett & Miller, 2010), in Aim 1b an equal sample size (e.g., N ~ 60^3) is chosen for each study

to compare how the similarity between sessions varies across studies. For all 360 pipelines, a

group-level (average) activation map is estimated for each session. In the case of the Jaccard

coefficient, the group maps are thresholded at *p* < .001. In the case of the Spearman coefficient,

the group maps are masked using a suprathreshold task-positive map from NeuroVault

(https://identifiers.org/neurovault.collection:4258; Image ID: 68843). Then, the paths for the

pipelines and sessions are called using the *pairwise_similarity* within the *similarity.py* script. The

resulting coefficients report the similarity between analytic pipelines and sessions for each study.

For each study, the coefficients are plotted to reflect the distribution and range of coefficients.

Both Jaccard's and Spearman correlation are reported separately. Like Aim 1a & Aim 2, two

HLMs are used to regress the Jaccard coefficients and Spearman correlation on the [four]

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[3] At Stage 1 the sample was based on an approximation. During Stage 2, we realized it would be more effective to take advantage of the complete available data by using standardized effect Cohen's *d* maps.

542 analytic decisions nested within study. Multiple comparisons corrections are applied using the
543 Tukey adjustment.

Results

Given the breadth of the analyses (see **Table 2**), the results in the main text focus on the
Session 1 between-run individual- and group-level reliability estimates for the supra-threshold
mask. Differences are briefly noted for between-session reliability estimates and sub-threshold
models and are reported in detail in the supplemental materials.

As permitted, aggregate and individual subjects' data are made publicly available on
NeuroVault (Gorgolewski et al., 2015) and/or OpenNeuro (Markiewicz et al., 2021). The
complete set of group-level and ICC maps are publicly available on Neurovault for ABCD (6180
images; https://identifiers.org/neurovault.collection:17171), AHRB (2400 images;
https://identifiers.org/neurovault.collection:16605) and MLS (2400 images;
https://identifiers.org/neurovault.collection:16606). For each run and session, the BIDS input
data and derivations for MRIQC v23.1.0 and fMRIPrep v23.1.4 are available on OpenNeuro for
AHRB (Demidenko, Huntley, et al., 2024) and MLS (Demidenko, Klaus, et al., 2024). Since the
ABCD data are governed by a strict data use agreement (March 2024), the processed data will be
made publicly available via the NDA at a later date as part of the ABCC release. The final code
for all analyses is publicly available on Github
(https://github.com/demidenm/Multiverse_Reliability[4]).

In the supplemental information of the Stage 1 submission, we stated that we would
adjust the smoothing weight for the MLS as its voxel size, 4 mm anisotropic, would result in
greater inherent smoothness of the data than ABCD/AHRB samples (2.4 mm isotropic voxel). A
weight of .50 was applied to the smoothing kernels of the MLS data. This resulted in 3.6, 4.8,
6.0, 7.2 and 8.4 mm smoothing kernels for the AHRB/ABCD data and 3.0, 4.0, 5.0, 6.0 and
7.0mm smoothing kernels for the MLS data (**Figure S4**). In the results, the MLS ordinal values
are relabeled to map onto the values used for AHRB/ABCD for reporting purposes.

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[4] Will revise with final Zenodo citation prior to Stage 2 acceptance.

568
Figure 2. Session 1 Between-runs and Between-sessions: Mean +/- 1 Standard Deviation (SD) of
Supra-threshold median Intraclass Correlation Coefficient (ICC), Jaccard and Spearman
Similarity Coefficients from 240 analytic models across ABCD, AHRB and MLS Samples.
Note: Estimates in supplemental **Table S5**

Deviations from Stage 1 Registered Report

There are one moderate and two minor deviations from the Stage 1 Registered Report

(https://doi.org/10.17605/OSF.IO/NQGEH). First, fieldmap-less distortion correction is not

applied on the MLS data because the data were collected using spiral acquisition. The ABCC

data selects a single fieldmap within a session to apply on *all* of the functional runs, so subjects

without a fieldmap folder are excluded and fieldmap-less distortion correction is not used on the

ABCD data. In AHRB, fieldmap-less distortion correction was used for only *one* subject.

Second, in Aim 1b we proposed to use thresholded images (e.g., $p < .001$, approx. $t > 3.2$) to

estimate the Jaccard/Spearman similarity between the model permutations for the estimated

group maps. However, this statistic is arbitrarily sensitive to differences in the number of model

permutations when subjects are excluded in cases of failed preprocessing features, such

aCompCor mask errors. To improve the interpretability of the similarity estimates across

analyses with different numbers of included observations (see supplemental **Figure S3**), we

converted all *t*-statistic group maps to Cohen's *d* effect size maps using the formula: $\frac{t-statistic}{\sqrt{N}}$.

Cohen's $d = .40$ is used as the alternative threshold for Aim 1b as for pre-registered N ~ 60 a

conversion of *t-statistic* = 3.2 would be near this threshold. Third, the analyses proposed to

evaluate 360 analytic decisions across the three samples. However, no subjects in the final

AHRB and MLS samples exceeded mean FD = .9 so it was not possible to perform Motion

option 5 (Motion option 3 + exclude mean FD ≥ .9) or Motion option 6 (Motion option 4 +

exclude mean FD ≥ .9). As a result, the model permutations are restricted to 240 permutations (5

= FWHM, 6 → 4 = Motion; 3 = Model Parameterization; 4 = Contrasts) with relevant data

across the three samples and are the focus of the below analyses.

Descriptive Statistics

The final sample for Aim 1 and Aim 2 for ABCD, AHRB and MLS samples (mean FD < .90) from the University of Michigan site that had two runs for at least two sessions, had behavioral data, and passed QC are Ns 119, 60 and 81, respectively. For $N = 15$ subjects in the ABCD sample aCompCor ROIs failed, but otherwise the data passed QC and so these subjects were not excluded in Motion option3 and option4 models that include the top-8 aCompCor components as regressors. The final random subsample from the Baseline ABCD data for Aim 3 is $N = 525$.

Demographic information across the three samples for Aim 1 and Aim 2 (ABCD = 119; AHRB = 60; MLS = 81) are reported in supplemental **Table S4**. The average number of days between sessions is largest for the MLS sample (1090 days), followed by ABCD (747 days) and AHRB (419 days; **Figure S5**). On average, mean FD was higher in the ABCD sample versus the AHRB and MLS samples (**Figure S6**; **Table S5**). The samples also differed on average response probe accuracy (%), whereby on average MLS participants had a higher and faster probe response accuracy than ABCD and AHRB samples.

The estimated model efficiency, defined as $Efficiency = \frac{1}{c(X\prime X)^{-1}c\prime}$, varied as a function of Model Parameterization and Contrast types across the three samples (see **Figure S7**). The Anticipation Model (i.e., onset times locked to Cue onset and duration the combined duration of Cue and Fixation cross) was consistently estimated to be the most efficient model across the three samples for the *Large Gain* versus *Neutral* and *Small Gain* versus *Neutral* contrasts.

Figure 3. Supra-threshold Median ICC Session 1 between-run reliability estimates for Contrast
(con) and Model Parameterization analytic options across the ABCD, AHRB and MLS samples.
Complete distribution across four analytic options in supplemental **Figure S9**.

Aim 1a: Effect of analytic decisions on median ICC estimates for individual continuous maps

Aim 1a proposed to evaluate the estimated individual map similarity between
measurement occasions (runs/sessions) using the ICC(3,1) across 240 pipeline permutations. In
Table S5 (**Figure 2**), the median between-run Session 1 ICCs are slightly lower than the
between-session ICCs (between-run: ABCD = .11 [range: -.04 - .43]; AHRB = .18 [range: .00 -
.52]; MLS = .18 [range: .04 - .55]; between-session: ABCD = .15 [range: .03 - .34]; AHRB = .21
[range: .04 - .53]; MLS = .21 [range: .06 - .47]). The mean and standard deviation of the 3D
volumes across the 240 analytic decisions are reported in supplemental **Figure S8**. Across the
three samples, a consistent pattern is observed, whereby the regions with the highest ICCs, on
average, are within the visual and motor regions. Notably, the lowest ICCs, on average, are
within the ventricles and white matter. The supra-threshold distribution of the median estimates
across the four model options and three samples are reported in Figure 3 and the specification

curve of the median ICC estimates are reported in Figure 4. Note, the sub-threshold reported in supplemental **Figure S10**.

Figure 4. The supra-threshold Specification Curve of the Session 1 Between-run Median ICC estimates across 240 pipeline permutations for the ABCD, AHRB and MLS samples. Full length of estimates reported in **Figure S11**.

639 A. The distribution of the point estimate (average) and distribution (error bars) across the three samples. B. The
640 model options (four) associated with each estimate.
641
The effects reported in **Figure 3** and **Figure 4** illustrate that the largest differences in the

median ICC estimate is associated with model parameterization and the contrast type. Even

though the Anticipation Model ('AntModel') has the highest estimated contrast efficiency within

each sample, contrary to our hypothesis the highest median ICC is associated with the Cue

Model ('CueMod') in which the onset and duration are locked to the cue stimulus. However,

using an interaction to probe the distributions in **Figure 3**, *post hoc* analyses suggest the Cue

Model finding is largely driven by the *Implicit Baseline* contrasts (see Aim 1b) and the plot of

the Model Parameterization-by-Contrast in supplemental **Figure S12** suggests negligible

differences between Model Parameterization for the contrast of the *Neutral* contrasts.

Independent of model parameterization and consistent with our hypothesis and previous

reports in the task fMRI literature (Han et al., 2022; Kennedy et al., 2022), the highest median

ICC is consistently observed for the *Large Gain* versus *Implicit Baseline* contrast. In line with

the reported estimates in **Figure 3** and **Figure 4**, the HLM model for the supra-threshold mask

shows a significant association between different FWHM, Motion, Model Parameterization and

Contrasts model options compared to their respective reference values (**Table 3**). Specifically,

the median ICC estimates increased with larger smoothing kernels and decreased with more

stringent motion correction. Additionally, primarily driven by the *Implicit Baseline* conditions,

median ICC for the 'CueMod' and 'FixMod' increased in comparison to the 'AntMod' (see

interaction plot in **Figure S12**). Last, median ICC decreased in comparison to the *Large Gain*

versus *Implicit Baseline* contrast. For example, the contrast *Large Gain* versus *Neutral* has an

median ICC that is .17 lower, on average, compared to the *Implicit Baseline* contrast when

holding other decisions constant (see marginal means comparisons in supplemental **Table S6**).

While most parameters are significant in **Table 3**, the effects vary in their relative importance in

the model. The variability in the median ICC estimate across 240 pipelines and three samples is

best explained by contrast (marginal ΔR2: .55) and model parameterization (marginal ΔR2: .10).

FWHM and motion had a smaller impact on ΔR2, .03 and .03 respectively. In fact, including

aCompCor components (Motion option 3) and aCompCor components + censoring high motion

volumes (Motion option 4) is associated with a slight decrease in the median ICC estimate as

compared to no motion correction (Motion option 1), b = -.05 and b = -.05, respectively. A

similar finding is observed for the sub-threshold mask, whereby the contrast (ΔR2: .56) and

model parameterization (ΔR2: .10) decision had a larger impact on ΔR2 than the FWHM (ΔR2:

.04) or motion (ΔR2: .02) decisions (see **Figure S14**; **Table S7).** In general, the voxelwise

distribution of ICC estimates tends to be higher for the supra-threshold mask than the sub-

threshold masks (see supplemental Figure S14). Interpretations are generally consistent for

between-session median ICC estimates across the 240 pipeline permutations (see **Table S9** and

Figure S18, **S19**).

We had hypothesized that the ICC estimates in the older samples (AHRB/MLS) would

meaningfully differ from the younger sample (ABCD). Overall, ICC estimates were higher in the

older than younger sample for *between-run*, $t(497.2) = 5.53$, p < .001, $d = .43$, and *between-*

session, $t(669.9) = 9.57$, p < .001, $d = .66$.

Table 3. Hierarchical Linear Model: (A) Linear associations between the analytic decisions and

the *Session 1 between-run median Intraclass Correlation Coefficient (ICC[3,1]), Between-*

subject (BS) and Within-subject variance (WS) from supra-threshold mask and (B) the impact of

the analytic category on the marginal R^2.

A. HLM Estimates for Supra-threshold Mask

	Median ICC(3,1)			Median BS			Median WS		
Predictors	*b*	*CI*	*p*	*b*	*CI*	*p*	*b*	*CI*	*p*
(Intercept)	.23	.20 – .26	<.001	.27	.18 – .35	<.001	.91	.72 – 1.10	<.001
Reference [3.6]									
fwhm [4.8]	.02	.01 – .04	.003	-.03	-.06 – .00	.09	-.23	-.28 – -.18	<.001
fwhm [6.0]	.04	.03 – .06	<.001	-.04	-.07 – -.01	.003	-.36	-.41 – -.31	<.001
fwhm [7.2]	.06	.04 – .07	<.001	-.06	-.09 – -.03	<.001	-.44	-.49 – -.39	<.001
fwhm [8.4]	.07	.05 – .08	<.001	-.07	-.10 – -.04	<.001	-.49	-.54 – -.44	<.001
Reference [opt1]									
motion [opt2]	-.01	-.03 – .00	.07	-.04	-.06 – -.01	.01	-.14	-.18 – -.09	<.001
motion [opt3]	-.05	-.06 – -.04	<.001	-.10	-.13 – -.08	<.001	-.23	-.28 – -.19	<.001
motion [opt4]	-.05	-.06 – -.03	<.001	-.10	-.13 – -.08	<.001	-.24	-.28 – -.20	<.001
Reference [AntMod]									
model [CueMod]	.10	.09 – .11	<.001	.15	.13 – .17	<.001	.26	.23 – .30	<.001
model [FixMod]	.05	.04 – .06	<.001	.12	.10 – .14	<.001	.27	.23 – .31	<.001
Reference [LgainBase]									
con [LgainNeut]	-.17	-.18 – -.16	<.001	-.22	-.25 – -.19	<.001	-.28	-.32 – -.23	<.001
con [SgainBase]	-.02	-.04 – -.01	<.001	-.02	-.05 – .00	.09	.00	-.04 – .05	.93
con [SgainNeut]	-.23	-.24 – -.22	<.001	-.24	-.27 – -.21	<.001	-.31	-.35 – -.26	<.001

B. Analytic Category Model Impact

Comparison	χ2	Orig R2	New R2	ΔR2	χ2	Orig R2	New R2	ΔR2	χ2	Orig R2	New R2	ΔR2
[Full] vs [New - fwhm]	95	.72	.69	.03	25	.47	.45	.02	384	.52	.31	.21
[Full] vs [New - motion]	81	.72	.69	.03	81	.47	.42	.05	138	.52	.46	.06
[Full] vs [New -	263	.72	.62	.10	162	.47	.37	.10	221	.52	.42	.10

model]											
[Full] vs [New - con]	864	.72	.17	.55	397	.47	.17	.30	285	.52	.38	.14

Summary of Findings for Aim 1a:

Overall, between-run ICCs are slightly lower than between-session ICCs. Across the

three samples, the highest ICCs, on average, are within visual and motor areas and the lowest

ICCs are within the ventricles and white matter. In Table 1, it was hypothesized that the optimal

analytic decisions would be: FWHM Smoothing 2.5x the voxel size, Motion correction that

includes translation/rotation, their derivatives, the first 8 aCompCor components and exclusion

of > .90 mFD subjects, the anticipation Model Parameterization, and Contrast *Large Gain >*

Implicit Baseline. Contrary to registered hypotheses: (1) smoothing had a small but linear effect

on ICC estimates, whereby the largest median ICC was for the largest FWHM smoothing kernel

(3.5x voxel size); (2) Motion correction had minimal and negative impact on median ICCs in

case of more rigorous corrections; and (3) the Cue and Fixation Models had higher estimated

median ICCs than the Anticipation model. *Post hoc* analyses illustrated Model Parameterization

is largely driven by the Implicit Baseline contrast, as Model Parameterization has a negligible

impact on between condition contrasts. Consistent with registered hypotheses, the *Large Gain*

versus *Implicit Baseline* had the highest estimated median ICC. Contrary to registered

hypotheses, there was little evidence to suggest that analytic decisions differentially impacted

estimated median ICCs between developmental samples (e.g., oldest MLS/AHRB versus

younger ABCD data). Finally, the older samples (AHRB/MLS) had higher between- and

between-session estimated ICCs than the younger sample (ABCD).

Aim 1b: Effect of analytic decisions on Jaccard (binary) and Spearman

(continuous) similarity estimates of group maps

Aim 1b proposed to evaluate the estimated group map similarity between measurement

occasions (runs/sessions) using a Jaccard similarity for thresholded binary maps and a Spearman

similarity for continuous measures across the 240 pipeline permutations. The distribution of the

estimates across the four model options and three samples are reported in **Figure 5** for Jaccard

and supra-threshold Spearman similarity. The specification curve of the Session 1 between-run

estimates are reported in **Figure 6** for Spearman similarity (see **Figure S21** for Jaccard)**.** Based

on the group-level Cohen's *d* maps, there is a high similarity between the *Small Gain* and *Large*

Gain versus *Implicit Baseline* (and *Large Gain*) contrasts that appears to be driven by the

Implicit Baseline condition and high similarity between Cue and Fixation models (see **Figure**

S22).

717
718 *Figure 5.* (**A**) *Jaccard* and (**B**) *supra-threshold Spearman Session 1 Between-run* similarity

719   estimates across [Four] analytic options for between-run reliability across the ABCD, AHRB and

720   MLS samples.

722
Figure 6. The supra-threshold Specification Curve of the *Session 1 Between-run Spearman*
similarity estimates across 240 pipeline permutations for the ABCD, AHRB and MLS samples.
A. The distribution of the point estimate (average) and distribution (error bars) across the three samples. B. The
model options (four) associated with each estimate.

Similar to Aim 1a (**Table S5**; **Figure 2**), on average the Session 1 between-run supra-

threshold Spearman similarity is slightly lower than the supra-threshold between-session

Spearman similarity (between-run: ABCD = .68 [range: .35 - .89]; AHRB = .73 [range: .22 -

.96]; MLS = .84 [range: .47 - .96]; between-session: ABCD = .80 [range: .40 - .94]; AHRB = .82

[range: .32 - .97]; MLS = .87 [range: .59 - .97]). A similar trend is observed for the Jaccard

Similarity coefficient. The effects reported in **Figure 5** illustrate that the analytic categories have

unique impacts on the estimated Jaccard and supra-threshold Spearman coefficients. While the

Jaccard coefficient varies most across contrast and model parameterization options (**Figure 5A**),

the Spearman similarity varies most across FWHM and contrast type (**Figure 5B**). The

specification curve for the Spearman similarity coefficients illustrate a near ceiling similarity for

estimates at the upper tail of the estimates and little variability across the three samples (**Figure**

6). The HLM estimates indicate that a change from 3.6 to 8.4 FWHM results in a $b = .08$

increase in Jaccard similarity and a $b = .13$ increase in Spearman similarity. Furthermore, the

change from the contrast *Large Gain* versus *Implicit Baseline* to *Large Gain* versus *Neutral*

results in a $b = -.09$ decrease in Jaccard Similarity and a $b = -.20$ decrease in Spearman

similarity. While most parameters are significant in **Table 4**, the effects vary in relative

importance in the model. The variability in the estimated coefficients across 240 pipelines and

three samples is best explained by Contrast (marginal ΔR^2: .21) and model parameterization

(marginal ΔR^2: .05) for Jaccard similarity coefficient, and Contrast (marginal ΔR^2: .66) and

FWHM (marginal ΔR^2: .08) for supra-threshold Spearman similarity coefficient. Surprisingly,

the motion regressor options had a near-zero impact on the variability on both Jaccard and

Spearman similarity coefficients. Similar to Aim 1a, *post hoc* analyses illustrate an interaction

between Contrasts and Model Parameterization (**Figure S23**), whereby the largest driver of

Model Parameterization differences in the Spearman *rho* similarity is as a function of the

contrasts included the *Implicit Baseline*.

Table 4. Hierarchical Linear Model: (A) Linear associations between the analytic decisions and

the *Jaccard and Spearman supra-threshold* mask Session 1 between-run similarity and (B) the

impact of the analytic category on the marginal R^2.

A. HLM Group-map Estimates						
		Jaccard			Spearman	
Predictors	*b*	*CI*	*p*	*b*	*CI*	*p*
(Intercept)	.20	.09 – .31	<.001	.76	.69 – .83	<.001
Reference [3.6]						
fwhm [4.8]	.03	.01 – .05	.004	.05	.04 – .07	<.001
fwhm [6.0]	.05	.03 – .07	<.001	.09	.07 – .10	<.001
fwhm [7.2]	.07	.05 – .09	<.001	.11	.10 – .13	<.001
fwhm [8.4]	.08	.06 – .10	<.001	.13	.12 – .15	<.001
Reference [opt1]						
motion [opt2]	.01	-.00 – .03	.13	.01	-.00 – .03	.05
motion [opt3]	.00	-.02 – .02	.85	.01	-.00 – .02	.20

motion [opt4]	.00	-.01 – .02	.69	.01	-.00 – .03	.08
Reference [AntMod]						
model [CueMod]	.05	.04 – .07	<.001	.02	.01 – .03	<.001
model [FixMod]	.08	.07 – .10	<.001	.01	-.00 – .02	.18
Reference [LgainBase]						
con [LgainNeut]	-.09	-.10 – -.07	<.001	-.20	-.21 – -.18	<.001
con [SgainBase]	-.03	-.05 – -.01	.001	-.01	-.02 – -.00	.17
con [SgainNeut]	-.18	-.20 – -.16	<.001	-.34	-.35 – -.32	<.001

B. Analytic Category Model Impact

Comparison	$\chi 2$	Orig R2	New R2	ΔR2	$\chi 2$	Orig R2	New R2	ΔR2
[Full] vs [New - fwhm]	78	.30	.26	.04	292	.74	.66	.08
[Full] vs [New - motion]	3	.30	.30	.00	5	.74	.74	.00
[Full] vs [New - model]	104	.30	.25	.05	14	.74	.73	.01
[Full] vs [New - con]	348	.30	.09	.21	1205	.74	.08	.66

The group-level maps indicate a notable difference in contrasts using the *Neutral* and

Implicit Baseline conditions (NeuroVault ABCD:

https://identifiers.org/neurovault.collection:17171 AHRB:

https://identifiers.org/neurovault.collection:16605

; MLS: https://identifiers.org/neurovault.collection:16606). As **Figure S22** shows, the *Large*

Gain versus *Neutral* contrast reflects a qualitatively comparable activation map across Cue,

Fixation and Anticipation Models. On the other hand, the *Large Gain* versus *Implicit Baseline*

contrast differs across models, where the most notable pattern is that the Cue model is negative

of the Fixation model across the samples. Specifically, in ABCD, AHRB and MLS there is

increased negative activity in the insular, visual, motor and visual areas, in the Cue Model, and

this pattern is mostly opposite of the Fixation Model. Meanwhile, in the Anticipation model there

is high positive activity in the dorsal striatal, SMA and Insular regions. This reflects the variable

meanings of *Implicit Baseline* across the models. The relative symmetry between the Cue and

Fixation models is consistent with the fact that each serves as the B_0 in the models, e.g.,

$B_{1[Condition\ A, Cue]} - B_{0[All\ Fixation\ +\ Probe\ Phase]}$ and

$B_{1[Condition\ A, fixation]} - B_{0[All\ Cue\ +\ Probe\ Phase]}$. The Anticipation model is more variable as it

is contrasted with a more narrow phase of the task, e.g., $B_{1[Condition\ A, Cue+Fixation]}$ −

$B_{0[Probe\ Phase]}$.

Summary of Findings for Aim 1b:

Similar to Aim 1a, on average, the supra-threshold Session 1 between-run Spearman and

Jaccard similarity is slightly lower between-session similarity. Spearman similarity meaningfully

differed across Contrast, Model Parametrization and Smoothing, and it is near the ceiling for the

upper tail of the Spearman similarity estimates. Like Aim 1a, Model Parametrization is driven by

the Implicit Baseline. Finally, mean-based group activity maps illustrate that the Cue and

Fixation models are opposite of each other when the contrast is a between condition and implicit

baseline comparison.

Aim 2: Effect of analytic decisions on median BS/WS estimates from individual

continuous maps

Aim 2 proposed to evaluate the changes in the Between-subject variance (BS) and

Within-subject variance (WS) components that differentially relate to the ICC(3,1) across the

240 workflow permutations. The supra- and sub-threshold distributions across the four model

options and three samples are reported in supplemental **Figure S24 & S25** and specification

curves for BS in supplemental **Figure S28** and WS in supplemental **Figure S29**. The HLM

estimates (**Table 3**) suggest that the Implicit Baseline contrasts increase BS variance and more

stringent motion correction decrease BS variance, and Implicit Baseline contrasts and larger

smoothing kernels reduce WS variance. The variability in the estimated BS coefficients across

240 pipelines and three samples is best explained by Contrast (ΔR^2: .30), model parameterization

(ΔR^2: .10) and then motion (ΔR^2: .04). The variability in the estimated WS coefficients across

240 pipelines and three samples is best explained by FWHM (ΔR^2: .21), Contrast (ΔR^2: .14) and

then model parameterization (ΔR^2: .10). A comparable trend is observed in the between-session

estimates (**Table S9**), with the exception of Contrast selection explaining more variability (ΔR^2:

.26) than FWHM (ΔR^2: .16). We avoid interpreting the sub-threshold mask as it includes regions

that are high-noise (e.g., white matter and ventricles) and drop-out areas (e.g. cerebellar and

medial orbital frontal cortex) which exaggerates the BS and WS components.

Aim 3: Stability of the ICC, BS and WS Components across Sample Size

As expected, based on sampling theory which demonstrates that variability decreases as a
function of the square root of N, the variability in estimates decreased as N increased.
Specifically, the bootstrapped estimates for the median ICC, BS and WS change slowly at higher
intervals of N (**Figure 7**). In *post hoc* comparisons of whole brain voxelwise ICC maps, the
largest variability occurs below N = 275. As reported in supplemental **Figure S36**, at $N = 25$ the
minimum and maximum median whole brain ICC maps have a wider voxelwise distribution of
ICC values which are notably different (Cohen's $d = 1.9$). With increasing $N,$ Cohen's d of the
whole brain voxelwise distributions between the minimum and maximum 3D ICC maps narrows,
$d = 1.4$ at $N = 225$ and $d = 1.0$ at $N = 525$, respectively.

Figure 7. Changes in the Supra- & Supra-threshold Median Intraclass Correlation (ICC), Between-subject variance (BS) and Within-subject variance (WS) estimate in the ABCD sample for *N* 25 to 525 with 100 bootstraps at each *N*

Note: Based on the top model from Figure 2: *Small Gain* vs *Implicit Baseline* Contrast, 'CueMod' Model, Motion option 1 and FWHM 8.4.

Post Hoc Analyses

An exploratory set of analyses were performed to evaluate 1) the effect of analytic decisions on ICC for the Left and Right Nucleus Accumbens and 2) the association between voxelwise Cohen's *d* estimates at the group-level and the voxelwise ICC maps. These are reported in supplemental **section 2.6.**

Discussion

Understanding the analytic decisions that may consistently increase individual- and/or group-level reliability estimates has implications for the study of individual differences using fMRI. The current study expands on previous work by simultaneously evaluating the effects of smoothing, motion correction, task parameterization and contrast selection on the continuous and binary reliability estimates of BOLD activity during the MID task for run- and session-level data across three independent samples. The five major findings are: (1) The ICC(3,1) test-retest reliability estimates in the MID task are consistently low; (2) Group-level estimates of reliability are higher than individual [ICC] estimates; (3) Contrast selection and Model Parameterization have the largest impact on median ICC estimates, and Smoothing and Contrast selection has the largest impact on similarity estimates; however, gains in reliability across different contrasts comes at the cost of interpretability and may differ; (4) Motion correction strategies in these analyses did not meaningfully improve individual or group similarity estimates and, in some cases, *reduced* estimates of reliability; and (5) the median ICC estimate varied across sample size but the variability decreased with increased sample size. Excluding some differences, the results are relatively consistent across the three samples, runs and sessions, providing a comprehensive overview of how analytic decisions at the GLM impact reliability of estimated BOLD in commonly used versions of the MID task.

The findings from these multiverse analyses confirm previous reports that ICC estimates are relatively low in univariate task-fMRI and in the current state are inadequate measures for use in individual differences research (Elliott et al., 2020; Kennedy et al., 2022). Consistent with Elloitt et al (2020), reliability estimates in the sub-threshold (or non-target mask) are lower than the supra-threshold of the MID task (target mask). The range of median ICCs varied across analytic decisions. Using commonly employed cut-offs (Cicchetti & Sparrow, 1981; Elliott et al., 2020; Noble et al., 2019), ICC estimates for *Large Gain* versus *Neutral* contrast are in the 'Poor'

range and the *Large Gain* versus *Implicit Baseline* contrast ranged between 'Poor' and 'Fair'
across the three samples. Test-retest reliability for the *Large Gain* (*Small Gain*) versus *Implicit*
Baseline contrast are modulated by Model Parameterization, whereby the Cue Model had a
meaningfully higher reliability than the Anticipation Model. However, this may come at the cost
of validity, which is discussed below. Nevertheless, based on voxelwise distributions from the
top performing model (Model: Cue Model, Contrast: *Small Gain* versus *Implicit Baseline*,
Motion Correction: None, Smoothing: 8.4 mm kernel), visual and motor regions had the highest
ICCs, in the 'Fair' to 'Good' range. *Post hoc* analyses of the bilateral NAc illustrate that, on
average, ICC estimates in this region of interest are in the 'Poor' range. Notably, ICCs in this
post hoc region were not meaningfully impacted by Model Parameterization but were impacted
by Contrast and Motion correction, suggesting that test-retest reliability may be uniquely
impacted by analytic strategy depending on the voxels under consideration. These findings
illustrate that the test-retest reliability of the MID task is relatively low, even in the most
common ROI such as the Left and Right NAc. While Kennedy et al. (2022, p. 13) speculated that
low reliabilities in the ABCD sample may be attributed to the participants' young age, our results
demonstrate that median ICC estimates are *higher* in older than younger samples but reliability
estimates in the MID task remain consistently low across early adolescents and late
adolescents/young adults. To understand how analytic strategies differentially impact ICCs in
different brain regions, we encourage future researchers to use the publicly available estimated
maps to probe this question further.

       Consistent with Fröhner et al. (2019), the group-level maps are not always representative
of the individual-level maps across analytic decisions. On average, the Spearman *rho*, Jaccard
coefficients and median ICC estimates are higher for the between-session than between-run
estimates. Consistently, Spearman *rho* estimates are meaningfully higher for supra-threshold
group maps than supra-threshold median ICC estimates derived from individual maps. This
suggests that across each of the three samples, the MID task is relatively effective at eliciting a
group-level activation map; however, the individual estimates are lower and more variable. In
the context of the MID task, the between-run and between-session effects may be the result of
within-session effects *decreasing* across runs (Demidenko, Mumford, et al., 2024). Notably, the
higher between-session than between-run reliabilities is inconsistent with values reported in
previous work (Fröhner et al., 2019), this is likely the result of those between-run estimates being

based on randomly split-half (within runs) which are inflated as a result of dependencies in the
model estimates within runs (Mumford et al., 2014). Nevertheless, the results here emphasize
that group-level maps and group similarity are not a good indicator of individual-level
reliabilities. This is unsurprising, considering that the MID task design was optimized to elicit
activity in anatomical regions at a group-level and for averaged time-courses within an
anatomical region (Knutson et al., 2003).

A major question of these analyses was: Are there decisions that *consistently* result in
higher individual- (continuous) and/or group-level reliability estimates (continuous/binary)? The
results across the analytic choices illustrate that reliability estimates are impacted most by
contrast, model parameterization and smoothing decisions. Across the three samples, for
between-run and between-session estimates, the contrast type had the largest influence of
individual and group reliability estimates. Consistent with previous reports (Baranger et al.,
2021; Han et al., 2022; Kennedy et al., 2022; Vetter et al., 2015, 2017), the contrast *Large Gain*
(and *Small Gain*) versus *Implicit Baseline* had meaningfully higher estimated ICC, Jaccard and
Spearman *rho* similarity estimates than the *Large Gain* versus *Neutral* contrast. The estimated
ICC and Spearman *rho* coefficients for contrasts are modulated by the model parameterization,
whereby the conditions including the *Implicit Baseline* are highest for the Cue Model
parameterization. Conversely, ICC and similarity estimates are relatively stable across the three
model parameterizations when comparisons are against the *Neutral* condition. Whether using
contrasts or percent signal changes, estimates of BOLD activity suffer from decreases in
reliability due to difference scores (Hedge et al., 2018). Where gains are observed from the less
reliable *Large Gain* versus *Neutral* to the more reliable *Large Gain* versus *Implicit Baseline*
contrast, it comes at the cost of interpretability and face validity that is expected in the estimated
BOLD activity. Finally, higher FWHM smoothing kernels positively impacted between-run and
between-session median ICC estimates and Spearman *rho* similarity estimates whereas motion
correction strategies had a smaller but negative impact on these estimates (i.e., more stringent
motion correction reduced reliability estimates). Decisions to smooth in the MID task are
especially important given that larger smoothing kernels have been reported to spatially bias
reward-related activity in the MID task (Sacchet & Knutson, 2013). In general, variability in
reliability estimates decreased with large sample sizes.

Improvements in estimated reliability as a function of contrast selection may come at the

cost of interpretability. For example, in the context of the *Large Gain* versus *Neutral* contrast,

despite differences in the estimated efficiencies the ICC estimates are relatively stable across the

model parameterizations in each of the three samples and the activation patterns are interpretable

at the group-level. In the context of the *Large Gain* versus *Implicit Baseline* contrast, there are

meaningful differences in the ICC estimates across model parameterizations, whereby the Cue

and Fixation models demonstrate a substantial improvement over the Anticipation model

parameterization, but the group-level activity patterns are less interpretable. As a researcher

looking for BOLD estimates that are consistent from run-to-run or session-to-session for

individual participants, the *Implicit Baseline* suggests a considerable and valuable improvement

on the reliability of estimated values. However, the difference of means for the *Implicit Baseline*

is complicated by the intercept in the GLM at the first level. For example, in the Cue Model

parameterization, the intercept takes on the average for the unmodeled phase of the task which

includes the fixation cross (between cue and probe phase) and the probe response phase. In this

instance, isolating the difference of [Cue *Large Gain*] - [Fixation + Probe phase] to a specific

cognitive function becomes especially challenging (Poldrack & Yarkoni, 2016; Price & Friston,

1997). It is well recognized that different definitions of "baseline", whether rest, passive or task-

related, in task-fMRI will result in different activation patterns (Newman et al., 2001). The use of

"neutral" or "fixation" is a cause for caution as it impacts interpretability in various fMRI task

designs (Balodis & Potenza, 2015; Filkowski & Haas, 2017). Here, we illustrated how contrasts

with the unmodeled phases of a task (*Implicit Baseline*) may improve reliability estimates but

may be heavily biased by the activity patterns throughout the task and diminish the validity of

the measure. It is reasonable to suspect that subtle modeling deviations between similar and

different task designs would further complicate comparisons between studies when using an

Implicit Baseline condition.

In the context of test-retest reliability of estimated BOLD activity, it is important to

consider alternative methods to improve reliability, estimation procedures and considerations of

what a 'reliable' BOLD estimate implies. In general, the evidence here illustrates that the test-

retest reliability for the modified version of the MID task is consistently low using the intraclass

correlation (ICC[3,1]), even at its maximum. The analytic decisions at the GLM modeling phase

demonstrated improvements in reliability from between-run to between-session. Higher between-

session reliability may be related to decreasing activity from early to later runs (Demidenko,

Mumford, et al., 2024) or based on the sessions being an average of two runs/increased trials

(Han et al., 2022; Ooi et al., 2024). In the current analyses, we focused on univariate maps and

the parametric, voxelwise ICC estimation procedures (ICC[3,1]). Parametric and non-parametric

multivariate methods are reported to improve reliability estimates over univariate estimates using

multi-dimensional BOLD data (Gell et al., 2023; Noble et al., 2021). For example, I2C2 is a

parametric method that pools variance across images to estimate a global estimate of reliability

using a comparable ratio as ICC (Shou et al., 2013) and the discriminability statistic is a non-

parametric statistic that is a global index of reliability testing whether the between-subject

distance between voxels is greater than the within-subject voxels (Bridgeford et al., 2021). Each

of these metrics uniquely summarizes the within- and between-subject variability of the

estimated BOLD data and so a consensus and definition of reliability in task-fMRI remains a

challenge (Bennett & Miller, 2010). In our analyses we used the ICC as it estimated the

reliability for each voxel in an easy-to-interpret coefficient that is useful in common brain-

behavior studies. Cut-offs from the self-report literature (Cicchetti & Sparrow, 1981) are often

leveraged in fMRI research (Elliott et al., 2020; Noble et al., 2019); however, these cut-offs

should depend on the optimal level of precision necessary for the question and reasonable for the

methods (Bennett & Miller, 2010; Lance et al., 2006). Some recommendations have been made

to use bias-corrections in developmental samples to adjust for suboptimal levels of reliability

(Herting et al., 2017), but these corrections should be used cautiously as they do not account for

the underlying problems of the measure or the complexities in the data that prevent accurate

measurement of the latent process (Nunnally, 1978).

Study Considerations

The analytic decisions in the current analyses focused primarily on a subset of decisions

at the First Level GLM model and its impact on estimates and supra/sub-threshold masks. As a

result, other decisions were not considered that may arise at the preprocessing (Li et al., 2021),

assumed hemodynamic response function (Kao et al., 2013; Lindquist et al., 2009), cardiac and

respiratory correction (Allen et al., 2022; Birn et al., 2006), and the effects of different methods

of signal distortion correction (Montez et al., 2023). Furthermore, we focused on voxelwise

estimates of reliability which are typically noisier than *a priori* anatomical regions. It is unclear

how much interpretation would change if ICC estimates were compared across variable

parcellations. Nevertheless, we shared all aggregate maps for the three samples and the

preprocessed data for the MLS/AHRB samples to facilitate reanalysis.

The results provide a comprehensive overview of individual and group reliability

estimates for the modified version of the MID task, but it is challenging to infer how reflective

these results are of alternate MID designs and different reward tasks. Based on prior reports of

low test-retest reliabilities in task fMR, if a sufficient sample size is used, we suspect that results

may be comparable to other MID and reward task designs. Future research should consider how

reliability estimates change as a function of modeling decisions in different task paradigms.

Conclusion

With the increasing interest in test-retest reliability in task fMRI and methods for

improving reliability estimates of BOLD, the current study evaluated which decisions at the

GLM model improved group and individual reliability estimates of reliability. In general, the

findings illustrate that the MID task group activation maps are more reliable than individual

maps across testing occasions and independent samples. Across group and individual models,

between-session estimates are consistently higher than between-run estimates of reliability.

Furthermore, estimates of reliability were more variable at the median fMRI sample size and

stabilized with N. While individual estimates of reliability are low (ICC[3,1]), contrasts and

model parameterization meaningfully improved test-retest reliability. However, the improvement

in reliability came at the cost of interpretability and may be region specific in the current version

of the MID task. This underscores the importance of evaluating reliability in larger samples sizes

and ensuring improved estimates reflect the neural processes of interest. While Model

Parameterization and Contrast selection had the largest impact on voxelwise ICCs, further work

is needed to expand on these findings by evaluating alternative brain regions and analytic

decisions that may result in improved test-retest reliability that may be meaningful in individual

differences research.

Data & Code Availability Statement

Adolescent Brain Cognitive Development (ABCD) data: The ABCD BIDS data, MRIQC v23.1.0 and fMRIPrep v23.1.4 derivatives can be accessed through the ABCD-BIDS Community Collection (ABCC) with an established Data Use Agreement (see https://abcdstudy.org/). The data used in these analyses will be available at a future release onto the National Institute of Mental Health Data Archive. The complete set of group-level and ICC maps are publicly available on Neurovault for ABCD (6180 images; https://identifiers.org/neurovault.collection:17171).

Michigan Longitudinal Study (MLS) and *Adolescent Health Risk Behavior* (AHRB) data: The BIDS inputs, fMRIPrep v23.1.4 and MRIQC v23.1.0 derivates are available on OpenNeuro.org (MLS: https://doi.org/10.18112/openneuro.ds005027.v1.0.1 AHRB: https://doi.org/10.18112/openneuro.ds005012.v1.0.1). The complete set of group-level and ICC maps are publicly available on Neurovault for MLS (2400 images; https://identifiers.org/neurovault.collection:16606) and AHRB (2400 images; https://identifiers.org/neurovault.collection:16605)

R and Python code: The *.html* and *.rmd* file containing the code to be run on extracted estimates from reliability maps are available on Github with the associated output files containing the estimates across the models and samples. Likewise, all of the code for first level, fixed effect, group and ICC models are available online at https://github.com/demidenm/Multiverse_Reliability.

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Author's Contribution

MID obtained data sharing agreements. MID conceptualized the study with critical input from RAP. MID defined the methodology with critical input from RAP and JAM. MID curated the

1063   analytic code and performed the formal analysis and interpretation with input from RAP and

1064   JAM. MID wrote the original draft and curated the visualizations. RAP and JAM reviewed,

1065   edited, and provided critical feedback on the draft and all revisions.

1066   **Conflicts of Interest**

1067   The authors declare that they have no conflicts of interest.

1068

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