

Key Variables Extracted from the Best Practices in Data Analysis and Sharing in Neuroimaging using MRI (COBIDAS) Document

Experimental Design		
<input type="checkbox"/> Age (mean, SD, range)	<input type="checkbox"/> Exclusion criteria	<input type="checkbox"/> Race/Ethnicity (if applicable)
<input type="checkbox"/> Sex	<input type="checkbox"/> Clinical criteria	<input type="checkbox"/> Matching strategy (if applicable)
<input type="checkbox"/> Education (Parents) / SES	<input type="checkbox"/> Clinical Instruments	<input type="checkbox"/> Subject scanning order
<input type="checkbox"/> Handedness	<input type="checkbox"/> Cognitive measures	<input type="checkbox"/> Pre-registration of study
Ethical Considerations		
<input type="checkbox"/> Ethical approval	<input type="checkbox"/> Informed consent	<input type="checkbox"/> Informed assent (if applicable)
Acquisition Reporting		
<input type="checkbox"/> Special accommodations	<input type="checkbox"/> Pulse sequence & parameters (<i>full set listed in COBIDAS document</i>)	<input type="checkbox"/> Fat suppression
<input type="checkbox"/> MRI model and field strength	<input type="checkbox"/> Parallel imaging / multiband	<input type="checkbox"/> Brain coverage
<input type="checkbox"/> Type of head coil		<input type="checkbox"/> Internal scanner pre-processing
Functional MRI		
<input type="checkbox"/> Task or resting-state fMRI	<input type="checkbox"/> “Dummy” scans acquired	<input type="checkbox"/> Motion monitoring / correction
<input type="checkbox"/> Length and number of runs	<input type="checkbox"/> Slice order and timing	<input type="checkbox"/> rs-fMRI: Eyes open or closed?
Task functional MRI		
<input type="checkbox"/> Design (Event related, block, etc.)	<input type="checkbox"/> Optimization of design?	<input type="checkbox"/> Ordering of events/blocks
<input type="checkbox"/> Length and type of baseline	<input type="checkbox"/> Presentation software used?	<input type="checkbox"/> Presentation hardware
<input type="checkbox"/> Condition and stimuli used	<input type="checkbox"/> Specifics of stimuli presented	<input type="checkbox"/> Responses collected
<input type="checkbox"/> Timing structure of events	<input type="checkbox"/> Unique number of stimuli	<input type="checkbox"/> Run order
<input type="checkbox"/> Random/Jittered pattern used	<input type="checkbox"/> Repetition of the stimuli used	
Behavioral Performance		
<input type="checkbox"/> Number of variables recorded	<input type="checkbox"/> Types of variables recorded	<input type="checkbox"/> Performance summary statistics
Preprocessing		
<input type="checkbox"/> Software used	<input type="checkbox"/> Spatial filtering	<input type="checkbox"/> Intensity correction
<input type="checkbox"/> Segmentation method	<input type="checkbox"/> Registration algorithm	<input type="checkbox"/> Physiologic noise removal
<input type="checkbox"/> Gradient distortion correction	<input type="checkbox"/> Native or Standard space	
DTI Preprocessing		
<input type="checkbox"/> Eddy current correction	<input type="checkbox"/> Voxel-wise or tract-based?	<input type="checkbox"/> Reference or atlas used
<input type="checkbox"/> Diffusion estimation models	<input type="checkbox"/> Threshold to define voxels of interest	<input type="checkbox"/> Algorithm used for tractography
fMRI Preprocessing		
<input type="checkbox"/> T ₁ Stabilization volumes	<input type="checkbox"/> Excluded due to excessive motion	<input type="checkbox"/> Volume censoring
<input type="checkbox"/> Slice timing correction	<input type="checkbox"/> Function/Structure coregistration	<input type="checkbox"/> Summary measures used for analyses
<input type="checkbox"/> Motion correction approaches	<input type="checkbox"/> Temporal filtering	<input type="checkbox"/> Masks or seed regions applied?
Statistical Modeling & Inference		
<input type="checkbox"/> Dependent variable	<input type="checkbox"/> Model settings	<input type="checkbox"/> Inference: Multiple testing correction
<input type="checkbox"/> Independent variables	<input type="checkbox"/> Inference: Contrast/Effect	Machine Learning
<input type="checkbox"/> Model type	<input type="checkbox"/> Inference: Search region	<input type="checkbox"/> Training procedure
<input type="checkbox"/> Feature extraction	<input type="checkbox"/> Inference: Statistic type	<input type="checkbox"/> Evaluation metrics
<input type="checkbox"/> Dimension reduction	<input type="checkbox"/> Inference: p-Value computation	<input type="checkbox"/> Significance and fit
Results Reporting		
<input type="checkbox"/> List of tested and omitted effects	<input type="checkbox"/> Unthresholded maps	<input type="checkbox"/> ICA analyses and # of components
<input type="checkbox"/> Extracted data	<input type="checkbox"/> Extracted data	<input type="checkbox"/> Graph Theory: Null hypothesis tested
<input type="checkbox"/> Table of Coordinates	<input type="checkbox"/> Spatial features	<input type="checkbox"/> Optimized evaluation metrics
Data Sharing & Reproducibility		
<input type="checkbox"/> Materials shared	<input type="checkbox"/> Ethics compliance	<input type="checkbox"/> Tools used (names, versions, URLs)
<input type="checkbox"/> URL and/or Access information	<input type="checkbox"/> Data format (i.e., DICOM, BIDS)	<input type="checkbox"/> Computational infrastructure

Note: For a complete description of each item, reporting of power analyses, and for alternative neuroimaging methods (i.e. arterial spin labelling, dynamic susceptibility contrast MRI, or combined MRI and EEG, etc.) please refer to the original COBIDAS document at the follow site: (<https://www.biorxiv.org/content/10.1101/054262v2>). This Table is open and can be copied, pasted, adapted, or re-used in any format.