The Artifact Subspace Reconstruction Method

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Background

- Originally created at SCCN in 2013
- Developed as part of Cognition and Neuroergonimics CTA program to support ambulatory neuroscience/-ergonomics research & applications



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Artifact Subspace Reconstruction

 Method for removing high-amplitude artifacts from EEG



 Reconstructs underlying EEG activity, recovers missing energy, leaves no pathologies in signal

visualization generated by vis_artifacts

Artifact Subspace Reconstruction

Designed to handle non-stationary / non-repeating artifacts



Prerequisites

- ASR requires data to be already high-pass filtered (signal mean should be *zero*)
- Data is assumed to be *full rank* (i.e., cannot use common average reference before ASR)
- Best to remove misc channels beforehand (e.g., trigger channel, packet counter, esp. flatline channels)
- Keeping EXG channels *should* work, but better to test with and without

Sliding-Window Processing

Applied independently sample-by-sample, using a sliding window for statistics



 Sliding window is typically short (to handle nonstationary / non-stereotypical artifacts), but longer windows can be used

Finding Artifact Components

- Want to find only *high-amplitude* signal components (potential artifact components) in the short statistics window
- Can be done using Principal Component Analysis
 (PCA) on signal in statistics window W [nCh x nWnd]:



Optional Spectral Weighting

• Statistics can be *spectrally weighted* by applying an IIR filter to the data in the statistics window (8-ord. Yule-Walker), used by default



- This filter is rarely touched, but can be used to make ASR "blind" to certain desired signals (e.g., SSVEP)
- Can also be used to cause ASR to be more sensitive to certain kinds of artifacts (e.g., muscle, EM)

Artifact Detection Threshold

- Applied to the variance for each detected component
- Needs to separate artifacts from non-artifacts (i.e., background EEG, ERPs)
 →Threshold is data-dependent
- Different brain areas will have different background activity levels (e.g., occipital alpha)
 →Threshold is direction-dependent
- Need statistics of "clean" EEG data to calibrate threshold

Finding Clean Calibration Data

- For real-time use, ASR must be calibrated on some initial reasonably clean resting-state data (can be eyes-closed), e.g. 1 minute
- On recorded data, ASR will optionally use clean_windows to find the cleanest data segments in a recording (e.g., 10% / few minutes), on by default
- Fine if these data *still* have artifacts in them
 → usually don't have to worry about it

Clean-Data Statistics

- We extract several statistics from clean data D:
 - Robust covariance matrix
 - Robust mixing matrix
 - Direction-dependent threshold matrix



Robust Estimators

• We use a robust estimator to estimate the covariance matrix (geometric median)

$$\mathcal{G}(\mathbf{X}) = \arg\min_{\mathbf{y}} \sum_{i=1}^{m} \|\mathbf{x}_{i} - \mathbf{y}\|_{2} \qquad y_{i+1} = \left(\sum_{j=1}^{m} \frac{x_{j}}{\|x_{j} - y_{i}\|}\right) / \left(\sum_{j=1}^{m} \frac{1}{\|x_{j} - y_{i}\|}\right)$$

Geometric Median

Iterative Formula (iteratively reweighted least squares)



x_i is a (vectorized)
covariance matrix of a
short window of *D*

Detection Threshold

- Artifacts components detected using a threshold function $th(v_k)$
- Can be thought of as a robust Mahalanobis distance estimated from clean data *D*

$$\operatorname{th}(v_k) \coloneqq \|\operatorname{diag}(z)\overline{P}^T v_k\|$$

• \overline{P} are the principal components derived from the robust covariance matrix



Detection Threshold

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$$\operatorname{th}(v_k) \coloneqq \|\operatorname{diag}(z)\overline{P}^T v_k\|$$

z_i are the thresholds along each clean data component's direction



Rejection Rule

 Reject if short-window PCA component's standard deviation exceeds threshold along its direction

$$\sqrt{\lambda_k} = \sigma_k > \operatorname{th}(v_k)$$

Now what do we do with rejected components?

Reconstructing Rejected Components

- Any two EEG channels are *almost always* correlated
- So are any two linear combinations of channels
- So are the components in our short statistics window
- Background EEG content in artifact PCs can be estimated from non-artifact PCs
- Analogy: estimating one channel from other channels
- Here not using channels but artifact / non-artifact subspaces on a short window



Reconstructing Rejected Components

- But isn't the artifact subspace uncorrelated from the non-artifact subspace?
- Only in the short-window decomposition
- Not in the average background EEG statistics (correlation structure)
- This correlation structure is described by our robust mixing matrix *M* (from clean data)

Performing the Reconstruction



Performing the Reconstruction



Overlapped-Window Composition

- Re-calculate R for every k'th sample (step size)
 (in real time can reconstruct at end of each chunk)
- Blend between successive *R* estimates for intermediate samples, e.g., using raised-cosine weighting



- Avoids sharp transients (jump discontinuities)
- Also maintains full-rank output data (i.e., no pathological subspaces)

Future Directions

- Calibrate in real time on expanding window or sliding window (difficult due to truncated generalized Gaussian est.)
- Use Riemannian estimator for calibration covariance
- Find good rules of thumb for threshold

Thanks!

Questions?

Figure References

- Brain model: Zeynep Akalin Acar
- Geometric median figure: Wikipedia
- Demo video: Cognionics