Supporting Information 4: Eliminating accidental deviations to minimize generalization error and maximize replicability: applications in connectomics and genomics

Eric W. Bridgeford¹, Shangsi Wang¹, Zeyi Wang¹, Ting Xu³, Cameron Craddock³, Jayanta Dey¹, Gregory Kiar¹, William Gray-Roncal¹, Carlo Colantuoni¹, Christopher Douville¹, Stephanie Noble⁴, Carey E. Priebe¹, Brian Caffo¹, Michael Milham³, Xi-Nian Zuo^{2,5}, Consortium for Reliability and Reproducibility, Joshua T. Vogelstein^{1,6*}

S4 Simulations The following simulations were constructed, where σ_{min} , σ_{max} are the variance ranges, and settings were run at 15 intervals in $[\sigma_{min}, \sigma_{max}]$ for 500 repetitions per setting. For a simulation setting with variance σ , the variance is reported as the normalized variance, $\bar{\sigma} = \frac{\sigma - \sigma_{min}}{\sigma_{max} - \sigma_{min}}$. Dimensionality is 2, the number of items is K, and the total number of measurements across all items is 128. Typically, i indicates the individual identifier, and j the measurement index. Notationally, in the below descriptions, we adopt the convention that z_i^j obeys the true distribution for a single observation j of item *i*, and x_i^j incorporates the controlled error term ϵ_i^j , which is the term which is varied the simulation. Further, each item features $\frac{n}{K}$ measurements.

Goodness of Fit Testing and Bayes Error

- 1. No Signal: K = 2 items, where the true distributions for class 1 and class 2 are the same.
 - $\boldsymbol{z}_i^j \stackrel{iid}{\sim} \mathcal{N}(\boldsymbol{0}, \boldsymbol{I}), i = 1, \dots, 2, t = 1, \dots, 64$. Note: $\boldsymbol{0} \in \mathbb{R}^2$ is $\boldsymbol{0}$, and likewise for \boldsymbol{I}

•
$$\boldsymbol{\epsilon}_{i}^{j} \stackrel{iid}{\sim} \mathcal{N}(\boldsymbol{0}, \sigma^{2}\boldsymbol{I}), \sigma \in [0, 20]$$

•
$$\boldsymbol{x}_{i}^{j} = \boldsymbol{z}_{i}^{j} + \boldsymbol{\epsilon}_{i}^{j} \stackrel{iid}{\sim} \mathcal{N}(\boldsymbol{0}, (1 + \sigma^{2})\boldsymbol{I})$$

- 2. Cross: K = 2 items, where the true distributions for class 1 and class 2 are orthogonal.
 - $\Sigma_1 = \begin{bmatrix} 2 & 0 \\ 0 & 0.1 \end{bmatrix}$, $\Sigma_2 = \begin{bmatrix} 0.1 & 0 \\ 0 & 2 \end{bmatrix}$
 - $\boldsymbol{z}_{i}^{j} \stackrel{iid}{\sim} \mathcal{N}(\boldsymbol{0}, \boldsymbol{\Sigma}_{i}), i = 1, 2$

•
$$\boldsymbol{\epsilon}_{i}^{j} \stackrel{iid}{\sim} \mathcal{N}(\boldsymbol{0}, \sigma^{2}\boldsymbol{I}), \sigma \in [0, 20]$$

•
$$\boldsymbol{x}_{i}^{j} = \boldsymbol{z}_{i}^{j} + \boldsymbol{\epsilon}_{i}^{j}$$

3. Gaussian: K = 16 items, where the true distributions are each gaussian.

- $\boldsymbol{\mu}_i \stackrel{iid}{\sim} \pi_1 \mathcal{N}(\mathbf{0}, 4\mathbf{I}), i = 1, \dots, 16$ $\boldsymbol{\Sigma} = \begin{bmatrix} 1 & 0.1 \\ 0.1 & 1 \end{bmatrix}$ • $\boldsymbol{z}_{i}^{j} \stackrel{iid}{\sim} \mathcal{N}(\boldsymbol{\mu}_{i}, \boldsymbol{\Sigma})$ • $\boldsymbol{\epsilon}_{i}^{j} \stackrel{iid}{\sim} \mathcal{N}(\boldsymbol{0}, \sigma^{2}\boldsymbol{I}), \sigma \in [0, 20]$
- $x_i^j = z_i^j + \epsilon_i^j$ 4. Ball/Circle: K = 2 items, where 1 item is uniformly distributed on the unit ball with gaussian error, and the second item is uniformly distributed on the unit sphere with gaussian error.
 - $z_1^t \stackrel{iid}{\sim} \mathbb{B}(r=1) + \mathcal{N}(\mathbf{0}, 0.1I)$ samples uniformly on unit ball of radius 2 with Gaussian
 - $z_2^{t} \stackrel{iid}{\sim} \mathbb{S}(r = 1.5) + \mathcal{N}(\mathbf{0}, 0.1 \mathbf{I})$ samples uniformly on unit sphere of radius 2 with Gaussian error

¹ Johns Hopkins University, Baltimore, Maryland, USA, ² Shanghai Jiaotong University, Shanghai, China ³ Child Mind Institute, New York, New York, USA ⁴ Yale University, New Haven, Connecticut, USA ⁵ Beijing Normal University, Beijing, China, Nanning Normal University, Nanning, China, University of Chinese Academy of Sciences, Beijing, China, ⁶ Progressive Learning, Baltimore, Maryland, USA. * jovo@jhu.edu.

Bayes error was estimated by simulating n = 10,000 points according to the above simulation settings, and approximating the Bayes error through numerical integration. The classification labels for K = 2 simulations were consistent with the individual labels, and for the K = 16, the first class consists of the 8 distributions whose means were leftmost, and the rest of the distributions were the other class.

Comparison Testing Items are sampled with the same true distributions z_i^j as before, with the following augmentation:

$$oldsymbol{x}_{i,k}^{j} = egin{cases} oldsymbol{z}_{i}^{j} & k = 1 \ oldsymbol{z}_{i}^{j} + oldsymbol{\epsilon}_{i}^{j} & k = 2 \end{cases}$$

That is, the observed data $\boldsymbol{x}_{i,k}^{j}$ for item *i*, observation *j*, and sample $k \in [2]$ is such that the first sample is distributed according to the true item distribution, and the second sample is distributed according to the true item distribution, where $\boldsymbol{\epsilon}_{i}^{j} \stackrel{iid}{\sim} \mathcal{N}(\mathbf{0}, \sigma^{2} \boldsymbol{I})$:

- 1. No Signal: K = 2 $\sigma \in [0, 10]$ 2. Cross: K = 2
- $\sigma \in [0, 1]$ 3. Gaussian: K = 16 $\sigma \in [0, 1]$

4. Ball/Circle:
$$K = 2$$

 $\sigma \in [0, 1]$

5. XOR: K = 2 $\boldsymbol{x}_{i,k}^{j} = \begin{cases} \boldsymbol{z}_{i}^{j} + \boldsymbol{\tau}_{i}^{j} & k = 1 \\ \boldsymbol{z}_{i}^{j} + \boldsymbol{\tau}_{i}^{j} + \boldsymbol{\epsilon}_{i}^{j} & k = 1 \end{cases}$ where $\tau_{i}^{j} \stackrel{iid}{\sim} \mathcal{N}(\boldsymbol{0}, 0.1\boldsymbol{I})$ $\sigma \in [0, 0.2]$

By construction, one would anticipate Discr of the first sample to exceed that of the second sample, as the second sample has additional error. Therefore, the natural hypothesis is:

$$H_0: D^{(1)} = D^{(2)}, \qquad H_A: D^{(1)} > D^{(2)}$$