Fast Threshold Tests for Detecting Discrimination Supplementary Information

1 Proofs

Proof of proposition 3.1 [Monotonicity] Given a discriminant distribution $disc(\phi, \mu_0, \sigma_0, \mu_1, \sigma_1)$, the mapping g from signal space to probability space is monotonic if and only if $\sigma_0 = \sigma_1$.

Proof. The mapping from signal space to probability space can be found with Bayes' rule:

$$g(x) = \Pr(Y = 1 \mid X = x) = \frac{\Pr(Y = 1)N(x; \mu_1, \sigma_1)}{\Pr(Y = 1)N(x; \mu_1, \sigma_1) + \Pr(Y = 0)N(x; \mu_0, \sigma_0)}$$
$$= \log it^{-1}(Ax^2 + Bx + C)$$

where

$$A = \frac{\sigma_1^2 - \sigma_0^2}{2\sigma_0^2 \sigma_1^2} \qquad B = \frac{\mu_1 \sigma_0^2 - \mu_0 \sigma_1^2}{\sigma_0^2 \sigma_1^2}$$
$$C = \frac{\sigma_1^2 \mu_0^2 - \sigma_0^2 \mu_1^2}{2\sigma_0^2 \sigma_1^2} - \log\left(\frac{1 - \phi}{\phi} \frac{\sigma_1}{\sigma_0}\right).$$

This is the composition of a quadratic in x and the inverse logit function, which will be monotonic if and only if the quadratic is monotonic, requiring $\sigma_0^2 - \sigma_1^2 = 0$.

Proof of proposition 3.2 [2-parameter representation]: Suppose disc $(\phi, \mu_0, \sigma, \mu_1, \sigma)$ and disc $(\phi', \mu'_0, \sigma', \mu'_1, \sigma')$ are two homoskedastic discriminant distributions. Let

$$\delta = \frac{\mu_1 - \mu_0}{\sigma}$$

and define δ' analogously. Then the two distributions are identical if $\phi = \phi'$ and $\delta = \delta'$. As a result, homoskedastic discriminant distributions can be parameterized by ϕ and δ alone.

Proof. We establish this result by explicitly deriving the density of g(X), where g is the usual mapping from signal to probability space. Doing so first requires computing the inverse transformation from probability space to signal space:

$$g^{-1}(p) = \frac{\mu_1^2 - \mu_0^2 - 2\sigma^2 \log\left(\frac{\phi}{1-\phi}\frac{1-p}{p}\right)}{2(\mu_1 - \mu_0)}$$

Now,

$$\frac{d}{dp}(g^{-1}(p)) = \frac{\sigma^2}{p(1-p)(\mu_1 - \mu_0)}.$$

The density of X is

$$f_X(x) = \phi N(x; \mu_1, \sigma_1) + (1 - \phi) N(x; \mu_0, \sigma_0)$$

We can accordingly compute the density of g(X) by the change of variables formula:

$$f_P(p) = f_X(g^{-1}(p)) \left| \frac{d}{dp}(g^{-1}(p)) \right|$$
$$= \frac{\sigma}{\sqrt{2\pi}p(1-p)(\mu_1 - \mu_0)} \left[\phi \exp\left(-\frac{\left(\alpha + \frac{(\mu_1 - \mu_0)^2}{\sigma^2}\right)^2}{8\frac{(\mu_1 - \mu_0)^2}{\sigma^2}}\right) + (1-\phi) \exp\left(-\frac{\left(\alpha - \frac{(\mu_1 - \mu_0)^2}{\sigma^2}\right)^2}{8\frac{(\mu_1 - \mu_0)^2}{\sigma^2}}\right) \right],$$

where

$$\alpha = 2\log\left(\frac{\phi}{1-\phi}\frac{1-p}{p}\right).$$

Without loss of generality we can set $\delta = (\mu_1 - \mu_0)/\sigma$, resulting in a 2-parameter family of densities:

$$f_P(p) = \frac{1}{\sqrt{2\pi}p(1-p)\delta} \left[\phi \exp\left(-\frac{\left(\alpha+\delta^2\right)^2}{8\delta^2}\right) + (1-\phi)\exp\left(-\frac{\left(\alpha-\delta^2\right)^2}{8\delta^2}\right) \right].$$
 (1)

Proof that discriminant distributions are a subset of logit-normal mixtures: Discriminant distributions approximate logit-normal distributions particularly well because they in fact form a subset of logit-normal mixture distributions. To see this, consider the following rearrangement of the first component in the mixture in Eq. (1) (ie, the component with weight ϕ):

$$\frac{\exp\left(-\frac{\left(2\log\left(\frac{\phi}{1-\phi}\frac{1-p}{p}\right)+\delta^2\right)^2}{8\delta^2}\right)}{\sqrt{2\pi}p(1-p)\delta} = \frac{\exp\left(-\frac{\left(\log i(p)-\left(\log i(\phi)+\frac{\delta^2}{2}\right)\right)^2}{2\delta^2}\right)}{\sqrt{2\pi}p(1-p)\delta}.$$

This is the density of a logit-normal with parameters $\mu = \text{logit}(\phi) + \frac{\delta^2}{2}$ and $\sigma = \delta$. Thus, we can express the discriminant distribution as a specific 2-parameter mixture of logit-normals (where $f_l(x; \mu, \sigma)$ is the density function of the logit-normal):

$$f_P(p) = \phi f_l\left(p; \operatorname{logit}(\phi) + \frac{\delta^2}{2}, \delta\right) + (1 - \phi) f_l\left(p; \operatorname{logit}(\phi) - \frac{\delta^2}{2}, \delta\right).$$

In this form, we can see that for small δ or ϕ close to 0 or 1, the distribution is almost equivalent to a (single) logit-normal.

2 Supplementary figures

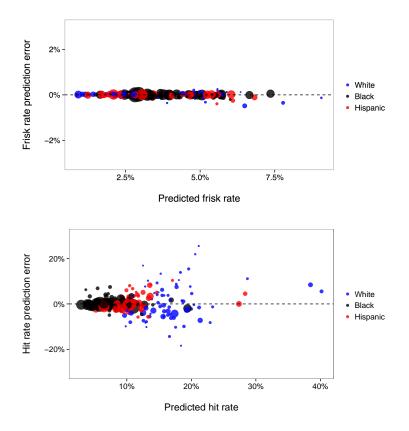


Figure 1: Posterior predictive checks for frisk rate and hit rate in stop-and-frisk data. Each point represents one precinct-race pair. Points are sized by the number of stops for that precinct and race.

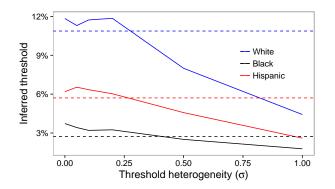


Figure 2: Inferred frisk thresholds when the model is fit to simulated data, where the threshold applied to each stop is perturbed by logit-normal noise. The dashed lines indicate the unperturbed thresholds. Racial disparities in thresholds persist even as large amounts of noise are introduced.

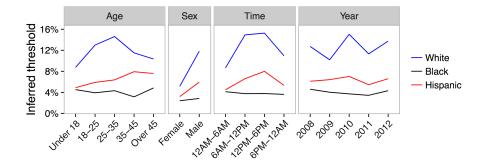


Figure 3: Frisk thresholds inferred on disaggregated subsets of the primary dataset. While thresholds vary across subsets, thresholds for whites are consistently higher than thresholds for blacks and Hispanics.

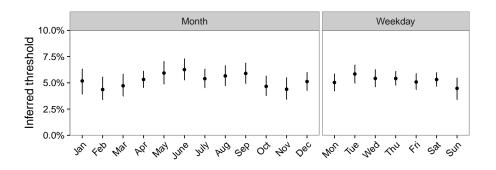


Figure 4: Placebo test: frisk thresholds inferred using month or weekday rather than race show only slight variation, as expected. Vertical lines denote 95% credible intervals.

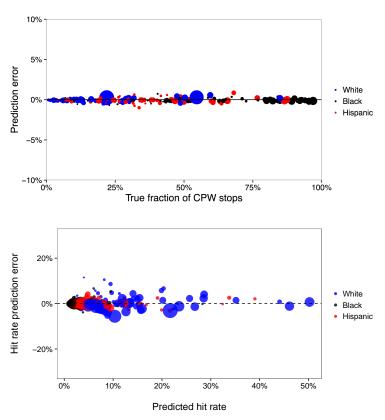


Figure 5: Posterior predictive checks for stop model. Each point represents one precinct-race pair. Points are sized by the population of each race in each precinct.

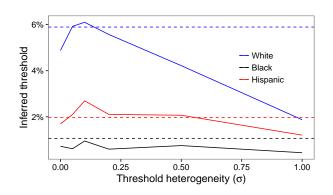


Figure 6: Inferred stop thresholds when the model is fit to simulated data, where the threshold applied to each stop is perturbed by logit-normal noise. The racial disparities in thresholds persist even as large amounts of noise are introduced.

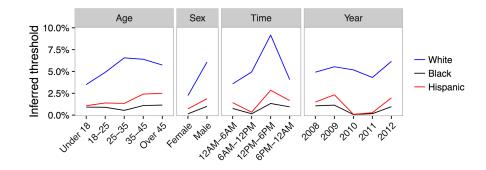


Figure 7: Stop thresholds inferred on disaggregated subsets of the primary dataset. While thresholds vary across subsets, thresholds for whites are consistently higher than thresholds for blacks and Hispanics.

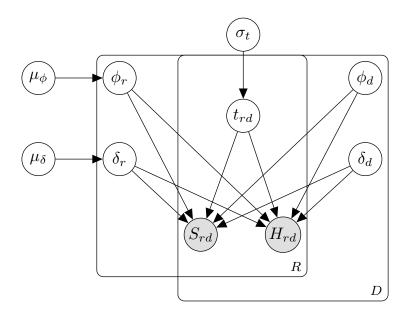


Figure 8: Graphical representation of the generative model for stop decisions. Observed outcomes are shaded, and unshaded nodes are latent variables inferred from data.