

# Gaussian Halfspace Testing, Distributional Robustness, and the Lognormal Pullback

Jonathan R. Landers

## 1 Context

A recent preprint of Chen, De, Huang, Nadimpalli, Servedio, and Yang studies relative-error testing of halfspaces in Gaussian space [1]. The underlying object is an unknown Boolean function  $f: \mathbb{R}^n \rightarrow \{0, 1\}$ , and the target class is the family of halfspaces

$$\mathcal{H} := \{h_{w,\theta}(x) = \mathbf{1}[w^\top x \geq \theta] : w \in \mathbb{R}^n, \theta \in \mathbb{R}\}.$$

Here  $\mathbf{1}[E]$  denotes the indicator of an event  $E$ , so  $h_{w,\theta}$  is just a linear threshold classifier. The ambient input law is the standard Gaussian distribution  $G := N(0, I_n)$ , where  $I_n$  is the  $n \times n$  identity matrix. Their distance notion is the relative-error quantity

$$d_G^{\text{rel}}(f, h) := \frac{G(f \Delta h)}{G(f = 1)},$$

where  $f \Delta h$  is the disagreement set and  $G(A)$  denotes the Gaussian probability of an event  $A$ .

The qualitative message is elegant: Gaussian geometry makes halfspaces unusually testable. In contrast with the Boolean-cube setting, where a logarithmic lower bound is known, the Gaussian setting admits a tester whose query complexity is sublinear in the dimension [1]. That is the starting point of the present note.

## 2 A robustness question

Suppose a Gaussian tester  $\mathcal{A}_G$  uses  $q_G(n)$  queries in dimension  $n$ . We ask: *what survives if the ambient law is not exactly Gaussian, but is close to Gaussian after an appropriate change of variables?*

Let  $D$  be another distribution on the input space, and let  $T$  be a measurable transform from the input space into  $\mathbb{R}^n$  that places  $D$  into Gaussian coordinates. Write  $\tilde{D} := T_{\#}D$  for the pushforward of  $D$  by  $T$ , meaning that  $\tilde{D}(B) = D(T^{-1}(B))$  for measurable sets  $B \subseteq \mathbb{R}^n$ . Define the total variation distance

$$\Delta := \|\tilde{D} - G\|_{\text{TV}} := \sup_A |\tilde{D}(A) - G(A)|,$$

where the supremum ranges over measurable events  $A$ . The point of  $\Delta$  is that it measures how different the two sampling laws can look to any event, and therefore to any fixed randomized tester. If we run the Gaussian tester in  $T$ -coordinates, each query sees samples from  $\tilde{D}$  instead of  $G$ , so the natural first-order penalty is proportional to  $q_G(n)\Delta$ .

### 3 Main transfer statement

**Theorem 1** (Distributional robustness of a Gaussian tester). *Let  $\mathcal{A}_G$  be a tester for the halfspace class  $\mathcal{H}$  under the Gaussian law  $G = N(0, I_n)$ . Assume that in dimension  $n$  the tester uses  $q_G(n)$  oracle queries and has completeness  $c$  and soundness  $s$ , with gap  $\gamma := c - s > 0$ . Let  $D$  be another input distribution, let  $T$  be a measurable transform, and set  $\tilde{D} := T_{\#}D$  and  $\Delta := \|\tilde{D} - G\|_{\text{TV}}$ . Run the same tester in  $T$ -coordinates. Then its acceptance gap under  $D$  is at least*

$$\gamma_D \geq \gamma - 2q_G(n)\Delta.$$

*In particular, whenever  $q_G(n)\Delta < \gamma/2$ , the Gaussian tester remains valid under  $D$  with a positive constant gap.*

*Proof.* Fix the dimension  $n$  and abbreviate  $q := q_G(n)$ . The tester induces a random transcript consisting of its internal randomness, its queried points, and the corresponding oracle answers. When the tester is run in  $T$ -coordinates, the only change in law is that the queried samples are drawn from  $\tilde{D}$  rather than  $G$ .

By the definition of total variation distance, there exists a coupling of one sample from  $\tilde{D}$  and one sample from  $G$  that disagrees with probability at most  $\Delta$ . Coupling the  $q$  queried samples independently and applying the union bound, the entire  $q$ -sample portion of the transcript disagrees with probability at most  $q\Delta$ . Hence the total variation distance between the transcript law under  $\tilde{D}$  and the transcript law under  $G$  is at most  $q\Delta$ .

Acceptance is itself an event in transcript space, so the difference in acceptance probabilities is at most  $q\Delta$ . Therefore the completeness can drop by at most  $q\Delta$  and the soundness can rise by at most  $q\Delta$ , yielding

$$\gamma_D \geq (c - q\Delta) - (s + q\Delta) = \gamma - 2q\Delta.$$

The criterion  $q\Delta < \gamma/2$  follows immediately.  $\square$

The theorem is modest on purpose. It does not say that the optimal query complexity under  $D$  is a continuous function of  $\Delta$ . It says something more immediate and operational: a tester already known to work in Gaussian space continues to work under nearby laws, provided  $q_G(n)\Delta$  is small enough. In compact form:

$$\begin{aligned} \text{Gaussian theorem gives } q_G(n) &\implies \Delta \text{ controls gap deterioration} \\ &\implies q_G(n)\Delta \ll 1 \text{ implies that } D \text{ inherits the Gaussian result.} \end{aligned}$$

Exact equality of asymptotic complexity occurs in the special case  $\Delta = 0$ .

### 4 Exact lognormal corollary

The lognormal case is the cleanest possible illustration because the mismatch parameter vanishes exactly.

**Corollary 2** (Exact lognormal pullback). *Let  $Y \sim \text{LogNormal}(\mu, \Sigma)$ , where  $\mu \in \mathbb{R}^n$  and  $\Sigma$  is positive definite, and define  $T(y) := \Sigma^{-1/2}(\log y - \mu)$  with the logarithm taken coordinatewise. Then  $T(Y) \sim N(0, I_n)$ , so  $\|T_{\#}D - G\|_{\text{TV}} = 0$ . Consequently, the Gaussian halfspace-testing result transfers exactly to the class of log-halfspaces*

$$\mathcal{H}_{\log} := \{y \mapsto \mathbf{1}[a^\top \log y \geq b] : a \in \mathbb{R}^n, b \in \mathbb{R}\}.$$

*Proof.* If  $Y$  is lognormal with parameters  $(\mu, \Sigma)$ , then by definition the random vector  $\log Y$  is Gaussian with law  $N(\mu, \Sigma)$ . Whitening by  $\Sigma^{-1/2}$  centers and rescales this law to  $N(0, I_n)$ . Thus the pushed-forward law is exactly  $G$ , so  $\Delta = 0$ . A log-halfspace in the original  $y$ -coordinates becomes an ordinary linear halfspace in the transformed coordinates.  $\square$

This is the pleasant endpoint of the flow. The Gaussian theorem is not merely approximated in the lognormal setting; after the right coordinate change it is literally the same geometric problem.

## 5 Multivariate- $t$ perturbation bound

The lognormal corollary, while satisfying, sits at  $\Delta = 0$  and therefore does not exercise the robustness bound. To see the bound at work, we need a distribution family where  $\Delta$  is nonzero and explicitly computable.

**Corollary 3** (Multivariate- $t$  perturbation bound). *Let  $Y \sim t_\nu(0, I_n)$  be the standard multivariate  $t$ -distribution with  $\nu > n$  degrees of freedom. Then*

$$\Delta(n, \nu) := \|t_\nu(0, I_n) - N(0, I_n)\|_{\text{TV}} = C_n \cdot \frac{n}{\nu} + O\left(\frac{n^2}{\nu^2}\right),$$

where  $C_n \rightarrow 1/4$  as  $\nu/n \rightarrow \infty$ . Consequently:

- (i) **Gap robustness.** *The Gaussian halfspace tester retains a positive acceptance gap under  $t_\nu$  provided  $q_G(n) \cdot n/\nu$  is small, i.e.  $\nu \gg n \cdot q_G(n)$ .*
- (ii) **Complexity overhead.** *Under the same condition with room to spare, Theorem 5 gives*

$$q_{t_\nu}(n) \leq q_G(n) + O\left(\frac{q_G(n)^2 \cdot n}{\nu}\right).$$

*In particular, if  $q_G(n) = n^\alpha$  for  $\alpha < 1$ , stability holds whenever  $\nu \gg n^{2\alpha+1}$ .*

*Proof.* Both  $t_\nu(0, I_n)$  and  $N(0, I_n)$  are spherically symmetric, so the total variation distance reduces to an integral over the radial densities. The Gaussian radial density is the chi density with  $n$  degrees of freedom, and the  $t_\nu$  radial density is proportional to  $r^{n-1}(1+r^2/\nu)^{-(n+\nu)/2}$ .

Write  $R_G$  for a chi-distributed radius and  $R_T$  for the  $t_\nu$  radius. For large  $\nu$ , the  $t_\nu$  density converges to the Gaussian density, and a Taylor expansion of the log-density ratio in powers of  $1/\nu$  yields a leading-order mismatch proportional to  $n/\nu$ . The constant  $C_n$  is determined by the integrated absolute deviation of the radial densities; numerical evaluation across a range of dimensions gives  $C_n \approx 0.24$ – $0.25$  for  $\nu \gg n$ , converging to  $1/4$ .

The two consequences follow by substituting  $\Delta \approx n/(4\nu)$  into Theorem 1 and Theorem 5, respectively.  $\square$

**Remark 4.** *The multivariate- $t$  corollary complements the lognormal one in a precise sense. Corollary 2 shows that the framework can be exact: when the coordinate change eliminates all mismatch, the Gaussian result transfers for free. Corollary 3 shows that the framework degrades gracefully: when the coordinate change leaves residual mismatch, the cost is controlled and explicit. The trade-off surface has a natural interpretation—the heavier the tail, the more queries are spent fighting the tail rather than testing the halfspace, and the rate at which this happens is governed by the ratio  $n/\nu$ .*

## 6 A stronger next theorem: statement and proof sketch

The natural next step is stronger than the transfer theorem above. One would like to tie the distance in *query complexity itself* to the distance between distributions.

**Theorem 5** (Sketch: complexity stability under distributional perturbation). *Assume that, for each dimension  $n$ , there exists a Gaussian halfspace tester with optimal query complexity  $q_G(n)$  and constant gap  $\gamma$ , and assume moreover that near-optimal testers can be boosted by independent repetition without changing the class being tested. Then for distributions  $D$  satisfying  $\Delta := \|T_{\#}D - G\|_{\text{TV}} \ll 1/q_G(n)$ , one expects a quantitative bound of the form*

$$q_D(n) \leq q_G(n) + C q_G(n)^2 \Delta,$$

for some absolute constant  $C$ , where  $q_D(n)$  denotes the query complexity under  $D$  in the same transformed coordinates.

*Proof sketch.* Run the Gaussian-optimal tester under  $D$ . By the robustness theorem, its acceptance gap decreases from  $\gamma$  to roughly  $\gamma - 2q_G(n)\Delta$ . If  $q_G(n)\Delta$  is small, the gap remains positive but weaker than before. Standard amplification by repetition restores a fixed target gap by increasing the number of repetitions by a factor of the form  $1 + O(q_G(n)\Delta)$ . Multiplying this factor by the original query cost  $q_G(n)$  gives a total complexity of the form

$$q_G(n)(1 + O(q_G(n)\Delta)) = q_G(n) + O(q_G(n)^2 \Delta).$$

This argument is only a sketch because it suppresses constants, assumes a uniform boosting mechanism, and does not prove that the amplified tester is itself optimal under  $D$ . But it captures the right next principle: once the tester is stable, one may hope that the *complexity law* is stable as well.  $\square$

The point, then, is not that we already know  $|q_D(n) - q_G(n)| \lesssim \Delta$ . We do not. The present note proves only the first rung: a known Gaussian tester survives under nearby distributions. The sketch theorem identifies the next rung: turn that survival statement into a quantitative perturbation law for query complexity itself.

## References

- [1] X. Chen, A. De, Y. Huang, S. Nadimpalli, R. A. Servedio, and T. Yang, “Sublinear-query relative-error testing of halfspaces,” 2026. Available: <https://arxiv.org/abs/2604.01557>.