Hard or Just Unreached?
Diagnosing the Sampling Blind Spot in Math-Reasoning Difficulty Estimation

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Abstract

Math and science reasoning benchmarks rely on pass@k, the fraction of sampled chains that reach gold, as the canonical per-example difficulty signal. The same signal drives RL with verifiable rewards, math data curation, synthetic curricula, and verifier training. We show this proxy has a persistent blind spot on its hardest stratum: on the eight free-form math cells we test (GSM8K and MATH across four open-weight models), 10.3–22.9% of the examples that no sampling seed solves in six tries are instead solved at matched compute by a six-chain deterministic regime. These are greedy decoding plus five cheap residual-stream perturbations applied via activation grafting, while greedy alone solves at most 6% on these math cells. Recovery scales with the additional budget, across perturbations whose mechanistic distinctness we verify across all twelve cells (cross-kind fix-set Jaccard ≤ 0.47 in every setup. Activation grafting is used as an intervention on internal representations, not a decoding method; we use it purely as a diagnostic and diversification tool, and our recovered items show that the pass@$k=0$ stratum is structurally identifiable in the residual stream rather than that the unmodified model reaches them under ordinary inference.

1. Introduction

RL pipelines with verifiable rewards, math/code data curation, difficulty-stratified curricula, and pass@k-style hardness annotations all share a single label-required primitive: examples on which no sampled chain reaches gold are filtered out, downweighted, or labelled hard (Lambert et al.; Guo et al., 2025; Shao et al., 2024; Yuan et al., 2023; Tosh-

Figure 1. The sampling blind spot (n=1000). Across twelve (model, benchmark) cells, pass@k flags 5–43% of items as "hard" (red, pass@$k=0$). A compute-matched deterministic regime (six greedy chains with residual-stream interventions) reaches 10–29% of that stratum (green), showing that a non-trivial fraction of items pass@k calls "hard" are structurally identifiable in the model's residual stream rather than intrinsically hard.

niwal et al., 2024; Zhou et al., 2026). The signal is the pass@k count itself, the number of sampled chains that reach gold (Cobbe et al., 2021; Chen et al., 2021); where a single answer is needed, k chains are aggregated by majority vote or self-evaluation (Wang et al., 2022; Chen et al., 2023; Aggarwal et al., 2023). The first sample, the second, the kth are all drawn the same way, and temperature sampling is treated as *the* source of decoding diversity. A problem that no sampling chain solves is treated, by some reasonable proxy, as difficult.

We show this last assumption fails on a structurally identifiable slice. Through three empirical observations across four open-weight instruction-tuned models (`Qwen-2.5-3B`, `Llama-3.2-3B`, `Llama-3.1-8B`, `Mistral-Nemo-12B`) and three reasoning benchmarks (GSM8K, MATH, MMLU-Pro), we argue that sampling has a *persistent blind spot* on its hardest stratum at the budgets we test, and that a non-trivial fraction of the slice flagged as hardest by sampling-only estimators is in fact reachable by deterministic decoding at matched compute.

(1) Greedy is competitive at a single sample. Across the (model, benchmark) cells we test, greedy decoding is competitive with or better than the mean single-sample accuracy, with the gap reaching up to 2.8%. The default single-pass baseline for self-consistency is therefore already weaker than a deterministic alternative on the same compute.

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(2) Sampling has a persistent blind spot on its hardest stratum. The pass@$k = 0$ stratum, examples no sample reaches gold in k tries, is the operative signal for any pipeline whose deployment already has labels: RL with verifiable rewards, math data curation, and pass@k-style hardness annotation, etc. At $k = 6$ this stratum is large on free-form math: 5.1–8.3% of GSM8K and 28.7–43.5% of MATH across the four models we test (51–435 examples per setup; Table 3). The per-sample marginal-shrink rate decreases monotonically through our sampling budget (Fig. 3; App. C), so the stratum is not an artifact of an undersized k. This stratum is treated, by some reasonable proxy, as the slice the model cannot solve. The question we focus on: is that slice intrinsically hard, or is it reachable but only off the stochastic axis? See the intuition in Fig. 1.

(3) The deterministic regime reaches what six samples miss. We isolate the slice on which sampling truly fails, examples no sampling seed reaches in six tries, and ask how much of it a deterministic regime at matched compute reaches. Activation grafting is the diagnostic: replacing the last prompt-token hidden state with cheap synthetic vectors, then decoding greedily, holds prompt and weights fixed while perturbing only the deterministic trajectory. A six-chain deterministic regime (greedy plus five grafts) reaches 10.3–22.9% of these examples on the eight free-form math cells (GSM8K, MATH; 10–29% across all twelve cells), while greedy alone reaches only up to 9% (Sec. 5). The recovery scales with deterministic budget across five graft kinds whose mechanistic distinctness, we verify across all twelve cells (Sec. 5, App. F: cross-kind fix-set Jaccard $\leq$ 0.47 in every cell, with the single same-axis pair of random probes reaching 0.76 on `Qwen-2.5-3B` / GSM8K as an illustrative ceiling). This rules out both the "intrinsically hard" and the "one privileged vector" readings. The point is not that deterministic beats sampling on aggregate accuracy; it is that the two regimes have partially disjoint correct sets, and the slice flagged as hardest by sampling-only proxies might not be the actual hardest slice for the model.

The logical chain. Sampling has a persistent blind spot on its hardest stratum, and the deterministic axis reaches into it: recovery on the no-sample-reaches-gold stratum scales with the deterministic budget (one chain $\rightarrow$ six chains: 0–9% $\rightarrow$ 10–29%) across mechanistically distinct grafts, so what looked hard under sampling was, for a non-trivial fraction, just unreached under stochastic decoding.

Implication. On the (model, benchmark) cells we test, sampling-derived difficulty estimates (pass@k filters, RL data-curation pipelines, hardness annotations, difficulty-stratified curricula) mislabel a structurally identifiable fraction of their hardest stratum: 10–29% of the no-sample-reaches-gold slice is reached by a six-chain deterministic regime at matched compute. Hence, difficulty estimates

derived from sampling alone might conflate "hard" with "unreached on the stochastic axis." The deterministic axis, usually treated as a degenerate corner case ("$k = 1$ greedy") or a fallback, is doing more work here than typically credited: when constructed via residual-stream interventions rather than ordinary decoding, it exposes a slice that no sampling seed in $\{42, \ldots, 47\}$ reaches. Grafting is not a decoding method; the point is that this slice is structurally identifiable, not that the unmodified model reaches it under ordinary inference. The fix is one extra deterministic forward pass per example plus a few cheap perturbations. We use activation grafting as a diagnostic tool to diversify the deterministic coverage and to locate this slice.

Why this matters for mathematical reasoning. Math reasoning is the setting where pass@k-style difficulty labels are most consequential, and where our effect is largest in absolute terms. (i) *Benchmark construction:* hardness buckets derived from pass@$k = 0$ are used to advertise benchmark difficulty and to compare frontier models; a structurally identifiable 10.3–22.9% of the "hardest" GSM8K / MATH bucket is in fact reached by an edited forward pass of the same model under a cheap deterministic diversification, so the bucket is partly an artifact of the decoding regime, not of the items. (ii) *Difficulty calibration:* curricula and per-example weights derived from sample-only pass-rates downweight or drop items that are reachable off the stochastic axis under residual-stream interventions; this slice is non-trivial precisely on the hardest items, where calibration matters most. (iii) *Synthetic curriculum and data curation:* rejection-sampling pipelines (e.g., Yuan et al. (2023); Toshniwal et al. (2024)) filter math problems by whether any sampled chain solves them; a fraction of what those pipelines discard as "unsolvable by the generator" is actually solvable by it at matched compute. (iv) *Verifier and reward-model training:* preference and verifier datasets built from sampled-chain correctness inherit the same blind spot; items with no positive sample contribute only negatives, even when a deterministic chain at matched compute would produce a positive. We do not propose to deploy activation grafting downstream; we use it to expose this slice so that math-reasoning pipelines account for it.

2. Related Work

Self-consistency and inference-time scaling. Self-consistency aggregates k sampled chains by majority vote (Wang et al., 2022) and is the default protocol for inference-time scaling on reasoning benchmarks (Wang et al., 2022; Chen et al., 2023; Aggarwal et al., 2023). Variants adaptively allocate the budget (Aggarwal et al., 2023), weight chains by self-evaluation (Chen et al., 2023), or combine sampling with verifier-guided search (Cobbe et al., 2021; Lightman et al., 2024). Across this line of work, $k = 1$

greedy is reported as a weak baseline that majority-voted sampling improves on. We do not dispute the aggregate ranking; we point out that the cell where greedy is correct and sampling-majority is not is non-trivial and persistent, and is hidden by aggregate accuracy.

Decoding diversity. Existing methods for decoding diversity operate in token space: nucleus and top-k sampling (Holtzman et al., 2020; Fan et al., 2018), diverse beam search (Vijayakumar et al., 2016; Kool et al., 2019), and contrastive decoding (Li et al., 2023); see Welleck et al. for a survey of inference-time decoding and meta-generation. All inject diversity through the per-step output distribution. Our diagnostic is orthogonal: the pass@$k = 0$ stratum persists at $k = 6$ under standard self-consistency hyperparameters, and is reached by deterministic perturbations applied to the residual stream rather than the output distribution.

Hidden-state and representation interventions. A growing body of work edits or analyzes internal activations: causal tracing for factual recall (Meng et al., 2022), representation engineering and steering (Zou et al., 2023; Turner et al., 2024; Rimsky et al., 2024), and cross-model communication via projected hidden states (Ramesh & Li, 2025). We borrow this primitive only as a probe, a minimal residual-stream perturbation whose deterministic outputs we compare against sampling, not as a method that should be deployed at inference.

Test-time compute and difficulty estimation. Recent work on test-time compute scaling reports favourable exchange rates between extra inference passes and accuracy on reasoning benchmarks (Snell et al., 2025; Brown et al., 2024; Wu et al., 2025). A common companion measure scores per-example difficulty by pass@k or sampling agreement (Cobbe et al., 2021; Brown et al., 2024), and the same signal is used to filter hard subsets out of a larger dataset (Zhou et al., 2026). Downstream RL-with-verifiable-rewards pipelines (Lambert et al.; Guo et al., 2025) consume this signal: prompts on which the model never samples a correct answer contribute no reward gradient. Our second observation directly affects this practice: a non-trivial fraction of examples that fail under sampling are solved by deterministic decoding, so sampling-only difficulty estimates conflate "hard" with "unreachable on the stochastic axis". Prior work on inference-time scaling has measured how coverage of correct answers grows with the sampling budget (Brown et al., 2024); we complement this by isolating the pass@$k = 0$ stratum and showing it persistence under additional stochastic samples.

3. Setup

We evaluate on the cross product of four open-weight instruction-tuned models, Qwen-2.5-3B (Qwen Team,

Figure 2. Constructing the deterministic regime. The frozen model is run on a prompt up to layer 26, producing the last-prompt-token residual-stream activation $h_{\text{last\_token}}$. It is then *replaced* with one of k cheap, parameter-free alternatives: $h_{\text{last\_token}}$ itself (the greedy baseline), the zero vector, a Gaussian sample, ..., the BOS-token activation. The same model continues its forward pass through layers 27 to num_layers under greedy decoding. The k chains differ only in this one residual vector, yielding k deterministic outputs $g_B, g_Z, g_R, \ldots, g_{\text{BOS}}$.

2024), Llama-3.2-3B, Llama-3.1-8B (Llama Team, 2024), and Mistral-Nemo-12B (Mistral AI Team, 2024), and three reasoning benchmarks, GSM8K (Cobbe et al., 2021), MATH (Hendrycks et al., 2021) and MMLU-Pro (Wang et al., 2024); $n = 1000$ matched prompts per cell, and max_new_tokens $= 2048$ to prevent truncation.

Decoding regimes. The *sampling regime* draws six chains $S_{1\ldots6}$ at $T = 0.7$, $p_{\text{top}} = 0.9$; we report sample-level quantities (pass@k, single-sample mean) directly, and where a sample aggregate is needed we use majority vote with ties broken uniformly at random over 50 seeds. A temperature sweep at $T \in \{0.3, 0.7\}$ appears in §8 (Table 5). The *deterministic regime* consists of greedy chains that differ only in a single last-prompt-token activation graft at $\ell = 26$, fired during prefill only; decoding is greedy throughout. All graft vectors are fixed across the dataset (no per-example optimisation, parameter-free, no extra forward passes beyond the prefill hook). Figure 2 provides an overview of grafting, while table 1 lists the eight chains used in the paper.

Matched compute. By *matched compute* we mean k deterministic chains against k sampled chains: the same number of prefill+decode forward passes per example. The graft is a single register_forward_hook that overwrites one residual-stream vector during prefill (no extra forward pass, no gradient, negligible wall-clock overhead), so we operationalise "compute" as the FLOP count of those k passes. The headline claim of Sec. 5 is at $k = 6$.

Chains were chosen to be mechanistically distinct in residual-stream geometry (the origin, a high-norm random direction, a token-position substitution, a hidden-dim permutation, a previous-layer state). We quantify the resulting fix-set diversity in §5 (cross-kind Jaccard ≤ 0.47 in every cell vs. a same-axis ceiling reaching 0.76 between

Table 1. Deterministic chains used in the paper. Short names are used interchangeably with the g. symbols from §5 onward.

Symbol / name	Description	Set
g_B	greedy, no graft	3
g_Z / ZERO	zero vector	3
g_R / RAND	norm-matched Gaussian	3
RUNIT	random unit direction	6
SHUF	hidden-dim permutation of g_B's activation	6
BOS	BOS-token activation	6
AVG	mean over prompt-token activations	8
PREV	same position from $\ell-1$	8

two random-direction probes; App. F). Layer and position choices shift the oracle ceiling by $\leq 4\text{pp}$ and reorder leading methods (Apps. B, H, F); the existence of the recovered slice is insensitive to these choices, the specific chain that recovers a given example is not.

4. Greedy is Competitive at $k=1$

At $k=1$, the sampling regime is one stochastic draw S_i, the deterministic regime is g_B. Running six independent sampling seeds (42–47), greedy is competitive with or better than the mean single-sample accuracy (Table 2), with the gap reaching up to $+2.8\%$ on `Llama-3.2-3B` / MATH. On six of twelve cells it exceeds the seed-to-seed range and beats the best seed. Greedy follows the local MAP trajectory while a single sample is a high-variance estimator of it; on harder problems the per-step distribution is flatter and MAP becomes a better single-pass bet than any draw.

Table 2. Single-pass head-to-head over six sampling seeds.

Setup	$1S$ Mean [Min,Max]	g_B	Δ
Qwen-2.5-3B/GSM8K	0.842 [0.838, 0.847]	0.843	+0.001
Qwen-2.5-3B/MATH	0.544 [0.539, 0.547]	0.557	+0.013
Qwen-2.5-3B/MMLU-Pro	0.403 [0.398, 0.411]	0.401	−0.002
Llama-3.2-3B/GSM8K	0.748 [0.736, 0.756]	0.770	+0.022
Llama-3.2-3B/MATH	0.461 [0.448, 0.469]	0.489	+0.028
Llama-3.2-3B/MMLU-Pro	0.345 [0.336, 0.361]	0.365	+0.020
Llama-3.1-8B/GSM8K	0.837 [0.826, 0.848]	0.847	+0.010
Llama-3.1-8B/MATH	0.485 [0.477, 0.492]	0.485	+0.000
Llama-3.1-8B/MMLU-Pro	0.483 [0.464, 0.502]	0.496	+0.013
Mistral-Nemo-12B/GSM8K	0.847 [0.842, 0.851]	0.857	+0.010
Mistral-Nemo-12B/MATH	0.321 [0.307, 0.340]	0.340	+0.019
Mistral-Nemo-12B/MMLU-Pro	0.431 [0.418, 0.437]	0.442	+0.011

The deterministic regime's clearest per-pass advantage is at $k=1$; at higher k, its contribution is better understood in terms of coverage (Sec. 5).

5. The Blind Spot is Structural, Not Hardness

The stratum. The pass@k=0 stratum, examples no sampled chain in k reaches gold, is the operative signal for label-required pipelines (RL with verifiable rewards, data curation, hardness annotation). At k=6 this stratum holds

Figure 3. Size of the pass@k=0 stratum as a function of sampling budget k, across the twelve (model, benchmark) cells ($n=1000$ each). The curve flattens early: most of the shrink from $k=1$ to $k=6$ happens by $k=3$, and a residual 5.1–43.5% of prompts remains unreached at $k=6$. The diminishing returns motivate spending budget on a different axis.

5.1–43.5% of prompts (51 to 435 examples per setup; see Table 3). This is the slice Sec. 5 attacks at matched compute.

The stratum persists at higher k**.** The per-sample marginal-shrink rate of pass@$k=0$ decreases monotonically across all twelve setups (Fig. 3; App. C). Within our sampling budget the shrinkage has already plateaued: pass@6=0 still contains 5.1–43.5% of prompts, and additional samples do not reliably close the gap. The stratum we attack is therefore not an artifact of an undersized sampling denominator.

Two readings of the residual slice are possible: those examples are intrinsically hard, or they are reachable but not on the stochastic axis. Section 5 adjudicates between them.

Anatomy. The recovered pass@6=0 examples share a recurring structure: the prompt admits two internally coherent readings of a discrete choice, sampling drifts toward one reading on all six seeds, and at least one deterministic chain commits to the gold reading. Which chain rescues a given example varies by example: no single graft dominates, but the union over a handful of mechanistically distinct chains catches a non-trivial fraction of the stratum. As a representative case, MATH Counting & Probability `0445` on `Qwen-2.5-3B` asks: *"A mother purchases 5 blue plates, 2 red plates, 2 green plates, and 1 orange plate. How many ways are there for her to arrange these plates for dinner around her circular table if she doesn't want the 2 green plates to be adjacent?"* (gold: 588). Greedy and the random graft both account for the identical-colour multiplicities and the circular symmetry, computing $10!/(5!\,2!\,2!\cdot 10) - 9!/(5!\,2!\cdot 9) = 756 - 168 = 588$. The zero graft instead treats the plates as 9 distinct objects and returns $9! - 8!\cdot 2 = 282{,}240$, ignoring both the identical-colour multiplicities and the circular division. All six sampled seeds in $\{42, \ldots, 47\}$ return non-gold answers, so the item is in pass@6=0, yet two of the three deterministic chains

recover it. Two further worked examples, spanning the other two rescuing pairs from $\{g_B, g_Z, g_R\}$, are in App. E.

Adjudicating the two readings. To adjudicate between the two readings of the residual sampling-failure slice, we work directly on the pass@6=0 stratum: examples for which no seed in $\{42, \ldots, 47\}$ reaches gold. On this slice, no sampled chain in six tries was correct: this is the strictest sampling-failure test. The slice contains 51 to 435 examples per setup (Table 3). We then ask: of these, how many does a deterministic regime at matched compute reach? Those would not actually be hard, but anyway flagged as hard by a sampling-based filter.

We define R_k as the number of pass@6=0 examples on which at least one of k deterministic chains is correct. R_1 is greedy alone. R_3 adds zero and norm-matched random grafts (g_Z, g_R). R_6 extends to greedy plus five grafts (zero, random, random unit direction, hidden-dim permutation, BOS-token replacement), matching the $k = 6$ sampling budget. R_8 adds mean-activation and previous-layer grafts. All grafts are at $\ell = 26$, last token position. The headline matched-compute claim is R_6 vs. pass@6=0 (six deterministic chains against six sampled chains, on the stratum where no sample reaches gold); R_1 and R_3 are reported at sub-matched budget and so strengthen, not weaken, the comparison (one or three deterministic chains reaching examples that six samples miss).

Three observations follow. (a) At matched compute, R_6 recovers 10–29% of examples that six sampling seeds collectively miss. The pass@6=0 stratum is therefore not intrinsically hard: a non-trivial fraction is reachable by deterministic decoding given enough representatives. (b) R_1 is still relatively small (0–9%), so a single deterministic chain is closer to "another sample" than to a separate regime; the regime emerges once multiple mutually disjoint perturbations are admitted. The recovery scales near-linearly with the deterministic budget (R_6 is several times R_1 on every setup where $R_1 > 0$). (c) The recovery is not driven by one privileged vector. Among the seven grafts surveyed, no single graft kind is the single largest individual contributor to R_6 on more than half the cells, and the leader rotates across cells (SHUFFLED on the math-style cells we inspected, RANDOM on Mistral-Nemo-12B / MMLU-Pro, multi-way ties on the smallest strata). Across all twelve cells, pairwise cross-kind fix-set Jaccard never exceeds 0.47 (range 0.06–0.47; App. F, Table 7), while the single same-axis pair (random vs. random-unit, two norm-matched Gaussians) reaches 0.76 on Qwen-2.5-3B / GSM8K as an illustrative ceiling for what resampling along a single random axis can contribute (see App. F for the per-cell breakdown). Different graft kinds therefore probe different subsets of the stratum in every cell we test, which is consistent with the recovery scaling with the deterministic budget.

These three readings together rule out both the "intrinsically hard" explanation and the "one privileged vector" explanation. Crucially, the recovery on pass@6=0 *scales with the deterministic budget*, admitting more mutually disjoint deterministic perturbations reaches more examples no sampling seed reached, and the dimensions on which it scales are mechanistically distinct from each other and from sampling's stochastic axis. What looked intrinsically hard under sampling was, for a non-trivial fraction, just unreached on the stochastic axis: the model was close, the missing ingredient was a different *kind* of diversity.

66–88% of pass@6=0 remains unadjudicated by an 8-chain deterministic regime: those examples may be intrinsically hard, or reachable on some other diversity axis we do not probe. Within the 3-chain set, each of $\{g_B, g_Z, g_R\}$ contributes some examples uniquely (Table 4): no single chain dominates across the twelve cells, and the deterministic budget pays back per chain. Per-setup unique contributions sit in the 1.3–8.9% band of $n = 1000$, with g_B leading in nine of twelve cells but never to the point of making either graft redundant, removing any one chain drops the 3-chain oracle by at least 13 examples on every setup, and the leading and trailing contributor differ by a factor of ≤ 2.2.

6. Mechanism: Residual-Stream Perturbation, Not Attention Rerouting

Counts and case studies establish that grafts reach examples sampling does not. They do not, on their own, explain why the deterministic chains in our table behave as multiple distinct axes rather than a single noisy one. This section gives the mechanistic answer.

We instrumented Qwen-2.5-3B with eager attention on 100 GSM8K examples and ran each twice: once unperturbed, once with a last-token graft at $\ell = 26$ (random and zero, separately). For every transformer block we recorded (i) the hidden state at the graft position before and after grafting, yielding cosine similarity and L_2 distance per layer, and (ii) the full attention-weight matrices, allowing direct comparison between baseline and grafted runs at identical token positions. For the random graft, $4/100$ examples exhibited a runaway-divergence mode in which L_2 grows to 10^4–10^5 in deeper layers; we exclude these from the per-layer summary to avoid heavy-tail contamination and report the trajectory on the remaining 96. Zero shows no such outliers.

Hidden-state divergence is shown in Figure 4 (per-layer numbers in App. A, Table 6). At $\ell = 26$, the random graft makes the hidden state essentially orthogonal to the baseline (cos = 0.001) at L_2 distance 121.8; the zero graft sets it to the origin ($L_2 = 86.2$, the baseline norm itself). Crucially the divergence does *not* decay through the remaining nine layers: L_2 grows monotonically to 466.7 (random) and

Table 3. Deterministic recovery on the pass@6=0 slice. R_k counts examples where no sampling seed in $\{42, \ldots, 47\}$ reaches gold but at least one of k deterministic chains does. Percentages are of the pass@6=0 slice (column 2); $n = 1000$ for all cells.

Setup	p@6=0	R_1	R_3	R_6	R_8
`Qwen-2.5-3B` / GSM8K	58	0 (0.0%)	5 (8.6%)	8 (13.8%)	10 (17.2%)
`Qwen-2.5-3B` / MATH	293	10 (3.4%)	23 (7.8%)	39 (13.3%)	48 (16.4%)
`Qwen-2.5-3B` / MMLU-Pro	323	14 (4.3%)	43 (13.3%)	81 (25.1%)	100 (31.0%)
`Llama-3.2-3B` / GSM8K	83	2 (2.4%)	12 (14.5%)	19 (22.9%)	21 (25.3%)
`Llama-3.2-3B` / MATH	287	11 (3.8%)	24 (8.4%)	42 (14.6%)	50 (17.4%)
`Llama-3.2-3B` / MMLU-Pro	336	30 (8.9%)	66 (19.6%)	96 (28.6%)	114 (33.9%)
`Llama-3.1-8B` / GSM8K	51	3 (5.9%)	7 (13.7%)	10 (19.6%)	11 (21.6%)
`Llama-3.1-8B` / MATH	291	9 (3.1%)	22 (7.6%)	30 (10.3%)	36 (12.4%)
`Llama-3.1-8B` / MMLU-Pro	252	19 (7.5%)	46 (18.3%)	72 (28.6%)	80 (31.7%)
`Mistral-Nemo-12B` / GSM8K	54	1 (1.9%)	8 (14.8%)	11 (20.4%)	13 (24.1%)
`Mistral-Nemo-12B` / MATH	435	16 (3.7%)	39 (9.0%)	64 (14.7%)	75 (17.2%)
`Mistral-Nemo-12B` / MMLU-Pro	322	11 (3.4%)	55 (17.1%)	93 (28.9%)	109 (33.9%)

Table 4. Per-source unique contribution to the 3-chain deterministic oracle $\det^{\oplus} = g_B \cup g_Z \cup g_R$ across all twelve cells. "Unique to x" counts examples whose gold answer is reachable through x but not the other two ($n = 1000$); equivalently, the drop in $\det^{\oplus}$ if x is removed. No single chain dominates: g_B leads on nine cells, g_R on two (both `Llama-3.1-8B`), and g_B is tied with g_Z on `Llama-3.2-3B` / GSM8K.

Setup	unique to g_B	unique to g_Z	unique to g_R
`Qwen-2.5-3B` / GSM8K	24 (2.4%)	21 (2.1%)	23 (2.3%)
`Qwen-2.5-3B` / MATH	33 (3.3%)	30 (3.0%)	21 (2.1%)
`Qwen-2.5-3B` / MMLU-Pro	68 (6.8%)	57 (5.7%)	58 (5.8%)
`Llama-3.2-3B` / GSM8K	37 (3.7%)	37 (3.7%)	31 (3.1%)
`Llama-3.2-3B` / MATH	79 (7.9%)	37 (3.7%)	53 (5.3%)
`Llama-3.2-3B` / MMLU-Pro	89 (8.9%)	61 (6.1%)	58 (5.8%)
`Llama-3.1-8B` / GSM8K	14 (1.4%)	13 (1.3%)	23 (2.3%)
`Llama-3.1-8B` / MATH	55 (5.5%)	39 (3.9%)	53 (5.3%)
`Llama-3.1-8B` / MMLU-Pro	49 (4.9%)	54 (5.4%)	57 (5.7%)
`Mistral-Nemo-12B` / GSM8K	35 (3.5%)	23 (2.3%)	21 (2.1%)
`Mistral-Nemo-12B` / MATH	71 (7.1%)	44 (4.4%)	45 (4.5%)
`Mistral-Nemo-12B` / MMLU-Pro	77 (7.7%)	52 (5.2%)	71 (7.1%)

415.1 (zero) by the final layer, while cosine similarity to baseline remains below 0.37. In this model, the forward computation does not project the perturbation back onto the baseline trajectory; it carries it forward.

By contrast, the graft's *direct* effect on attention is negligible (attention over the divergent generation itself is not what we measure here; see footnote). We report two quantities: (a) the mean absolute difference in `prefill_attn_received` (per-token attention received during the prompt forward pass, averaged over heads and query positions) between baseline and grafted runs, and (b) the mean absolute difference in the full attention-weight matrix under teacher forcing on *identical* token sequences (prompt + baseline output for both runs, so any difference is attributable to the graft and not to divergent generation).[1] On the 100 examples, (a) $= (1.1 \pm 0.3) \times 10^{-5}$ and

(b) $= (3.5 \pm 0.8) \times 10^{-5}$ for the random graft; the zero graft is within rounding $((1.2 \pm 0.4) \times 10^{-5}$ and $(3.5 \pm 0.9) \times 10^{-5})$. This is four to five orders of magnitude smaller than the residual-stream perturbation the same intervention induces.

From mechanism to behaviour. The two measurements above pin down how the graft acts on the forward pass: it does not reroute attention but injects a content vector that the remaining nine blocks carry forward without correction (Figure 4, L_2 grows monotonically; attention deltas are 10^{-5}). Mechanistically, each graft vector is therefore a different initial condition for the same nine-block computation producing the first generated token, and because the forward pass does not contract these perturbations back onto the baseline trajectory, distinct graft kinds produce distinct downstream trajectories that the autoregressive decode then locks in. This is the link to the behavioural table: different residual-stream perturbations propagate into different generated answers, and the eight chains in Table 3 are eight non-collinear initial conditions, not eight redraws from one.

The deterministic axis is therefore multi-dimensional in a behaviourally measurable sense: cross-kind fix-set Jaccard ≤ 0.47 in every one of the twelve cells (App. F) shows the chains act on largely disjoint subsets of the input distribution. That is why R_k scales with the deterministic budget rather than saturating after one graft: more graft kinds are more independent initial conditions, each reaching its own slice of pass@6=0.

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[1] Teacher forcing the grafted run on the baseline output is nec-

essary to isolate the graft's direct effect on attention from indirect effects mediated by divergent generation: the grafted chain typically produces a different output, and attention over that different output would conflate the two. Comparing on the *same* token sequence asks the cleaner question: does the graft, at layer $\ell = 26$, change which tokens subsequent attention heads attend to?

Figure 4. Hidden-state divergence at the graft position over layers $\ell \geq 26$ (Qwen-2.5-3B, GSM8K, last token, $\ell = 26$ graft; random excludes 4 runaway-divergence outliers, $n = 96$). Cosine similarity to the unperturbed run stays low and L_2 distance grows monotonically through the remaining nine blocks: the perturbation is carried forward, not absorbed. Baseline norm at the graft position is 86.2 ± 0.9 at $\ell = 26$ and 309.2 ± 14.5 at $\ell = 35$.

7. Practical Utility

The headline finding of §5 is qualitative: a six-chain deterministic regime reaches 10–29% of pass@6 = 0. This section turns it into two *deployable* recipes that need neither gold labels nor per-cell tuning. First, a matched-cost substitution (§7.1): a fixed-policy rule that spends one of B forward passes on the mean-prompt-activation graft AVG rather than an extra sample covers more of the dataset than B samples on the majority of (cell, budget) combinations, with AVG chosen once and never tuned. Second, a label-free curation flag (§7.2): chain disagreement among $\{g_B, S_0, S_1\}$ ranks recoverable items above the base rate without gold labels, with 3–5× lift on free-form math at three forward passes per item. Both recipes are deployable directly on top of an existing pass@k curation pipeline, at the cost of a small constant overhead per item.

7.1. Matched-cost coverage

A sharper test of the §5 finding is direct substitution at matched compute: at total budget B forward passes per item, does $(B-1)$ samples plus *one* deterministic chain cover more of the dataset than B samples? We propose the mean-prompt-activation graft AVG as a fixed, no-tuning rule: at every budget and every cell, spend one of the B

forwards on AVG rather than an extra sample. Across the 96 (cell, B) slots with $B \in \{2, \ldots, 9\}$, the fixed-AVG policy reduces residual pass@$B = 0$ count by $+1.66$ items per slot on average, and is net-positive on 8 of 12 cells: all four MMLU-Pro cells (mean $\Delta +4.4$ to $+7.0$ per slot), Qwen-2.5-3B / MATH ($+7.9$), Mistral-Nemo-12B / MATH ($+4.5$), Llama-3.1-8B / GSM8K ($+2.0$), and Mistral-Nemo-12B / GSM8K ($+0.4$). As an upper bound, we also report the oracle in which the best chain at each (cell, B) is picked from the eight in Table 1; this is an analysis, not a deployable rule. The oracle wins on 80 of 96 slots and is particularly large on MMLU-Pro and Mistral-Nemo cells (e.g. Mistral-Nemo-12B / MMLU-Pro: $379 \to 357$ at $B = 4$ and $277 \to 269$ at $B = 9$; Llama-3.2-3B / MMLU-Pro: $336 \to 317$ at $B = 6$; Qwen-2.5-3B / MATH: -10 at $B = 6$). The unconstrained best mix even goes det-heavy on three cells at $B = 9$ ($s = 1, d = 8$ on Qwen-2.5-3B / MATH; $s = 2, d = 7$ on Mistral-Nemo-12B / MMLU-Pro; $s = 4, d = 5$ on Llama-3.2-3B / GSM8K). The gap between fixed-AVG and the oracle is modest at the slot level (median best-chain wins $\approx$ 1–2 items more than AVG when they differ), so AVG reaches near-oracle coverage with no per-cell tuning. Per-cell, per-B oracle tables are in App. I. The headline is operational: when budgeting forward passes for coverage on a curation task, spending one of them on the AVG graft rather than an extra sample is the better choice on the majority of (cell, budget) combinations. The choice of a *single* graft slot is a proof-of-concept simplification: the same matched-cost substitution extends to $(B-d)$ samples plus d deterministic chains for any $d \leq B$, and a curation pipeline with a fixed audit budget per item can tune d to its own sample/deterministic exchange rate.

7.2. Toward label-free identification

A natural follow-up is whether the recoverable slice can be flagged *without* gold labels, purely from chain agreement. As a deployable probe we use only three forward passes per item, the greedy chain g_B and two sampling seeds S_0, S_1, and score each item by how many of the three pairs disagree. We then rank items by this score and ask whether the top-K% most-suspect items contain a higher density of recoverable items (pass@6 = 0 *and* some deterministic chain correct) than random selection would. The base rate of recoverable items is 0.8–9.6% across the twelve cells. On the eight free-form math cells (GSM8K, MATH) the probe concentrates recoverable items at the top of the ranking, with lift 3–5× at $K = 2$–5% on those cells and precision up to 20% at $K = 1$% on the strongest cell (Llama-3.1-8B / MATH); lift declines with K, as expected for a ranker. The signal lowers with MMLU-Pro cells, where the lift lies within 1.0–1.5× (see Limitations). Aggregate "3–5×" is therefore a majority claim, not universal. Per-setup numbers

are in App. G, Table 9; Fig. 6 shows precision-recall curves. This reframes the deployment from an accuracy intervention into a curation tool: given a fixed budget to re-examine $K\%$ of items with deterministic perturbations, the probe identifies the $K\%$ most likely to be *mislabelled* as intrinsically hard, at three forward passes per item and no gold labels. A calibrated precision guarantee (rather than rank-only lift), however, would require a small labelled dev set per pipeline, which we leave to future work.

8. Discussion

Robustness. Mechanism is residual-stream perturbation, not attention rerouting (§6); the diagnostic holds at lower sampling temperature (Table 5). On `Qwen-2.5-3B` / GSM8K with $T = 0.3$ across four seeds, the pass@4 = 0 stratum *grows* ($6.7\% \rightarrow 8.2\%$ of n) and the deterministic recovery on that stratum grows with it: R_6 rises from 20.9% at $T = 0.7$ to 31.7% at $T = 0.3$. Lowering T peaks the per-step distribution and reduces stochastic exploration, so the sampling-only blind spot widens while the deterministic regime continues to reach into it. The blind spot is therefore not a high-temperature artifact; if anything, the diagnostic is more visible at lower T.

Table 5. Pass@k = 0 and deterministic recovery on `Qwen-2.5-3B` / GSM8K at two sampling temperatures ($n = 1000$, four matched seeds each). R_k counts examples in the pass@4 = 0 stratum on which at least one of k deterministic chains is correct; percentages are of the stratum. Deterministic chains are temperature-independent; the only thing that changes between columns is the sampling stratum.

	$T = 0.3$	$T = 0.7$
pass@4=0 stratum size	82 (8.2%)	67 (6.7%)
R_1 on stratum	3 (3.7%)	1 (1.5%)
R_3 on stratum	15 (18.3%)	9 (13.4%)
R_6 on stratum	26 (31.7%)	14 (20.9%)

Intervention vs decoding. Activation grafting is an intervention on internal representations, not a decoding method: it overwrites the activation the model actually computed at layer 26 with a fixed vector and reads off predictions from the edited forward pass. The two regimes we compare are therefore not two ways of decoding the same system; one is the model running, the other is an internally edited model at matched compute. Our claim is correspondingly narrow: of the examples that pass@k = 0 classifies as too hard, a non-trivial fraction is reached by an edited forward pass at matched compute, so the slice is structurally identifiable in the residual stream. We do not claim those items are "truly easy" or that the unmodified model reaches them under ordinary inference; only that they are not unreachable in the broader sense, and that their reachability is decoding-regime dependent rather than an intrinsic property of the

items. Relatedly, there is no difficulty oracle: any operational notion of difficulty (pass@k, sampling agreement, intervention-based reachability) is a choice, not a measurement. The reverse asymmetry is real and we do not quantify it: some items the deterministic regime misses are reachable by sampling. We do not probe this direction because it is not what downstream pipelines act on: sampling is the de facto default for difficulty estimation, and pass@k-style proxies, RL-with-verifiable-rewards filters, and curation pipelines are often built on it. The deterministic regime, in current practice, is collapsed to greedy alone — a single chain, on which a pass@k hardness signal is undefined — so no operational pipeline currently labels items hard based on deterministic misses. What needs auditing is therefore the direction we report: items the sampling-based proxy currently flags as hard, on a stratum that downstream pipelines treat as such.

Limitations. Our study covers four open-weight models (3B, 8B and 12B parameters) on three reasoning benchmarks (two free-form math, GSM8K and MATH, plus one multiple-choice, MMLU-Pro), with seven graft vectors at a single layer and position and a default sampling temperature of $T = 0.7$ (plus one $T = 0.3$ check). Even with eight deterministic chains, 66–88% of the pass@6 = 0 stratum remains unrecovered; those examples may be genuinely hard, or reachable along a diversity axis we do not test. We report conditional recovery as point estimates across the twelve cells, without bootstrap confidence intervals or repeated-seed-set variance; the qualitative direction holds on every cell, but per-cell rates should be read with this uncertainty in mind. We also do not study whether the recovered slice can be identified *without* gold labels at deployable precision; we treat that as a separate question. The cheap cross-chain disagreement probe in App. G works well on free-form numeric and symbolic answers (lift@5% up to $8.0\times$ on GSM8K / MATH cells) but degrades on multiple-choice benchmarks (lift@5% collapses to 1.0–$1.5\times$ across the four MMLU-Pro cells): with a bounded answer vocabulary (A–J), two sampled chains and greedy frequently agree by chance even on recoverable items, so chain disagreement loses its signal. A deployable label-free flag would need a different signal on MCQ-style outputs. Finally, on the three smallest pass@6 = 0 strata (`Llama-3.1-8B` / GSM8K, 51 examples, `Mistral-Nemo-12B` / GSM8K, 54, and `Qwen-2.5-3B` / GSM8K, 58), absolute recovery counts are small ($R_6 \in \{10, 11, 8\}$); the recovery *rates* line up with the other cells, but the noise floor is higher.

9. Conclusion

Pass@k is the canonical sampling-based primitive for per-example difficulty, driving RL data curation, hardness annotation, and curricula across reasoning-LM evaluation. On

the twelve (model, benchmark) cells we test (four open-weight models spanning 3B, 8B and 12B parameters across three reasoning benchmarks), we show it has a persistent blind spot on its hardest stratum at the $k = 6$ budget: of examples no sampling seed in $\{42, \ldots, 47\}$ solves, 10–29% are reached by a six-chain deterministic regime at matched compute, while greedy alone reaches 0–9%. The recovery scales with the deterministic budget across five graft kinds whose mechanistic distinctness we verify across all twelve cells (cross-kind fix-set Jaccard ≤ 0.47 in every cell; App. F). The decoder is unchanged: stochastic decoding and sample majority remain the right default at inference, and grafting is purely a diagnostic tool for locating the slice.

The implication is that sampling-derived difficulty annotations conflate "hard" with "unreached on the stochastic axis" on a non-trivial fraction of their hardest stratum, which is what RL filters, curricula, and pass@k-based labels build on. For AI-for-math pipelines specifically, benchmark hardness buckets, pass@k-filtered synthetic curricula, and verifier datasets, this means an identifiable fraction of items currently labeled as too hard for the generator are, at matched compute, not. The slice is structurally locatable and can be audited at a few extra deterministic forward passes per item.

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Impact Statement

This work is a diagnostic study of how pass@k-based hardness labels behave on math and science reasoning benchmarks; it does not propose a new inference method or a more capable model, and the activation-grafting intervention is used only as an analysis tool rather than as something to deploy. The main downstream consequence we anticipate is more careful construction of math-reasoning evaluation suites, curricula, and verifier datasets, where treating pass@$k = 0$ as intrinsic hardness can silently and erroneously drop a structurally identifiable slice of samples the model can in fact solve. Better labels of this kind should reduce wasted compute in rejection-sampling pipelines and reduce miscalibrated difficulty signals in reinforcement-learning-with-verifiable-rewards training. We do not foresee specific dual-use risks: the diagnostic does not enable new generation capabilities, does not require any training, and is run entirely on existing open-weight models and publicly released benchmarks.

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A. Per-Layer Hidden-State Divergence (Numerical)

Table 6 reports the per-layer mean and standard deviation of cosine similarity and L_2 distance to the unperturbed run, summarised graphically in Figure 4.

Table 6. Hidden-state divergence at the graft position, mean $\pm$ std over 100 GSM8K examples (Qwen-2.5-3B, last token, $\ell = 26$; random excludes 4 runaway-divergence outliers, $n = 96$). Layers below 26 are not shown (cos $= 1$, $L_2 = 0$ identically). The baseline norm at the graft position is 86.2 ± 0.9 at $\ell = 26$ and 309.2 ± 14.5 at $\ell = 35$.

ℓ	g_R (random) cos	L_2	g_Z (zero) cos	L_2
26	0.001 ± 0.021	121.8 ± 2.0	0.000	86.2 ± 0.9
27	0.130 ± 0.047	155.2 ± 8.2	0.136 ± 0.074	147.7 ± 29.9
28	0.095 ± 0.047	186.8 ± 10.8	0.071 ± 0.056	177.1 ± 23.8
30	0.003 ± 0.048	258.4 ± 16.2	0.021 ± 0.057	238.3 ± 20.3
32	-0.005 ± 0.094	341.3 ± 27.5	0.136 ± 0.165	301.2 ± 34.3
34	0.205 ± 0.130	425.7 ± 37.3	0.361 ± 0.139	370.9 ± 38.9
35	0.001 ± 0.108	466.7 ± 33.4	0.017 ± 0.073	415.1 ± 28.1

B. Layer-Selection Sweep

We computed the $\det^{\oplus}$ oracle ceiling on Qwen-2.5-3B / GSM8K at four layers $\ell \in \{2, 13, 26, 32\}$, spanning early, middle, late, and final-block positions in the 36-layer stack. The ceilings are 0.902, 0.903, 0.908, and 0.905, with corresponding $G_1 = \det^{\oplus} - g_B$ values of $+5.9\%$, $+6.0\%$, $+6.5\%$, and $+6.2\%$. G_1 remains within a 0.6% band across this 30-layer range: the diversification effect is broadly layer-insensitive, consistent with a perturbation propagating forward through the residual stream rather than acting at a privileged depth. We adopt $\ell = 26$ as the marginal best across the 3B main-text cells (sweep done on Qwen-2.5-3B); the same layer is used for the 8B cells, and the choice is not load-bearing.

C. Pass@$k$$=$$0$ Shrink-Rate Diagnostics

A natural concern is that six sampling chains undersample the sampling regime, and that the pass@$k$$=$$0$ stratum would collapse with more samples. Going from $k$$=$$4$ to $k$$=$$6$ shrinks pass@$k$$=$$0$ substantially in absolute terms, but the stratum remains 5.1–43.5% of prompts at $k$$=$$6$, and the per-sample marginal-shrink rate is monotonically decreasing across the twelve cells (Fig. 3). Within our $k$$\leq$$6$ budget the shrinkage has already plateaued, so the R_6 recoveries reported in Sec. 5 (10–29% of pass@6$=$0) are not a quirk of an undersized sampling denominator. Whether the stratum would eventually vanish at much larger k (e.g. $k$$=$$50, 100$) is outside the budget of this study.

D. Recovery Scaling with the Deterministic Budget

Figure 5 visualises Table 3: how R_k grows from $k$$=$$1$ to $k$$=$$8$ across the twelve cells. Recovery scales near-linearly with budget on every cell, with no saturation by $k$$=$$8$.

E. Examples of Recovered Items

The R_k table is a counting argument. To make the mechanism concrete, we examine three pass@6 $= 0$ examples (Qwen-2.5-3B) on which exactly two of $\{g_B, g_Z, g_R\}$ are correct, with the rescuing pair differing between examples. No single chain is the privileged recoverer. We characterise each case by reading the generated chains and identifying the apparent reasoning step on which the failing chain diverges from the gold-reaching one; we do not claim these descriptions are recovered from inspecting the model's internals.

GSM8K 0798 (Rescued by g_Z, g_R). *"In the next 3 years, Billy will save \$X. He's saving up for a raise from \$10/hr to \$11/hr at his current job. Meanwhile, his co-worker Sally is going to retire in 3 years at \$12/hr. If Sally earns \$1.50 more than Billy after she retires and Billy gets his raise, what's the difference between their hourly wages?"* Gold: \$20. The intended reading puts Billy at \$11/hr after his raise and Sally at \$10.50/hr after retirement (Sally earns \$1.50 more than Billy's pre-raise wage), so Billy is \$20/hr higher in scenarios that double the difference, both grafts arrive at \$20. Greedy

Figure 5. Deterministic recovery on the pass@6=0 slice as the deterministic budget grows from 1 to 8 chains. R_k = number of pass@6=0 examples on which at least one of k deterministic chains is correct. Numbers above bars are raw counts; percentages are of column 2 of Table 3. Recovery scales near-linearly with budget across all twelve cells.

collapses "raise" and "retire" into an averaged rate $10.025/hr and returns $439.50, an arithmetically clean but semantically off-axis answer; all six sampled chains return non-gold answers.

MATH Counting & Probability `0445` (Rescued by g_B, g_R). *"A mother purchases 5 blue plates, 2 red plates, 2 green plates, and 1 orange plate. How many ways are there for her to arrange these plates for dinner around her circular table if she doesn't want the 2 green plates to be adjacent?"* Gold: 588. Total circular arrangements of 10 plates with multiplicities $(5, 2, 2, 1)$ are $10!/(5! \, 2! \, 2! \cdot 10) = 756$; arrangements with the two greens adjacent (treat them as one block of identical greens) are $9!/(5! \, 2! \cdot 9) = 168$, giving $756 - 168 = 588$. Greedy and the random graft both execute this argument cleanly. The zero graft instead treats the plates as 9 distinct objects in a circle and returns $9! - 8! \cdot 2 = 282{,}240$, dropping both the identical-colour multiplicities and the circular division. All six sampled chains return non-gold answers.

MATH Counting & Probability `0542` (Rescued by g_B, g_Z). *"Bob rolls a fair six-sided die each morning. If Bob rolls a composite number, he eats sweetened cereal. If he rolls a prime number, he eats unsweetened cereal. If he rolls a 1, then he rolls again. In a non-leap year, what is the expected number of times Bob will roll his die?"* Gold: 438. The expected rolls per morning solves $E = 1 + \frac{1}{6}E$, giving $E = 6/5$; over 365 days this is 438. Greedy and the zero graft both execute this recursion. The random graft instead sets $E = 2$ (a common slip from treating the re-roll as a single additional draw) and returns 730. All six sampled chains return non-gold answers.

Pattern. The three cases instantiate three different rescuing pairs from $\{g_B, g_Z, g_R\}$: $\{g_Z, g_R\}$, $\{g_B, g_R\}$, $\{g_B, g_Z\}$. The chain that fails is not the same chain across examples, and the chain that rescues is not the same either. The behavioural payoff of admitting multiple mechanistically distinct chains comes precisely from this lack of a privileged direction: any single chain, greedy included, misses a non-trivial slice of pass@6=0 that another chain in the set catches (Table 4).

F. Extra Graft Vectors

As a partial empirical check on `Qwen-2.5-3B` / GSM8K, augmenting $\{g_B, g_Z, g_R\}$ with five additional grafts, mean activation, BOS-token replacement, previous-layer activation, random unit direction, and shuffled activation, all at $\ell = 26$, raises $\det^{\oplus}$ from 0.908 to 0.940, suggesting that the diversification effect is not specific to zero and random vectors.

Fix-Set Diversity. We further check that distinct grafts rescue *structurally different* examples, not the same ones in slightly different forms. For each graft g_x, define its fix-set $F_x = \{i : g_B(i) \neq \text{gold}_i, \; g_x(i) = \text{gold}_i\}$ (examples baseline gets wrong but the graft rescues). We report pairwise Jaccard over fix-sets in two views: a per-cell summary across all twelve (model, benchmark) cells (Table 7) and a full per-pair matrix on `Qwen-2.5-3B` / GSM8K as a worked example (Table 8).

Across all twelve cells, the maximum cross-kind Jaccard never exceeds 0.47, with the per-cell maximum landing in $[0.26, 0.47]$. The single same-axis pair, RANDOM vs. RANDOM_UNIT (two norm-matched Gaussian probes that by con-

Table 7. Per-cell fix-set Jaccard summary across all twelve (model, benchmark) cells. *cross-kind J*: range of pairwise Jaccard over all unordered pairs of distinct graft kinds (ZERO, RAND, RUNIT, SHUF, BOS, AVG, PREV at $\ell = 26$, last-token). *same-axis J*: Jaccard between the two random-direction probes (RAND vs. RUNIT), both norm-matched Gaussians targeting the same axis by construction. $\max |F|$: largest fix-set across the seven grafts. The Qwen-2.5-3B / GSM8K row corresponds to the full per-pair matrix in Table 8.

Setup	n	cross-kind J	same-axis J	$\max |F|$
Qwen-2.5-3B / GSM8K	1000	0.28–0.45	**0.76**	55
Llama-3.2-3B / GSM8K	1000	0.20–0.45	0.43	75
Llama-3.1-8B / GSM8K	1000	0.26–**0.47**	0.46	55
Mistral-Nemo-12B / GSM8K	1000	0.07–0.44	0.43	47
Qwen-2.5-3B / MATH	1000	0.21–0.43	0.46	77
Llama-3.2-3B / MATH	1000	0.22–0.36	0.37	91
Llama-3.1-8B / MATH	1000	0.22–0.42	0.31	91
Mistral-Nemo-12B / MATH	1000	0.08–0.27	0.24	83
Qwen-2.5-3B / MMLU-Pro	1000	0.19–0.29	0.45	112
Llama-3.2-3B / MMLU-Pro	1000	0.12–0.28	0.42	124
Llama-3.1-8B / MMLU-Pro	1000	0.17–0.29	0.27	97
Mistral-Nemo-12B / MMLU-Pro	1000	0.06–0.26	0.21	114

Table 8. Pairwise Jaccard of fix-sets on Qwen-2.5-3B / GSM8K, last-token, $\ell = 26$. Each cell is $|F_a \cap F_b|/|F_a \cup F_b|$, where F_x is the set of examples baseline gets wrong but graft g_x rescues. RAND=norm-matched Gaussian (=g_R), RUNIT=random unit direction, SHUF=hidden-dim permutation, BOS=BOS-token replacement, AVG=mean prompt activation, PREV=same position from layer $\ell - 1$.

	ZERO	RAND	RUNIT	SHUF	BOS	AVG	PREV
ZERO	–	0.32	0.36	0.40	0.45	0.33	0.32
RAND	0.32	–	**0.76**	0.29	0.32	0.28	0.31
RUNIT	0.36	**0.76**	–	0.30	0.35	0.31	0.32
SHUF	0.40	0.29	0.30	–	0.43	0.39	0.34
BOS	0.45	0.32	0.35	0.43	–	0.40	0.44
AVG	0.33	0.28	0.31	0.39	0.40	–	0.43
PREV	0.32	0.31	0.32	0.34	0.44	0.43	–

struction target the same axis), ranges from 0.21 to 0.76 across cells and exceeds the per-cell cross-kind maximum on five of twelve cells (Qwen-2.5-3B / GSM8K, Qwen-2.5-3B / MATH, Llama-3.2-3B / MATH, Qwen-2.5-3B / MMLU-Pro, Llama-3.2-3B / MMLU-Pro), reaching 0.76 on Qwen-2.5-3B / GSM8K as an illustrative ceiling for what resampling along a single random axis can contribute. On the remaining seven cells (the larger Llama-3.1-8B and Mistral-Nemo-12B rows), the same-axis pair sits within the cross-kind band rather than above it; this reflects compression of all overlaps under the bounded MCQ output space (MMLU-Pro, A–J) and the smaller per-cell fix-sets at the larger model scales, not a violation of the cross-kind ceiling, which remains ≤ 0.47 everywhere. The universal statement is the cross-kind ceiling: distinct graft kinds rescue largely disjoint slices of the pass@$k = 0$ stratum in every cell we test.

The Qwen-2.5-3B / GSM8K matrix in Table 8 illustrates the pattern at full resolution: most pairs overlap on only 28–45% of their union, including $J(F_Z, F_R) = 0.32$ between the two main-text grafts, and the single high-overlap pair ($J = 0.76$) is the same-axis RANDOM vs. RANDOM_UNIT comparison. This justifies treating g_Z and g_R as independent diversification channels rather than redundant perturbations.

G. Label-Free Flag: Per-Setup Numbers

Table 9 reports the per-setup precision, recall, and lift of the cheap label-free probe described in §7.2. The probe scores each item by the number of disagreeing pairs in the three-chain set $\{g_B, S_0, S_1\}$ (three forwards per item) and ranks items in descending order. Recoverable items are those in the pass@6 = 0 stratum on which at least one chain in $\{g_B, g_Z, g_R\} \cup \{\text{RUNIT}, \text{SHUF}, \text{BOS}\}$ is correct. Lift is precision at the operating point divided by the base rate.

Table 9. Cheap label-free probe (3 forwards per item) at flagging recoverable items. $n_{p6} = 0$: pass@6 = 0 count. n_{rec}: recoverable count. Base: n_{rec}/n. AUPRC: rank-only area under PR. P@K%: precision among the top K% ranked. L@K%: lift over the base rate. Top-1% cells with P=0 reflect noise: on the two smallest recoverable strata ($n_{rec} = 8, 10$), the single highest-scoring item happens not to be recoverable; precision stabilises by P@2%.

Setup	Stratum sizes					Precision (%)				Lift (× base)			
	n	$n_{p6=0}$	n_{rec}	base	AUPRC	P@1	P@2	P@5	P@10	L@1	L@2	L@5	L@10
`Qwen-2.5-3B` / GSM8K	1000	58	8	0.8%	0.020	0.0	0.0	4.0	3.0	0.0	0.0	5.0	3.8
`Qwen-2.5-3B` / MATH	1000	293	39	3.9%	0.106	10.0	15.0	12.0	10.0	2.6	3.8	3.1	2.6
`Qwen-2.5-3B` / MMLU-Pro	1000	323	81	8.1%	0.106	30.0	20.0	12.0	9.0	3.7	2.5	1.5	1.1
`Llama-3.2-3B` / GSM8K	1000	83	19	1.9%	0.054	10.0	10.0	6.0	4.0	5.3	5.3	3.2	2.1
`Llama-3.2-3B` / MATH	1000	287	42	4.2%	0.080	10.0	15.0	12.0	6.0	2.4	3.6	2.9	1.4
`Llama-3.2-3B` / MMLU-Pro	1000	336	96	9.6%	0.140	0.0	0.0	10.0	16.0	0.0	0.0	1.0	1.7
`Llama-3.1-8B` / GSM8K	1000	51	10	1.0%	0.050	0.0	5.0	8.0	5.0	0.0	5.0	8.0	5.0
`Llama-3.1-8B` / MATH	1000	291	30	3.0%	0.083	20.0	15.0	8.0	8.0	6.7	5.0	2.7	2.7
`Llama-3.1-8B` / MMLU-Pro	1000	252	72	7.2%	0.098	0.0	10.0	10.0	11.0	0.0	1.4	1.4	1.5
`Mistral-Nemo-12B` / GSM8K	1000	54	11	1.1%	0.032	0.0	5.0	4.0	4.0	0.0	4.6	3.6	3.6
`Mistral-Nemo-12B` / MATH	1000	435	64	6.4%	0.103	0.0	10.0	10.0	10.0	0.0	1.6	1.6	1.6
`Mistral-Nemo-12B` / MMLU-Pro	1000	322	93	9.3%	0.091	10.0	10.0	10.0	8.0	1.1	1.1	1.1	0.9

Table 10. Oracle ceilings $\det^\oplus$ on `Qwen-2.5-3B` / GSM8K when the graft is applied at three different prompt positions; $\ell = 26$ throughout. Each ceiling is the union of g_B and the per-position graft set described in the text. Last-token grafting both reaches the highest ceiling and most cleanly diversifies the ensemble.

position p	ceiling $\det^\oplus$	Δ over g_B	# grafts
last (-1)	**0.940**	$+9.7\%$	7
first (0)	0.915	$+7.2\%$	6
penultimate (-2)	0.903	$+6.0\%$	7

H. Graft Position Ablation

The main-text grafts replace the hidden state at the *last prompt token* ($p = -1$). We ablate this choice on `Qwen-2.5-3B` / GSM8K by repeating the seven-graft sweep of Appendix F at three positions: last, penultimate ($p = -2$), and first ($p = 0$); per-position oracle ceilings appear in Table 10. The BOS-token graft at $p = 0$ is a no-op (it would replace BOS by itself) and is omitted there.

The last-token position dominates by $+2.5$–3.7% of oracle ceiling and is the position at which a single graft already realises the largest gain (g_{BOS} alone reaches 0.898, i.e. $+5.5\%$). First-token grafting is competitive ($\det^\oplus = 0.915$) but is dominated by the AVERAGE_ACT graft alone ($+4.2\%$), suggesting that early-position perturbations act through a single mechanism rather than via several disjoint channels. Penultimate is uniformly worst. We adopt the last-token position throughout the paper on this basis.

I. Matched-Cost Sample-vs-Graft Substitution

Per-cell, per-budget numbers backing the matched-cost coverage result in §7.1; Figure 7 visualises the same data, sweeping B for both the fixed-AVG policy and the oracle upper bound. At total budget B forward passes per item, the canonical sampling-only baseline ("pure S") runs the first B sample seeds (canonical order $S_{42}, \ldots, S_{50}$); the substitution ("S+D") runs the first $B-1$ samples plus the single deterministic chain that minimises residual pass@$B = 0$ count, picked from $\{g_B, g_Z, g_R, \text{RUNIT}, \text{SHUF}, \text{BOS}, \text{AVG}, \text{PREV}\}$. Lower is better; the best chain is shown in parentheses. Across the 96 (cell, B) combinations summarised below, substitution wins on 80, ties on 4, and loses on 12.

The strongest substitution gains concentrate on MMLU-Pro and on the larger models within each benchmark, where the sampling axis saturates earlier (cf. Fig. 3) and the deterministic axis still contributes fresh coverage. The AVG chain is the single most-frequently-chosen substitution (32/96 slots), followed by RAND (25), SHUF (10), BOS (7), B (6), Z (6), RUNIT (6), and PREV (4); no single chain dominates, consistent with the fix-set diversity reported in App. F.

Figure 6. Precision–recall of the cheap label-free probe (g_B plus two sampling seeds, three forwards) at flagging recoverable items (pass@6 = 0 *and* at least one deterministic chain correct). Dashed line is the base rate (random selection). The probe ranks recoverable items above the base rate across all twelve cells; precisions look low absolutely because the recoverable slice itself is small (0.8–9.6% of n). Per-setup top-K% numbers and lift are in Table 9.

Figure 7. Matched-cost coverage: pass@B = 0 count under pure sampling (gray dashed, "S": first B sample seeds), the proposed *fixed-policy* substitution (blue solid, "S+avg": $B-1$ samples plus the mean-prompt activation chain), and the oracle upper bound (red dotted, "$S+D^\star$": $B-1$ samples plus the single best deterministic chain at that (cell, B), picked from $\{g_B, g_Z, g_R, \text{RUNIT}, \text{SHUF}, \text{BOS}, \text{AVG}, \text{PREV}\}$), swept over $B \in \{2, \ldots, 9\}$. Lower is better. The fixed avg policy wins 59/96 (cell, B) slots with 8 ties and net mean $+1.66$ items per slot; the oracle, an upper bound that picks the best chain per slot, wins 80/96. Both lines drop below gray on every MMLU-Pro cell. Per-cell, per-B counts in Table 11.

Table 11. Pass@$B\!=\!0$ count under pure sampling ("S") vs. one-chain substitution ("S+D", best deterministic chain in parentheses) at matched compute $B \in \{2, \dots, 9\}$. Lower is better; **bold** marks the better column at each B (ties not bolded). Per-cell unconstrained best mix $(s^\star, d^\star)$ at $B\!=\!9$ in the last column ($n = 1000$). "M.-Pro" denotes MMLU-Pro

	Setup	B=2 S	B=2 S+D	B=3 S	B=3 S+D	B=4 S	B=4 S+D	B=5 S	B=5 S+D	B=6 S	B=6 S+D	B=7 S	B=7 S+D	B=8 S	B=8 S+D	B=9 S	B=9 S+D	best mix at B=9
GSM8K	Qwen-2.5-3B	**95**	99 (BOS)	**74**	75 (PREV)	67	**65** (RUNIT)	62	**61** (R)	58	**57** (R)	56	**54** (AVG)	55	**53** (Z)	53	53 (Z)	$s=2, d=7$
	Llama-3.2-3B	**154**	157 (PREV)	**118**	117 (Z)	100	**98** (SHUF)	92	**86** (SHUF)	83	**82** (Z)	80	**75** (SHUF)	75	**73** (SHUF)	71	**69** (SHUF)	$s=4, d=5$
	Llama-3.1-8B	92	**89** (R)	74	**68** (B)	66	**62** (B)	59	**57** (AVG)	**51**	52 (AVG)	**44**	46 (Z)	42	**40** (AVG)	42	**39** (SHUF)	$s=7, d=2$
	Mistral-Nemo-12B	94	**90** (R)	71	**70** (R)	62	**57** (R)	57	**53** (R)	54	**50** (R)	51	**48** (SHUF)	46	**45** (SHUF)	44	**42** (SHUF)	$s=4, d=3$
MATH	Qwen-2.5-3B	372	**368** (Z)	339	**330** (AVG)	314	**308** (AVG)	303	**292** (AVG)	293	**283** (AVG)	287	**276** (AVG)	282	**273** (AVG)	272	**269** (AVG)	$s=1, d=8$
	Llama-3.2-3B	414	**406** (B)	**358**	359 (PREV)	**322**	326 (RUNIT)	310	**301** (RUNIT)	**287**	290 (PREV)	279	**273** (RUNIT)	269	**265** (RUNIT)	263	**256** (RUNIT)	$s=8, d=1$
	Llama-3.1-8B	**398**	408 (BOS)	354	**352** (R)	**321**	325 (R)	303	303 (R)	291	**290** (R)	283	**279** (R)	**270**	271 (R)	262	**260** (R)	$s=9, d=0$
	Mistral-Nemo-12B	582	**560** (B)	531	**523** (B)	490	**488** (AVG)	462	**459** (AVG)	**435**	438 (AVG)	422	**416** (SHUF)	409	**405** (R)	402	**395** (B)	$s=8, d=1$
M.-Pro	Qwen-2.5-3B	486	**474** (BOS)	427	**415** (BOS)	377	**376** (BOS)	347	**341** (AVG)	323	**317** (AVG)	**298**	300 (AVG)	285	**278** (AVG)	275	**268** (AVG)	$s=7, d=2$
	Llama-3.2-3B	527	**524** (BOS)	456	**447** (AVG)	398	**393** (AVG)	355	**352** (AVG)	336	**317** (AVG)	304	**300** (AVG)	286	**277** (AVG)	266	**262** (AVG)	$s=7, d=2$
	Llama-3.1-8B	389	**388** (AVG)	324	324 (AVG)	300	**282** (R)	271	**265** (R)	252	**241** (AVG)	233	**226** (AVG)	214	**211** (AVG)	204	**195** (AVG)	$s=9, d=0$
	Mistral-Nemo-12B	467	**452** (BOS)	407	**395** (R)	379	**357** (R)	344	**337** (R)	322	**311** (R)	305	**295** (R)	293	**281** (R)	277	**269** (R)	$s=2, d=7$