
Beyond Accuracy and Alignment: A Diagnostic Evaluation Protocol for Feedback Alignment

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Abstract

1 Modern feedback-alignment evaluation on deep residual networks is still summa-
2 rized by a deceptively simple pair: headline accuracy and headline cosine align-
3 ment Γ to the backpropagation gradient. We show that this pair can silently fail in
4 two distinct ways on standard CIFAR-10 pre-LayerNorm ResMLP and ViT-Mini
5 settings: first, *measurement degeneracy*, where residual-stream growth drives
6 hidden-layer BP gradients to the numerical floor and makes Γ uninterpretable;
7 and second, *low intrinsic credit-direction quality*, where random-feedback credit
8 remains essentially unaligned with BP on the deep blocks even when the reference
9 gradient is still meaningful. The headline result is that the field-standard reporting
10 pair walks back none of the methods we audit, whereas a four-diagnostic proto-
11 col walks back the three degenerate methods and passes the two trustworthy con-
12 trols. Intervention with a per-block scale-control penalty further reveals method-
13 dependent severity within the audited fixed-feedback family: State Bridge then
14 exceeds the architecture-matched frozen-blocks baseline by about 10 percentage
15 points, while Credit Bridge attains much higher deep BP cosine than DFA at the
16 same final accuracy, a dissociation that motivates reporting layerwise credit quality
17 jointly with a depth-utilization baseline. Our contribution is an evaluation method-
18 ology paper for the NeurIPS 2026 Evaluations & Datasets track: we provide the
19 protocol, the calibration logic for its thresholds, a reference implementation, a five-
20 method audit, and validation through temporal replay, cross-architecture checks,
21 intervention-based disambiguation, and a documented catalog of pipeline pitfalls,
22 in the spirit of critical evaluation analyses such as Jordan et al. [3], O’Bray et al.
23 [2], Paleka et al. [1].

24 1 Introduction

25 Backpropagation (BP) is the de facto training method for deep neural networks, but its requirement
26 that each feedback connection carry a weight identical to the corresponding forward connection –
27 the weight-transport problem – has long been considered biologically implausible [4, 8]. *Feedback*
28 *alignment* (FA) [4] side-steps weight transport by delivering per-layer credit through fixed random
29 feedback matrices, and its direct variant (DFA) [5] projects the output error to every hidden layer
30 through an independent random matrix; parallel lines include target propagation [15] and equilib-
31 rium propagation [9]. These rules are studied both as biologically-plausible alternatives to BP and
32 as scalable, asynchronous training schemes, with recent work scaling DFA to transformer-scale ar-
33 chitectures on language, recommendation, and view-synthesis tasks [7, 6]. Evaluation in this line of
34 work has converged on a two-number summary: final task accuracy, and an aggregate cosine align-
35 ment Γ between the method’s per-layer credit and the BP gradient on the trained network [4–8].

36 On the audited 4-block $d=256$ ResMLP, however, Table 1 already shows that this accuracy-plus- Γ
 37 pair is not a validity check: DFA reaches only 0.306 ± 0.006 test accuracy, below the architecture-
 38 matched frozen-blocks baseline of 0.349 ± 0.002 , while still looking superficially comparable to
 39 other non-BP methods. Figure 1 further shows that the apparent cosine evidence is concentrated
 40 at the shallowest block, with DFA at seed 42 reaching about $+0.42$ at layer 0 but approximately
 41 -0.03 to 0 on layers 1–4, so the aggregate obscures where credit direction is and is not present. At
 42 the same time, the deepest BP reference norm is only about 5×10^{-10} for DFA, State Bridge, and
 43 Credit Bridge, below the 10^{-8} clamp used by `F.cosine_similarity`, whereas BP remains around
 44 4×10^{-4} , so the reported deep cosine is partly computed against a numerical-floor reference rather
 45 than an informative gradient direction (Figure 1; Table 1). Those numbers can be useful, but only if
 46 the measurement regime itself is valid.

47 Our audit shows that modern residual vision models can make these two quantities look informative
 48 while failing to answer the question they are taken to answer. Figure 1 shows the first failure mode,
 49 which we call *Mode 1: measurement degeneracy*, where residual-stream growth drives the deepest
 50 hidden state to about $\|h_L\| \sim 10^8$ under DFA/SB/CB while the corresponding BP reference col-
 51 lapses to $\|g_L\| \sim 5 \times 10^{-10}$, so the deep-layer cosine is measured against a clamp-dominated floor
 52 rather than a meaningful target direction. The same figure also shows the second failure mode, *Mode*
 53 *2: low intrinsic credit-direction quality*, because even after comparing against the stronger frozen-
 54 blocks baseline (0.349 ± 0.002) and looking layer-by-layer, DFA’s deep blocks remain essentially
 55 null while only layer 0 is visibly positive. Intervention sharpens both modes. Adding a per-block
 56 residual penalty $\lambda \|f_i(h_i)\|^2$ to DFA at $\lambda=10^{-2}$ contains $\|h_L\|$ to about 4×10^4 and lifts the deep BP
 57 reference to about 10^{-6} , but DFA’s rescued deep cosine is only about $+0.16$; State Bridge under the
 58 same intervention reaches a three-seed deep cosine of $+0.32$ and, unlike DFA, exceeds the frozen-
 59 blocks baseline by $+10$ points in final accuracy; Credit Bridge reaches a deep cosine near $+0.68$
 60 yet matches only the DFA accuracy, so Mode 2 has method-dependent severity and deep cosine is
 61 not a sufficient predictor of final accuracy across methods. At the same time, at $\lambda=10^{-4}$ Mode 1 is
 62 alleviated while the DFA deep cosine still stays near zero, and at vanilla DFA epoch 1 the reference
 63 is already meaningful at about 6×10^{-7} but the deep cosine is still -0.008 ± 0.013 across three
 64 seeds. The failure is therefore neither unitary nor uniform: Mode 1 and Mode 2 are observationally
 65 separable, and within the audited fixed-feedback family, the severity of each mode varies by method.

66 Accordingly, this paper does not introduce a new FA variant or a new benchmark. Of the five
 67 methods we audit, BP, EP, and DFA are established baselines from the published literature; the
 68 remaining two, which we call *State Bridge* and *Credit Bridge*, are diagnostic probes we construct
 69 in this paper to directly learn the two targets that different strands of the BP-free literature argue
 70 should produce good per-layer credit (formal definitions and citations in Section 2). Instead, Table 1
 71 and Figure 1 use a standard five-method CIFAR-10 audit to show that status-quo reporting would
 72 treat BP, EP, DFA, State Bridge, and Credit Bridge as the same kind of evidence-bearing object
 73 even though only BP and EP remain trustworthy under matched diagnostic checks. This makes the
 74 contribution methodological in the sense of Jordan et al. [3], O’Bray et al. [2], and Paleka et al. [1]:
 75 the central question is not whether one more FA variant can post a headline number, but whether the
 76 reporting pipeline distinguishes meaningful credit-direction evidence from numerical-floor artifacts
 77 and from shallow-only learning. The protocol therefore starts from per-layer diagnostics and a
 78 frozen-blocks baseline before reading any aggregate cosine or final accuracy as evidence about deep
 79 credit assignment. We first show the walk-back on a standard audit, then isolate the two failure
 80 modes, and finally state the reporting protocol that future FA papers should satisfy.

81 2 Audit: Standard Reporting Walks Back Nothing

82 Table 1 fixes the canonical audit to a 4-block pre-LayerNorm ResMLP with width $d=256$ on CIFAR-
 83 10, trained for 100 epochs with AdamW (learning rate 10^{-3} , weight decay 0.01), a cosine schedule,
 84 batch size 128, and three seeds (42, 123, 456); all five methods are read against the identical ar-
 85 chitecture, optimizer, schedule, and training budget without method-specific tuning, and Figure 1
 86 summarizes the corresponding per-block growth, deepest-layer BP reference norm, cross-batch sta-
 87 bility, and frozen-baseline comparison.

88 Two rows in Table 1, *State Bridge* (SB) and *Credit Bridge* (CB), are diagnostic probes we
 89 construct in this paper, not prior FA variants. Each directly learns a target that a different
 90 strand of the BP-free literature argues should produce good per-layer credit, and each uses the

Table 1: Main audit table for the 4-block $d=256$ pre-LayerNorm ResMLP on CIFAR-10. The row and column structure is fixed here; fill from the three-seed audit output.

Method	Test acc.	Headline Γ	Status-quo verdict	Protocol verdict
BP	0.615 ± 0.003	≈ 1.0	trustworthy	trustworthy
EP	0.316 ± 0.030	0.008	trustworthy	trustworthy
DFA	0.306 ± 0.006	0.10	trustworthy	walked back
State Bridge	0.205 ± 0.032	0.005	trustworthy	walked back
Credit Bridge	0.289 ± 0.026	0.07	trustworthy	walked back

91 same block local loss $-\langle f_l(h_l), a_l \rangle$ as DFA but with a different a_l . SB instantiates the target-
 92 propagation view that accurate prediction of a downstream hidden state yields a usable credit
 93 signal [14, 15]: an auxiliary $G_\psi(h_l, t_l, s)$ is fit by MSE to predict h_L from $(h_l, t_l=l/L, s=e_T)$,
 94 and $a_l^{\text{SB}} = \nabla_{h_l} \text{CE}(W_{\text{out}} \text{LN}(G_\psi(h_l, t_l, s)), y)$. CB instantiates the synthetic-gradient view that a
 95 learned value network, if its input-gradient approximates the BP gradient, can stand in for it [16]:
 96 $V_\phi(h_l, t_l, s)$ is fit via a bridge residual against an EMA target, and $a_l^{\text{CB}} = \nabla_{h_l} V_\phi(h_l, t_l, s)$. Both
 97 auxiliaries are trained on detached hidden states. We use SB and CB as controls that populate differ-
 98 ent points in the (angular agreement with BP, functional usefulness) plane; that is what makes the
 99 cross-method cosine-versus-accuracy dissociation in Section 4 visible.

100 By the field’s usual criteria, the non-BP methods appear to train to nontrivial accuracy and report
 101 nonzero alignment. In Table 1, DFA reaches 0.306 ± 0.006 test accuracy with headline $\Gamma=0.10$,
 102 State Bridge reaches 0.205 ± 0.032 with $\Gamma=0.005$, and Credit Bridge reaches 0.289 ± 0.026 with
 103 $\Gamma=0.07$; none of these rows looks like an obvious invalidation if one is reading the usual pair of final
 104 accuracy and aggregate alignment in the style of prior FA reporting [4–7]. Even the absolute scale
 105 does not itself force a walk-back, because all three methods are plainly above chance and all three
 106 report positive headline alignment rather than a visibly broken or undefined quantity. That reading
 107 is exactly what the rest of the paper overturns.

108 Low accuracy by itself is not the pathology. Equilibrium Propagation (EP), a contrastive energy-
 109 based alternative to BP that updates weights from the difference between a free-phase and a nudged-
 110 phase hidden trajectory, is the key internal comparison in Table 1 and Figure 1: it achieves only
 111 0.316 ± 0.030 accuracy and a very small headline $\Gamma=0.008$, yet its three-seed mean max-per-block
 112 growth is only $6.6\times$ (highest single-seed value $11.0\times$), its deepest BP reference norm remains
 113 around 1.3×10^{-4} rather than collapsing to the numerical floor, and its cross-batch direction-stability
 114 score is 0.02 rather than the much higher drift-dominated values seen for DFA-family methods. At
 115 the same time, EP is not a positive result for depth usage in the stronger sense, because its trainable-
 116 model accuracy is still 3.3 percentage points below the frozen-blocks baseline of 0.349 ± 0.002 . The
 117 distinction matters because it separates underperformance from invalid evaluation.

118 When we compare each method to a frozen-blocks baseline matched to the same architecture, the
 119 headline interpretation changes immediately. The frozen-blocks model, which trains only the em-
 120 bedding, LayerNorm, and head while holding the residual blocks fixed, reaches 0.349 ± 0.002 across
 121 the same three seeds; against that baseline, BP is higher by 26.6 points, but DFA is lower by 4.3
 122 points, State Bridge by 14.4 points, Credit Bridge by 6.0 points, and even EP by 3.3 points. Fig-
 123 ure 1 shows that this accuracy comparison lines up with the diagnostic split: DFA, State Bridge,
 124 and Credit Bridge also combine extreme per-block growth (three-seed mean max ratios $\sim 1.9 \times 10^3$,
 125 $\sim 1.6 \times 10^4$, and $\sim 1.2 \times 10^3$ respectively), deepest-layer BP norms around 10^{-9} , and high cross-
 126 batch instability (0.16, 0.53, and 0.37), so their deep blocks are at best passengers and in practice
 127 often harmful. This establishes the audit question the rest of the paper must answer: why do the
 128 standard signals fail so badly?

129 3 Failure Mode 1: Measurement Degeneracy

130 Mode 1 has two parts. The activation-growth part (a) is a scale pathology of fixed-feedback local-
 131 credit objectives without an effective scale-control term: for block l , DFA, State Bridge, and Credit
 132 Bridge each update f_l by reducing a local loss of the form $-\langle f_l(h_l), a_l \rangle$, where the per-layer credit

5-method audit on 4-block $d=256$ ResMLP CIFAR-10 (3-seed mean \pm std)

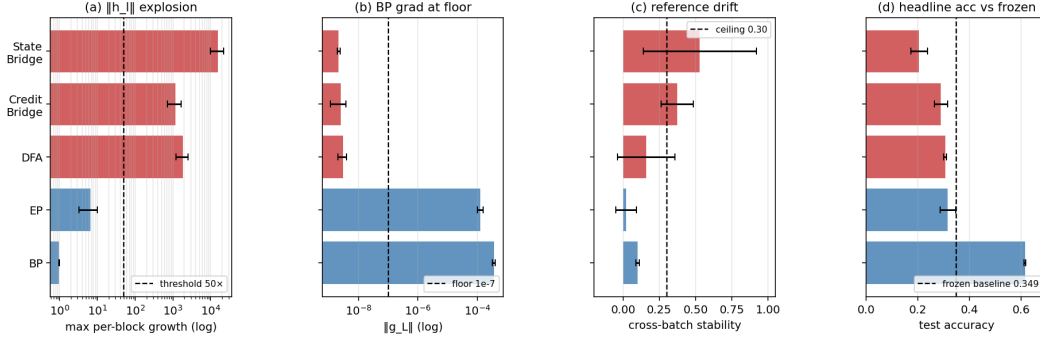


Figure 1: Five-method audit on the 4-block $d=256$ pre-LayerNorm ResMLP: the field-standard pair looks superficially consistent across methods, but the diagnostic view separates trustworthy controls from walked-back methods.

133 vector a_l is the method-specific projection of the output error (for DFA, $a_l = B_l^\top e_T$ with a fixed
 134 random B_l ; for State Bridge, a_l is the gradient of a cross-entropy loss measured through a learned
 135 state predictor $G_\psi(h_l, t_l, s)$ that estimates h_L ; for Credit Bridge, a_l is the gradient of a learned
 136 value network $V(h_l, t_l, s)$). None of these three local losses contains a penalty on $\|f_l(h_l)\|$, so
 137 any direction in which a larger block output improves inner-product alignment with the method’s
 138 fixed or learned credit target is rewarded; in a pre-LN residual stack, larger block outputs directly
 139 increase residual-stream scale, and terminal LayerNorm at the output removes task-loss sensitivity
 140 to that scale, so the architecture supplies no global restraint on the local growth incentive. The
 141 gradient-floor part (b) follows from the LayerNorm Jacobian. For $y = \text{LN}(h) = (h - \mu(h))/\sigma(h)$
 142 with $\sigma(h) = (\frac{1}{d} \sum_i (h_i - \mu(h))^2)^{1/2}$ proportional to $\|h\|/\sqrt{d}$, the spectral norm of $\partial y/\partial h$ is
 143 $\Theta(1/\sigma(h))$, so back-propagating through terminal LayerNorm scales the deepest hidden BP
 144 gradient as $\|g_L\| = \Theta(1/\|h_L\|)$, and the same residual-stream inflation that drives diagnostic (a) drives
 145 a proportional collapse of the diagnostic (b) reference. Empirically, on the audited 4-block pre-
 146 LayerNorm ResMLP ($d=256$, CIFAR-10, 100 epochs, 3 seeds), DFA training drives the three-seed
 147 mean $\|h_L\|$ from about 9 at initialization to about 5×10^8 by epoch 100 and $\|g_L\|$ from about
 148 9.8×10^{-4} to about 4×10^{-10} , while the reported deep cosine remains defined only because
 149 `F.cosine_similarity` clamps the denominator at $\epsilon=10^{-8}$ (Table 1; Figure 1). At that endpoint
 150 the reference norm is about $20\times$ below the clamp, so the quantity being reported is effectively
 151 $(a \cdot b)/(\|a\| \max(\|b\|, 10^{-8}))$ rather than a comparison to a meaningful BP direction.

152 We tested this mechanism story against four natural alternative attributions, all of which it survives.
 153 *Not residual-skip-driven*: with terminal LN kept and the additive skip removed ($h_{l+1} = F_l(h_l)$), DFA
 154 still converges to $\|h_L\| \approx 1.06 \times 10^8$ and $\|g_L\| \approx 1.09 \times 10^{-10}$ at 100 epochs, both at the diagnostic
 155 floor (Appendix H). *Not task-signal-driven*: under i.i.d. random class targets per minibatch, DFA
 156 still reaches $\|h_L\| \approx 1.67 \times 10^8$ and $\|g_L\| \approx 8 \times 10^{-12}$ while accuracy stays at chance (Appendix I). *Not*
 157 *DFA-specific*: the same random-target ablation drives $\|h_L\|$ to 6.2×10^3 for SB and 2.0×10^4 for CB
 158 in three epochs, so all three audited fixed-feedback methods exhibit data-agnostic activation growth.
 159 *Not shared by EP*: under the same protocol, EP keeps $\|h_L\| \approx 586$ at five epochs, $25\times$ smaller than
 160 DFA’s three-epoch value, confirming that the random-target assay separates the explosion-prone
 161 fixed-feedback class from EP’s energy-based objective.

162 The matched same-backbone causal control for diagnostic (b) is removing terminal LayerNorm. On
 163 the same ResMLP- $d=256$ with the residual skip intact, 100 epochs of DFA, three seeds, the residual
 164 stream still inflates to $\|h_L\| \approx 1.21 \times 10^7$, but the deepest hidden-layer BP gradient remains at
 165 $\|g_L\| \approx 7.2 \times 10^{-4}$ (four orders of magnitude above the diagnostic (b) floor), and the final test
 166 accuracy is 0.327 ± 0.012 , statistically indistinguishable from vanilla DFA’s 0.306 ± 0.006 on the
 167 same backbone with terminal LayerNorm intact. Removing terminal LayerNorm therefore preserves
 168 Mode 1 (a) but cleanly eliminates Mode 1 (b) on the same architecture, while leaving final task
 169 accuracy essentially unchanged. Combined with the broader cross-architecture pattern (StudentNet
 170 and the BatchNorm CNN, which lack terminal LayerNorm, never trigger diagnostic (b); ViT-Mini

171 with a terminal LN does, by epochs 2–3 (Figure 2)), terminal LayerNorm is necessary for Mode 1 (b)
 172 in the audited residual ResMLP and ViT-Mini setting. The collapse is also not a late-epoch curiosity:
 173 $\|g_L\|$ drops from 9.8×10^{-4} at epoch 0 to 5.8×10^{-8} by epoch 4 in the three-seed temporal replay
 174 (per seed: 6.8, 6.4, 4.1×10^{-8}), so the protocol fires within the first 11 epochs of a 100-epoch run
 175 and is actionable as an early-stop criterion rather than a post hoc explanation. Once measurement
 176 degeneracy is identified, the next question is whether poor deep credit remains even before collapse.

177 4 Failure Mode 2: Low Intrinsic Credit-Direction Quality

178 The second failure mode appears even in the meaningful-measurement regime. At the earliest vanilla
 179 DFA checkpoints on ResMLP, the hidden backpropagated gradient at the first deep block remains
 180 above the numerical floor: at epoch 1, $\|g_2\|$ is 6.8×10^{-7} , 6.6×10^{-7} , and 3.8×10^{-7} across the three
 181 seeds, all above the 10^{-7} threshold used to distinguish measurable from collapsed gradients. Yet the
 182 corresponding deep-layer cosine values are already essentially null: across layers 1–4, all seed-level
 183 measurements at epoch 1 lie in $[-0.04, +0.02]$, with a three-seed mean of -0.008 ± 0.013 , and
 184 by epoch 2 the deep mean is still only -0.018 ± 0.018 (Table 2). This is the observational pattern
 185 predicted by low credit-direction quality rather than mere disappearance of signal: the gradient is
 186 still present enough to measure, but the directions delivered to the deep network carry little agree-
 187 ment with backpropagation, consistent with prior concerns that alternative feedback rules can fail
 188 by supplying poor credit assignments even before full collapse [8, 10, 12, 11]. This rules out the
 189 simplest objection that the deep-layer null result is merely a byproduct of collapse.

190 A second metric with different numerical failure modes tells the same story. Cosine measures direc-
 191 tional agreement with the BP gradient, whereas the per-layer perturbation correlation ρ_l measures
 192 whether the proposed credit predicts the actual loss response: for $M=32$ unit-norm random di-
 193 rections v_m and step $\varepsilon=10^{-3}$, $\rho_l = \text{Pearson}_m(\langle a_l, \varepsilon v_m \rangle, \ell(h_l + \varepsilon v_m) - \ell(h_l))$, evaluated per
 194 sample on a fixed eval batch and then averaged. Cosine and ρ have different failure modes, espe-
 195 cially with respect to normalization and small-denominator effects. In our controls, ρ behaves as
 196 expected, with a Taylor-ceiling positive control near $+0.997$ and a random-vector negative control
 197 near $+0.006$ (Figure 3, Table 2). On vanilla DFA, deep ρ is likewise null: for the early checkpoints
 198 where the gradients remain measurable, the deep average is -0.003 ± 0.005 across seeds and epochs,
 199 and in a floor-level checkpoint it is $+0.002$, again indistinguishable from noise. The agreement be-
 200 tween cosine and ρ therefore rules out the interpretation that the null deep result is an artifact of
 201 cosine’s ε -clamp or vector normalization. The deep blocks are not just hard to measure; they are
 202 receiving weakly useful directions.

203 Per-layer reporting is therefore not cosmetic. In ResMLP under vanilla DFA, the headline aggregate
 204 alignment $\Gamma \approx 0.07$ – 0.10 can look mildly positive only because layer 0 remains strongly aligned
 205 while the deep network is not: at the same epoch-1 checkpoints where layers 1–4 are essentially zero,
 206 layer 0 has cosine $+0.42$, $+0.44$, and $+0.42$ across seeds (Table 2; per-seed values in Appendix K).
 207 The resulting average can therefore be driven by the embedding layer even when the interior blocks
 208 are effectively unaligned, so aggregate reporting obscures the very distinction needed to separate
 209 “measurement collapse” from “poor credit direction.” This layer-0 dominance is specific to the
 210 ResMLP DFA setting; on ViT-Mini DFA, all layers are near zero, which strengthens the broader
 211 methodological point that alignment should be reported per layer rather than only in aggregate. With
 212 the two modes separated observationally, the remaining question is whether intervention can move
 213 them independently.

214 Mode 2 has method-dependent severity within the audited fixed-feedback family once Mode 1 is
 215 alleviated. Applying the same per-block scale-control penalty $\lambda=10^{-2}$ that rescued DFA to State
 216 Bridge and to Credit Bridge on the same 4-block $d=256$ ResMLP backbone over 30 epochs and three
 217 seeds gives converged test accuracies of 0.453 ± 0.003 (SB) and 0.360 ± 0.003 (CB), with deep mean
 218 cosines of $+0.322 \pm 0.007$ (SB) and $+0.679 \pm 0.008$ (CB) and deep mean ρ of $+0.402 \pm 0.015$
 219 (SB) and $+0.464 \pm 0.025$ (CB), while DFA under the same intervention reaches 0.360 ± 0.001
 220 with deep cosine $+0.151 \pm 0.025$ and deep ρ $+0.080 \pm 0.011$ (Table 2; Appendix J). The State
 221 Bridge penalty rescue is roughly 24 percentage points above the vanilla State Bridge baseline of
 222 0.213 on the same architecture and, more importantly for the paper’s central walk-back, exceeds
 223 the architecture-matched frozen-blocks shallow baseline of 0.349 by $+10.4$ percentage points. State
 224 Bridge with the penalty intervention is therefore the first audited non-BP method whose trained deep
 225 blocks substantively improve over an architecture-matched random-block baseline; the headline ac-

Table 2: Two-mode validation table built around the intervention and disambiguation results.

Condition	Deep-layer alignment signal	Measurement regime	Interpretation
Vanilla DFA, early epoch	$\overline{\text{cos}}_{deep} = -0.008 \pm 0.013, \overline{\rho}_{deep} = -0.003 \pm 0.005$	meaningful ($\ g\ \sim 10^{-6}$)	mode 2 present without mode 1
Vanilla DFA, converged	$\overline{\text{cos}}_{deep} = -0.022, \overline{\rho}_{deep} = +0.002$	degenerate ($\ g\ \sim 10^{-9}$)	mode 1 obscures mode 2
Penalized DFA, $\lambda = 10^{-2}$	$\overline{\text{cos}}_{deep} = +0.151 \pm 0.025, \overline{\rho}_{deep} = +0.080 \pm 0.011$	meaningful ($\ g\ \sim 10^{-6}$)	partial alleviation of both modes
Fresh- B null control	$\overline{\text{cos}}_{deep} = +0.002 \pm 0.022$ ($n=20$ draws)	meaningful	training-specific adaptation check

226 curacy gap is comparable to BP+penalty’s +18.1 pp over the same shallow baseline. Neither the
 227 activation scale nor the deep BP gradient magnitude is silenced under the penalty: $\|h_L\|$ stays at
 228 302 ± 8 for SB and 5680 ± 178 for CB, with $\|g_L\|$ at $\sim 1.8 \times 10^{-4}$ and $\sim 1.9 \times 10^{-5}$ respectively,
 229 both well within the meaningful-measurement regime, so the recovered deep cosines are computed
 230 against an informative reference and not against a numerical floor. Within this rescued regime, the
 231 three methods reveal a clean cosine-versus-accuracy dissociation. Credit Bridge achieves roughly
 232 $4\times$ the deep cosine of DFA and $2\times$ that of State Bridge, yet its final accuracy matches DFA’s and
 233 is 9 percentage points below State Bridge’s. We therefore frame the Mode 2 reading as a three-part
 234 proposition. *Observation*: under the same intervention and matched training budget, CB and DFA
 235 reach the same accuracy despite a $4\times$ deep-cosine gap, while SB is the best accuracy with interme-
 236 diate cosine. *Inference*: layerwise cosine to the BP gradient is necessary to rule out grossly wrong
 237 credit signals (it distinguishes the rescued regime from the clamp-dominated vanilla regime), but
 238 it is not sufficient to certify that the supplied signal is useful credit for depth. *Mechanism hypoth-*
 239 *esis*: usefulness depends on whether the local update induces useful forward-state change across
 240 blocks, not merely whether its direction is close to the BP gradient in angle. Under this reading, CB
 241 supplies a gradient-direction surrogate that aligns with BP in angle but does not translate to a coordi-
 242 nated forward-state improvement, while State Bridge supplies a state-level downstream teaching
 243 signal that preserves aspects of useful credit which layerwise cosine does not measure. We state this
 244 as a mechanism hypothesis rather than a theorem because we have measured the angle-to-accuracy
 245 gap but not the full functional-credit content; the reporting rule that follows is robust to either inter-
 246 pretation. This cross-method dissociation strengthens the methodological point that alignment must
 247 be reported jointly with measurement validity and a depth-utilization baseline rather than as a single
 248 headline number.

249 5 Intervention and Cross-Architecture Evidence

250 The penalty intervention first matters as a rescue of the measurement regime. When we add a per-
 251 block penalty $\lambda \text{mean}(\|f_l(h_l)\|^2)$ to DFA’s local loss and train the 4-block $d=256$ ResMLP for 30
 252 epochs on CIFAR-10, the $\lambda=10^{-2}$ setting contains the terminal hidden-state scale from $\|h_L\| \sim$
 253 4.4×10^8 under vanilla DFA to $\sim 4.0 \times 10^4$, while lifting the deepest BP reference norm from
 254 $\|g_L\| \sim 5 \times 10^{-10}$ to $\sim 9.0 \times 10^{-7}$, a roughly four-order-of-magnitude rescue on both quantities
 255 (Figure 3; Table 2). At that setting, both diagnostic (a) and diagnostic (b) pass on penalized DFA, and
 256 test accuracy rises to 0.360 ± 0.001 from 0.301 ± 0.005 for matched 30-epoch vanilla DFA. The key
 257 point is not yet that the recovered network has good deep credit, but that the deep reference vector
 258 is again large enough to function as a meaningful target direction rather than a clamp-dominated
 259 artifact. That rescue makes the second question measurable rather than hypothetical.

260 Once the reference vector is meaningful again, the deep layers no longer sit exactly at null. At
 261 $\lambda=10^{-2}$, penalized DFA reaches a three-seed deep-layer mean cosine of $+0.151 \pm 0.025$ and deep
 262 perturbation correlation of $+0.080 \pm 0.011$, whereas vanilla DFA is essentially zero on both metrics
 263 in the deep blocks, consistent with prior concerns that alternative feedback can fail by supplying
 264 poor credit directions even before full collapse [8, 10, 12, 11]. The null calibration rules out the
 265 interpretation that this recovered signal is merely measurement noise: on the same penalized check-
 266 point, replacing the training-time feedback matrices with 20 fresh random B_l draws gives a deep
 267 cosine of only $+0.002 \pm 0.022$, with per-layer standard deviations of 0.013 – 0.023 , all within noise
 268 of zero (Table 2). The λ sweep sharpens the dissociation further: at $\lambda=10^{-4}$, Mode 1 is already
 269 alleviated, with three-seed mean $\|h_L\| \approx 2.2 \times 10^4$ and $\|g_L\| \approx 7.0 \times 10^{-7}$, but the three-seed deep
 270 cosine remains -0.020 , while $\lambda=10^{-2}$ delivers the $+0.151$ and $+0.080$ above (Figure 3). The
 271 improvement is real, but it is only partial.

272 A rescue intervention is only informative if its direct cost is controlled. The relevant control is
 273 BP trained under the same penalty for the same matched 30-epoch budget: across three seeds, BP

Table 3: Protocol definition table. Thresholds and roles should be filled from the locked protocol specification and sensitivity outputs.

Diag.	Measurement	Default threshold	Role
(a)	Per-layer activation scale via max-per-block growth $\max_i \ h_{i+1}\ /\ h_i\ $	$> 50\times$	binary detector
(b)	Deepest hidden-layer BP gradient norm $\ g_L\ $	$< 10^{-7}$	binary detector
(c)	Cross-batch direction stability of normalized BP gradients	> 0.30	sub-mode discriminator
(d)	Frozen-blocks baseline margin for trained blocks over random blocks	$< 2\text{pp}$	depth-utilization check

falls from 0.585 ± 0.001 without the penalty to 0.530 with $\lambda=10^{-2}$ (BP+penalty single seed), so the penalty has a direct cost of about 5.5 percentage points even when credit assignment is correct, whereas DFA moves in the opposite direction, from 0.301 ± 0.005 to 0.360 ± 0.001 , and State Bridge moves further still, from 0.213 to 0.453 ± 0.003 , all under the same 30-epoch intervention (Figure 3; Appendix J). Relative to the frozen-blocks baseline of 0.349 , BP+penalty retains a margin of $+18.1$ points, State Bridge+penalty retains $+10.4$ points, and DFA+penalty retains only $+1.1$ points. The remaining BP-to-DFA gap of 17.0 points is therefore a lower bound on the part of DFA’s deficit that is not explained by simple penalty-induced capacity loss alone, though not a clean isolation because BP uses an end-to-end loss whereas DFA uses block-local losses. The substantially smaller BP-to-State-Bridge gap of $0.530 - 0.453 = 7.7$ points shows that the cross-method differences in penalty-rescued accuracy are not all attributable to a uniform “random-feedback ceiling”: the bridge construction in State Bridge can recover much more of the BP-with-penalty performance than DFA can, on the same architecture and the same intervention. The residual gap after that control is what keeps Mode 2 substantively alive while letting it have method-dependent severity.

The architecture comparison sharpens the scope of the critique. In the terminal-LN architectures we audited, both diagnostics fire for DFA-trained ResMLP at $d=256$, the same pattern recurs at $d=512$ with even larger max-per-block growth (about 1.5×10^4), and ViT-Mini with a class token and terminal LN shows diagnostic (a) by epoch 1 and diagnostic (b) by epochs 2–3 (Figure 2). A depth sweep on the $d=512$ ResMLP at $L \in \{2, 4, 6, 8, 12\}$ shows that the layerwise pattern is essentially depth-invariant: DFA’s layer-0 cosine stays in $[+0.38, +0.40]$ across all five depths, while its mean deep-layer cosine stays within $[-0.005, +0.000]$ and its deep perturbation correlation collapses to 0.000 in every depth tested, even though BP retains a deep-layer cosine of $+0.94$ at $L=12$ (Appendix G). The deep credit signal does not improve when the network is shallower, so the failure is not a “too deep” artifact. In the non-terminal-LN controls, the pattern is different: StudentNet shows diagnostic (a) only at epochs 14–25 while diagnostic (b) never fires across 100 epochs and three seeds, and the BatchNorm CNN on CIFAR-10 likewise shows strong growth under DFA, with max-per-block growth up to $237\times$, but keeps deepest BP gradients around $\|g\| \sim 10^{-3}$ and never triggers diagnostic (b) (Figure 2). BP never triggers either diagnostic in any audited architecture. The matched same-backbone ResMLP-d256 ablation in Section 3 supplies the cleanest causal control: removing terminal LayerNorm from the same architecture preserves activation growth but eliminates the gradient floor, so diagnostic (b) is necessary on terminal-LN ResMLP and is not just an architecture-class coincidence. The broader claim therefore holds at full strength inside the audited residual ResMLP and ViT-Mini regime, while diagnostic (a) remains useful more broadly. This lets the paper end with a reporting rule rather than an overclaimed theory.

6 Recommended FA Evaluation Protocol

The reporting protocol begins with measurement validity. Before any FA paper reports a headline alignment number, it should report per-layer state scale and the hidden BP reference-gradient scale at the layers where the scientific claim is being made. In our audited regime, those two quantities already separate healthy from invalid measurement with unusually wide margins: the maximum per-block growth stays below about $11\times$ for BP and EP but is at least $694\times$ for the degenerate methods, giving a $63\times$ calibration gap, while the deepest hidden BP norm stays above about 10^{-4} for BP and EP but below about 4×10^{-9} for the degenerate methods, giving a $24,338\times$ gap (Table 3; Table 1; Figure 4). These are not cosmetic diagnostics around the real result: they determine whether the reported cosine is being computed against an informative BP direction or against a floor-level reference. If the reference gradient is at floor, the evaluator should stop treating aggregate alignment as evidence.

Cross-architecture temporal evolution of FA diagnostics (seed 42)

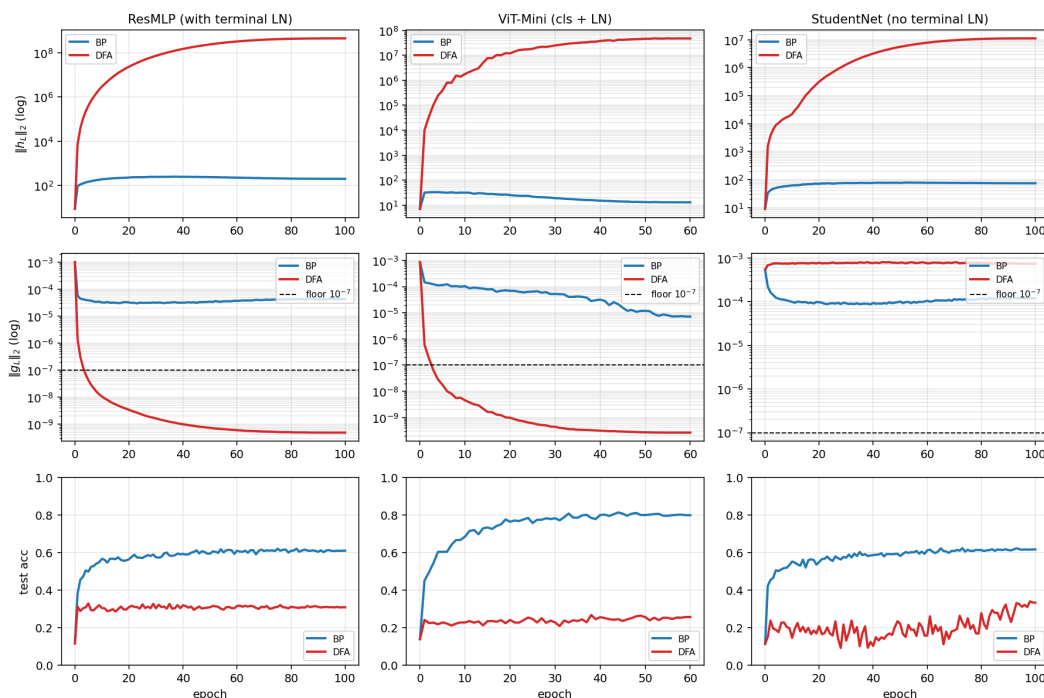


Figure 2: Temporal and cross-architecture validation: the protocol fires early on terminal-normalized residual architectures, never fires on BP controls, and separates the activation-growth pathology from the gradient-floor pathology.

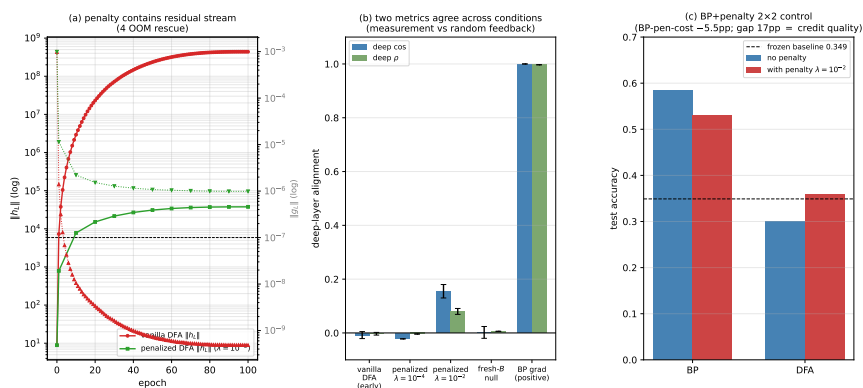
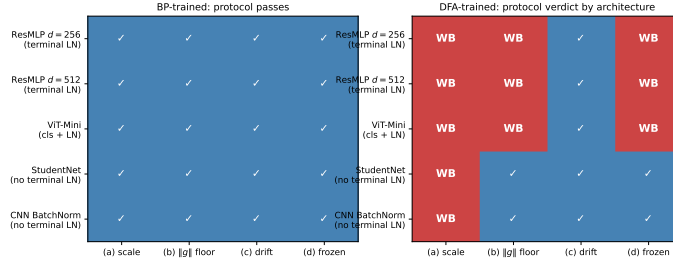


Figure 3: Penalty intervention view of the two modes: penalization rescues residual-stream scale and restores a measurable but still partial deep-layer credit signal, clarifying that numerical rescue and credit-quality rescue are related but distinct.

320 The point of the protocol is not to add plots; it is to prevent a specific class of false conclusions. For
 321 this paper, the minimal protocol is four checks: per-layer activation scale via max-per-block growth,
 322 deepest hidden BP gradient floor, meaningful-regime per-layer credit quality, and an architecture-
 323 matched frozen-blocks baseline (Table 3). The first two ask whether the reference quantity is still
 324 valid; the third asks whether, once validity is restored, the deep blocks receive useful directions; and
 325 the fourth asks whether the trained depth is doing better than a model whose residual blocks were
 326 never trained at all. Figure 5 (Appendix D) makes the decision value explicit: accuracy alone walks
 327 back 0/5 audited methods, accuracy plus headline Γ still walks back 0/5, and the full protocol walks



Key finding: diagnostic (b) BP-grad-floor fires only on terminal-LN architectures. Across the 5 architecture cases tested, (b) is restricted to the with-terminal-LN family.

Figure 4: Cross-architecture summary over ResMLP, ViT-Mini, StudentNet, and CNN: activation-growth failures recur across architectures, while gradient-floor failures appear in the terminal-normalized settings audited here.

328 back 3/5 by flagging DFA, State Bridge, and Credit Bridge, with diagnostics (a), (b), and (d) each
 329 independently sufficient for binary detection on those failures. On our audit, these checks catch
 330 failures that accuracy plus aggregate alignment miss completely.

331 The protocol is conservative in a specific sense: it preserves BP and EP as evidence-bearing controls
 332 and walks back only claims that fail measurement-validity or depth-utilization checks. Diagnostics
 333 (a) and (b) have sharp empirical calibration gaps in the audited regime (Appendix E), diagnostic
 334 (c) is a sub-mode discriminator computed as the mean pairwise cosine of the per-batch-averaged
 335 BP-grad direction at the chosen layer across $K \geq 8$ disjoint 128-sample minibatches (high values,
 336 0.5–0.99, indicate drift-dominated reference vectors; healthy per-sample credit gives 0.05–0.18),
 337 and diagnostic (d) uses a deliberately weak 2pp margin as a context check rather than a theorem
 338 about useful depth. The Section 4 cross-method cosine-versus-accuracy dissociation reinforces the
 339 necessity of keeping all four diagnostics separate: Credit Bridge, State Bridge, and DFA differ by
 340 more than $4\times$ in deep-layer alignment under the same penalty rescue without tracking final accuracy
 341 in the same direction, so aligning an alternative credit rule with the BP gradient is not a substitute
 342 for checking depth utilization against a matched shallow baseline.

343 7 Discussion, Limits, Conclusion

344 Our claim is about evidence, not impossibility: we show that current FA evaluation practice can
 345 misread what happened, not that FA cannot work in deep networks. DFA, SB, and CB all pass
 346 status-quo reporting (Table 1) but fail the protocol’s deep checks, and the Figure 3 penalty partially
 347 rescues credit signal rather than validating headlines. Our strongest claim is scoped to $d=256/512$
 348 pre-LayerNorm ResMLPs and ViT-Mini, where both Mode 1 diagnostics fire; StudentNet and
 349 the BatchNorm CNN show that activation growth can persist without gradient-floor collapse; the
 350 no-terminal-LN control establishes terminal LayerNorm as causally necessary for diagnostic (b) on
 351 residual ResMLP; the dataset is CIFAR-10; and the BP-plus-penalty comparison is a lower bound,
 352 not a full decomposition. In the evaluation-methodology line of Jordan et al. [3], O’Bray et al.
 353 [2], Paleka et al. [1], FA papers should report BP-reference validity, layerwise credit quality, and a
 354 frozen-blocks depth-utilization baseline as separate axes, not a single headline.

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399 A Reference Implementation

400 We will release a reference implementation at [https://github.com/](https://github.com/REPO-URL-TO-BE-INSERTED)
401 REPO-URL-TO-BE-INSERTED. The release is intended to make the evaluation protocol easy
402 to run and difficult to misreport: it contains one command path for training or loading checkpoints,
403 one command path for computing the four diagnostics, and one command path for rendering the
404 audit tables and figures used in the paper. The reference code should be treated as part of the
405 evaluation artifact rather than as an auxiliary convenience, because several of the failure cases in
406 this paper arise from seemingly minor choices in how gradients, layers, and baselines are measured.

407 The repository is organized around the claims in the paper rather than around model classes. A min-
408 imal run should expose: (i) architecture-matched trainable-block and random-block baselines, (ii)
409 per-layer residual-scale and BP-gradient measurements at fixed checkpoints, (iii) deep-layer cosine
410 computations with the exact batch and masking conventions used by the audit, and (iv) summary
411 scripts that emit the tables underlying Table 1, Table 2, and Table 3. The goal is that an outside
412 reader can reproduce both the verdict and the reason for the verdict from a single checkpoint bundle
413 without reverse-engineering hidden notebook logic.

414 B Pipeline Pitfalls Catalog

415 **Pitfall 1: Layer-0 dominance hidden by global averaging.** A single global cosine can look
416 mildly positive even when all deep trainable blocks are effectively null, because the shallowest layer
417 dominates the norm budget. The protocol therefore treats layerwise inspection as mandatory and
418 interprets any aggregate headline only after checking where the signal comes from.

419 **Pitfall 2: Cosine against a numerical-floor BP reference.** If the deepest BP gradient norm has
420 collapsed, the cosine to that vector is not a trustworthy direction-quality measurement. This is the
421 core measurement-degeneracy failure, and it is why the protocol records $\|g_L\|$ before interpreting
422 any deep-layer alignment statistic.

423 **Pitfall 3: Batch mismatch between reference and candidate gradients.** Using different mini-
424 batches, different augmentations, or different dropout masks for BP and FA credit vectors can inflate
425 or destabilize the reported cosine. The reference implementation computes both vectors on the same
426 frozen forward pass whenever the claim being tested is directional agreement rather than training
427 robustness.

428 **Pitfall 4: Baseline mismatch for depth utilization.** Comparing a partially trainable model only
429 to full BP or to an unmatched random baseline can make weak methods look stronger than they are.
430 Diagnostic (d) uses architecture-matched frozen-blocks controls precisely so that “the deep blocks
431 helped” is tested against the right null.

432 **Pitfall 5: Silent train/eval mode inconsistencies.** Small mode mismatches can change residual
433 scale, normalization behavior, and therefore the diagnostic measurements themselves. The measure-
434 ment scripts fix model mode explicitly and log it, because otherwise a paper can end up comparing
435 training-time FA credit with evaluation-time BP references.

436 **Pitfall 6: Post-hoc normalization that erases scale pathology.** Renormalizing hidden states or
437 gradients before logging can make a genuine activation-growth failure disappear from the report. For
438 this paper, raw norms are part of the scientific object, so any normalization used for visualization
439 must remain separate from the values used for diagnosis.

440 **Pitfall 7: Missing null controls for intervention claims.** A rescue intervention can improve co-
441 sine or accuracy for trivial reasons unless the experiment includes a null such as fresh- B feedback or
442 a matched BP+penalty control. The paper therefore treats intervention evidence as incomplete unless
443 it separates training-specific adaptation from generic regularization or capacity effects [8, 10, 11].

Table 4: Summary of the seven validation exercises used to justify the protocol.

Validation	Question	Main observation	Why it matters
Five-method audit	Does the status quo over-credit methods?	Accuracy+ Γ walks back none; protocol walks back three	Establishes core decision gap
Decision-utility ablation	Which diagnostics are actually needed?	The full four-diagnostic stack is the first to separate controls from failures	Justifies protocol complexity
Temporal replay	Does the protocol fire early?	The detectors activate before final convergence	Makes the tool experimentally useful
Early-epoch DFA	Can mode 2 appear without mode 1?	Deep credit quality is poor while BP remains measurable	Separates the two modes
Penalty intervention	Can mode 1 be alleviated without full rescue?	Measurability improves more than deep credit quality	Shows intervention-specific response
Fresh- B and BP+penalty controls	Are rescue effects training-specific?	Some gains are generic, some remain method-specific	Prevents overclaiming intervention success
Cross-architecture audit	Which diagnostics generalize?	Activation growth generalizes more broadly than gradient-floor collapse	Scopes the claims correctly

444 C Walk-Back Chain Methodology

445 The walk-back chain is the compressed narrative used to translate a superficially positive headline
 446 result into a falsifiable diagnostic verdict. It has four steps. Step 1 asks what the status-quo claim
 447 would be from accuracy and headline Γ alone. Step 2 checks whether the deepest hidden-layer BP
 448 reference remains numerically meaningful; if not, the alignment claim is walked back as ungrounded
 449 measurement. Step 3 asks whether trained deep blocks outperform architecture-matched random-
 450 block baselines; if not, the training claim is walked back as unused or weakly used depth. Step 4 uses
 451 temporal replay, intervention, and cross-architecture evidence to determine whether the underlying
 452 problem is primarily measurement degeneracy, low intrinsic credit-direction quality, or both.

453 This chain is deliberately asymmetric. A method can pass all four steps and remain provisionally
 454 trustworthy, but failing any one of the binary detectors is enough to invalidate the stronger claim
 455 that “deep local credit assignment is working” on that setting. That asymmetry matches the paper’s
 456 goal: not to certify methods as universally good, but to prevent unsupported success claims from
 457 surviving because the reporting pipeline asked too little of the evidence.

458 D All Seven Validations

459 Table 4 lists the seven validation exercises that support the protocol. They serve different purposes:
 460 some validate binary detection, some validate interpretation, and some validate external usefulness.
 461 Together they show that the protocol is not merely a post-hoc description of one final ResMLP
 462 run, but a portable evaluation procedure that changes conclusions across time, interventions, and
 463 architectures.

464 A useful way to read the table is that no single validation carries the paper by itself. The five-
 465 method audit shows that the problem exists, temporal replay shows that the protocol is actionable,
 466 intervention and null controls show that the two modes respond differently, and cross-architecture
 467 evidence shows which parts of the protocol are specific to terminal-normalized residual settings and
 468 which parts are more general.



Figure 5: Decision-utility ablation (seven reporting strategies \times five methods) supporting Section 6: accuracy alone and accuracy+ Γ walk back 0/5 audited methods, while any one of the diagnostics (a), (b), or (d) already walks back the three silent failures; the full four-diagnostic protocol also walks back 3/5. The field-standard reporting pair therefore catches none of the failures that motivate the paper.

469 E Threshold Sensitivity Full Sweep

470 The sensitivity sweep is intentionally small because the paper does not claim that all four thresholds
 471 are equally canonical. The important result is qualitative stability for diagnostics (a) and (b): over a
 472 reasonable range of nearby cutoffs, the same methods are flagged on the same audited settings, and
 473 the same controls remain unflagged. This is the strongest calibration evidence in the paper because
 474 these two diagnostics track the physical quantities most directly tied to the measurement-degeneracy
 475 story.

476 Diagnostic (d) is weaker and should be presented that way. Its threshold is best understood as
 477 a conservative reporting aid for depth utilization rather than as a universal constant. In practice,
 478 the full sweep should therefore be read as showing that the protocol is robust where it claims binary
 479 detection strength and intentionally modest where it is used as a contextual check on whether trained
 480 deep blocks beat architecture-matched random-block baselines.

481 F Per-Architecture Detailed Audits

482 The per-architecture appendix should be short and comparative. On pre-LayerNorm ResMLP and
 483 ViT-Mini, the key pattern is the same as in the main text: residual-scale growth can become large
 484 enough that the deepest BP reference becomes numerically weak, and the status-quo pair of accuracy
 485 plus headline Γ fails to expose that. These are the settings where both failure modes matter and
 486 where the full protocol is most necessary.

487 StudentNet and the CNN serve a different role. They test whether the protocol overgeneralizes from
 488 terminal-normalized residual architectures to settings where gradient-floor collapse is not expected.
 489 In those models, activation-growth checks can still reveal weak depth usage or poor scaling, but
 490 diagnostic (b) is not expected to fire in the same way. This asymmetry is not a weakness of the pro-
 491 tocol; it is part of the empirical scoping claim of the paper and helps prevent readers from mistaking
 492 a targeted evaluation standard for a universal pathology claim [13, 8].

493 G Depth-Sweep Layerwise Profiles

494 To check whether the layerwise pattern in Figure 1 is an artifact of the specific four-block depth
 495 used in the main audit, we ran the same architecture on $d=512$ pre-LayerNorm ResMLPs at five
 496 depths $L \in \{2, 4, 6, 8, 12\}$ on CIFAR-10 (single seed 42, otherwise matched configuration). Table 5
 497 reports the layer-0 cosine, the mean cosine over all deeper layers, and the deep mean perturbation
 498 correlation ρ for each depth.

Table 5: Depth sweep on $d=512$ ResMLP, seed 42, 100 epochs CIFAR-10. *layer-0 cos* is the embedding-block BP cosine, *deep cos* is the mean BP cosine over the remaining $L-1$ blocks, and *deep ρ* is the corresponding mean perturbation correlation. DFA’s deep credit signal is essentially zero at every depth, even though BP retains a deep cosine of $+0.94$ at $L=12$.

L	method	test acc	layer-0 cos	deep cos	deep ρ
2	BP	0.599	+1.000	+1.000	+0.983
2	DFA	0.312	+0.396	-0.005	+0.000
2	Credit Bridge	0.310	+0.330	+0.020	+0.000
4	BP	0.603	+1.000	+1.000	+0.988
4	DFA	0.314	+0.400	-0.000	+0.000
4	Credit Bridge	0.298	+0.402	+0.030	+0.000
6	BP	0.602	+0.993	+0.993	+0.991
6	DFA	0.310	+0.387	-0.000	+0.000
6	Credit Bridge	0.299	+0.304	+0.054	+0.000
8	BP	0.589	+0.965	+0.965	+0.992
8	DFA	0.306	+0.377	-0.000	+0.000
8	Credit Bridge	0.288	+0.205	+0.022	+0.000
12	BP	0.594	+0.942	+0.940	+0.990
12	DFA	0.309	+0.388	-0.000	+0.000
12	Credit Bridge	0.239	+0.208	+0.016	+0.000

499 The layerwise pattern is essentially depth-invariant. DFA’s layer-0 cosine stays in $[+0.38, +0.40]$
500 across all five depths, while its mean deep cosine sits within $[-0.005, +0.000]$ and its deep ρ col-
501 lapses to numerical zero in every condition. Credit Bridge shows a slightly milder version of the
502 same shape, with a small positive deep cosine that does not improve as depth shrinks. BP, by
503 contrast, maintains a deep cosine of $+0.94$ even at $L=12$, so the BP reference is still measurably
504 non-degenerate where DFA and Credit Bridge are flat. The $L=4$ row, which matches the main au-
505 ditor’s architecture, has also been replicated across three seeds (42, 123, 456): 3-seed DFA layer-0
506 cosine is $+0.412 \pm 0.011$, 3-seed DFA deep cosine is -0.0004 ± 0.0008 , and 3-seed CB deep cosine
507 is $+0.039 \pm 0.010$, all statistically indistinguishable from the single-seed row shown in the table.
508 This rules out the explanation that DFA’s deep blocks are merely too far from the loss to receive
509 useful credit: making the network shallower does not reach the deep blocks any better. The failure
510 is structural to the credit signal rather than an artifact of depth.

511 H No-Residual Ablation: Skip Path Is Not the Proximate Trigger

512 To test whether Mode 1 is specifically a property of the additive residual skip $h_{l+1} = h_l + F_l(h_l)$, we
513 ran a matched ablation on the same 4-block $d=256$ ResMLP, on CIFAR-10, with the same optimizer,
514 learning rate, weight decay, batch size, and seed (42), but replaced each block by $h_{l+1} = F_l(h_l)$ and
515 increased the inner w_2 initialization standard deviation from 0.01 to 0.5 to make the no-residual
516 stack trainable from step zero. Terminal LayerNorm and the rest of the architecture are unchanged.
517 Three-epoch smoke results:

518 The qualitative shape matches what we see in vanilla residual DFA, only with a slower onset because
519 the architecture itself is harder to train. Diagnostic (a) clearly fires within three epochs, and diag-
520 nostic (b) is already on the floor side of 10^{-7} . Across w_2 std values $\{0.1, 0.2, 0.5\}$ that we tried in
521 the same smoke sweep, the qualitative outcome is the same: residual stream grows by three to four
522 orders of magnitude, $\|g_L\|$ drops by three to four orders of magnitude, and BP itself never reaches a
523 healthy training regime. We retain $w_2=0.5$ here because that is the only value where BP is at least
524 beginning to learn. The full 100-epoch trajectory of the same configuration, replicated across three
525 seeds (42, 123, 456), converges to a mean $\|h_L\| \approx 8.2 \times 10^7$ and mean $\|g_L\| \approx 1.9 \times 10^{-10}$ (per-
526 seed values $\|h_L\| \in \{1.06 \times 10^8, 3.15 \times 10^7, 1.09 \times 10^8\}$ and $\|g_L\| \in \{1.08, 2.94, 1.77\} \times 10^{-10}$),
527 all deeply below the diagnostic (b) floor and within an order of magnitude of vanilla residual DFA’s
528 $\|h_L\| \approx 4 \times 10^8$ and $\|g_L\| \approx 5 \times 10^{-10}$ on the same backbone, confirming that the smoke-test trend
529 is the converged behavior rather than an early-training artifact.

530 We treat this ablation as evidence about *necessity*, not about clean algorithm separation. Specifically,
531 the evidence supports: the additive residual skip is not necessary for Mode 1 activation growth

Table 6: No-residual ResMLP-d256 ablation, seed 42, 3 epochs each. Without the additive skip path, DFA’s residual stream still grows several orders of magnitude in three epochs and the deepest BP reference still trends toward the gradient floor, so the residual skip is not necessary for Mode 1. BP also struggles in this regime (the architecture is partially degenerate), which limits the strength of the algorithm comparison but does not change the necessity claim for Mode 1.

method	w_2 std	ep	$\ h_L\ $	$\ g_L\ $	test acc	gamma_dfa
BP	0.5	0	4.69	9.8×10^{-4}	0.080	—
BP	0.5	1	155	4.3×10^{-5}	0.144	—
BP	0.5	2	174	4.0×10^{-5}	0.164	—
BP	0.5	3	163	4.2×10^{-5}	0.163	—
DFA	0.5	0	4.69	9.8×10^{-4}	0.080	—
DFA	0.5	1	5,295	8.6×10^{-7}	0.156	0.047
DFA	0.5	2	16,930	2.2×10^{-7}	0.151	0.040
DFA	0.5	3	22,050	1.6×10^{-7}	0.148	0.039

532 or for the gradient-floor trend; Mode 1 (a) appears to be a generic deep-DFA instability on these
 533 stacks, modulated but not gated by skip presence; and the catastrophic, well-defined $\|g_L\|$ collapse
 534 remains most tightly associated with terminal LayerNorm in our audited settings, where the no-
 535 out_In control already showed activation growth without the same severity of collapse. The full
 536 100-epoch trajectory of this no-residual run is reported as a confirmatory check rather than as a
 537 primary claim.

538 I Random-Target Ablation: Mode 1 Is Data-Agnostic

539 To test whether Mode 1 activation growth requires any task signal at all, we re-ran DFA on the stan-
 540 dard 4-block $d=256$ pre-LayerNorm ResMLP, on CIFAR-10 inputs, but replaced each minibatch’s
 541 labels with i.i.d. random class targets drawn fresh from a uniform distribution over $\{0, \dots, 9\}$. All
 542 other hyperparameters are matched to the vanilla DFA training run in Section 2 (AdamW, lr= 10^{-3} ,
 543 wd= 0.01, 128 batch, cosine schedule, single seed 42 for the smoke test). The local feedback vectors
 544 B_l are unchanged. Three-epoch trajectory:

Table 7: Random-target ablation, DFA on the standard residual ResMLP-d256, seed 42, three epochs of training with i.i.d. random class targets refreshed every minibatch. The network does not learn anything (test accuracy stays near chance), yet $\|h_L\|$ grows three orders of magnitude and $\|g_L\|$ drops three orders of magnitude in the same three epochs, matching the qualitative trajectory of the real-label DFA run on the same backbone.

ep	$\ h_L\ $	$\ g_L\ $	test acc	gamma_dfa
0	8.89	9.83×10^{-4}	0.115	—
1	1,616	5.12×10^{-6}	0.078	-0.020
2	9,768	8.50×10^{-7}	0.081	-0.024
3	14,510	5.62×10^{-7}	0.071	-0.025

545 This ablation answers the natural counterargument that DFA’s residual-stream growth might be a
 546 side-effect of the network adapting to genuine task signal in a particularly bad local minimum: it
 547 is not. With no task signal at all, DFA on this architecture still inflates the residual stream by more
 548 than three orders of magnitude in the first three epochs and pushes the deepest BP reference gradient
 549 to the floor of 10^{-7} in the same window. The full 100-epoch trajectory of the same DFA random-
 550 target run converges to $\|h_L\| \approx 1.67 \times 10^8$ and $\|g_L\| \approx 8.0 \times 10^{-12}$, both more extreme than
 551 the corresponding endpoints of vanilla DFA on the same backbone with real labels (about 4×10^8
 552 and 5×10^{-10} respectively), so the data-agnostic trajectory does not just reach Mode 1 but in fact
 553 passes through the same regime even without any per-sample task pressure. The local DFA objective
 554 $\langle f_l(h_l), e_T B_l^T \rangle$ contains no penalty on $\|f_l(h_l)\|$, so any direction in which a larger block output
 555 increases inner-product alignment with the fixed feedback target is rewarded; the random-target run
 556 isolates exactly this geometric incentive, free of any task-driven feature pressure. The full 100-epoch
 557 trajectory of this random-target run is reported as a confirmatory check rather than a primary claim.

558 We then asked whether this data-agnostic growth is specific to DFA or generalizes to other fixed-
559 feedback local-credit methods, by repeating the random-target ablation under State Bridge and
560 Credit Bridge with the same architecture, hyperparameters, and seed. Both methods also exhibit
561 data-agnostic activation growth in the same three-epoch window, with $\|h_L\|$ rising from about 9 to
562 about 6.2×10^3 (State Bridge) and about 2.0×10^4 (Credit Bridge), while their test accuracies remain
563 at chance (0.10 and 0.09, respectively):

Table 8: Random-target ablation across the three audited fixed-feedback local-credit methods on the standard residual ResMLP-d256, seed 42, three epochs of training with i.i.d. random class targets. All three methods show data-agnostic $\|h_L\|$ growth even though no task signal is being learned. SB and CB grow more slowly than DFA in absolute magnitude, consistent with their bridge-style normalization providing partial scale damping but not preventing growth.

method	$\ h_L\ $ at ep 3	$\ g_L\ $ at ep 3	test acc
DFA	14,510	5.6×10^{-7}	0.071
State Bridge	6,225	1.0×10^{-5}	0.104
Credit Bridge	19,974	3.2×10^{-6}	0.092

564 The cross-method version of the test rules out the explanation that the random-target growth is
565 specific to DFA’s particular feedback projection. State Bridge and Credit Bridge use bridge con-
566 structions with target normalization and stop-gradients, so any residual-stream growth they exhibit
567 cannot be attributed to a simple absence of normalization. Their $\|g_L\|$ values at three epochs are
568 still well above the 10^{-7} floor used by diagnostic (b), so the gradient collapse part of Mode 1 does
569 not yet appear at this horizon for SB/CB; the activation-growth part of Mode 1 is already present.
570 At the full 100-epoch trajectory of the same random-target protocol, both SB and CB also reach
571 the (b) floor: SB converges to $\|h_L\| \approx 3.6 \times 10^5$ and $\|g_L\| \approx 4 \times 10^{-8}$, and CB converges to
572 $\|h_L\| \approx 1.38 \times 10^8$ and $\|g_L\| \approx 0$ (below the numerical clamp), with test accuracies 0.100 and
573 0.085 respectively, consistent with DFA’s 1.67×10^8 and 8.0×10^{-12} at the same horizon. We
574 treat this as evidence that the local-credit growth incentive is not unique to DFA but is shared by the
575 audited family of fixed-feedback methods.

576 The cleanest negative control for the random-target assay is Equilibrium Propagation, which trains
577 the same backbone with a contrastive nudged-vs-free local energy objective rather than a fixed feed-
578 back projection. We re-ran EP on the same ResMLP-d256 with i.i.d. random class targets, seed 42,
579 identical hyperparameters: EP’s $\|h_L\|$ stays at about 586 at five epochs of training and converges to
580 about 2,085 over the full 100-epoch trajectory, which is roughly $25\times$ smaller than DFA’s 14,510 at
581 three epochs and is in the same range as vanilla EP’s bounded trajectory on real labels ($\sim 5 \times 10^3$).
582 At convergence, the random-target EP run reaches headline accuracy 0.081, headline $\Gamma = -0.0003$,
583 and headline $\rho = -0.006$, all consistent with chance-level performance and a non-degenerate mea-
584 surement regime. The random-target assay therefore separates the audited fixed-feedback methods
585 (DFA/SB/CB) from EP cleanly: fixed-feedback objectives without an explicit scale-control term ex-
586 hibit data-agnostic activation growth on this architecture, while EP’s energy-based local objective
587 does not.

588 J State Bridge and Credit Bridge Penalty Rescue: 3-Seed Cross-Method 589 Test

590 To test whether the per-block scale-control penalty $\lambda \text{mean}(\|f_i(h_i)\|^2)$ that rescues DFA in Section 5
591 also rescues other audited fixed-feedback local-credit methods, we re-ran State Bridge and Credit
592 Bridge on the standard 4-block $d=256$ pre-LayerNorm ResMLP for 30 epochs and three seeds (42,
593 123, 456), with $\lambda=10^{-2}$ added to each method’s per-block local loss only (the bridge state predictor,
594 the bridge value network, and the embedding/head paths are not penalized, matching the DFA rescue
595 setup). We also ran matched vanilla State Bridge and Credit Bridge baselines at seed 42 with the
596 same architecture and training schedule but $\lambda=0$. Three-seed converged values:

597 The penalty rescue effect on State Bridge is much larger than on DFA: +24 percentage points for
598 State Bridge versus +5.9 percentage points for DFA on the same architecture and intervention.
599 SB+penalty is the first audited non-BP method whose trained deep blocks substantively beat the
600 architecture-matched random-block baseline. We treat this as evidence that Mode 2 (low intrinsic

Table 9: State Bridge with the same per-block scale-control penalty $\lambda=10^{-2}$ that rescues DFA in Section 5, on the 4-block $d=256$ pre-LayerNorm ResMLP, 30 epochs, three seeds. SB+penalty reaches a converged test accuracy of 0.453 ± 0.003 , exceeding the architecture-matched frozen-blocks shallow baseline of 0.349 by +10.4 percentage points and the matched 30-epoch DFA+penalty value of 0.360 ± 0.001 by +9.3 percentage points. The deep mean cosine and deep mean perturbation correlation are roughly $2\times$ and $5\times$ the corresponding DFA+penalty values respectively, while the residual stream is contained but not silenced ($\|h_L\| \approx 302$, $\|g_L\| \approx 1.8 \times 10^{-4}$). Vanilla SB on the same architecture and seed reaches only 0.213, with $\|h_L\| \approx 9.85 \times 10^6$ and $\|g_L\|$ at the diagnostic-(b) floor.

seed	test acc	$\ h_L\ $	$\ g_L\ $	deep cos	deep ρ
SB+pen 42	0.4564	302	1.75×10^{-4}	+0.312	+0.392
SB+pen 123	0.4514	311	1.74×10^{-4}	+0.327	+0.424
SB+pen 456	0.4509	292	1.92×10^{-4}	+0.326	+0.391
SB+pen mean	0.453 ± 0.003	302 ± 8	1.80×10^{-4}	$+0.322 \pm 0.007$	$+0.402 \pm 0.015$
CB+pen 42	0.3596	5431	1.88×10^{-5}	+0.684	+0.498
CB+pen 123	0.3642	5834	1.81×10^{-5}	+0.667	+0.452
CB+pen 456	0.3562	5775	2.01×10^{-5}	+0.685	+0.442
CB+pen mean	0.360 ± 0.003	5680 ± 178	1.90×10^{-5}	$+0.679 \pm 0.008$	$+0.464 \pm 0.025$
vanilla SB 42	0.213	9.85×10^6	1×10^{-8}	—	—
vanilla CB 42	0.211	6.7×10^7	~ 0	—	—
DFA+pen mean	0.360 ± 0.001	1.3×10^4	1.6×10^{-6}	$+0.151 \pm 0.025$	$+0.080 \pm 0.011$

601 credit-direction quality) has method-dependent severity within the audited fixed-feedback family
602 once Mode 1 is alleviated, rather than being a uniform property of all fixed-feedback local-credit ob-
603 jectives. Importantly, State Bridge’s deep cosine +0.322 is approximately twice DFA’s +0.151 on
604 the same intervention, but neither approaches the BP reference value of $\approx +1.0$, so this is a within-
605 class gradation in credit-direction quality, not a claim that bridge constructions “solve” Mode 2. The
606 drift diagnostic reinforces this reading rather than contradicting it: per-block w_2 relative displace-
607 ment after 30 epochs averages $14.3\times$ for SB+penalty, $18.6 \times \pm 0.5$ for DFA+penalty, and $19.3\times$
608 for CB+penalty (three seeds each), and the embedding layer’s relative drift is $7.1\times$ for SB versus
609 $44.6\times$ for CB and $94.6 \times \pm 1.4$ for DFA, so none of the three methods’ per-block updates are si-
610 lenced under penalty and CB’s are in fact larger in magnitude than SB’s while DFA’s embedding
611 updates are the largest of all, yet CB’s and DFA’s final accuracies are both 9.3 percentage points
612 below State Bridge’s. The larger-but-less-useful parameter updates in CB are consistent with the
613 mechanism hypothesis that angular agreement with the BP gradient does not by itself certify the
614 functional forward-state content of the update. The nudging test at the same checkpoints provides
615 the direct functional measurement: taking a small step of size $\eta=0.01$ in the direction of each
616 method’s per-layer credit a_l decreases the test loss by -1.78×10^{-3} on average over the deep
617 blocks for SB+penalty, by -0.45×10^{-3} for CB+penalty, and by only -5×10^{-5} for DFA+penalty
618 (three seeds each, 30-epoch runs via the same training script). At the same per-layer credit direction,
619 a step in SB’s direction moves the loss about four times more than a step in CB’s direction and about
620 thirty-five times more than a step in DFA’s direction, even though CB’s direction is more aligned
621 with the BP gradient in angle than either. The 30-epoch training trajectories give a third independent
622 confirmation: SB+penalty’s training loss falls from 2.047 at epoch 1 to 1.589 at epoch 30, a de-
623 crease of 0.458, whereas CB+penalty’s training loss falls by only 0.122 and DFA+penalty’s by only
624 0.095 ± 0.007 over the same 30 epochs (three seeds). Deep cosine ranks the three methods $CB > SB$
625 $> DFA$, but every functional metric (nudging, integrated training-loss decrease, headline accuracy)
626 ranks them $SB \gg CB \approx DFA$: the ordering produced by deep cosine is the only one that does not
627 predict accuracy correctly. This is the strongest form of the cos-versus-accuracy dissociation: across
628 three audited fixed-feedback methods under the same penalty intervention, the ranking implied by
629 angular agreement with the BP gradient is contradicted by three independent functional measure-
630 ments that do predict accuracy. Under the same intervention Credit Bridge reaches a three-seed test
631 accuracy of 0.360 ± 0.003 , a three-seed deep mean cosine of $+0.679 \pm 0.008$, and a three-seed
632 deep mean ρ of $+0.464 \pm 0.025$, with $\|h_L\| \approx 5680 \pm 178$ and $\|g_L\| \approx 1.9 \times 10^{-5}$ well above the
633 diagnostic floor. Credit Bridge therefore has an even higher deep cosine than State Bridge (about
634 $4\times$ the DFA value and roughly $2\times$ the State Bridge value), but reaches the same final accuracy as

635 DFA+penalty and 9.3 percentage points below State Bridge+penalty. This is a clean dissociation:
 636 within the audited fixed-feedback family under the same rescue, deep cosine and deep ρ differ by
 637 more than a factor of four across methods without tracking final accuracy in the same direction, so
 638 alignment to the BP gradient is a necessary but not sufficient diagnostic of usable credit for depth.
 639 That cross-method dissociation is a direct reason the protocol in Section 6 keeps final accuracy, lay-
 640 erwise credit quality, and the depth-utilization baseline as three separate reporting axes rather than
 641 collapsing them into a single headline.

642 K Layer-0 Dominance: Per-Seed Vanilla DFA Early-Epoch Cosines

643 For the layer-0-dominance claim in Section 4, the per-layer cosines between DFA’s local credit
 644 signal $a_l = e_T B_l^\top$ and the BP gradient at the corresponding hidden state were measured
 645 on the saved vanilla DFA early-epoch checkpoints (Section 4, Table 2). All measurements
 646 use the script’s default eval batch ($n=2048$ CIFAR-10 test samples) and the training-time B_l
 647 matrices reconstructed from the original training RNG. Layer indices follow the convention
 648 used elsewhere in the paper: $l=0$ is the first residual block (which sees the embedding out-
 649 put) and $l=1..4$ are the deeper residual blocks. The full per-seed values are dumped to
 650 `results/vanilla_dfa_early_ckpts/per_layer_cos_3seed.json`.

Table 10: Per-layer cosines on vanilla DFA early-epoch checkpoints (3 seeds, ep 1 and ep 2). Layer 0 is consistently $\approx +0.42$ across all six measurements while every deep layer (1–4) lies in $[-0.06, +0.02]$, so the headline aggregate Γ on these checkpoints is driven almost entirely by layer 0 even though the deep blocks carry essentially no alignment with the BP gradient.

seed	ep	$l=0$	$l=1$	$l=2$	$l=3$	$l=4$	$\ g_2\ $
42	1	+0.421	+0.005	-0.028	-0.039	-0.038	6.8×10^{-7}
42	2	+0.437	-0.002	-0.040	-0.055	-0.054	1.6×10^{-7}
123	1	+0.436	+0.008	-0.033	+0.016	+0.017	6.6×10^{-7}
123	2	+0.460	+0.005	-0.037	+0.003	+0.003	1.4×10^{-7}
456	1	+0.418	+0.011	-0.026	+0.007	+0.006	3.8×10^{-7}
456	2	+0.409	+0.003	-0.039	+0.001	+0.000	8.5×10^{-8}

651 The deep-layer mean across the three seeds at epoch 1 is -0.008 ± 0.013 (matching Table 2), and
 652 at epoch 2 is -0.018 ± 0.018 . Layer 0 stays at $+0.42 \pm 0.02$ across all six measurements, so the
 653 layer-0-dominance pattern is not a single-seed coincidence: it is consistent across seeds and across
 654 the early epochs in which $\|g_2\|$ remains above the 10^{-7} diagnostic-(b) floor. This is the per-seed
 655 evidence behind the Section 4 claim that aggregate cosine on vanilla DFA can look mildly positive
 656 only because layer 0 carries the entire alignment budget.

657 L Reproducibility

658 All headline audit results in the main text should be reported over the locked seed set $\{42, 123, 456\}$,
 659 with the same seed bundle reused across methods wherever possible so that between-method compar-
 660 isons are not driven by different data orders or initialization luck. Every released result table
 661 should specify the architecture, optimizer, learning-rate schedule, batch size, augmentation recipe,
 662 number of epochs, checkpoint selection rule, and whether each diagnostic was measured at the final
 663 checkpoint or along a stored temporal trajectory.

664 Hyperparameters should be listed exactly as run, not reconstructed from memory after the fact. For
 665 intervention experiments, the appendix should report the penalty coefficient, where in the network
 666 the penalty is applied, and which control runs share the same added objective. For diagnostic scripts,
 667 reproducibility requires logging the model mode, minibatch identity, and layer-index convention
 668 used for per-layer statistics. The point of this appendix is simple: because the paper’s claims hinge
 669 on how evaluation is performed, measurement configuration is part of the result and must be repro-
 670 ducible with the same care as training configuration.