

Hadronic τ decays at higher orders in QCD

What four coefficients can, and cannot, support

Gauhar Abbas

Department of Physics, Indian Institute of Technology (BHU), Varanasi

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Outline

1. Motivation

$\delta^{(0)}$, $c_{5,1}$, and why it limits $\alpha_s(m_\tau)$

2. Formalism

The three-weight test is a tolerance

A no-go for a whole class of methods

Cross-validated renormalon closure (CVRC)

3. Validation

Moments, pole mass, all orders in large- β_0

... and the honest negative

4. Global comparison

Every approximant, one code path, one figure

5. Final predictions

$c_{5,1}$, $c_{6,1}$, and the calibrated Borel model

6. Summary and conclusions

One methodological thread runs through all of it:
declare the test before making the forecast, and
report the tests that fail as prominently as those that
pass.

Why inclusive τ decay is still the sharpest low-energy probe

and where the theory error actually sits

Optical theorem + analyticity:

$$R_{\tau,V+A} = N_c |V_{ud}|^2 S_{EW} \left[1 + \delta'_{EW} + \delta^{(0)} + \sum_{D \geq 2} \delta^{(D)} \right]$$

- $\delta^{(0)}$ is the massless perturbative term — the piece that carries α_s .
- Everything else (EW factors, condensates, duality violations) is *not* touched here.
- $\delta^{(0)}$ is known to $\mathcal{O}(\alpha_s^4)$: four Adler coefficients.

$$\hat{D}(Q^2) = \sum_{n \geq 1} c_{n,1} a^n, \quad a = \frac{\alpha_s}{\pi}$$

$$c_{1,1} = 1, \quad c_{2,1} = 1.6398, \quad c_{3,1} = 6.3710, \quad c_{4,1} = 49.076$$

The practical question

At fifth order, $c_{5,1}$ fixes the first omitted fixed-order term. It does *not* resolve the asymptotic FOPT–CIPT separation, which also depends on contour running and Borel singularities.

As Belle II and STCF reduce spectral-moment errors, perturbative control becomes increasingly important.

Can four coefficients support a **renormalon-resolved** extrapolation, not merely a locally successful one?

Why the FOPT–CIPT difference matters

Two perturbative organizations

- FOPT expands the running coupling at a fixed reference scale before performing the contour integral.
- CIPT evolves the coupling locally along the complex contour and does not re-expand it at a fixed scale.

At finite order,

$$\delta_W^{\text{FOPT}}(N) \neq \delta_W^{\text{CIPT}}(N), \quad (1)$$

and the difference has traditionally been included as an important theory uncertainty in extractions of α_s from hadronic τ decays.

Why the FOPT–CIPT difference matters

The modern renormalon interpretation

In the presence of infrared renormalons, the Borel representation underlying standard CIPT is not compatible with the usual local-operator form of the OPE. Consequently,

$$\delta_W^{\text{CI,Borel}} - \delta_W^{\text{FO,Borel}} = \Delta_W^{\text{Asymptotic-Separation}} \neq 0, \quad (2)$$

where Δ_W^{AS} is the asymptotic separation.

Interpretation

- The difference is not merely caused by the unknown coefficient $c_{5,1}$.
- Knowing $c_{5,1}$ reduces the first omitted fixed-order uncertainty, but does not remove the all-order separation.
- A renormalon-subtracted OPE scheme can strongly reduce the numerical FOPT–CIPT discrepancy. [See Pich and Rodriguez-Sanchez 2026](#)

Why $c_{5,1}$ is an important missing theory input

a qualitative statement — no α_s extraction is performed in this work

- FOPT and CIPT use the same Adler coefficients but organize contour running differently. Their truncated numerical sums differ, although the CIPT result re-expanded in the reference coupling reproduces FOPT through the stated order.
- $c_{5,1}$ fixes the next unknown fixed-order contribution and is therefore an important stability diagnostic. It does not make the all-order prescription separation vanish.
- The all-order separation involves analytic continuation and the Borel singularity structure. A credible continuation must be *renormalon-aware*, not merely a successful local fit.

Literature values of $c_{5,1}$

283	geometric ratio (not the model!) (Beneke–Jamin 2008)
277 ± 51	Borel–Padé (Boito, Masjuan, Oliani 2018)
278 ± 27	Levin (Abbas, Singh 2026)

The spread is not small, and — as we will see — different extrapolation methods applied to the *same four numbers* give anything from -22 to 3328 .

So the interesting question is not “what number?” but “which methods are entitled to answer?”

The rules, declared before any forecast

a validation hierarchy, not a post-hoc justification

Every method in this work must face the same four tests:

1. **Leave-one-order-out on three moments.** Withhold $c_{N,1}$, forecast it, and check the implied $d_N[W]$ for

$$W_\tau = (1-x)^3(1+x), \quad W_{GC} = (1-x)^3, \\ W_{P2} = (1-x)^2(1+x).$$

Criterion: below 10% errors on *all three*.

2. **Nearby renormalon.** Blind forecast of the four-loop pole- $\overline{\text{MS}}$ mass coefficient ($u = 1/2$).
3. **All orders.** Strictly recursive blind forecasting in the large- β_0 limit, where every coefficient is known exactly.
4. **A non-observable control.** Withhold the known five-loop $\overline{\text{MS}}$ β -function coefficient.

Two ground rules

Accuracy through order. A method must reproduce every supplied coefficient exactly. No fitting away the input.

No withheld coefficient may enter a forecast. Blind means blind, including in the calibration of any free parameter.

W_{GC} and W_{P2} are both $u = 2$ -sensitive but weight the Adler series very differently. Passing one and failing the other is diagnostic.

From moments back to a single number

the triangular weight map has unit diagonal

Analyticity turns the weighted spectral integral into a contour integral,

$$\delta_W^{(0)}(s_0) = \frac{1}{2\pi i} \oint_{|x|=1} \frac{dx}{x} W(x) \hat{D}(-s_0 x) = \sum_{N \geq 1} d_N[W] a^N.$$

In fixed order this is an exact **triangular** map from Adler coefficients to moment coefficients,

$$d_N[W] = \sum_{n=1}^N M_{Nn}[W] c_{n,1}, \quad M_{NN}[W] = 1 \text{ for every normalised } W.$$

Transfer identity

Because the diagonal is unity, an error in the forecast of $c_{N+1,1}$ passes into every moment with unit coefficient:

$$d_{N+1}[W] - \hat{d}_{N+1}[W] = c_{N+1,1} - \hat{c}_{N+1,1} \quad \textit{independently of } W.$$

The three-weight test is really a tolerance on one coefficient

For *any* transformation acting on the common Adler sequence, the declared ten-percent three-weight criterion is exactly

$$\left| \widehat{c}_{N+1,1} - c_{N+1,1} \right| \leq \min_{W \in \{W_\tau, W_{GC}, W_{P2}\}} 0.1 \left| d_{N+1}[W] \right|,$$

and with the exact moments,

$$\left| \widehat{c}_{3,1} - c_{3,1} \right| \leq 1.4523, \quad \left| \widehat{c}_{4,1} - c_{4,1} \right| \leq 2.1509$$

- The withheld- d_4 “ten-percent” test is in fact a 1.69% requirement on $d_4[W_\tau] = 127.08$.
- The withheld- d_3 test is a 5.51% requirement on $d_3[W_\tau] = 26.37$.

This single reformulation is what makes the rest of the analysis possible: a multi-observable criterion collapses to a one-dimensional window, and the whole problem becomes “*where can $\widehat{c}_{5,1}$ be?*”

Not a fixed-order artefact: in CIPT the map is *diagonal*, not triangular, and the error is no longer weight independent — but the collapse survives, and the same criterion is $\sim 35\times$ weaker (± 80 vs ± 2.15 at $N = 4$).

A no-go: why most methods cannot possibly work here

Define the one-step Padé/Shanks **defect** $\Delta_n = c_{n,1} - c_{n-1,1}^2/c_{n-2,1}$. From the four known coefficients,

$$(\Delta_2, \Delta_3, \Delta_4) = (+0.640, +3.682, +24.323).$$

Positive, and growing by a factor of six per order. But the geometric baseline at the next order is $c_{4,1}^2/c_{3,1} = 378.028$, so *any* literature-range $c_{5,1}$ requires $\hat{\Delta}_5 = \hat{c}_{5,1} - 378.028 < 0$: a **sign reversal**.

Proposition

Let $\mathcal{T}$ act on the common Adler sequence and be *defect-sign-preserving* (whenever $\Delta_2, \dots, \Delta_N > 0$ and increasing, it returns $\hat{\Delta}_{N+1} \geq 0$). Then

- (i) at $N = 4$ it returns $\hat{c}_{5,1} \geq 378.028$;
- (ii) the withheld- $c_{4,1}$ three-weight test does *not* exclude it, because the test window $\hat{\Delta}_4 \in [22.17, 26.47]$ lies entirely at $\hat{\Delta}_4 > 0$.

Hence: it can pass the weight test and still never return a $c_{5,1}$ in $[229, 305]$. The two are incompatible inside this class.

Ordinary Shanks/Padé, ARTS/PORT with a positive transported defect, and any single positive infrared factorial mode all live inside this class. Their mixed record is **forced, not accidental**.

Escaping the obstruction: declare the Borel structure

In the Borel plane, $u = \beta_0 t$, a singularity $N_p(p - u)^{-\alpha_p}$ contributes

$$c_{n,1}\Big|_p = N_p p^{-\alpha_p - n + 1} \frac{\Gamma(\alpha_p + n - 1)}{\Gamma(\alpha_p)} \beta_0^{n-1},$$

and a Borel monomial u^j contributes $\beta_0^j j!$ to $c_{j+1,1}$ *and nothing to any other order*.

The exponents are *declared*, not fitted:

$$\alpha_p = 1 + \frac{p\beta_1}{\beta_0^2} \quad (p > 0), \quad \alpha_{-k} = 2 - \frac{k\beta_1}{\beta_0^2} \quad (k > 0)$$

$$\alpha_2 = 2.5802, \quad \alpha_3 = 3.3704, \quad \alpha_{-1} = 1.2099$$

- $\alpha_{-1} = 2 - \beta_1/\beta_0^2$: the large- β_0 *double pole* at $u = -1$, RG-dressed. This is the Beneke–Jamin exponent 1.21.

Nesting rule at order N

N *constrained* columns, seeded by the leading IR singularity and the analytic term, then alternating:

$$\{u=2, u^0\}, \{u=3\}, \{u^1\}, \{u=4\}, \{u^2\}, \dots$$

plus **one more** column — the **closure**: the leading UV singularity $u = -1$, amplitude ρ .

$N + 1$ parameters for N constraints —
deliberately **one parameter richer than the data**.

Cross-validated renormalon closure (cvrc)

the closure family is exactly one-dimensional and affine

Imposing $c_{1,1}, \dots, c_{N,1}$ on $N + 1$ parameters leaves a one-dimensional affine set. Every member reproduces all supplied coefficients exactly (**accuracy through order for every ρ**), and every forecast is affine:

$$\widehat{c}_{N+1,1}(\rho) = A_N + B_N \rho$$

For the physical $n_f = 3$ series:

$$\widehat{c}_{3,1}(\rho) = 6.60484 + 24.500 \rho$$

$$\widehat{c}_{4,1}(\rho) = 31.6788 - 287.960 \rho$$

$$\widehat{c}_{5,1}(\rho) = 395.340 + 2439.209 \rho$$

$$\widehat{c}_{6,1}(\rho) = 3447.716 + 5720.039 \rho$$

- The slopes are *structural*: they depend on the declared singularity list and β_0 , not on the data.
- They **alternate in sign** with the order.

This is exactly the missing degree of freedom

One negative ρ simultaneously *lowers* $\widehat{c}_{3,1}$, *raises* $\widehat{c}_{4,1}$, and *lowers* $\widehat{c}_{5,1}$ relative to the pure infrared model.

A defect-sign-preserving transformation has no such handle. CVRC is not a better fit — it is **outside the class the Proposition forbids**, as is the Beneke–Jamin model, for the same reason. The new element is not the model but the **criterion** that can reject it.

The escape is the alternating ultraviolet mode at $u = -1$ — a physical statement, not a numerical trick.

Fixing the one free parameter — blind

There is exactly one free direction, so use exactly one leave-one-order-out fold. The model built from $c_{1,1}, \dots, c_{N-1,1}$ must reproduce the known $c_{N,1}$:

$$\boxed{\rho_\star = \frac{c_{N,1} - A_{N-1}}{B_{N-1}}} \implies \boxed{\widehat{c}_{N+1,1}^{\text{CVRC}} = A_N + B_N \frac{c_{N,1} - A_{N-1}}{B_{N-1}}}$$

A **closed-form nonlinear** transformation — nonlinear because $A_{N-1}, B_{N-1}, A_N, B_N$ themselves depend on the supplied data. **No withheld coefficient enters it.**

At physical order $N = 4$:

$$\rho_\star = \frac{49.07570 - 31.67880}{-287.96007} = -0.060414$$

- The *negative* sign is a statement about the physical series: the UV mode must enter with the sign that *suppresses* $c_{5,1}$.

Structural reading: closure = null
newest column

$\rho_\star$ is the unique value for which the column added at order N carries zero amplitude — i.e. the order- N and order- $(N-1)$ models coincide.

Verified numerically: at $N = 4$ the newest column is the monomial u^1 and its fitted amplitude is $\sim 10^{-16}$.

cvrc is the Beneke–Jamin model — stated explicitly

same singularities, same exponents, same closure, same data

Identical, ingredient by ingredient

Converting their convention, $\beta_k^{\text{BJ}} = 2\beta_{k-1}^{\overline{\text{MS}}}$:

$1 + \tilde{\gamma}$	BJ (2008)	this work
$u = 2$	2.580247	2.580247
$u = 3$	3.370370	3.370370
$u = -1$	1.209877	1.209877

- same singularity list $\{u=-1, 2, 3\}$;
- same parameter count (four);
- their $d_1^{\text{PO}} = 0$ is our **closure = null newest column**;
- same four coefficients.

Running our leave-one-out closure on *their* columns returns $\rho_\star = -0.016089$ — exactly their fitted d_1^{UV} .

We claim **no new model class** — the class that escapes the Proposition is theirs. What is new is the **acceptance criterion** that can reject a declared structure.

The only difference = the whole gap

Identical protocol, one ingredient at a time:

declared content	$\widehat{c}_{5,1}$	shift
pure powers (CVRC)	247.98	—
+ $\tilde{b}_1, \tilde{b}_2$ cuts	257.53	+9.6
+ $c_1[\langle aG^2 \rangle]$ (=BJ)	280.12	+22.6

The 32-unit gap is **not** a disagreement about renormalon content, closure, or data. It is 9.6 units of subleading cut structure plus 22.6 of a gluon-condensate Wilson coefficient we do not declare (backup: why not).

283 is *not* their model's output — it is their Eq. (3.10) geometric ratio, used as an **input**. Their own four-coefficient forecast is 280.

Test 1 — the three-weight leave-one-out test

fold	weight	exact	error
d_3	W_τ	26.366	-4.73%
d_3	W_{GC}	34.484	-3.61%
d_3	W_{P2}	14.523	-8.58%
d_4	W_τ	127.079	0.00%
d_4	W_{GC}	214.140	0.00%
d_4	W_{P2}	21.509	0.00%

All six inside the declared 10% band. No other construction examined in this work does that.

What this actually certifies — read carefully

The three d_4 entries are the **calibration fold** and are exact *by construction*. The three d_3 entries are a self-consistency reconstruction.

So this is a **consistency certificate for the declared structure**, *not* six blind forecasts. We say so in the paper, and we say so here.

The blind tests are the *external* ones — pole mass and large- β_0 — and they are next.

Is the certificate discriminating? — misspecification scan

is the admissible closure set nonempty for a *wrong* declared structure?

declared structure	admissible ρ window	$\widehat{c}_{5,1}$
baseline: $u_{\text{IR}} = 2$, $\alpha_{-1} = 2 - \beta_1/\beta_0^2$	$[-0.0679, -0.0529]$	247.98
$\alpha_{-1} = 1 - \beta_1/\beta_0^2$ (value used earlier in this work)	EMPTY — rejected	—
$\alpha_{-1} = 1$ (simple UV pole)	$[-0.0879, -0.0710]$	258.39
$\alpha_{-1} = 2$ (undressed double pole)	$[-0.0292, -0.0228]$	210.52
UV singularity at $u = -3/2$	EMPTY — rejected	—
UV singularity at $u = -2$	EMPTY — rejected	—
leading IR singularity at $u = 1$	EMPTY — rejected	—
leading IR singularity at $u = 3/2$	$[-0.0585, -0.0460]$	290.40
leading IR singularity at $u = 5/2$	EMPTY — rejected	—
leading IR singularity at $u = 3$	EMPTY — rejected	—

Six of ten declared structures are **rejected outright**: no value of ρ passes the three-weight criterion. The test has teeth — it is not satisfiable by construction, and it prefers the renormalization-group exponents.

Tests 2 and 3 — genuinely blind, external

Pole- $\overline{\text{MS}}$ mass, $u = 1/2$ renormalon

Blind forecast of the four-loop coefficient
 r_4 from the first three:

n_l	exact	CVRC	error
3	1702.7	1742.4	+2.33%
4	1235.6	1306.7	+5.75%
5	839.2	939.0	+11.90%

ARTS/PORT does *better* here
(1.60–3.84%) — and that is consistent
with the Proposition, because the
pole-mass defects do not require a sign
reversal.

Large- β_0 : every coefficient known, so test all orders

N	CVRC-min	CVRC-full	bal. Padé
4	SINGULAR	−108.3%	−68.7%
5	+23.7%	+264.1%	−11.6%
6	+9.5%	+4.6%	−45.7%
7	+13.0%	+3.4%	+787.0%
8	+9.6%	+0.8%	−22.7%
10	+8.0%	+1.2%	−9.5%
12	+6.6%	+0.7%	−3.7%
14	+5.6%	+0.6%	−1.4%

CVRC-complete: within 10% at every $N \geq 6$, within 1.5%
at every $N \geq 8$. The renormalon annihilator with *exact*
input needs 11–13 coefficients for the same persistent
control.

Two things the method does *not* do

1. At $N = 4$ in large- β_0 it *abstains*

At $N = 4$ in the minimal large- β_0 model, the design matrix is rank deficient. Two or more declared templates produce indistinguishable low-order coefficient patterns, so their amplitudes cannot be separated. The method returns no forecast rather than a pseudoinverse-dependent number.

Below $N = 6$ the procedure is not merely inaccurate — at $N = 4$ it is **not defined**.

2. Run strictly recursively, it misses $c_{4,1}$

Fix ρ one order lower still (from the $c_{3,1}$ fold) and forecast $c_{4,1}$ blind:

ρ from withheld- $c_{3,1}$ fold	-0.0095
ρ from withheld- $c_{4,1}$ fold	-0.0604
closure drift $ \Delta\rho $	0.0509
blind $\hat{c}_{4,1}$	34.43
exact $c_{4,1}$	49.08
error	-29.9%

This is reported as prominently as the successes. The closure drift is our **quantitative measure of how far four coefficients are from the asymptotic regime**.

We therefore do *not* claim that four coefficients *determine* $c_{5,1}$.

Every approximant, one set of inputs, one code path

how to read the next slide

Each section of the paper introduced a method and reported its continuation in its own normalisation. This figure recomputes **all of them** from

$$c_{1,1} = 1, \quad c_{2,1} = 1.6398, \quad c_{3,1} = 6.3710, \quad c_{4,1} = 49.076$$

through a *single* inverse renormalization-group map, so the entries are commensurable. No new method, no new assumption.

Status labels :

- CVRC — passes the acceptance criterion
- BASE — conditional baseline
- COND — accuracy through order, admissible as a conditional forecast
- DIAG — diagnostic only
- ABST — the method's own certificate abstains

Reference lines on the figure

- **grey band** — Borel–Padé literature range 277 ± 51
- dashed — Beneke–Jamin 283
- **solid red** — the **Shanks bound**
 $c_{4,1}^2/c_{3,1} = 378.03$ from the Proposition
- **blue band** — CVRC admissible closure window

The red line is a *prediction of the no-go theorem*: every defect-sign-preserving row must sit on or to the right of it. Watch whether it does.

Only one row is blue.

CVRC, top row:

$$c_{5,1} = 248 \pm 70$$

(the blue band drawn is the narrower *structural* one)

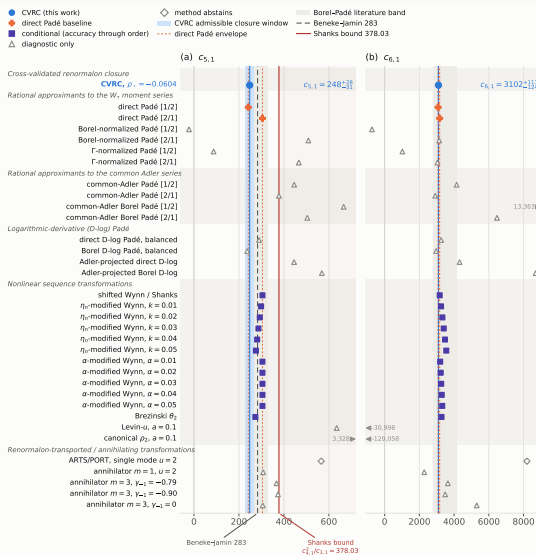
Red line: 378.03

The Shanks bound

$c_{4,1}^2/c_{3,1}$. Every defect-sign-preserving Adler-space row sits on or to the *right* of it — the Proposition, visible.

Grey band: 277 ± 51

Borel–Padé literature range; dashed line is Beneke–Jamin 283.



ARTS/PORT → 566

Passes the weight test, fails the range — exactly as the Proposition requires.

The cond block

Sequence transformations cluster tightly at 275–305, but none is validated: accuracy through order is necessary, not sufficient.

The diag scatter

Diagnostic rows run from −21.7 to 3327.8. That spread is the honest measure of how underdetermined four coefficients leave the problem.

Marker shape carries status as well as hue.

What the comparison shows

- The **fourteen** accuracy-through-order rows span

$$241.70 \leq c_{5,1} \leq 304.71.$$

- Admitting the *diagnostic* rows widens this to $[-21.73, 3327.76]$.
- So with four coefficients, **method choice dominates the answer** — by two orders of magnitude if one is not selective.

Two nonstatistical summaries, kept separate:

	$c_{5,1}$	$c_{6,1}$
direct- <i>Padé</i> baseline	273 ± 32	3125 ± 46
seq.-transformation grid	289 ± 16	3375 ± 175

Why CVRC is kept *outside* every average

It is the only row that passes the acceptance criterion the others fail. Averaging it with rows that fail that criterion would *destroy* the information the criterion carries.

We also state explicitly which combinations of rows are *not* admissible — e.g. direct *Padé* and shifted Wynn/Shanks coincide algebraically for the $[N-1/1]$ row and are not independent evidence.

The Proposition is confirmed empirically: every defect-sign-preserving row in the figure returns $c_{5,1} \geq 378.03$.

Final predictions

Physical-order result

$$c_{5,1}^{\text{CVRC}} = 248 \pm \underbrace{30}_{\text{struct}} \pm \underbrace{62}_{\text{method}} = 248 \pm 70$$

The **method band** is the *demonstrated* one-order-ahead accuracy at this order — not a parameter scan:

large- β_0 at $N = 4$ (the physical order)	undefined
large- β_0 at $N = 5$ (first defined order)	+23.7%
physical recursive test (one order lower)	-29.9%
pole mass (easier: nearby $u = 1/2$)	2.3–11.9%
adopted	25% = 62

Consistent with 277 ± 51 , with 273 ± 32 , and with **280**, the Beneke–Jamin four-coefficient forecast — from which we differ only by subleading cut structure (+9.6) and a gluon-condensate Wilson coefficient (+22.6) we omit. The criterion *forbids* adding them at this order — backup.

$c_{6,1}$ is only a two-order rolled value (3102^{+111}_{-124} , structural); it is not advanced as a prediction.

... and it is a single Borel function

Closure = null newest column, so this is not an N -dependent recipe:

$$\hat{B}(u) = \frac{6.463246}{(2-u)^{2.580247}} - \frac{26.662923}{(3-u)^{3.370370}} - \frac{0.060414}{(1+u)^{1.209877}} + 0.637084$$

- Positions and exponents: from the renormalization group; amplitudes from $c_{1,1}, \dots, c_{4,1}$.
- The one direction the data cannot fix — the $u = -1$ amplitude — is $\rho_* = -0.060414$. *Nothing is tuned.*

Summary

The structural result

- The triangular weight map has unit diagonal $\Rightarrow$ the three-weight 10% criterion is exactly $|\widehat{c}_{3,1} - c_{3,1}| \leq 1.4523$, $|\widehat{c}_{4,1} - c_{4,1}| \leq 2.1509$.
 - Known defects are positive, but a literature-range $c_{5,1}$ needs $\Delta_5 < 0$.
- $\Rightarrow$ **No defect-sign-preserving Adler-space transformation can pass the weight test *and* return a literature-range $c_{5,1}$.** The mixed record of the standard toolkit is forced.

The construction

- Declared Borel singularities with RG exponents, one parameter richer than the data $\Rightarrow$ forecast affine in a single closure amplitude ρ ; accuracy through order for all ρ .
- ρ fixed by one leave-one-order-out fold $\Rightarrow$ closed-form map. *Nothing tuned.*

What survived the hierarchy

- All six moment entries pass; six of nine alternatives (six of ten total structures) are rejected outright.
- Blind pole-mass forecast: 2.3–11.9%.
- Blind large- β_0 : complete tower $\leq 10\%$ for every $N \geq 6$ and $< 1.5\%$ for every $N \geq 8$; the minimal tower reaches persistent 10% accuracy only for $N \geq 8$.
- Physical order:

$$c_{5,1} = 248 \pm 30_{\text{struct}} \pm 62_{\text{meth}} = 248 \pm 70$$

$c_{6,1}$: rolled, ROLL status — not advanced as a prediction.

What did not

- Strictly recursive at physical order: misses $c_{4,1}$ by -29.9% ; closure drift $|\Delta\rho| = 0.0509$.
- At $N = 4$ in large- β_0 the minimal design is rank deficient — the method abstains.
- Every method fails the β -function control.

Conclusions

1. With four coefficients, **method choice dominates the answer**: admissible continuations span $[241.7, 304.7]$, diagnostics $[-21.7, 3327.8]$; validation is essential.
2. The failure of the standard toolkit is **a theorem, not bad luck**. Escaping it requires a transformation that can reverse the defect sign — physically, an **alternating ultraviolet renormalon**.
3. That class **already exists** (Beneke–Jamin 2008); we claim no new one. CVRC adds an **acceptance criterion**: affine in $\rho \Rightarrow$ the three-weight test selects an interval or *none*, so a declared Borel structure becomes **falsifiable**: six of nine alternatives are rejected. It gives $c_{5,1} = 248 \pm 70$, consistent with 277 ± 51 and the Beneke–Jamin forecast 280.
4. The claim is *weaker* than determination: a single closure value is consistent with everything testable. The -29.9% recursive miss measures the distance still to go.

Outlook for Belle II and STCF

- Higher-precision spectral data can justify a smaller accepted relative tolerance ϵ and a broader weight set. At fixed ϵ and fixed weights, they do not alter the unit-diagonal algebra.
- Weights with different $u = 2$ sensitivity are the most discriminating, and are exactly what a high-statistics spectral function enables.
- A genuine $c_{5,1}$ computation remains the decisive input.

Thank you.

Backup

Rolling the same model forward: $c_{5,1}$ to $c_{12,1}$

n	CVRC $c_{n,1}$	closure window	declared- α_3 range	$\pm(\text{rel.})$	status
5	247.98	[229.76, 266.20]	[235.26, 257.75]	7%	COND
6	3 102.1	[3 059.4, 3 144.9]	[3 021.3, 3 169.9]	2%	ROLL
7	13 003	[10 549, 15 458]	[10 952, 14 644]	19%	ROLL
8	3.7215×10^5	$[3.685, 3.758] \times 10^5$	$[3.638, 3.801] \times 10^5$	2%	ROLL
9	-2.5999×10^5	$[-7.736, 2.537] \times 10^5$	$[-6.555 \times 10^5, 5.7 \times 10^4]$	198%	INDET
10	8.7153×10^7	$[8.233, 9.198] \times 10^7$	$[8.701, 8.783] \times 10^7$	6%	ROLL
11	-9.0648×10^8	$[-1.098, -0.715] \times 10^9$	$[-1.030, -0.810] \times 10^9$	21%	ROLL
12	3.8699×10^{10}	$[3.506, 4.234] \times 10^{10}$	$[3.769, 4.018] \times 10^{10}$	9%	ROLL

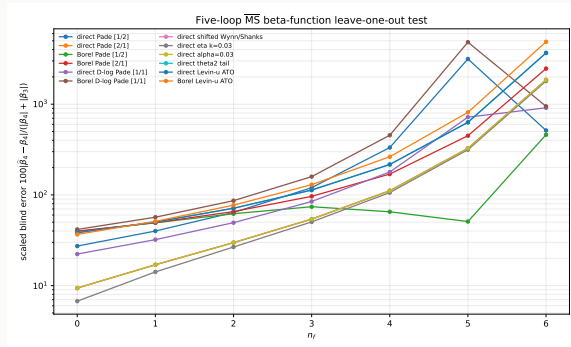
- **No controlled values are claimed** for $c_{7,1}$ - $c_{12,1}$. These are what the *single calibrated Borel model* returns.
- The roll is a *fixed point* of the closure rule: recomputing ρ at each step returns the same value to $\sim 10^{-17}$.
- Cost of each rolled order, measured *blind* in large- β_0 : about half a percent per order, on top of the one-order-ahead error.
- $c_{9,1}$ is flagged **sign-indeterminate**: its envelope contains zero.

A control that everything fails — and why we report it

Withhold the *known* five-loop $\overline{\text{MS}}$ β -function coefficient at $n_f = 3$ and forecast it from $\beta_0, \dots, \beta_3$.

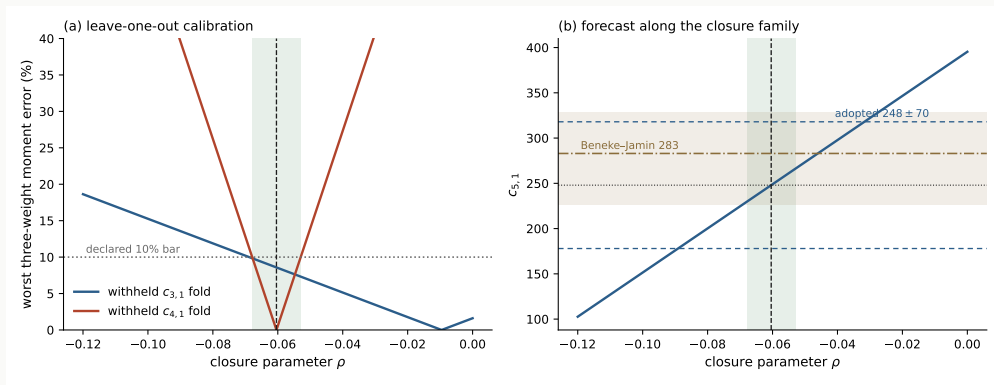
- Exact: $\beta_4 = 127.322$.
- **Every** tested continuation fails the declared ten-percent criterion.
- The least inaccurate member of the predeclared grid is still **65.8% high**.
- Direct and Borel Levin- u miss by 155.0% and 177.8%; their higher-order continuations change sign.

Accordingly β_5 and β_6 are retained only as *method diagnostics* and **no average is quoted**.



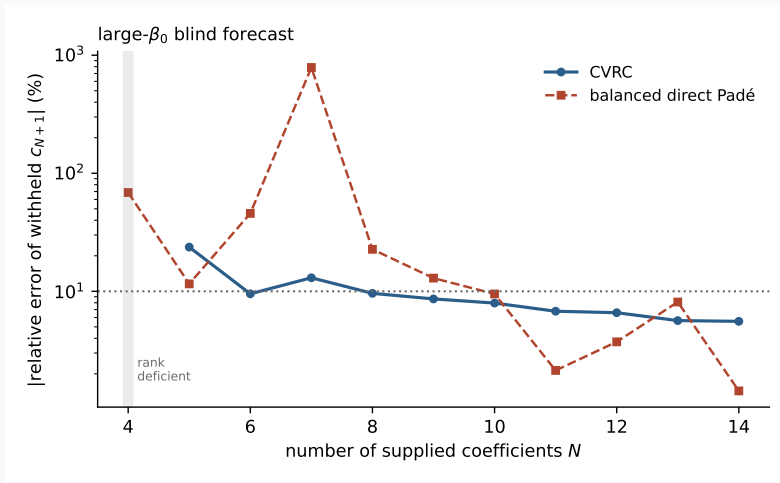
The β function has no renormalon interpretation, so this is a genuine negative control: it catches methods that “work” for reasons unrelated to the physics they claim to capture.

Backup — the closure family and its calibration



(a) Worst three-weight relative error on each leave-one-order-out fold as a function of ρ ; dashed line is $\rho_* = -0.060414$, shaded band the admissible window $[-0.0679, -0.0529]$. (b) $\hat{c}_{5,1}(\rho)$ along the same family, with the literature band 277 ± 51 , the geometric-ratio value 283, and the adopted result 248 ± 70 as the dashed blue pair — note how much wider the adopted band is than the vertical closure window, which is only its structural part.

Backup — blind large- β_0 forecast error



Strictly recursive error for the withheld large- β_0 Adler coefficient: CVRC-minimal versus balanced direct Padé on identical input. The dotted line is the ten-percent criterion; CVRC abstains at the shaded $N = 4$ point.

Backup — why we do *not* add the gluon condensate

The obvious objection: $c_1[\langle aG^2 \rangle]$ is known short-distance physics, it is worth **+22.6** on $c_{5,1}$, and adding it would move $248 \rightarrow 280$. Why not declare it?

Because it enters *only* inside the asymptotic bracket of the $u = 2$ cut, and that bracket is not a correction at the orders we actually have:

n	0	1	2	3	4	5	6
bracket	-2.90	-0.26	+0.27	+0.49	+0.61	+0.69	+0.74

Bold = the four orders that are known. It reaches 30% of unity only at $n = 6$, i.e. at $c_{7,1}$.

- At $n = 0$ it *reverses the sign* of the pure power and nearly triples it.
- Cut-corrected structure: admissible set **empty** (gap 0.206), closure drift 0.236 vs 0.051.
- Its blind postdiction of the known $c_{4,1}$: **244.5** against 49.076, an error of +398%.

The criterion rejects a *more complete* structure because the extra physics is not yet numerically operative at $N = 4$. Beneke–Jamin use their model from $c_{6,1}$ upward — exactly where the bracket is meaningful. No disagreement; a statement about the order.