

## Peptide COA Checklist

What to look for — and what to walk away from — before accepting a vial for laboratory use.

*A COA without a batch number is not a certificate of analysis — it is a portrait of a stranger who happens to share the molecule's name.*

### A — IDENTITY

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Confirms the molecule is the one on the label. Purity does not establish identity.

**Method named**

Mass spectrometry (MS/ESI-MS/MALDI-MS) is the standard. Method should be explicit, not implied.

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**Theoretical molecular weight shown**

Calculated from the target amino-acid sequence; allows you to check the arithmetic independently.

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**Observed mass shown and compared**

The instrument result next to

the  
theoretical  
value,  
within  
stated  
tolerance.  
"Confirmed"  
with  
no  
numbers  
is  
insufficient.

---

**Orthogonal data (if available)**

Amino-  
acid  
analysis,  
HPLC  
retention  
time  
vs  
reference  
standard,  
or  
sequence  
confirmation  
strengthens  
identity  
significantly.

**B – PURITY (HPLC)**

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Quantifies the main-peak fraction. Does not identify what the remaining percentage contains.

**Method is HPLC, peak-area %**

Reverse-  
phase  
HPLC  
(RP-  
HPLC)  
is  
standard.  
"Purity"  
without  
naming  
the  
technique  
is  
unverifiable.

---

**Result  $\geq 99\%$**

The  
field  
benchmark  
for  
research-  
grade  
synthetic  
peptides.

Below  
this,  
the  
impurity  
breakdown  
becomes  
decisive.

---

**Chromatogram or representative trace included**

The  
visual  
record  
of  
peaks.  
A  
single  
figure  
with  
no  
trace  
cannot  
be  
cross-  
checked.  
Dominant  
main  
peak  
+  
clean  
baseline  
is  
the  
target.

---

**Column and wavelength stated**

Conditions  
define  
what  
the  
method  
detects.  
Different  
wavelengths  
(UV  
210  
nm  
vs  
220  
nm)  
can  
shift  
apparent  
purity  
for  
some  
sequences.

---

**Run date on the chromatogram matches COA date**

Chromatograms  
from

an  
earlier  
or  
generic  
batch  
can  
be  
borrowed.  
Date  
consistency  
is a  
basic  
authenticity  
check.

## C — IMPURITY PROFILE

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The most informative — and most often skipped — section of a peptide COA.

**Individual impurities listed at  $\geq 0.1\%$**

The  
EMA  
2026  
synthetic-  
peptide  
guideline  
and  
ICH  
Q3A  
operate  
around  
a  
reporting  
threshold  
near  
0.1%.  
Impurities  
above  
this  
level  
should  
appear  
individually,  
not  
vanish  
into  
"other".<sup>12</sup>

---

**Nature of major impurities described where possible**

Deletion  
sequences,  
truncated  
chains,  
oxidised  
or  
deamidated  
variants,  
residual  
protecting  
groups

—  
these  
are  
the  
predictable  
cousins  
in  
synthetic  
peptide  
chemistry.

---

**No single impurity dominates unexpectedly**

A  
"99%  
pure"  
vial  
with  
a  
single  
0.9%  
impurity  
differs  
in  
risk  
profile  
from  
one  
with  
nine  
0.1%  
impurities.

---

**Residual solvents reported (if relevant)**

Particularly  
for  
peptides  
where  
the  
synthesis  
or  
lyophilisation  
process  
uses  
solvents  
that  
could  
carry  
over  
(e.g.  
acetonitrile,  
TFA).

**D — BATCH / LOT TRACEABILITY**

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The most diagnostic section for real-versus-recycled certificates.

**Lot / batch number present on COA**

A  
COA

without  
a  
lot  
number  
describes  
no  
specific  
batch.  
It  
is a  
generic  
document,  
not  
a  
release  
certificate.

---

**COA lot number matches the vial label**

Cross-  
check  
physically.  
A  
mismatch  
means  
the  
certificate  
was  
issued  
for  
different  
material.

---

**Analysis date is plausible**

A  
COA  
dated  
years  
before  
your  
order  
or  
after  
your  
delivery  
date  
is a  
red  
flag.  
Dates  
should  
precede  
or  
closely  
accompany  
dispatch.

---

**Net weight or quantity stated**

Relates  
the  
certificate

to  
the  
actual  
amount  
in  
the  
vial;  
makes  
concentration  
calculations  
traceable.

## E — WATER CONTENT & APPEARANCE

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### Water content measured and stated

Method:  
Karl  
Fischer  
titration  
or  
loss  
on  
drying.  
Lyophilised  
peptides  
are  
hygroscopic;  
5–  
10%  
water  
in  
the  
vial  
means  
5–  
10%  
less  
peptide  
per  
net-  
weight  
label.  
This  
directly  
affects  
reconstitution  
concentration  
calculations.

### Appearance described and consistent

Standard:  
white  
to  
off-  
white  
lyophilised  
powder.  
Discolouration  
or

clumping  
is a  
free  
integrity  
indicator  
worth  
checking  
against  
the  
COA  
description.

## F — ENDOTOXIN (CRITICAL FOR CELL-BASED & IN VITRO WORK)

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Invisible to HPLC. The most common silent confound in immunology and cell-culture models.

### Endotoxin test performed and named

Bacterial  
endotoxin  
test  
(BET/LAL  
—  
gel-  
clot,  
turbidimetric,  
or  
chromogenic)  
or  
recombinant  
Factor  
C  
(rFC).  
The  
method  
name  
must  
appear;  
"tested"  
alone  
is  
not  
sufficient.

---

### Result expressed in endotoxin units (EU/mg or EU/mL)

A  
specific  
measured  
value,  
referenced  
to  
USP  
<85>  
or  
Ph.  
Eur.  
2.6.14.  
Acceptable  
limits  
depend  
on

the  
intended  
research  
application.

---

**If not present: supplier can provide on request**

For  
applications  
where  
LPS  
contamination  
could  
confound  
results  
(cytokine  
assays,  
NF-  
κB  
reporters,  
macrophage  
models),  
treat  
an  
absent  
endotoxin  
section  
as  
incomplete  
QC  
data.

## **G – ISSUING LABORATORY**

---

**Laboratory name and location stated**

Anonymous  
"third-  
party  
testing"  
without  
identifying  
the  
lab  
is  
unverifiable.  
Condor  
Research  
uses  
an  
independent  
analytical  
laboratory  
in  
the  
Czech  
Republic  
(EU).

---

**Independent of the manufacturer**

Self-  
certified  
COAs  
(manufacturer  
tests  
own  
product  
without  
external  
verification)  
carry  
an  
inherent  
conflict  
of  
interest.

---

**Accreditation referenced (ISO 17025 or equivalent)**

Not  
universally  
present  
on  
peptide  
COAs  
but  
the  
strongest  
quality  
signal  
available.  
Even  
a  
named,  
non-  
accredited  
laboratory  
is  
better  
than  
no  
laboratory.

---

**Authorised signatory or stamp**

A  
real  
laboratory  
report  
carries  
a  
traceable  
signature  
or  
stamp.  
A  
document  
that  
could  
have  
been  
edited

in  
a  
word  
processor  
offers  
weaker  
assurance.

## RED FLAGS — WALK AWAY

WHAT YOU SEE	WHAT IT MEANS	VERDICT
No batch / lot number	Certificate describes no specific production run	Reject
Lot number does not match vial label	Certificate was issued for a different (or fictional) batch	Reject
Purity stated; no method named	Unverifiable claim — what instrument, what conditions?	Insufficient
"Confirmed" identity with no mass data	Identity not actually demonstrated; just asserted	Insufficient
No chromatogram or trace	Peak profile cannot be cross-checked; figure could be fabricated	Insufficient
Impurity percentage unaccounted for	What is the other 1–5%? If unlisted, it is uncharacterised	Insufficient
No laboratory named	No independent verification; self-certification at best	Insufficient
COA dated after your delivery	Temporal impossibility — document was backdated or generic	Reject
No water content or appearance data	Incomplete specification per ICH Q6A; affects concentration accuracy	Incomplete

## QUICK REFERENCE — WHAT GOOD LOOKS LIKE

COA SECTION	METHOD	MINIMUM ACCEPTABLE
<b>Identity</b>	Mass spectrometry (ESI-MS / MALDI)	Observed mass vs theoretical, both values shown, within stated tolerance
<b>Purity</b>	RP-HPLC, peak-area %	≥99%, chromatogram included, run date consistent
<b>Impurities</b>	HPLC profile vs ICH / EMA thresholds	Individual impurities listed at ≥0.1%; not hidden in "other" <sup>12</sup>
<b>Water content</b>	Karl Fischer / loss on drying	Stated value; relevant for concentration calculations
<b>Appearance</b>	Visual / physical description	Matches vial contents; white lyophilised powder is standard
<b>Batch / Lot</b>	Traceability	Number present on COA, matches vial; date plausible
<b>Endotoxin</b>	BET/LAL or rFC (USP <85> / Ph. Eur. 2.6.14)	Named method, EU/mg value stated; critical for cell-based assays <sup>34</sup>
<b>Laboratory</b>	Independent, named, EU preferred	Lab identified; ideally ISO 17025; not the manufacturer itself

## FREQUENTLY ASKED QUESTIONS

### What purity should a research peptide COA show?

The field benchmark is ≥99% by HPLC peak-area percentage. Below that, the fraction and identity of what accounts for the remainder becomes the deciding question — a 97% purity with a fully characterised 3% impurity profile may be more informative than a 99% figure with no breakdown.

### Can a COA be faked?

Yes. Red flags include no batch/lot number, no named analytical method, no observed mass next to theoretical, and a purity figure with no chromatogram or impurity breakdown. Independent re-testing by an accredited laboratory is the only absolute defence. Ask your supplier if they support third-party re-analysis.

### Why is the batch number so important?

A COA describes one specific manufacturing lot. Two batches of the same peptide can differ in purity and impurity profile even from the same manufacturer. Without a batch number that matches the one on your vial, the certificate describes a different — or entirely hypothetical — batch.<sup>5</sup>

### Does HPLC purity cover endotoxin contamination?

No. HPLC is essentially blind to lipopolysaccharide (LPS) endotoxins. A separate bacterial endotoxin test (BET/LAL or rFC) is required. High HPLC purity with no endotoxin data leaves a major confound completely uncharacterised for any cell-based or immunological assay.

### What does "independent laboratory" mean?

The testing laboratory is separate from the peptide manufacturer — ideally accredited to ISO 17025 — so the result is not self-certified. Condor Research uses an independent EU analytical laboratory based in the Czech Republic. COAs are issued per batch, not per product line.

### How do I read the mass spec result if I am not a chemist?

Look for two numbers: theoretical MW (calculated from the sequence) and observed MW (what the instrument measured). If they match within  $\pm 1-2$  Da (or the stated tolerance), the backbone is the right sequence. If only one number appears, or the result is written as "confirmed" without showing the observed value, the identity check cannot be independently verified.

## FURTHER READING — CONDOR RESEARCH

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- How to Read a Certificate of Analysis (COA) for a Research Peptide — full editorial guide
- What "99% Pure" on a Peptide COA Really Means: HPLC, Mass Spec and the Limits of a Purity Number
- Endotoxins and Sterility: The COA Section Almost Nobody Reads
- How to Choose a Research Peptide Supplier in Europe (2026)
- Why Peptide Experiments Fail: Reproducibility and the Reagent Problem

**How to use this checklist.** Work through sections A–G against the COA supplied with each batch. Keep a signed copy with your laboratory notebook entry for the lot. If an item cannot be confirmed, contact the supplier before use — a credible supplier will be able to provide documentation or direct you to the issuing laboratory.

## REFERENCES

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2. ICH Harmonised Guideline. *Impurities in New Drug Substances Q3A(R2)*. International Council for Harmonisation, 2006. [ich.org](https://www.ich.org)
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5. ICH Harmonised Guideline. *Specifications: Test Procedures and Acceptance Criteria for New Drug Substances and New Drug Products Q6A*. 1999. [ich.org](https://www.ich.org)

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