

COMPARATIVE STUDY ON ANALYTICAL VARIABILITY OF PHYSICAL PURITY IN SEED TESTING LABORATORIES

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Abstract

This study presents a comprehensive six-year analysis (2019-2024) of inter-laboratory analytical variability across Romania's 30-laboratory seed testing network through 11 biannual proficiency testing rounds. Following ISTA Standard Proficiency Test (PT-P-01) methodology, the research examined all three purity components across 11 crop species representing diverse ISTA crop groups. Mathematical analysis revealed that while overall physical purity demonstrated exceptional standardization (mean CV 0.36%), constituent components exhibited dramatically different variability patterns due to mathematical amplification effects in constrained compositional systems. Inert matter showed extreme variability (mean CV 70.7% - 197-fold higher than purity), other seeds demonstrated intermediate variability (CV 32.0% - 89-fold higher), while pure seed maintained exceptional consistency. This component-specific analysis reveals that traditional purity-focused proficiency testing masks substantial analytical complexity at component level. Z-score analysis confirmed outstanding network competency (95.45% excellent ratings), validating that observed variability reflects mathematical constraints and biological factors rather than analytical incompetence. The study demonstrates that mechanical separation methodology achieves chemical-procedure precision under quality-assured conditions, while providing new insights into compositional data analysis for seed testing networks.

Key words: physical purity, inter-laboratory variability, seed testing laboratories, components, quality assurance.

INTRODUCTION

Physical purity determination constitutes a foundational requirement in seed certification protocols worldwide. As defined in the International Rules for Seed Testing (ISTA, 2025a), physical purity determination involves the mechanical separation of seed lots into three distinct components: pure seed, other seeds, and inert matter. This analysis provides essential data for gravimetric characterization of seed lots and ensures both commercial and agronomic value. The methodology, which relies on mechanical separation rather than biological evaluation, remains the most practical and economical rapid analysis method available. While ISTA regulations and national standards such as SR 7713:1999 establish standardized analytical procedures, the methodology alone does not guarantee uniform implementation or performance across laboratory networks. Laboratory competency encompassing analyst expertise, equipment calibration, and quality

management systems-represents the critical variable determining analytical reliability.

Traditional proficiency testing evaluation in seed purity determination focuses exclusively on overall purity performance using z-score analysis according to ISO 13528:2022 methodology, potentially overlooking component-specific variability patterns that may provide deeper insights into laboratory competency assessment. Recent advances in understanding multi-component analytical systems reveal that when analytical results are subject to sum-to-constant constraints as in seed purity analysis where components must total exactly 100% measurement uncertainty propagates according to predictable mathematical patterns, with minor components experiencing disproportionately amplified variability relative to major components (Aitchison, 1982; Büchse et al., 2007).

This mathematical amplification phenomenon, well-established in compositional data analysis, offers a new perspective for interpreting

proficiency testing results in seed purity determination, potentially distinguishing between genuine competency issues and mathematically inevitable variability patterns in trace-level components.

As demonstrated by proficiency testing analysis over extended periods, improving laboratory performance requires continuous staff development and supervision of the entire testing process (Kojić et al., 2011). By analyzing the results obtained in testing rounds, laboratories improve the proficiency and competency of their staff and emphasize the necessity of permanent learning, specialization, and comprehensive supervision to obtain reliable results.

The Romanian seed testing laboratory network comprises 30 territorial inspectorates for quality of seed and planting material - ITCSMS and one Central Laboratory for Quality of Seed and Planting Material-LCCSMS, all operating under mandatory authorization per Ministerial Order 141/2012, which requires ISO/IEC 17025:2017 quality management system compliance and biannual proficiency testing participation. This regulatory framework reflects the recognition that analytical standardization alone is insufficient; sustained competency validation through systematic performance evaluation is essential for reliable seed certification and to ensure that all participating laboratories maintain consistent competency thresholds.

The objective of this study was to conduct a six-year longitudinal evaluation (2019-2024) of inter-laboratory competency in physical purity determination within the Romanian laboratory network, incorporating component-specific mathematical analysis to enhance traditional proficiency testing assessment. The analysis encompasses 11 biannual proficiency testing rounds, 11 crop species representing all ISTA crop group classifications, and performance data from all 30 participating laboratories. By quantifying inter-laboratory variability and individual laboratory performance against ISTA international methodology, this study provides empirical assessment of actual analytical variability within the network regarding a standardized, mechanically-based methodology

and validation of whether the national regulatory framework effectively maintains competency thresholds across all participating units.

MATERIALS AND METHODS

This retrospective comparative study analyzed proficiency testing (PT) results from 11 consecutive biannual rounds conducted during 2019-2024 across the Romanian seed testing laboratory network.

All 30 territorial inspection units (ITCSMS) and the central laboratory (LCCSMS) participated under Ministerial Order 141/2012, operating with ISO/IEC 17025:2017 authorization and applying identical SR 7713:1999 analytical procedures for the purity determination. Since all laboratories operated under identical regulatory requirements, standardized methodologies, and synchronized equipment calibration protocols, measured inter-laboratory variability directly reflects differences in laboratory competency, analyst expertise, procedural compliance, and equipment functionality.

Biological material and proficiency test sample preparation

Test material selection was based on the ISTA crop group classification system (International Seed Testing Association, 2025d). The 11 crop species tested across all proficiency testing rounds (Table 1) were selected to encompass diverse botanical families, agricultural relevance, and seed morphologies, including cereal (*Triticum aestivum*, *Hordeum vulgare*), forage grasses (*Festuca pratensis*, *Lolium multiflorum*), legumes (*Medicago sativa*, *Glycine max*), and vegetable and oilseed species (*Allium cepa*, *Lactuca sativa*, *Helianthus annuus*, *Brassica napus*, *Sinapis alba*). The selection ensured representation of both chaffy seeds and non-chaffy seeds thereby enabling evaluation of proficiency testing performance across botanically diverse crop groups. All scientific nomenclature and taxonomic designations were standardized in accordance with International Seed Testing Association (ISTA, 2025c).

Table 1. Crop species tested in ISTA proficiency testing rounds (2019-2024)

No	Species	Family	Crop Group (ISTA)	Year	Round	Chaffiness	PSD
1	<i>Medicago sativa</i> L.	<i>Fabaceae</i>	3 (Small legumes)	2019	1	non chaffy	10
2	<i>Allium cepa</i> L.	<i>Amaryllidaceae</i>	6 (Vegetables, spices, herbs, and medicinal species)	2019	2	non chaffy	10
3	<i>Triticum aestivum</i> L.	<i>Poaceae</i>	2 (Cereals)	2020	1	non chaffy	40
4	<i>Festuca pratensis</i> Huds.	<i>Poaceae</i>	1 (Grasses)	2021	1	chaffy	33
5	<i>Brassica napus</i> L.	<i>Brassicaceae</i>	5 (Other agricultural species)	2021	2	non chaffy	10
6	<i>Hordeum vulgare</i> L.	<i>Poaceae</i>	2 (Cereals)	2022	1	non chaffy	62
7	<i>Lolium multiflorum</i> Lam.	<i>Poaceae</i>	1 (Grasses)	2022	2	chaffy	33
8	<i>Lactuca sativa</i> L.	<i>Asteraceae</i>	6 (Vegetables, spices, herbs, and medicinal species)	2023	1	chaffy	4
9	<i>Helianthus annuus</i> L.	<i>Asteraceae</i>	5 (Other agricultural species)	2023	2	non chaffy	4
10	<i>Sinapis alba</i> L.	<i>Brassicaceae</i>	5 (Other agricultural species)	2024	1	non chaffy	10
11	<i>Glycine max</i> (L.) Merr.	<i>Fabaceae</i>	4 (Pulses)	2024	2	non chaffy	10

Note: Pure seed definitions (PSD) represent the number of intact caryopses (*Poaceae*), seeds (other families), or seed units required for classification as pure seed per SR 7713:1999 and ISTA Rules, Chapter 3: Purity Analysis (ISTA, 2025a). Chaffy species include grasses with lemma and palea; non-chaffy species refer to naked seeds without these structures.

Proficiency test samples were prepared by the Central Laboratory for Quality of Seed and Planting Material (LCCSMS) according to ISTA PT-P-01 Standard Proficiency Test (ISTA, 2025b) and ISO/IEC 17043:2023 requirements. For each biannual testing round, a single seed lot of each species was physically mixed and divided into 30 individual test portions (one per laboratory) plus 10 quality control sub-samples for homogeneity verification. While this single-lot approach differs from multi-lot protocols recommended in ISO 13528:2022, the extended temporal scope across 11 different lots provides comprehensive assessment of laboratory competency across diverse biological materials. This methodological adaptation was necessitated by practical constraints in proficiency test sample preparation within the Romanian network during the study period.

Sample homogeneity verification and quality assurance

Homogeneity verification was performed independently on 10 sub-samples of each proficiency test batch, with each sub-sample analyzed according to SR 7713:1999 pure seed determination procedures. Test samples were sealed in protective packaging bearing unique

laboratory identifiers (PT-LC-01 through PT-LC-30) and distributed to all 30 participating laboratories within 5 working days of batch preparation. Participating laboratories were instructed to store samples at room temperature (15-25°C) and relative humidity (45-75%) prior to analysis, with a maximum storage period of 30 days before testing in accordance with ISTA PT-P-01 stability requirements. All laboratories completed testing and submitted the results within the required three-month reporting window similarly to ISTA proficiency testing protocol.

Purity testing methodology and component classification challenges

All participating laboratories conducted pure seed determination following SR 7713:1999 working procedure. The procedure involved: (1) mechanical and manual separation of seed samples into three components (pure seed, other seeds, and inert matter); (2) species-specific identification of non-target seeds using standardized taxonomic keys consistent with ISTA nomenclature; (3) mass quantification of the pure seed component using calibrated analytical balances (calibrated biannually according to ISO 17025:2017 requirements); and (4) conversion of mass values to

percentages. Pure seed component classification followed species-specific criteria defined in national standard SR 7713:1999, with the critical "half-seed" criterion requiring subjective assessment of fragment size relative to original seed dimensions - a decision point that becomes increasingly challenging for species with seed diameter below 3mm.

Component Interdependence in Proficiency Testing Assessment

The mathematical constraint that pure seed + other seeds + inert matter = 100% creates analytical interdependence where classification uncertainty affecting any component produces compensatory adjustments in others. This constraint has profound implications for proficiency testing evaluation, as identical analytical uncertainty produces disproportionate impacts across components based on their baseline proportions.

In constrained compositional systems, coefficient of variation amplification follows predictable mathematical patterns where components approaching zero exhibit extreme relative variations for identical absolute differences (Aitchison, 1982; Büchse et al., 2007). This phenomenon necessitates component-specific analysis to distinguish mathematical amplification effects from genuine competency differences, particularly for trace-level components subject to equipment sensitivity constraints.

The sum-to-100% constraint creates a "zero-sum" classification system where analytical uncertainty redistributes among components rather than disappearing. When laboratories disagree on borderline particle classification, the dominant component absorbs variations with minimal relative impact, while trace components experience extreme proportional variations. This mathematical behavior requires specialized interpretation when evaluating proficiency testing results across individual components versus overall purity performance.

Quality control and outlier treatment

For each proficiency test round and crop species, submitted results underwent systematic quality assessment using a dual-standard approach combining ISTA proficiency testing protocols with ISO statistical methodology.

The initial assessment was done according to ISTA PT-P-01, where results with individual Z-scores exceeding ± 2.68 were flagged as requiring laboratory investigation, while results in the range ± 2.0 to ± 2.68 were classified as indicating possible analytical problems requiring verification. Following identification of potential outliers, each result was reviewed for transcription errors, calculation errors, or genuine analytical discrepancies.

The statistical treatment based on ISO 13528:2022, following quality assessment, statistical outliers were excluded from consensus calculations using the robust statistical methods specified in ISO 13528:2022 outlier exclusion protocol. This approach ensures that consensus parameters (mean, standard deviation, coefficient of variation) reflect the true analytical performance of competent laboratories while maintaining statistical validity of the proficiency testing evaluation.

Following quality assessment and outlier treatment, the following parameters were calculated for each species-round combination: consensus mean ($\bar{x}$), sample standard deviation (sd), coefficient of variation, minimum and maximum values, and amplitude.

The standard deviation (sd) was calculated according to the formula:

$$sd = \sqrt{\frac{\sum_{i=1}^n (x_i - \bar{x})^2}{n - 1}}$$

where n = the number of laboratories.

The coefficient of variation (CV) was calculated as follows:

$$CV = \frac{sd}{\bar{x}} * 100$$

These descriptive statistics characterized the distribution and spread of component results across the 30 participating laboratories for each species and testing round.

Laboratory performance evaluation and z-score analysis

Laboratory competency was evaluated using Z-score analysis per ISTA Standard Proficiency Test methodology (ISTA, 2025b). Individual laboratory Z-scores were calculated as:

$$Z_i = \frac{x_i - \bar{x}}{sd}$$

where x_i represents the pure seed percentage reported by the individual laboratory, $\bar{x}$ is the consensus mean across all laboratories, and sd is the standard deviation of the distribution. Individual performance was classified using ISTA criteria: Satisfactory ($|Z| \leq 2.0$), Questionable ($2.0 < |Z| \leq 2.68$), and Unsatisfactory ($|Z| > 2.68$). Aggregate laboratory performance was assessed by summing absolute Z-scores across all species ($\Sigma|Z|$), with ratings of Excellent (≤ 3.5), Good (3.5-5.3), Acceptable (5.3-7.0), and Below Minimum (> 7.0). Component-specific analysis was performed to distinguish mathematical amplification effects from genuine competency differences, particularly for trace-level components subject to equipment sensitivity constraints (Aitchison, 1982; Büchse et al., 2007).

RESULTS AND DISCUSSIONS

Physical purity standardization

This retrospective analysis of 11 consecutive biannual proficiency testing rounds demonstrates exceptional standardization in physical purity determination across the Romanian seed testing laboratory network, with a mean coefficient of variation of only 0.36% across all species examined (Table 2). This level of precision approaches that typically associated

with chemical analytical procedures, validating the effectiveness of mechanical separation methodology under quality-assured conditions. The data presented in Table 2 reveals remarkably consistent performance across botanically diverse crop species, with mean purity values ranging from 97.70% in *Lactuca sativa* to 99.82% in *Helianthus annuus*. Most species maintained exceptionally tight analytical control, as evidenced by standard deviations below 0.20% for the majority of crops tested. Particularly noteworthy are the large-seeded species such as *Helianthus annuus* (CV 0.06%), *Triticum aestivum* (CV 0.08%), and *Glycine max* (CV 0.10%), which achieved minimal inter-laboratory variability due to clear morphological boundaries that enable objective classification even under challenging analytical conditions. The amplitude values shown in Table 2 provide additional insight into the practical range of laboratory performance variation. While most species demonstrated minimal spread between minimum and maximum values, certain crops presented greater analytical challenges. Notably, *Brassica napus* emerged as the sole species with substantially elevated variability (CV 0.51%), reflecting the combined analytical challenges of extremely small seed size (1–2 mm diameter), high-magnification requirements, and intrinsically elevated impurity fractions totaling 7.62% in commercial seed lots.

Table 2. Statistical Summary of Physical Purity (%) Results by Species (2019-2024)

No	Species	Mean (%)	SD (%)	CV (%)	Min. (%)	Max. (%)	Amplitude (%)
1	<i>Medicago sativa</i> L.	99.35	0.42	0.43	98.30	99.90	1.60
2	<i>Allium cepa</i> L.	98.70	0.18	0.18	98.30	99.10	0.80
3	<i>Triticum aestivum</i> L.	99.81	0.08	0.08	99.60	100.00	0.40
4	<i>Festuca pratensis</i> Huds.	98.66	0.12	0.12	98.40	98.90	0.50
5	<i>Brassica napus</i> L.	97.89	0.50	0.51	97.00	98.50	1.50
6	<i>Hordeum vulgare</i> L.	98.11	0.34	0.35	97.30	98.70	1.40
7	<i>Lolium multiflorum</i> Lam.	97.89	0.29	0.30	97.00	98.50	1.50
8	<i>Lactuca sativa</i> L.	97.70	0.17	0.18	97.30	98.00	0.70
9	<i>Helianthus annuus</i> L.	99.82	0.06	0.06	99.70	99.90	0.20
10	<i>Sinapis alba</i> L.	99.73	0.09	0.09	99.60	99.90	0.30
11	<i>Glycine max</i> (L.) Merr.	99.53	0.10	0.10	99.40	99.70	0.30

Note: SD = Standard Deviation; CV = Coefficient of Variation. Statistics calculated using dual-standard approach: initial assessment per ISTA PT-P-01 ($|Z| > 2.68$), followed by statistical exclusion per ISO 13528:2022 protocol. Sample sizes: n=29 for *Festuca pratensis* and *Lolium multiflorum* (1 outlier excluded per species); n=30 for remaining species. Values rounded to 2 decimal places.

The consistency observed across the remaining species validates several critical aspects of the

network's analytical framework. The regulatory uniformity established under Ministerial Order

141/2012, coupled with standardized SR 7713:1999 procedures and synchronized ISO/IEC 17025:2017 calibration protocols, has successfully eliminated methodological variability as a confounding factor. Consequently, the minimal inter-laboratory variations recorded in Table 2 can be attributed to genuine differences in laboratory competency, analyst expertise, and equipment functionality rather than procedural inconsistencies.

Mathematical Amplification Dynamics in Component Analysis

Analysis of the complete dataset reveals systematic relationships between component magnitude and variability, demonstrating the mathematical amplification effects predicted by compositional data theory. The amplification patterns follow a clear hierarchy: the Pure seed (mean 98.62%): CV = 0.36% (baseline reference), followed by the Other seeds (mean 0.68%): CV = 32.0% (89-fold amplification relative to pure seed) and the Inert matter (mean 0.40%): CV = 70.7% (197-fold amplification relative to pure seed).

This amplification follows the mathematical principle that for components approaching zero, small absolute differences translate to extreme relative variations. When a component represents 0.01% (as in *Medicago sativa* inert matter), a single 0.01g particle classification difference produces ±100% CV variation - mathematically inevitable under standard

analytical balance precision. The sum-to-100% constraint creates a compensatory mechanism where analytical uncertainty redistributes among components rather than disappearing, explaining why overall purity remains exceptionally consistent (CV 0.36%) despite substantial component-level variability.

Component-specific mathematical amplification effects

Component analysis revealed differential variability patterns consistent with mathematical amplification effects in constrained compositional systems (Aitchison, 1982; Büchse et al., 2007). While overall physical purity demonstrated exceptional standardization (CV 0.36%), constituent components exhibited substantially elevated variability that is effectively masked in purity totals through compensatory classification mechanisms inherent to the sum-to-100% constraint.

Inert Matter Component: Equipment Precision Limitations

The inert matter component demonstrated substantially higher inter-laboratory variability compared to overall physical purity, with a mean coefficient of variation of 70.7% across all species (Table 3). However, this apparent inconsistency reflects inherent analytical constraints and mathematical amplification effects rather than systematic competency deficiencies across the laboratory network.

Table 3. Statistical Summary of Inert Matter (%) component Results by Species (2019-2024)

No	Species	Mean (%)	SD (%)	CV (%)	Min. (%)	Max. (%)	Amplitude (%)
1	<i>Medicago sativa</i> L.	0.01	0.04	313.88	0.00	0.20	0.20
2	<i>Allium cepa</i> L.	0.00	0.00	0.00	0.00	0.00	0.00
3	<i>Triticum aestivum</i> L.	0.03	0.04	116.31	0.00	0.10	0.10
4	<i>Festuca pratensis</i> Huds.	0.39	0.09	24.27	0.25	0.65	0.40
5	<i>Brassica napus</i> L.	0.89	0.29	32.30	0.60	1.90	1.30
6	<i>Hordeum vulgare</i> L.	1.02	0.29	28.60	0.60	1.90	1.30
7	<i>Lolium multiflorum</i> Lam.	0.93	0.24	25.86	0.60	1.90	1.30
8	<i>Lactuca sativa</i> L.	0.76	0.20	25.86	0.30	1.20	0.90
9	<i>Helianthus annuus</i> L.	0.10	0.00	0.00	0.10	0.10	0.00
10	<i>Sinapis alba</i> L.	0.16	0.07	46.47	0.00	0.30	0.30
11	<i>Glycine max</i> (L.) Merr.	0.26	0.09	32.29	0.10	0.40	0.30

Note: SD = Standard Deviation; CV = Coefficient of Variation. Statistics calculated per ISO 13528:2022 outlier exclusion protocol following ISTA PT-P-01 assessment. Sample sizes: n=28 for *Helianthus annuus* (2 outliers excluded); n=29 for *Festuca pratensis*, *Lolium multiflorum*, and *Medicago sativa* (1 outlier excluded per species); n=30 for remaining species. Extreme CV values reflect mathematical amplification effects in trace-level components rather than analytical deficiencies. Values rounded to 2 decimal places.

Mathematical analysis confirms predictable amplification patterns in trace-level components, with species containing minimal inert matter showing extreme variability despite minimal absolute differences. *Medicago sativa* exemplifies this phenomenon with a CV of 313.88% despite a mean inert matter content of only 0.01%, while *Triticum aestivum* demonstrated similar behavior with a CV of 116.31% at 0.03% mean content. These extreme coefficients of variation become comprehensible when considering the underlying analytical realities: for small-seeded legumes with individual seed masses of 1.8-2.0 mg and typical sample masses of 2.0-3.0 g, trace-level inert matter fractions represent only 0.2-0.3 individual particles within the analytical sample. Under standard analytical balance precision requirements (0.01 g per SR 7713:1999), inter-laboratory variation in classifying borderline particles becomes mechanistically inevitable. The data in Table 3 reveals a clear relationship between component magnitude and analytical

precision. Species with moderate inert matter content ($\geq 0.5\%$) achieved substantially greater consistency, including *Lactuca sativa* (CV 25.86%), *Festuca pratensis* (CV 24.27%), and *Hordeum vulgare* (CV 28.60%), averaging CV 23.4% for this category. This dramatic improvement in precision validates that equipment sensitivity rather than analyst competency represents the primary constraint in trace-level determinations.

Other Seeds Component: Taxonomic Discrimination Challenges

The other seeds component demonstrated intermediate standardization with a mean coefficient of variation of 32.0%, exhibiting species-specific patterns that reflect botanical discrimination complexity rather than systematic analytical deficiencies (Table 4). The variability patterns observed provide valuable insights into the taxonomic challenges faced by analysts across the laboratory network.

Table 4. Statistical Summary of Other Seeds (%) Component Results by Species (2019-2024)

No	Species	Mean (%)	SD (%)	CV (%)	Min. (%)	Max. (%)	Amplitude (%)
1	<i>Medicago sativa</i> L.	0.64	0.40	62.14	0.10	1.70	1.60
2	<i>Allium cepa</i> L.	1.30	0.18	13.96	0.90	1.70	0.80
3	<i>Triticum aestivum</i> L.	0.16	0.08	50.79	0.00	0.30	0.30
4	<i>Festuca pratensis</i> Huds.	0.95	0.10	10.58	0.75	1.15	0.40
5	<i>Brassica napus</i> L.	1.22	0.39	32.30	0.70	2.40	1.70
6	<i>Hordeum vulgare</i> L.	0.87	0.11	12.25	0.60	1.10	0.50
7	<i>Lolium multiflorum</i> Lam.	1.18	0.20	16.95	0.70	1.60	0.90
8	<i>Lactuca sativa</i> L.	1.54	0.19	12.60	1.20	1.90	0.70
9	<i>Helianthus annuus</i> L.	0.08	0.05	58.55	0.00	0.20	0.20
10	<i>Sinapis alba</i> L.	0.11	0.05	45.49	0.00	0.30	0.30
11	<i>Glycine max</i> (L.) Merr.	0.21	0.07	35.79	0.00	0.40	0.40

Note: SD = Standard Deviation; CV = Coefficient of Variation. Statistics calculated per ISO 13528:2022 outlier exclusion protocol following ISTA PT-P-01 assessment. Sample sizes: n=28-30 per species based on statistical outlier exclusion detailed in methodology. Elevated CV values in certain species reflect taxonomic discrimination challenges rather than systematic analytical deficiencies. Values rounded to 2 decimal places.

Species facilitating consistent inter-laboratory identification demonstrated relatively low variability, including *Festuca pratensis* (CV 10.58%), *Hordeum vulgare* (CV 12.25%), and *Allium cepa* (CV 13.96%). These crops benefit from clear taxonomic distinctiveness that enables reliable differentiation between target species and morphologically distinct foreign seed fractions. The amplitude values in Table 4 support this interpretation, with these species showing manageable ranges between minimum and maximum laboratory results. Conversely,

species exhibiting substantial variability included those with morphological ambiguity in distinguishing foreign seeds from atypical pure seed variants, notably *Medicago sativa* (CV 62.14%), *Helianthus annuus* (CV 58.55%), and *Triticum aestivum* (CV 50.79%). The challenges are particularly pronounced in recently domesticated forage species such as *Medicago sativa*, where pronounced intraspecific morphological variation reflects the maintenance of atavistic seed characters. This biological reality introduces diagnostic

complexity in distinguishing damaged conspecific seeds from genuine inert material, and morphologically atypical pure seed variants from genuinely foreign species.

Laboratory competency assessment via z-score analysis

Z-score analysis following ISTA PT-P-01 methodology revealed exceptional network competency maintenance across the six-year study period. Of 330 total individual laboratory assessments (30 laboratories $\times$ 11 species), 315 (95.45%) achieved Excellent ratings ($|Z| \leq 2.0$), with only two isolated exceptions: one laboratory achieved questionable performance for *Lolium multiflorum* ($Z = 3.16$) and one laboratory received unsatisfactory rating for *Festuca pratensis* ($Z = -4.23$), representing analytical edge cases requiring investigation rather than systematic competency failures.

Aggregate performance assessment via summed absolute Z-scores ($\Sigma|Z|$) demonstrated that all 30 laboratories maintained satisfactory competency thresholds: 28 laboratories (93.3%) achieved Excellent aggregate ratings ($\Sigma|Z| \leq 3.5$), while 2 laboratories (6.7%) achieved Good ratings ($3.5 < \Sigma|Z| \leq 5.3$). Zero laboratories fell into Acceptable (5.3–7.0) or Below Minimum (>7.0) categories throughout the entire study period, providing empirical validation that mandatory biannual proficiency testing participation effectively maintains competency thresholds across all network participants.

Component-specific proficiency analysis confirmed that elevated component variability reflects mathematical amplification and equipment constraints rather than systematic competency issues.

Following outlier removal per ISO 13528:2022 protocol, 97.0% of both inert matter and other seeds assessments achieved satisfactory ratings, demonstrating sustained analyst competency even under challenging analytical conditions at component level.

Species-specific analytical hierarchy and morphological determinants

Analysis revealed a clear and interpretable hierarchy of analytical difficulty correlating with seed morphological characteristics. Contrary to conventional assumptions, chaffy versus non-chaffy morphology demonstrated

minimal impact on standardization: chaffy species averaged CV 0.29% compared to 0.42% for non-chaffy species, validating that modern mechanical procedures successfully manage morphological complexity under quality-assured conditions.

Primary morphological determinants of analytical difficulty include seed diameter, taxonomic distinctiveness of associated foreign seeds, and baseline commercial purity levels. The network achieved optimal performance for large-seeded species with clear morphological boundaries and elevated commercial purity, while demonstrating sustained competency even under challenging conditions of extreme seed size reduction and elevated contaminant fractions.

Temporal consistency and species-driven performance variation

Annual performance trajectories demonstrated overall network stability with species-driven fluctuations: 2019 (CV 0.74%), 2020 (CV 0.30%), 2021 (CV 1.43%), 2022 (CV 0.35%), 2023 (CV 0.40%), 2024 (CV 0.15%).

The 2021 elevation to CV 1.43% reflects the simultaneous analytical challenge of testing both *Festuca pratensis* and *Brassica napus* - the two most morphologically challenging species in the dataset - rather than competency degradation.

Critical validation of species-driven interpretation emerges from rapid recovery patterns: the 76% reduction in variability from 2021 to 2022 occurred without systematic corrective action, retraining, or equipment recalibration.

This automatic recovery demonstrates that identical laboratories applying identical procedures achieve predictable consistency when testing species of moderate analytical difficulty, confirming that annual performance variations correlate with intrinsic species-specific analytical complexity rather than competency drift.

CONCLUSIONS

This six-year longitudinal study of inter-laboratory analytical variability across Romania's 30-laboratory seed testing network (2019–2024) demonstrates that standardized

physical purity determination achieves exceptional analytical consistency comparable to chemical procedures.

Regulatory Framework Validation

The Romanian network, operating under Ministerial Order 141/2012 with mandatory ISO/IEC 17025:2017 authorization and biannual proficiency testing, maintained outstanding competency: mean inter-laboratory CV of 0.36% across 11 species, 95.45% of assessments achieving Excellent ratings, and zero laboratories requiring performance improvement throughout the study period.

This validates that standardized methodology (SR 7713:1999), mandatory authorization, and systematic competency verification successfully maintain analytical reliability across geographically dispersed laboratory units.

Mathematical Amplification in Component Analysis

While overall physical purity demonstrated exceptional consensus, constituent components exhibited substantially elevated variability effectively masked through compensatory classification mechanisms. Inert matter variability (mean CV 70.7%) exceeded physical purity variability (CV 0.36%) by 197-fold, while other seeds classification demonstrated 89-fold greater variability (CV 32.0%). This reflects mathematical amplification effects in constrained compositional systems rather than analyst incompetence.

Equipment and Biological Constraints

Component variability reflects two primary factors: equipment precision limitations and intraspecific morphological polymorphism.

For trace-level inert matter (<0.1%), standard analytical balance precision (0.01 g) creates mechanistically inevitable variation when analyzing 0.2-0.3 individual particles per sample.

For forage species (*Medicago sativa*, *Festuca pratensis*, *Lolium multiflorum*), pronounced morphological polymorphism introduces legitimate diagnostic complexity in botanical classification.

Species-Specific Analytical Hierarchy

Analysis revealed morphological determinants of analytical difficulty: large-seeded species (>3 mm) achieved minimal variability (*Helianthus annuus* CV 0.06%), while small-seeded species demonstrated elevated but manageable

variability (*Brassica napus* CV 0.51%). Importantly, chaffy versus non-chaffy morphology showed minimal impact (mean CV 0.29% versus 0.42%), validating modern analytical procedures.

Temporal Stability and Species-Driven Variation

The 2021 CV elevation (1.43%) reflected testing of morphologically challenging species (*Festuca pratensis*, *Brassica napus*) rather than competency degradation. Rapid recovery to 0.35% CV in 2022 without corrective action confirms species-difficulty interpretation and validates regulatory framework effectiveness.

Component Compensation Mechanism

This study documents a previously undescribed phenomenon whereby analyst classification decisions in ambiguous cases result in compensatory adjustments between inert matter and other seeds allocations, maintaining overall purity consensus while masking component-level precision variations.

Recommendations and Future Directions

Enhanced component-specific proficiency monitoring would improve botanical composition reliability, particularly for small-seeded legumes. Investment in higher-precision balances (0.001 g sensitivity) could substantially reduce trace-component variability.

Current regulatory protocols successfully maintain network competency, with future enhancements focusing on component-specific rather than purity-global performance monitoring.

Study Limitations and Future Protocol Enhancements

The demonstrated network stability validates current proficiency testing protocols while highlighting opportunities for methodological enhancement.

Future implementation of multi-lot protocols per ISO 13528:2022 recommendations would further strengthen statistical robustness, complementing the already exceptional competency maintenance documented across the six-year study period.

Final Validation

The Romanian seed testing network demonstrates that mechanical component separation, conducted under quality-assured conditions, represents a highly reliable analytical procedure. Zero below-minimum

proficiency classifications across six years provide empirical validation that systematic competency verification maintains analytical standards across diverse laboratory units, supporting the effectiveness of current regulatory approaches.

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