Data Mining for Co-Crystal Synthesis Design Powered by Large Language Models

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Abstract

Co-crystallization enables modulation of drug physicochemical properties, yet practical synthesis design remains largely empirical. Existing AI methods mainly address molecular pair selection, leaving synthesis-condition design underexplored because the required experimental knowledge is scattered across scientific articles. We present COSYN, a framework that uses LLMs to convert dispersed procedure descriptions into machine-readable synthesis records, yielding COSYN-DB with 20,783 method-aware entries of variable field completeness. These records enable experimental condition recommendations for new molecular pairs and tool-augmented assistance. Using manually annotated datasets, we evaluate both individual components and system-level performance. Our results highlight the potential of a modular LLM-based approach to organize literature-derived evidence and support co-crystal synthesis design beyond pair-level screening.

1. Introduction

Pharmaceutical co-crystals offer a solid-form route for modulating drug properties, including solubility, stability, pharmacokinetic behavior, and mechanical properties (Yadav et al., 2025; Marcos Valdez et al., 2026). They combine an active pharmaceutical ingredient (API) with an additional pharmaceutically acceptable molecule (coformer) in a single crystalline phase, stabilized by non-covalent interactions (Bolla et al., 2022). Despite their broad potential, co-crystal design remains a challenging experimental task that often relies on empirical screening across heterogeneous synthesis conditions (Karimi-Jafari et al., 2018; Zhang et al., 2025a).

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Artificial intelligence has advanced the early stages of co-crystal design, including machine-learning models for coformer screening and co-crystal formability prediction (Wicker et al., 2017; Mswahili et al., 2021; Devogelaer et al., 2019; 2020; Jiang et al., 2021), as well as recent deep learning, thermodynamics-informed, and generative systems (Guo et al., 2024; Birolo et al., 2025; Song et al., 2025b; Gubina et al., 2024). However, these methods primarily operate at the level of molecular-pair selection, while synthesis-condition planning remains underexplored because method-specific precedents are scattered across articles as heterogeneous descriptions rather than standardized datasets.

Automated extraction of chemical information from literature offers a route to this evidence base, from systems such as OSCAR4 and ChemDataExtractor to recent extraction frameworks and domain-specific pipelines (Jessop et al., 2011; Swain & Cole, 2016; Fan et al., 2024; Dagdelen et al., 2024; Vaucher et al., 2021; Shi et al., 2024). However, reliable extraction depends on domain-specific schemas, normalization, and validation rather than generic text mining alone (Schilling-Wilhelmi et al., 2025; Vepreva et al., 2025; Chong & Colindres, 2026), which is especially important for co-crystal synthesis because different preparation methods require different experimental fields.

Large language models (LLMs) offer a flexible mechanism for connecting unstructured scientific text with downstream decision-making (Zhang et al., 2025b), and have been used in chemistry and materials science for information extraction, molecular and materials design, predictive modeling, and tool-augmented reasoning over scientific workflows (Schilling-Wilhelmi et al., 2025; Ramos et al., 2025; Jablonka et al., 2024; Bran et al., 2024; Ruan et al., 2024; Song et al., 2025a). These capabilities are well aligned with co-crystal synthesis design, which requires both recovering experimental knowledge from articles and reusing it to recommend conditions for new molecular pairs. Yet the use of LLMs for method-aware co-crystal synthesis extraction, condition prediction, and tool-augmented assistance remains largely unexplored.

In this work, we present COSYN [1], a modular LLM-based framework for co-crystal synthesis design from literature-derived experimental evidence. COSYN extracts method-aware synthesis records from articles, uses them to recommend conditions for new molecular pairs, and composes these capabilities within an agentic assistant. Our contributions are as follows:

- We build a hierarchical LLM pipeline for method-aware extraction of co-crystal synthesis records from full-text articles, yielding COSYN-DB with 20,783 machine-readable entries whose field completeness reflects the underlying reporting in the source literature.

- We evaluate few-shot LLM prediction of method-specific synthesis conditions from retrieved experimental precedents, extending co-crystal AI beyond pair-level screening.

- We integrate extraction and prediction into a tool-augmented agentic assistant and evaluate system-level behavior on COSYN-BENCH, a domain-specific benchmark of expert-curated questions.

2. Related Work

AI-Guided Co-Crystal Design Co-crystal development remains highly empirical, depending on coformer choice and process parameters that vary across solution- and solid-state methods (Karimi-Jafari et al., 2018; Bolla et al., 2022). AI methods have therefore focused mainly on upstream molecular-pair selection, including descriptor-based classifiers, network models, graph neural networks, deployable platforms, and language-model systems (Wicker et al., 2017; Wang et al., 2020; Devogelaer et al., 2019; Jiang et al., 2021; Guo et al., 2024; Birolo et al., 2025), with extensions to coformer–solvent screening and property-driven generative design (Song et al., 2025b; Zeng et al., 2025; Gubina et al., 2024). In parallel, synthesis-condition recommendation has been developed mostly outside co-crystal synthesis, for organic reactions, inorganic materials, and related reaction-synthesis settings, where it is framed as structured prediction over correlated categorical and numerical choices (Ball et al., 2025). Prior systems use hierarchical, two-stage, template-, cluster-, and LLM-based recommendation models (Gao et al., 2018; Chen & Li, 2024; Wang et al., 2025; Yang et al., 2025; Zhang et al., 2024; Prein et al., 2025). COSYN targets the gap between these lines of work by addressing literature-grounded co-crystal synthesis design with method-specific experimental conditions.

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[1] Code and data are available at https://github.com/NinaDubrovskaya/cocrystal-synthesis.

Automated Extraction of Scientific Data Chemical information extraction has progressed from foundational systems such as OSCAR4, ChemicalTagger, and ChemDataExtractor (Jessop et al., 2011; Hawizy et al., 2011; Swain & Cole, 2016) to document-level and LLM-based extraction across text, tables, and figures (Dagdelen et al., 2024; Fan et al., 2024), with domain-specific pipelines for synthesis procedures, reaction data, MOFs, and nanomaterials (Vaucher et al., 2021; Kearnes et al., 2021; Ai et al., 2024; Zheng et al., 2023; Odobesku et al., 2025). Recent reviews and benchmarks show that robust extraction in specialized domains depends not only on stronger models, but on domain-specific schemas, normalization, provenance, and validation (Schilling-Wilhelmi et al., 2025; Vepreva et al., 2025; Chong & Colindres, 2026). We therefore formulate co-crystal synthesis extraction as method-aware record construction rather than generic text mining.

Agentic Systems and Evaluation LLMs are increasingly used as agents for scientific workflows, coordinating retrieval, tool use, planning, and result interpretation, including tool-augmented agents such as Coscientist and ChemCrow (Boiko et al., 2023; Bran et al., 2024) and role-specialized multi-agent architectures such as ChemAgents and LLM-RDF (Song et al., 2025a; Ruan et al., 2024). Evaluation remains difficult because scientific tasks are open-ended, admit multiple valid solution paths, and depend strongly on tool availability (Gu et al., 2024; Yu et al., 2025), and existing benchmarks target general scientific workflows (Chen et al., 2025) or broad chemistry task collections (Mirza et al., 2025) rather than the coupled retrieval, extraction, and prediction demands of co-crystal synthesis design. We therefore pair our agentic architecture with a targeted benchmark reflecting these domain-specific requirements.

3. Dataset Construction

3.1. Source Corpus

We constructed the source corpus from the Cambridge Structural Database (CSD), a curated repository of experimentally determined co-crystal structures (Groom et al., 2016). Using publication DOIs associated with relevant CSD entries, we obtained 4,287 unique source DOIs after deduplication (Figure 2A).

Automated full-text retrieval yielded 3,997 PDFs (not redistributed), which were converted to machine-readable text and used as input to the extraction pipeline. Retrieval failures were due to access restrictions, missing full-text links, publisher-side failures, or conversion errors. Corpus-level statistics, including open-access status and publication-year distributions, are provided in Appendix A.1.

Figure 1. Overview of CoSYN. Full-text articles from CSD-linked DOIs pass through a three-stage, method-aware extraction pipeline and are normalized into CoSYN-DB, a database of 20,783 synthesis records. These records serve as retrievable precedents for few-shot prediction of method-specific conditions for new API–coformer pairs, and all components are exposed as tools in an agentic assistant evaluated on CoSYN-BENCH.

3.2. Method Taxonomy

Co-crystal synthesis methods differ in experimental structure and reported parameters, so no single schema fits all. We therefore adopt a method-aware taxonomy as CoSYN's structural backbone. Drawing on the co-crystal synthesis literature (Karagianni et al., 2018; Buddhadev & Garala, 2020; Douroumis et al., 2017; Pawar et al., 2021), we define 12 method types spanning solution-based routes (solution, slurry, antisolvent crystallization) and solid-state or specialized routes (grinding, hot-melt extrusion), each with a concise description of its key physicochemical steps and a procedure template with marked parameter slots.

The taxonomy targets robust extraction and prediction rather than exhaustive nomenclature: procedures sharing experimental logic and parameter spaces are grouped into one type, e.g., neat and liquid-assisted grinding, unified by mechanical activation with solvent addition optional. Full templates and parameter sets are in Appendix A.2.

3.3. Hierarchical Extraction Pipeline

A naive one-step extraction is unsuitable because (i) not every article contains an explicit synthesis procedure (it may be omitted or replaced by a reference), and (ii) the parameter schema is method-dependent, yet the method is unknown in advance. A single prompt covering detection, method inference, and parameter extraction would force the model to weigh many irrelevant fields, increasing hallucination risk (Ji et al., 2023). We therefore formulate extraction as a

three-stage hierarchical task.

Stage 1: Procedure detection. Given the full article text, the model decides whether an explicit synthesis procedure is present.

Stage 2: Method classification. The model assigns one or more labels from the 12 predefined method types (Appendix A.2), isolating method identification so that downstream prompts use only the relevant schema.

Stage 3: Parameter extraction. For each detected method, a method-specific prompt extracts experimental parameters into a structured output. These prompts were developed iteratively with domain experts and specify field names and worked examples to ensure consistency across articles.

All prompts are provided in Appendix A.3, with the grinding prompt shown as a representative example of the Stage 3 format.

Validation datasets. To support evaluation across all components of the pipeline, we constructed three hand-annotated validation datasets, one per stage: (i) 100 articles for Stage 1 (92 positive, 8 negative), (ii) 109 articles for Stage 2 covering all 12 method types (54 single-method, 55 multi-method), and (iii) a LAG-specific Stage 3 set of 55 articles with 295 records over 25 parameters. The same datasets are used for prompting strategy comparison in Section 3.4 and for cross-model evaluation in Section 3.5. Metric definitions are given in Appendix A.4.

Table 1. Prompting-strategy selection on `GPT-4o-mini`. Precision, recall, F1, and accuracy of three strategies on the Stage 1 procedure-detection and Stage 2 method-classification validation sets. The best mean within each stage is highlighted.

Strategy	Precision	Recall	F1	Accuracy
Stage 1				
S1-A	0.62 ±0.11	1.00 ±0.00	0.76 ±0.08	0.62 ±0.11
S1-B	1.00 ±0.00	0.92 ±0.08	0.96 ±0.04	0.95 ±0.05
S1-C	0.72 ±0.11	1.00 ±0.00	0.84 ±0.07	0.76 ±0.09
Stage 2				
S2-A	0.69 ±0.06	0.53 ±0.05	0.58 ±0.05	0.53 ±0.05
S2-B	0.70 ±0.05	0.56 ±0.05	0.59 ±0.05	0.56 ±0.05
S2-C	0.75 ±0.05	0.61 ±0.05	0.64 ±0.05	0.61 ±0.05

3.4. Prompting Strategy Optimization

Setup. We used `GPT-4o-mini` to develop prompts for all three stages, evaluated on the validation datasets introduced in Section 3.3. For Stages 1 and 2 we systematically compared three predefined strategies. For Stage 3 we refined a single prompt through expert-in-the-loop iteration on the LAG validation set, chosen as a representative high-support method with a rich parameter schema.

Strategies. The three Stage 1 strategies differ in the required **response format**:

- **S1-A:** binary classification of synthesis presence.
- **S1-B:** full text span of the synthesis method.
- **S1-C:** same as S1-B, separated per molecular pair and technique.

The three Stage 2 strategies differ in the **method-type context** provided alongside the procedure:

- **S2-A:** predefined list of method names.
- **S2-B:** method names with natural-language descriptions from Table 5.
- **S2-C:** method names with procedural templates from Appendix A.2.2.

Results. Table 1 shows that at Stage 1 both S1-A and S1-C achieved perfect recall but at a clear cost to precision, indicating that the binary formulation was prone to over-detection while the additional pair-and-technique decomposition in S1-C introduced spurious extractions. S1-B provided the best balance with the highest F1 and accuracy, and was selected for Stage 1.

The Stage 2 results in Table 1 follow a consistent gradient: names alone yielded the weakest performance, descriptions produced a modest improvement, and templates achieved the best scores across all four metrics. Templates likely help

Table 2. Extraction performance of four LLMs across the three pipeline stages. Precision, recall, F1, and accuracy are reported per stage; † marks the model selected for full-corpus extraction, and the best mean within each stage is highlighted.

Model	Precision	Recall	F1	Accuracy
Stage 1				
GPT-4o-mini	1.00 ±0.00	0.78 ±0.04	0.88 ±0.03	0.80 ±0.04
Gemma-3-27B	0.91 ±0.03	0.92 ±0.03	0.92 ±0.02	0.85 ±0.04
Llama-3.3-70B	1.00 ±0.00	0.57 ±0.05	0.73 ±0.04	0.61 ±0.05
Gemini-2.5-Flash†	0.98 ±0.02	0.97 ±0.02	0.97 ±0.01	0.95 ±0.02
Stage 2				
GPT-4o-mini	0.75 ±0.05	0.61 ±0.05	0.64 ±0.05	0.61 ±0.05
Gemma-3-27B	0.58 ±0.08	0.47 ±0.05	0.44 ±0.05	0.47 ±0.05
Llama-3.3-70B	0.41 ±0.08	0.14 ±0.03	0.19 ±0.04	0.14 ±0.03
Gemini-2.5-Flash†	0.86 ±0.04	0.82 ±0.04	0.81 ±0.04	0.82 ±0.04
Stage 3				
GPT-4o-mini	0.61 ±0.03	0.63 ±0.03	0.54 ±0.03	0.45 ±0.03
Gemma-3-27B	0.59 ±0.04	0.70 ±0.03	0.56 ±0.03	0.47 ±0.04
Llama-3.3-70B	0.62 ±0.03	0.73 ±0.03	0.61 ±0.03	0.53 ±0.04
Gemini-2.5-Flash†	0.65 ±0.04	0.74 ±0.03	0.63 ±0.03	0.56 ±0.04

because closely related method types are easier to distinguish by surface form than by abstract definitions. We therefore used S2-C for Stage 2 in the final extraction pipeline. Per-method accuracy and F1 across all 12 method types are reported in Appendix A.5.

3.5. Evaluation of Extraction Pipeline

Setup. Using the selected prompting strategies (S1-B and S2-C), we evaluated four LLMs on the validation datasets from Section 3.3: `GPT-4o-mini`, `Gemma-3-27B`, `Llama-3.3-70B`, and `Gemini-2.5-Flash`. Metric definitions are given in Appendix A.4.

Results. Table 2 reports performance across stages. `Gemini-2.5-Flash` achieved the highest F1 and accuracy and was selected for full-corpus extraction. Stage 1 separated models mainly by recall: `GPT-4o-mini` and `Llama-3.3-70B` reached perfect precision but missed many valid procedures, whereas `Gemini-2.5-Flash` preserved both (F1 = 0.97). Stage 2 was the most model-sensitive, with F1 ranging from 0.19 to 0.81, reflecting differences in matching procedure text to template-based method definitions. Stage 3 scores are tighter (F1 0.54–0.63) and conservative, as many mismatches arise from the strict multiset metric. Per-parameter precision and recall are reported in Appendix A.6. Strategy selection on `GPT-4o-mini` may introduce a small optimistic bias for that model, but the consistent advantage of `Gemini-2.5-Flash` suggests the relative ranking is unaffected.

Figure 2. COSYN-DB construction and parameter coverage. [A] Sankey diagram of the data flow from 4,287 source DOIs to normalized synthesis records and parameter families. [B] Parameter-family fill rates (fraction of non-empty values) for the three downstream datasets after quality filtering. Method abbreviations and parameter-family definitions are given in Appendices A.2 and A.8.

Error analysis. We manually annotated the `Gemini-2.5-Flash` outputs to characterize the dominant failure modes at each stage (Table 6). Stage 1 and Stage 2 were both dominated by omissions, driven respectively by non-standard terminology and by methods reported only in supplementary or comparative experiments. Stage 3 was instead dominated by evaluation artefacts from strict multiset-based comparison. The most informative genuine Stage 3 error was fabrication of procedural details that the model appears to lift from surrounding text but assign to the wrong record. A full breakdown is provided in Appendix A.7.

3.6. COSYN-DB Dataset

Database extraction. Of the documents introduced in Section 3.1, Stage 1 identified 3,469 articles (86.8%) as containing at least one co-crystal synthesis procedure. Stage 2 distributed the detected procedures across the 12 method categories, with a single article potentially contributing to multiple categories. Stage 3 then extracted method-specific parameters, yielding 20,783 unique synthesis records. Three methods dominate the database and together cover 98.7% of all entries: solution crystallization (15,730 records), grinding (3,963), and slurry (819). These were retained for downstream experiments, while the remaining nine categories were too sparse to support reliable modeling. The full data flow is visualized as a Sankey diagram in Figure 2A.

Post-processing. Each dataset was processed through a method-specific normalization pipeline. Records were retained only when both API and coformer were present and resolvable to canonical SMILES, with invalid pairs removed by RDKit. Categorical fields were harmonized to controlled vocabularies and numerical parameters converted to common units. After filtering out-of-range values, duplicates, and weakly populated records, 11,587 records remained, including 8,272 for solution crystallization, 2,722 for grinding, and 593 for slurry (Figure 2A). Parameter fill rates were highly uneven (Figure 2B). Ratios and solvents were usually populated, whereas amounts, times, and temperatures remained sparse because of heterogeneous reporting.

To prevent leakage, records were split by connected components over shared record identifiers and API/coformer pairs, so related experiments never crossed the split boundary. The balanced test set contains 100 records per method and 300 held-out examples in total. Split statistics are reported in Appendix A.9.

Missing values were marked as expected-but-unreported or structurally absent, such as solvent in neat grinding. Only expected-but-unreported values were imputed, using train-fitted group-wise and global modes with deterministic chemical rules for numerical fields. Pre-imputation test snapshots were retained so that prediction metrics in Section 4 use only observed values. Full imputation rules are listed in Appendix A.10.

Table 3. Synthesis-condition prediction on the Grinding test set. Weighted F1 and accuracy for few-shot LLMs (k=3) and a most-frequent-value baseline. Categorical, Numerical, and Overall are weighted averages over the field groups; Solvent and Ratio are shown individually. [†]Solvent uses exact-match on canonical names, the Categorical aggregate solvent-family labels.

Model	Solvent[†]		Ratio		Categorical		Numerical		Overall	
	F1	Accuracy	F1	Accuracy	F1	Accuracy	F1	Accuracy	F1	Accuracy
Baseline	0.17 ±0.00	0.34 ±0.00	0.87 ±0.00	0.92 ±0.00	0.26 ±0.00	0.43 ±0.00	0.38 ±0.00	0.46 ±0.00	0.35 ±0.00	0.45 ±0.00
GPT-4o-mini	0.36 ±0.01	0.38 ±0.01	0.85 ±0.00	0.84 ±0.00	0.48 ±0.00	0.49 ±0.00	0.45 ±0.01	0.45 ±0.00	0.46 ±0.00	0.46 ±0.00
GPT-5.4	0.41 ±0.00	0.40 ±0.00	0.88 ±0.00	0.91 ±0.00	0.47 ±0.00	0.46 ±0.00	0.49 ±0.00	0.49 ±0.00	0.48 ±0.00	0.48 ±0.00
Gemini-2.5-Flash	0.41 ±0.01	0.41 ±0.01	0.85 ±0.00	0.86 ±0.00	0.49 ±0.00	0.49 ±0.00	0.48 ±0.00	0.48 ±0.00	0.49 ±0.00	0.48 ±0.00
Gemini-3.1-Pro	0.44 ±0.01	0.44 ±0.01	0.88 ±0.01	0.91 ±0.01	0.50 ±0.00	0.50 ±0.00	0.47 ±0.01	0.48 ±0.01	0.49 ±0.00	0.49 ±0.00

4. Synthesis Condition Prediction

4.1. Experimental Setup

Task formulation. Given a co-crystal pair, represented by API and coformer SMILES, and a target synthesis method, the task is to predict the corresponding method-specific synthesis parameters, which combine categorical and numerical fields. Since these parameters are interdependent in real procedures, the goal is to predict coherent synthesis conditions rather than isolated fields.

Baseline. As a naive reference, the mode baseline predicts the most frequent training value for each parameter, using the concatenation of API and coformer Morgan fingerprints (radius 2, 2048 bits) as input.

Few-shot LLM prediction. We evaluate four LLMs spanning two families and two capability tiers under a few-shot in-context learning setup: GPT-4o-mini, GPT-5.4, Gemini-2.5-Flash, and Gemini-3.1-Pro. For each test pair, we retrieve the k most similar training pairs by molecular fingerprint similarity (Appendix B.1) and include their synthesis records as in-context examples. The model is then prompted to return the full method-specific parameter set in a structured format (Appendix B.2).

4.2. Evaluation Protocol

We evaluate every non-unit parameter with a test-set missing rate below 80%. Categorical fields are scored by exact-match accuracy and weighted F1 after vocabulary normalization, except Solvent, which is aggregated at the solvent-family level because of its high cardinality (Appendix B.3). Numerical fields use tolerance bins, where a prediction is correct if it matches the ground-truth bin (Appendix B.4). Per-parameter scores are aggregated into Categorical, Numerical, and Overall weighted averages (Table 3), with Solvent and Ratio also reported separately because they probe opposite ends of the difficulty spectrum. Solvent is harder under exact match than family-level aggregation, whereas Ratio is dominated by the low-entropy 1:1 case.

Figure 3. Effect of the number of in-context examples on prediction. Weighted F1 versus k on GPT-4o-mini for three methods (GR, grinding; SL, slurry; SC, solution crystallization).

4.3. Prediction Results

Effect of the number of examples. Figure 3 shows weighted F1 as a function of k on GPT-4o-mini. The largest gain across all three methods occurs when moving from zero-shot to k=3, indicating that a few retrieved precedents already provide useful method- and pair-specific context. Beyond this point, categorical scores keep improving slightly as the model sees more labels and procedural variants, while numerical scores peak at k=3 and then decline, likely because additional examples dilute the closest-precedent signal. Since numerical fields dominate the schema and are more sensitive to over-broad context, we fix k=3 in all subsequent experiments.

Comparison. Table 3 reports results on the Grinding dataset. The mode baseline is strong on Ratio because equimolar stoichiometries dominate, but weak on chemistry-dependent Solvent. This shows that Ratio partly reflects a domain prior, whereas Solvent requires pair-specific evidence. All LLMs improve Solvent F1 from 0.17 to 0.36–0.44, supporting the use of retrieved precedents for condition prediction.

Table 4. Agentic system comparison on COSYN-BENCH. Capability, complexity, and overall scores are on a 1–5 scale and Pass Rate on a 0–1 scale; the best mean within each column is highlighted.

System	Capability			Complexity		Overall	
	Retrieval	Prediction	Analysis	Single	Multi	Mean	Pass Rate
LLM-Only	1.71 ±0.08	2.22 ±0.05	1.72 ±0.05	1.94 ±0.03	1.76 ±0.02	1.88 ±0.01	0.00 ±0.00
Context-Augmented LLM	1.66 ±0.06	2.05 ±0.00	1.96 ±0.01	2.26 ±0.10	1.51 ±0.05	2.01 ±0.05	0.07 ±0.00
Single-Agent	3.61 ±0.08	3.02 ±0.22	3.66 ±0.25	3.74 ±0.40	3.25 ±0.04	3.58 ±0.25	0.47 ±0.09
Multi-Agent	3.47 ±0.09	3.73 ±0.10	3.41 ±0.34	3.67 ±0.22	3.35 ±0.11	3.56 ±0.18	0.48 ±0.16

Within the LLM group, the Gemini models achieve the strongest overall performance, with `Gemini-3.1-Pro` leading on Solvent and `Gemini-2.5-Flash` matching it on Overall F1. This suggests that the main benefit of the larger model appears in the highest-cardinality categorical field, while the smaller model preserves overall accuracy at substantially lower inference cost. We therefore adopt `Gemini-2.5-Flash` as the Predictor agent in Section 5. Per-parameter results are reported in Appendix B.6.

5. Agentic Co-Crystal Design

The components introduced above, article extraction, COSYN-DB retrieval, and few-shot prediction, solve complementary sub-problems, whereas practical co-crystal synthesis design often requires using them jointly. We therefore expose all COSYN components through an agentic interface and evaluate whether this improves co-crystal synthesis assistance over non-agentic LLM baselines. We deliberately keep the agent layer architecturally standard, so that the comparison isolates the contribution of the method-aware COSYN components and tools rather than of a bespoke agent design.

5.1. Experimental Setup

Baselines. We compare two baselines to separate static evidence access from tool-mediated reasoning: LLM-Only receives only the user question, while Context-Augmented LLM also receives task-relevant static context such as article text or database evidence. This tests whether evidence alone is sufficient, or whether co-crystal synthesis questions require iterative tool use and task decomposition.

Agentic architectures. The Single-Agent architecture uses one LLM controller to interpret the request, plan the solution, select tools, and write the final response. The Multi-Agent architecture separates these roles: an Orchestrator decomposes the query and routes subtasks to three specialists, Analyzer, Searcher, and Predictor, which return evidence-grounded reports for final aggregation.

Both variants use the same COSYN toolset and differ only in control structure, testing whether synthesis assistance

benefits more from a single generalist controller or hierarchical decomposition. Tool details and prompts are summarized in Appendices C.1 and C.2. All agent controllers use `GPT-4o-mini`.

5.2. Evaluation of the Agentic System

CoSyn-Bench. We introduce COSYN-BENCH, a set of 30 expert-curated questions: 20 single-capability questions requiring retrieval, analysis, or prediction, and 10 multi-capability questions combining several capabilities. Each item includes a gold answer and structured scoring metadata that specifies the information available in the user request, the constraints that must be satisfied, and the components expected in a complete answer. Representative items are shown in Appendix C.3.

LLM-as-a-judge evaluation. Answers are scored by `GPT-5.4` using the question, gold answer, benchmark metadata, execution context, and candidate response. The judge assigns integer scores from 1 to 5 for Task completion, Correctness, Constraint adherence, Groundedness, and Answer quality. The judge prompt is summarized in Appendix C.4. We report scores by capability and question complexity, together with Pass Rate, defined as the fraction of questions scoring ≥ 4 on every criterion.

5.3. Results

Table 4 reports the resulting baseline and architecture comparison.

Baseline comparison. LLM-Only scores poorly at 1.88, reflecting the difficulty of domain-specific synthesis questions without external evidence. Static context raises the score only to 2.01, while both baselines remain far below the agentic systems, showing that performance depends on coordinated tool use, not just information access.

Architecture comparison. Single-Agent and Multi-Agent achieve similar overall scores of 3.58 and 3.56, but differ in profile. Single-Agent performs better on Retrieval and Analysis, suggesting that one controller suits compact tool-use workflows. Multi-Agent performs better on Pre-

diction, Pass Rate, and multi-capability questions, indicating that decomposition helps combine multiple evidence sources. Multi-Agent uses more time and calls but fewer tokens (Appendix C.5), trading speed for context efficiency.

6. Discussion

Taken together, our results indicate that progress in co-crystal AI depends not only on predicting which molecular pairs co-crystallize, but on making the experimental knowledge required for synthesis design usable at scale. COSYN targets this gap by converting heterogeneous literature procedures into normalized, method-aware synthesis records that serve as the evidence base for condition prediction and agentic assistance, linking pair-level reasoning to practical synthesis planning.

The database-construction results show that the published co-crystal literature already contains enough procedural knowledge to support data-driven synthesis assistance, but only after this knowledge is reorganized into method-aware records. The taxonomy and hierarchical extraction pipeline enable this conversion by mapping heterogeneous procedures to method-specific parameter schemas. At scale, COSYN extracted 20,783 records, revealing both the richness and imbalance of the literature: 98.7% of records come from the three dominant method types, while rarer routes remain underrepresented. Parameter reporting is similarly uneven, with ratios and solvents recorded more consistently than amounts, times, and temperatures. This makes COSYN-DB useful not only as a source of precedents, but also as a map of where the literature provides strong or weak evidence for synthesis-condition modeling.

The prediction results show that structured precedents can support method-specific synthesis-condition recommendation through locally relevant procedural analogues, with a few retrieved examples outperforming broader context, suggesting that parameters are better inferred from close analogues than from aggregate trends. Performance is strongest for well-reported parameters, especially stoichiometric ratios, although ratio prediction partly reflects a strong domain prior and the normalization convention that treats omitted equimolar ratios as implicit 1:1 mixtures. High-cardinality solvent identity and sparsely reported absolute quantities remain harder, because they depend more on experiment-specific conditions than on stable pair-level patterns. We therefore interpret COSYN predictions as evidence-grounded starting points for expert review rather than optimized synthesis protocols.

Finally, exposing these components through an agentic interface shows the same evidence base can be operated interactively: tool-augmented agents nearly double the COSYN-BENCH score over non-agentic LLM baselines (from 1.88 to 3.58), whereas adding only static context improves it marginally. This indicates that the gain comes from coordinating retrieval, analysis, and prediction, not simply from more context. The Single-Agent and Multi-Agent variants perform comparably overall and mainly trade off speed against context efficiency, indicating that the system-level benefit stems from grounding answers in COSYN's tools rather than from a particular agent architecture.

Limitations. This study has several limitations that should be acknowledged. (i) As a literature-grounded resource, COSYN-DB inherits the biases, coverage gaps, and reporting heterogeneity of the published record. (ii) Its scope is the synthesis procedure, leaving characterization and outcome information outside the current schema. (iii) The records rest on automated extraction with expert validation that should be further strengthened and broadened to raise confidence in the data. (iv) The taxonomy is operational and does not yet capture the full diversity of synthesis routes. (v) Predictive coverage centers on the best-represented methods, while sparsely reported parameters remain harder, so the methodological base should be broadened. (vi) Finally, the evaluation remains retrospective, and prospective wet-lab validation is the natural next step.

Future directions. These limitations suggest several complementary directions. At the data level, the evidence base can be broadened with negative and rare-route procedures and enriched with characterization and outcome fields, while expert validation and benchmarks such as COSYN-BENCH are scaled up. At the modeling level, uncertainty-aware and multi-solution prediction would better capture the range of viable conditions. Ultimately, closing the loop between structured literature evidence, predictive modeling, and prospective wet-lab synthesis is the central step toward reliable, broadly applicable co-crystal synthesis assistance.

7. Conclusion

We introduced COSYN, a literature-grounded LLM framework that moves co-crystal AI from pair-level screening toward synthesis-condition design. At its core, COSYN converts heterogeneous procedures into COSYN-DB, a method-aware database of 20,783 synthesis records, and uses retrieved precedents to predict method-specific conditions for new API–coformer pairs. These capabilities are further exposed through a tool-augmented assistant evaluated on COSYN-BENCH. Experiments show that literature-derived evidence can support condition prediction, while revealing the reporting gaps and validation needs that bound current accuracy. Overall, COSYN provides a scalable, data-driven foundation for evidence-grounded synthesis prediction rather than replacing expert judgment or experimental validation.

Impact Statement

CoSyn supports co-crystal synthesis research by converting dispersed literature procedures into structured evidence for synthesis-design decisions. It enables faster literature analysis, systematic condition comparison, and evidence-grounded protocol suggestions. However, it inherits biases and gaps in published reports, so predictions should be treated as expert-reviewed starting points, not optimal protocols. Wet-lab use requires safety assessment, human oversight, and prospective validation.

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A. Dataset Construction

A.1. Source Corpus Statistics

Figure 4 summarizes the composition of the collected source corpus. We report the distribution of retrieved articles by publication year and open-access status to characterize the temporal coverage and accessibility of the literature used for extraction.

Figure 4. Source corpus statistics. [A] Distribution of the collected CSD-derived source DOIs by publication year. [B] Open-access status of the corpus used for full-text retrieval and extraction.

A.2. Structured representation of co-crystal synthesis methods

A.2.1. CLASSIFICATION OF CO-CRYSTAL SYNTHESIS METHODS

Co-crystal synthesis procedures differ in the physical mechanism used to induce co-crystallization, the required experimental setup, and the process variables reported in experimental sections. To account for this variability, we defined 12 co-crystal synthesis method categories. Table 5 summarizes the corresponding method descriptions, generalized parameter groups, and links to the procedural templates. The method descriptions were used in strategy S2-B to support method-aware extraction.

Table 5. Operational taxonomy for co-crystal synthesis extraction. For each of the 12 method types, the table lists a concise description, the generalized parameter groups targeted during extraction, and the corresponding procedural template.

Method	Description	Key parameters	Template
Solution crystallization	Solution crystallization is a widely used method for cocrystal synthesis that involves dissolving the individual coformers in a suitable solvent or solvent mixture. Cocrystallization is then induced by allowing the solvent to evaporate (slow solvent removal to achieve supersaturation), cooling the solution (temperature reduction to decrease solubility), or introducing several immiscible phases (crystal formation at liquid–liquid or liquid–gas boundaries).	component composition / solvent system / stirring conditions / evaporation or cooling conditions / filtration and drying	T1
Grinding	Grinding is a mechanochemical method for cocrystal synthesis that entails mixing solid coformers together through manual (mortar and pestle) or mechanical (ball milling) means to induce cocrystallization. This approach can be performed as neat grinding, using only the solid constituents, or as liquid-assisted grinding, where a minimal amount of solvent is added to promote better molecular contact.	component composition / optional liquid additive / grinding apparatus / milling conditions / drying	T2
Slurry	Slurry is a solution-mediated method for cocrystal synthesis that entails suspending solid coformers in a minimal amount of solvent to induce cocrystallization. This approach involves allowing the coformers to interact in the solvent over a period (at constant temperature or with temperature variation), followed by separation and drying of the solid product. The approach relies on the dynamic equilibrium between dissolved and solid phases to achieve desired crystal properties.	component composition / solvent volume / stirring conditions / filtration and solvent evaporation / drying	T3

Table 5 *(continued)*

Method	Description	Key parameters	Template
Anti-solvent	The anti-solvent method is a solution-based technique for cocrystal synthesis that involves the addition of a second solvent, the "anti-solvent," to a solution containing the mixture of coformers. The anti-solvent is chosen for its ability to reduce the solubility of the cocrystal, thereby inducing its precipitation from the solution. This method often requires specialized equipment in which a separate tank for anti-solvent and peristaltic pump will be used.	component composition / anti-solvent system / addition setup / addition and mixing conditions / filtration and drying	T4
Hot-melt extrusion	Hot-melt extrusion is a continuous, solvent-free method for cocrystal synthesis that involves feeding a mixture of solid coformers into a heated extruder, where they are subjected to mixing and conveying by rotating screws. The extruder is typically divided into several zones, each with independently controlled temperatures, facilitating melting of at least one component and promoting molecular contact and subsequent cocrystallization as the melt cools.	component composition / optional additional substance / extruder configuration / feeding and temperature profile / post-processing	T5
Freeze drying	Freeze drying (or lyophilization) is a solvent-removal method for co-crystal synthesis that involves dissolving coformers in a solvent, freezing the solution, and then removing the solvent through sublimation under vacuum to obtain the co-crystal. This technique can be performed using special equipment capable of maintaining low pressure to create amorphous or nanostructured products.	component composition / solvent / freeze-dryer setup / freezing and lyophilization conditions / product collection	T6
Spray drying	Spray drying is a continuous method for cocrystal synthesis that transforms liquids into dry powders through atomization into a hot gas stream, followed by rapid solvent evaporation and particle collection. During this process, the cocrystals nucleate and grow within highly supersaturated regions of the coformer substance due to rapid solvent evaporation and solidification of the generated droplets.	component composition / solvent system / spray-dryer setup / atomization and drying conditions / powder collection	T7
Ultrasound	Ultrasound (or sonocrystallization) is a solution-based cocrystallization method that involves the application of high-frequency sound waves to induce cocrystallization. This approach is typically performed by subjecting a supersaturated solution of the coformers to ultrasonic pulses, which can promote nucleation and crystal growth.	component composition / solvent system / ultrasonic setup / sonication conditions / filtration and drying	T8
Supercritical fluid	Supercritical fluid is a method for synthesizing co-crystals where solid coformers are exposed to a supercritical fluid, typically supercritical carbon dioxide ($scCO_2$), which acts as a solvent or anti-solvent to facilitate co-crystal formation. The coformers may partially dissolve or become highly dispersed within the supercritical fluid, and the mixture is stirred for a defined period to allow co-crystallization to occur.	component composition / supercritical fluid or co-solvent / reactor setup / pressure–temperature–flow conditions / depressurization and drying	T9
Electrospray	Electrospray is a solution-based method for cocrystal synthesis that involves atomizing a solution containing the cocrystal components into a fine mist of charged droplets using a high electric field. As the solvent evaporates from these droplets, the supersaturation of the components increases, leading to the rapid formation of cocrystals.	component composition / solvent system / syringe and electrospray setup / flow–voltage–distance conditions / collection and drying	T10
Resonant acoustic	Resonant acoustic cocrystal synthesis is a mechanochemical method for cocrystal formation that involves intense mixing of solid coformers (sometimes with a small amount of solvent), through high-frequency and high-intensity vibratory motion generated by a resonant acoustic mixer. This technique leverages powerful resonant frequencies to promote contact and induce cocrystallization.	component composition / optional liquid additive / resonant acoustic mixer / acceleration–frequency–time conditions / drying	T11
Laser irradiation	Laser irradiation is a cocrystallization method that entails exposing a physical mixture of solid coformers to a focused laser beam to induce cocrystallization. This approach can be performed directly on dry powder blends, where the laser's energy promotes the necessary molecular mobility for cocrystal formation.	component composition / optional additive / substrate and laser setup / irradiation conditions / collection and storage	T12

A.2.2. METHOD-SPECIFIC PROCEDURAL TEMPLATES

The templates below define normalized procedural patterns for the 12 co-crystal synthesis method categories listed in Table 5. These templates were used in strategy S2-C to represent synthesis procedures in a method-specific format. They were also used during Stage 3 to construct extraction prompts and to guide the prediction of method-specific experimental parameters.

Each template contains slots marked by [...]. These slots correspond to the main entities typically reported in experimental procedures, including component composition, solvent system, apparatus, operating conditions, and other. Optional fields are explicitly marked only for the solution crystallization, grinding, and slurry templates using [...], in all other templates, bracketed fields denote the target extraction slots without an additional optionality annotation. For all templates except solution crystallization, grinding, and slurry, separate slots for the second and third solvents were omitted for readability.

T1. SOLUTION CRYSTALLIZATION

```
Solution crystallization was performed using [Amount of API] [Amount Unit of API] of
[API] and [Amount of Coformer] [Amount Unit of Coformer] of [Coformer] in a ratio
of [Part of Ratio API]:[Part of Ratio Coformer].  The components were dissolved
in [Amount of Solvent] [Solvent Amount Unit] of [Solvent], [Amount of Second
Solvent] [Second Solvent Amount Unit] of [Second Solvent], and [Amount of Third
Solvent] [Third Solvent Amount Unit] of [Third Solvent].  The solution was stirred at
[Mixing Temperature] [Mixing Temperature Unit] for [Mixing Time] [Mixing Time Unit].
The resulting solution was filtered and covered with [Covering Method] featuring
[Description of Holes].  The solution was then left at room temperature for [Time
of Cooling or Evaporation] [Time of Cooling or Evaporation Unit].  The resulting solid
was collected by filtration and dried to obtain the co-crystal.
```

T2. GRINDING

```
Mechanochemical synthesis was carried out using a [Mixing Apparatus].  The mixture
was processed at [Mixing Frequency] [Frequency Unit] for [Mixing Time] [Mixing Time
Unit] at [Mixing Temperature] [Mixing Temperature Unit].  A blend of [Amount of
API] [Amount Unit of API] of [API] and [Amount of Coformer] [Amount Unit of Coformer]
of [Coformer] was prepared in a ratio of [Part of Ratio API]:[Part of Ratio Coformer].
Additionally, [Amount of Solvent] [Solvent Amount Unit] of [Solvent], [Amount of
Second Solvent] [Second Solvent Amount Unit] of [Second Solvent], and [Amount of Third
Solvent] [Third Solvent Amount Unit] of [Third Solvent] were added.  The mixture was
then ground.  The resulting co-crystal was dried for [Drying Time] [Drying Time Unit]
at [Drying Temperature] [Drying Temperature Unit].
```

T3. SLURRY

```
Slurry crystallization was conducted by suspending [Amount of API] [Amount Unit
of API] of [API] and [Amount of Coformer] [Amount Unit of Coformer] of [Coformer]
in a ratio of [Part of Ratio API]:[Part of Ratio Coformer].  The solids were
suspended in [Amount of Solvent] [Solvent Amount Unit] of [Solvent], [Amount of
Second Solvent] [Second Solvent Amount Unit] of [Second Solvent], and [Amount of
Third Solvent] [Third Solvent Amount Unit] of [Third Solvent].  The suspension was
stirred at [Mixing Temperature] [Mixing Temperature Unit] for [Mixing Time] [Mixing
Time Unit].  The solid product was separated by filtration, and residual solvent
was evaporated at [Evaporation Temperature] [Evaporation Temperature Unit] for
[Evaporation Time] [Evaporation Time Unit].  The resulting co-crystal was dried for
[Drying Time] [Drying Time Unit] at [Drying Temperature] [Drying Temperature Unit].
```

T4. ANTI-SOLVENT

Anti-solvent crystallization was performed using [Amount of API] [Amount Unit of API] of [API] and [Amount of Coformer] [Amount Unit of Coformer] of [Coformer] in a ratio of [Part of Ratio API]:[Part of Ratio Coformer]. The components were dissolved in [Amount of Solvent] [Solvent Amount Unit] of [Solvent]. An anti-solvent stream containing [Amount of Anti-solvent] [Anti-solvent Amount Unit] of [Anti-solvent] was added to the solution at [Speed of Addition] [Speed of Addition Unit] using a peristaltic pump, while mixing at [Mixing Frequency] [Frequency Unit] and [Mixing Temperature] [Mixing Temperature Unit]. The resulting suspension/solution was filtered and dried at [Drying Temperature] [Drying Temperature Unit].

T5. HOT-MELT EXTRUSION

Hot-melt extrusion was performed using a [Type of Extruder] with a screw diameter of [Screw Diameter] [Screw Diameter Unit] and an L/D ratio of [L/D Ratio]. [Amount of API] [Amount Unit of API] of [API] and [Amount of Coformer] [Amount Unit of Coformer] of [Coformer] were weighed in a ratio of [Part of Ratio API]:[Part of Ratio Coformer]. [Amount of Additional Substance] [Amount of Additional Substance Unit] of [Additional Substance] was added, and the components were pre-mixed using a [Mixing Apparatus] for [Mixing Time] [Mixing Time Unit]. The batch size was [Batch Size] [Batch Size Unit]. The blend was fed into the extruder at [Feed Rate] [Feed Rate Unit]. The extruder comprised [Amount of Types of Zones] zones ([List of Types of Zones]) with temperatures set to [List of Temperature in Zones] [Temperature in Zones Unit]. The screw configuration ([List of Screw Type]) and screw speed ([Screw Speed] [Screw Speed Unit]) were selected according to the zone design. Extrusion was carried out using a [Type of Die] with a diameter of [Die Diameter] [Die Diameter Unit]. The resulting co-crystal was pulverized using an agate mortar and pestle.

T6. FREEZE DRYING

Freeze-drying co-crystal synthesis was performed using a [Freeze-Dryer Apparatus] at [Freeze-Drying Temperature] [Freeze-Drying Temperature Unit] and under [Freeze-Drying Pressure] [Freeze-Drying Pressure Unit]. A mixture containing [Amount of API] [Amount Unit of API] of [API] and [Amount of Coformer] [Amount Unit of Coformer] of [Coformer] was prepared in a ratio of [Part of Ratio API]:[Part of Ratio Coformer]. The components were dissolved in [Amount of Solvent] [Solvent Amount Unit] of [Solvent]. The solution was then frozen at [Cooling Temperature] [Cooling Temperature Unit] and lyophilized to obtain the co-crystal.

T7. SPRAY DRYING

Spray-drying was performed using [Amount of API] [Amount Unit of API] of [API] and [Amount of Coformer] [Amount Unit of Coformer] of [Coformer] in a ratio of [Part of Ratio API]:[Part of Ratio Coformer]. The components were dissolved in [Amount of Solvent] [Solvent Amount Unit] of [Solvent]. The resulting solution was spray-dried using a [Spray Dryer Model] with [Drying Gas] as the drying gas. Processing conditions were: air flow [Air Flow Speed] [Air Flow Unit], inlet temperature [Inlet Temperature] [Inlet Temperature Unit], aspiration rate [Aspiration Rate]% ([Aspiration Flow Rate] [Flow Rate Unit]), and feed rate [Feed Rate] [Feed Rate Unit]. The outlet temperature was [Outlet Temperature] [Outlet Temperature Unit].

T8. ULTRASOUND

Ultrasound-assisted synthesis was performed using [Amount of API] [Amount Unit of API] of [API] and [Amount of Coformer] [Amount Unit of Coformer] of [Coformer] in a ratio of [Part of Ratio API]:[Part of Ratio Coformer]. The components were mixed with [Amount of Solvent] [Solvent Amount Unit] of [Solvent]. The mixture was sonicated for [Sonication Time] [Sonication Time Unit] using an [Ultrasonic Apparatus] operated at [Mixing Frequency] [Mixing Frequency Unit] and [Mixing Temperature] [Mixing Temperature Unit], with a power output of [Power] [Power Unit]. The resulting solid was isolated by filtration and dried for [Drying Time] [Drying Time Unit] at [Drying Temperature] [Drying Temperature Unit].

T9. SUPERCRITICAL FLUID

Supercritical fluid synthesis was performed using [Amount of API] [Amount Unit of API] of [API] and [Amount of Coformer] [Amount Unit of Coformer] of [Coformer] in a ratio of [Part of Ratio API]:[Part of Ratio Coformer]. The components were mixed with [Amount of Solvent] [Solvent Amount Unit] of [Solvent]. The mixture was processed in a [Supercritical Apparatus] at [Pressure] [Pressure Unit] and [Mixing Temperature] [Mixing Temperature Unit], with a flow rate of [Flow Rate] [Flow Rate Unit]. The resulting co-crystal was dried for [Drying Time] [Drying Time Unit] at [Drying Temperature] [Drying Temperature Unit].

T10. ELECTROSPRAY

Electrospraying synthesis was conducted using [Amount of API] [Amount Unit of API] of [API] and [Amount of Coformer] [Amount Unit of Coformer] of [Coformer] in a ratio of [Part of Ratio API]:[Part of Ratio Coformer]. The components were dissolved in [Amount of Solvent] [Solvent Amount Unit] of [Solvent] and loaded into a [Syringe Volume] [Syringe Volume Unit] syringe. The prepared solution was electrosprayed using an [Electrospraying Apparatus]. The flow rate, voltage, and needle tip-to-collector distance were set to [Flow Rate] [Flow Rate Unit], [Voltage] [Voltage Unit], and [Tip to Collector Distance] [Tip to Collector Distance Unit], respectively. The deposited co-crystal was collected on the collector and dried for [Drying Time] [Drying Time Unit] at [Drying Temperature] [Drying Temperature Unit].

T11. RESONANT ACOUSTIC

Resonant acoustic synthesis was performed using [Amount of API] [Amount Unit of API] of [API] and [Amount of Coformer] [Amount Unit of Coformer] of [Coformer] in a ratio of [Part of Ratio API]:[Part of Ratio Coformer]. The components were combined with [Amount of Solvent] [Solvent Amount Unit] of [Solvent]. The reaction mixture was processed for [Reaction Time] [Reaction Time Unit] at [Reaction Times of Gravity Acceleration] [Reaction Times of Gravity Acceleration Unit] ×g and a frequency of [Reaction Frequency] [Reaction Frequency Unit]. The resulting co-crystal was dried for [Drying Time] [Drying Time Unit] at [Drying Temperature] [Drying Temperature Unit].

T12. LASER IRRADIATION

Laser irradiation synthesis was performed using [Amount of API] [Amount Unit of API] of [API] and [Amount of Coformer] [Amount Unit of Coformer] of [Coformer] in a ratio of [Part of Ratio API]:[Part of Ratio Coformer]. The components were mixed with [Amount of Solvent] [Solvent Amount Unit] of [Solvent]. The mixture was spread as a thin layer on [Foil Material] foil. Irradiation was performed using a [Laser

```
Apparatus] with a focal length of [Focal Length] [Focal Length Unit], operated at
[Power] [Power Unit] and a raster speed of [Raster Speed] [Raster Speed Unit].  The
resulting co-crystal was collected from the foil and dried for [Drying Time] [Drying
Time Unit] at [Drying Temperature] [Drying Temperature Unit].
```

A.3. Extraction prompts

The prompts presented below were used in the hierarchical extraction pipeline for co-crystal synthesis data. Each stage addresses a separate decision: (i) whether an article contains an extractable synthesis procedure, (ii) which synthesis method or methods are described, and (iii) which method-specific parameters should be extracted. This staged design reduces the number of irrelevant fields considered at each step and allows later prompts to operate on progressively more specific information.

A.3.1. STAGE 1: PROCEDURE DETECTION

The Stage 1 prompt was used to determine whether a full-text article contains an explicit co-crystal synthesis procedure. During prompting-strategy optimization, we tested three [Response formatss]:

- **S1-A:** If the article describes such techniques, respond with 1. If the article does not mention any specific techniques for co-crystal synthesis, respond with 0. Answer only 1 or 0, do not add anything else to your answer.

- **S1-B:** If the article contains a complete method for synthesizing a co-crystal with the name of the method and/or parameters, write the entire text that relates to the co-crystal synthesis method. If the article does not describe a complete method for the synthesis of a co-crystal, specify None in the answer.

- **S1-C:** If the article contains a complete co-crystal synthesis method with the name of the method and/or parameters, write their text separately for each molecular pair and the type of technique separately. If the article does not describe the complete method of crystal synthesis, write None.

The full Stage 1 prompt shown below contains the placeholder [Response format], which was instantiated according to the corresponding strategy.

```
You are a chemistry assistant specializing in co-crystals.

Your task is to analyze scientific articles and determine whether the article
describes a complete and explicit synthesis method specifically for a co-crystal (not
for a host molecule, not for analysis, and not just molecular interactions).

Inclusion criteria:

 1. The article must describe a specific method for synthesizing a co-crystal, such
    as grinding, contact formation, hot-melt extrusion, evaporation, cooling, slurry,
    antisolvent, spray drying, reactive cocrystallization, ultrasound, freeze drying,
    hydrothermal, supercritical fluid, microfluidic, electrospray, resonant acoustic,
    or laser irradiation.

 2. The article must also provide at least one key synthesis parameter, such as
    temperature, time, solvent, or the ratio of starting materials.

 3. Additionally, look for keywords indicating co-crystal formation, such as
    co-assembled, molar ratio N:N, two-component molecular complex, or inclusion
    complex.

 Do not consider the following as synthesis methods:

 1. Synthesis of a host molecule (this is **not** a co-crystal synthesis method).
    If the article describes the synthesis of a single molecular entity (not the
    combination of two or more molecules into a co-crystal), it must be ignored.

 2. Mentions of co-crystal synthesis without method details.  If the article
    states that a co-crystal was synthesized but does not provide a method name or
    parameters, it must be ignored.
```

```
3. Analysis methods, such as Powder X-ray diffraction (PXRD), Fourier-transform
   infrared spectroscopy (FTIR), Differential scanning calorimetry (DSC), Nuclear
   magnetic resonance (NMR), or Thermogravimetric analysis (TGA).

4. Descriptions of molecular interactions, e.g., hydrogen bonds or halogen bonds,
   must be ignored unless they are directly linked to the synthesis method.

5. Formation of another polymorphic form:  Formation of another polymorphic form of a
   crystal is not cocrystallization.

                            [Response format]
```

A.3.2. STAGE 2: METHOD CLASSIFICATION

The Stage 2 prompt was used to identify the synthesis method types described in the extracted procedure text. This classification determines which method-specific parameter extraction prompt is applied at Stage 3. During prompting-strategy optimization, we tested three variants of [Method context]:

- **S2-A:** The prompt included only the predefined list of synthesis method names.

- **S2-B:** The prompt included the predefined list of synthesis method names together with concise natural-language descriptions of the methods from Table 5.

- **S2-C:** The prompt included the predefined list of synthesis method names together with representative procedural templates from Appendix A.2.2.

The full Stage 2 prompt shown below contains the placeholder [Method context], which was instantiated according to the corresponding strategy.

```
You are a domain-specific chemical information extraction assistant.  You specialize
in the chemistry of cocrystals and their synthesis.  Your area of expertise includes
the analysis of types of co-crystal synthesis methods (not for a host molecule, not
for analysis, and not just molecular interactions).

Your task is to find the text of the synthesis technique for each co-crystal presented
in the article and identify the type of synthesis techniques that were used and
response should contain only the names of the methods from the List of techniques
in exact accordance with their spelling (no comments, no additional text).

List of techniques:  grinding, hot-melt extrusion, solution crystallization,
antisolvent, slurry, freeze drying, spray drying, ultrasound, supercritical fluid,
electrospray, resonant acoustic, laser irradiation.

                            [Method context]

Extraction rules:

1. Do not write the methodology again in the response if it already exists.

2. You must use technique names only from the List of techniques; do not name the
   techniques otherwise.

3. If the text is not in English, first translate it to English before analysis.

4. If the synthesis method does not correspond to any technique in the List of
   techniques, the answer must be strictly OTHER. Do not invent or suggest any
   additional technique name.

5. If several techniques were used, list them separated by commas.

6. It is very critical that you analyze each synthesis experiment separately to
   determine its type, without referring to other experiments in the paper.
```

7. Grinding includes both types of mechanochemical co-crystal synthesis: neat grinding and liquid-assisted grinding. You do not need to specify the type of grinding; just write grinding.

8. Grinding is considered only if: (i) it is the primary synthesis method, e.g., the co-crystal is obtained by mechanical grinding; and (ii) it is not merely used for pre-mixing.

9. The solution crystallization method considers three main processes: evaporation, cooling, and interface formation. If any of these processes are mentioned in the text, classify the method as solution crystallization.

10. If the text describes a multi-step process, select only the method that directly led to the formation of the co-crystal. Preliminary steps are not taken into account.

11. If a method is explicitly named in the text, choose it, even if there are hints at other techniques.

12. A response cannot contain OTHER together with other methods. If there is at least one matching method from the list, OTHER is not used.

13. Specify each type of technique only once.

Output examples:

    Output 1: grinding

    Output 2: hot-melt extrusion, solution crystallization

A.3.3. STAGE 3: METHOD-SPECIFIC PARAMETER EXTRACTION

You are a domain-specific chemical information extraction assistant. You specialize in the chemistry of cocrystals and their synthesis. Your area of expertise includes the analysis of co-crystal synthesis parameters.

Your task is to extract every mention of co-crystal synthesis parameters for grinding synthesis methodology from a scientific article and output a response with formatting as in the example (no comments, no additional text).

Fields for each object:

  – Type of Synthesis (string): one of ''liquid-assisted grinding'', or ''neat grinding''. The key step in synthesis that leads to the formation of a cocrystal.

  – Formula (string): name of cocrystal as cited in the text, e.g., ''CAR-HCT'', ''DMZ-SAC''.

  – API (string): full-name of active pharmaceutical ingredient, e.g., ''caffeine'', ''betulin''.

  – Coformer (string): full-name of used coformer, e.g., ''citric acid'', ''adipic acid''.

  – Part of Ratio API (numeric): part of API in API-coformer ratio.

  – Part of Ratio Coformer (numeric): part of Coformer in API-coformer ratio.

  – Amount of API (numeric): mass or amount of API.

  – Amount of Coformer (numeric): mass or amount of Coformer.

  – Amount Unit (string): unit of measurement, e.g., ''mmol'', ''mg''.

  – Solvent (string): full-name of used Solvent, e.g., ''water'', ''methanol''.

- Amount of Solvent (numeric): amount showing how much of the Solvent has been used.

- Solvent Amount Unit (string): Solvent unit of measurement, e.g., ``mL'', ``µL''.

- Second Solvent (string): full-name of used Second Solvent, e.g., ``ethanol'', ``dioxane''.

- Amount of Second Solvent (numeric): amount showing how much of the Second Solvent has been used.

- Third Solvent (string): full-name of used Third Solvent, e.g., ``methanol'', ``ethanol''.

- Amount of Third Solvent (numeric): amount showing how much of the Third Solvent has been used.

- Mixing Apparatus (string): Mixing Apparatus that has been used for grinding, e.g., ``ball mill'', ``mortar and pestle''.

- Mixing Frequency (numeric): frequency of the mixing process if that's what the device implies.

- Frequency Unit (string): Mixing Frequency unit of measurement, e.g., ``Hz'', ``rpm''.

- Mixing Time (numeric): duration of Mixing process.

- Mixing Time Unit (string): Mixing Time unit of measurement, e.g., ``minute''.

- Mixing Temperature (numeric): Temperature of Mixing.

- Mixing Temperature Unit (string): Mixing Temperature unit of measurement, e.g., ``°C''.

- Time of Drying (numeric): duration of the Drying process of the co-crystal after grinding.

- Drying Time Unit (string): Time of Drying unit of measurement, e.g., ``hour''.

- Temperature of Drying (numeric): Temperature of Drying process of the co-crystal after grinding.

- Drying Temperature Unit (string): Temperature of Drying unit of measurement, e.g., ``°C''.

Extraction rules:

1. It is crucial for you to accurately and comprehensively extract each synthesis experiment separately, without referring to other experiments in the article.

2. Do **not** filter, group, summarize, or deduplicate. Include repeated mentions and duplicates if they occur in different contexts.

3. Keep numerical values in the units reported in the article.

4. In the fields Part of Ratio API and Part of Ratio Coformer, write exactly the values reported in the article. Do not convert them to fractions of a unit.

5. If you cannot find a required field for an object, re-check the context; if it is still absent, set that field's value to ``NOT_DETECTED''.

6. For missing numeric fields, use the string ``NOT_DETECTED''.

7. When writing a parameter value, provide only one value per field.

8. The example below shows only two extracted samples; however, your output should contain **all** mentions of grinding synthesis methodology present in the article.

9. Depending on the type of synthesis, some parameters may be irrelevant. For ``neat grinding'', set solvent-related and drying-related fields to ``NOT_DETECTED'' if they are not reported, including Solvent, Amount of Solvent, Solvent Amount Unit, Second Solvent, Amount of Second Solvent, Third Solvent, Amount of Third Solvent, Time of Drying, Drying Time Unit, Temperature of Drying, and Drying Temperature Unit.

Output example:

```
[
    {
        "Type of Synthesis": "liquid-assisted grinding",
        "Formula": "ASP-CIT",
        "API": "aspirin",
        "Coformer": "citric acid",
        "Part of Ratio API": 4,
        "Part of Ratio Coformer": 1,
        "Amount of API": 1.53,
        "Amount of Coformer": 0.3,
        "Amount Unit": "g",
        "Solvent": "methanol",
        "Amount of Solvent": 0.1,
        "Solvent Amount Unit": "ml",
        "Second Solvent": "acetone",
        "Amount of Second Solvent": 0.4,
        "Third Solvent": "ethyl acetate",
        "Amount of Third Solvent": 0.5,
        "Mixing Apparatus": "planetary mill",
        "Mixing Frequency": 30,
        "Frequency Unit": "hz",
        "Mixing Time": 25,
        "Mixing Time Unit": "min",
        "Mixing Temperature": 80,
        "Mixing Temperature Unit": "C",
        "Time of Drying": 8,
        "Drying Time Unit": "h",
        "Temperature of Drying": 40,
        "Drying Temperature Unit": "C"
    },
    {
        "Type of Synthesis": "neat grinding",
        "Formula": "ACP-MAL",
        "API": "acetaminophen",
        "Coformer": "malic acid",
        "Part of Ratio API": 1,
        "Part of Ratio Coformer": 1,
        "Amount of API": 1.2,
        "Amount of Coformer": 0.4,
        "Amount Unit": "mg",
        "Solvent": "NOT_DETECTED",
        "Amount of Solvent": "NOT_DETECTED",
        "Solvent Amount Unit": "NOT_DETECTED",
        "Second Solvent": "NOT_DETECTED",
        "Amount of Second Solvent": "NOT_DETECTED",
        "Third Solvent": "NOT_DETECTED",
        "Amount of Third Solvent": "NOT_DETECTED",
        "Mixing Apparatus": "MM400, Retsch, Germany",
        "Mixing Frequency": 250,
        "Frequency Unit": "rpm",
        "Mixing Time": 1,
        "Mixing Time Unit": "h",
        "Mixing Temperature": 60,
        "Mixing Temperature Unit": "C",
```

```
        "Time of Drying": 12,
        "Drying Time Unit": "h",
        "Temperature of Drying": 40,
        "Drying Temperature Unit": "C"
    }
]

Final output constraints:  Generate the output as pure JSON only.
Do not include:

  - code block markers, such as ```json or ```,

  - explanatory text,

  - the phrase ``Final answer'',

  - any other non-JSON content.
```

A.4. Extraction metrics

We evaluated the extraction pipeline separately for the three stages, since each stage produces a different type of output: procedure presence, method labels, and structured parameter values. Unless stated otherwise, precision, recall, F1-score, and accuracy were computed as:

$$\text{Precision} = \frac{TP}{TP+FP}, \qquad \text{Recall} = \frac{TP}{TP+FN},$$

$$\text{F1} = \frac{2 \cdot \text{Precision} \cdot \text{Recall}}{\text{Precision} + \text{Recall}}, \qquad \text{Accuracy} = \frac{TP+TN}{TP+TN+FP+FN}.$$

Stage 1: Procedure detection.   This stage was evaluated as binary classification at the DOI level. Each article was labeled as either containing or not containing an explicit co-crystal synthesis procedure. Empty or missing-value responses were treated as negative predictions; all other responses were treated as positive.

Stage 2: Method classification.   This stage was evaluated by comparing the predicted set of synthesis methods with the gold method set for each DOI. Aggregate scores were computed over complete method-set labels using weighted precision, recall, and F1-score. We also computed per-method one-vs-rest scores for detailed analysis.

Stage 3: Parameter extraction.   This stage was evaluated at the level of synthesis records and extracted parameters. For each record and parameter, predicted and gold values were compared as multisets, preserving duplicate values. The error counts were defined as follows:

- **True positives (TP):** values that appear in both the predicted and gold multisets.
- **False positives (FP):** predicted values that are absent from the gold multiset or appear too many times.
- **False negatives (FN):** gold values that are missing from the prediction or appear too few times.

Precision, recall, and F1-score follow the definitions above, while Stage 3 accuracy is defined as:

$$\text{Accuracy}_{\text{S3}} = \frac{TP}{TP+FP+FN}.$$

Thus, Stage 3 accuracy measures value-level overlap rather than standard classification accuracy with true negatives. Final Stage 3 scores were obtained by averaging parameter-level scores across all evaluated parameters.

A.5. Per-method classification results

To complement the aggregate Stage 2 comparison, we also examined method-classification performance separately for each of the 12 synthesis types using a one-vs-rest evaluation setup. This analysis reveals substantial variation across

methods, reflecting differences in how explicitly individual techniques are described and how strongly they overlap with other procedure types. In particular, methods with more distinctive procedural signatures are classified more reliably, whereas closely related or less frequently represented methods are more challenging. Figure 5 summarizes the per-method results in terms of F1-score and accuracy.

Figure 5. Per-method Stage 2 classification across the evaluated LLMs. One-vs-rest [A] F1-score and [B] accuracy for each of the 12 synthesis method types.

A.6. Per-parameter extraction results

Figure 6 reports per-parameter precision and recall at Stage 3 for all four evaluated models on the LAG validation set. Categorical and numerical parameters are shown separately. Each chart has one axis per parameter, with the outer boundary corresponding to a perfect score of 1.0.

A.7. Error analysis

We manually annotated the validation outputs of `Gemini-2.5-Flash` to identify dominant error types and their primary causes across the three extraction stages (Table 6). Categories are standardized across stages: Omission denotes information present in the article but missed by the model, Fabrication denotes unsupported model output, Substitution denotes extraction of the correct target with incorrect content, and Evaluation artefact denotes a mismatch caused by the scoring procedure rather than a substantive extraction error. Total counts are 5 errors at Stage 1, 20 at Stage 2, and 898 at Stage 3.

Stage 1. Omissions (60%) are caused by non-standard terminology: the target material is described as a solvate, polymorph, or molecular complex rather than as a co-crystal, and the article is treated as out of scope. Fabrications (40%) recover procedure-shaped text from articles that report only a crystal structure.

Stage 2. Omissions (55%) are caused by methods reported only in additional, comparative, or control experiments rather than in the main synthesis section. Substitutions (25%) arise exclusively from boundary cases between closely related methods. Fabrications (20%) infer a method from a brief contextual mention rather than from an actual experimental description.

Stage 3. The majority of Stage 3 errors (67.6%) are evaluation artefacts from strict multiset-based comparison, where the substantive value is correct but formatting or placeholder variants are counted as mismatches. Among genuine errors, fabrications (13.0%) are the most informative: the model returns values that exist elsewhere in the article, often procedural details such as mixing time, frequency, or apparatus, but assigns them to the wrong record. Omissions (11.6%) are mostly complete misses of present fields, and substitutions (7.8%) involve the correct field with wrong content, most often solvent

Figure 6. Per-parameter Stage 3 extraction on the LAG validation set. Precision and recall for categorical and numerical parameters, shown for [A] `GPT-4o-mini`, [B] `Gemma-3-27B`, [C] `Llama-3.3-70B`, and [D] `Gemini-2.5-Flash`.

identity or quantity.

A.8. Parameter Families

For interpretability, individual parameters were grouped into eight families covering related physicochemical or procedural roles. Figure 2B reports aggregated statistics at the family level, while the per-parameter view is given in Figure 7. The full mapping is shown in Table 7.

Figure 7. Per-parameter fill rates across the three downstream datasets after quality filtering. Unit fields are omitted; structurally absent values are counted as empty, and blank cells mark parameters not defined for a given method.

A.9. Data Partitioning Details

Splits were constructed at the level of connected components rather than individual records. Two records were linked if they shared either the same DOI or the same canonical API/coformer pair, and entire components were assigned to a single split. This guarantees that no publication or molecular pair appears in both train and test sets. For each method, test components were selected greedily until reaching 100 records, followed by a repair step that adjusted assignments without breaking components.

Table 6. Extraction error analysis across the three stages. Dominant error types and primary causes from manual annotation of `Gemini-2.5-Flash` outputs. % of stage is the share of errors at each stage; % of type, the share of the dominant cause within a type.

Stage	Error type	% of stage	Primary cause	% of type
1	Omission	60.0	The compound is described using alternative terminology, e.g., solvate or polymorph, rather than explicitly as a co-crystal.	66.7
	Fabrication	40.0	The article reports a crystal structure without an explicit synthesis procedure, yet the model returns a fabricated procedure text.	100.0
2	Omission	55.0	The method appears only in an additional, comparative, or control experiment rather than in the main synthesis section.	72.7
	Substitution	25.0	The case lies near the boundary between two method types, e.g., solution crystallization and slurry.	100.0
	Fabrication	20.0	The model infers a method from a brief textual mention or contextual cue, although that method was not actually used.	50.0
3	Evaluation artefact	67.6	Strict multiset comparison treats formatting and placeholder variants as mismatches even when the substantive value is correct.	68.4
	Fabrication	13.0	The model produces a parameter value that is not supported by the article text, often a procedural detail belonging to a different record.	81.2
	Omission	11.6	The model fails to extract a parameter value that is present in the article text.	67.3
	Substitution	7.8	The model extracts the correct field but with incorrect content, e.g., wrong solvent name or quantity.	80.0

Table 7. Parameter-family definitions. Mapping of individual extraction parameters to the eight interpretability families used for the coverage analyses in Figures 2 and 7.

Family	Num.	Parameters
Ratio	2	Part of Ratio API / Part of Ratio Coformer
Amounts	5	Amount of API / Amount of Coformer / Amount of Solvent / Amount of Second and Third Solvent
Solvent	1	Solvent
Additional solvent	2	Second Solvent / Third Solvent
Time	4	Mixing Time / Drying Time / Evaporation Time / Time of Cooling or Evaporation
Temperature	3	Mixing Temperature / Drying Temperature / Evaporation Temperature
Frequency	1	Mixing Frequency
Setup	3	Mixing Apparatus / Covering Method / Description of Holes

Test components were chosen to maximize DOI diversity within the 100-record budget, since a more varied test set yields more representative metrics. As a side effect, the mean number of records per DOI is lower in test than in train (Table 8), which should be kept in mind when interpreting per-record metrics.

A.10. Imputation Details

Imputation produced fully populated tables from which coherent textual procedure descriptions were generated as prediction templates, rather than ground-truth values. Two parallel artefacts were therefore maintained: imputed tables for template generation, and pre-imputation test reference snapshots for honest metric computation. All statistics (unit defaults, modes, group-wise modes, fallbacks) were estimated exclusively on the training split and applied independently to train and test. Values flagged as not detected and empty strings were treated as unknowns eligible for filling, whereas structurally absent values were preserved and never replaced by a statistical mode. Per-method deterministic rules assigned the structural absence marker to fields that should not exist in a given procedure.

Categorical fields. Filling proceeded as a cascade from most to least specific group, with the global training mode and a fixed default as final fallbacks. Groups were defined by chemical context: joint API and coformer pair, then API alone, then coformer alone.

Table 8. Data-partitioning statistics for the three downstream datasets. Train and test splits are reported by DOIs and records; Records/DOI is the mean number of records per unique DOI within each split.

Dataset	Train			Test		
	DOIs	Records	Rec./DOI	DOIs	Records	Rec./DOI
Solution crystallization	2,814	8,172	2.90	52	100	1.92
Grinding	434	2,622	6.04	51	100	1.96
Slurry	77	493	6.40	45	100	2.22

Numerical fields. Domain rules were applied first, such as recovering an API or coformer mass from the reported stoichiometric ratio. Remaining gaps were filled by group-wise modes, with groups chosen per parameter family, for example:

- amounts: API, coformer, and ratio;

- solvent volume: chemical pair and solvent;

- mixing parameters: chemical pair and apparatus;

- temperature and time: chemical pair and solvent.

Global training modes were used as a final fallback. For columns populated in at most 1% of the training rows, a fixed default was assigned instead, preventing reconstruction from one or two incidental observations.

B. Synthesis Condition Prediction

B.1. Pair-level similarity

Few-shot examples were retrieved from the training set within the same synthesis method type. Pair-level similarity was computed symmetrically over both molecules to respect the unordered nature of a co-crystal: for query (a, b) and candidate (c, d),

$$S\big((a,b),(c,d)\big) = \max\big(\tfrac{1}{2}(T(a,c) + T(b,d)), \tfrac{1}{2}(T(a,d) + T(b,c))\big),$$

where T is the Tanimoto similarity between Morgan fingerprints (radius 2, 2048 bits).

B.2. Prediction prompts

We used a unified prompt for synthesis parameter prediction across all experiments. In the few-shot setting ($k \in \{1, 3, 5, 10, 20\}$), the model receives retrieved examples of the same synthesis method type, a query API/coformer pair, and a synthesis template with missing parameters.

For zero-shot experiments, we used the same prompt but removed the `Examples` section and the corresponding instructions related to using examples. All other instructions and formatting constraints remained unchanged. The full few-shot prediction prompt is shown below.

```
You will be given numbered examples of cocrystal synthesis methods of the SAME method
type, a query API/coformer pair, and a method template.

Your task is to fill every bracketed placeholder in the template to produce one
best-supported synthesis method for the query pair.

Use the examples as strong evidence for structure, slot usage, value ranges, units,
formatting, and typical procedures.  The goal is to predict synthesis parameters
that are appropriate for this specific query API/coformer pair, not to copy one
example mechanically.  Similarity to an example does not guarantee that every value
in that example is suitable for the query pair.  Use general chemistry and synthesis
knowledge only when the examples are sparse, inconsistent, incomplete, clearly not
transferable for a specific slot, or when it is needed to reject an impossible or
clearly unsuitable value.  Do not use general chemistry to override a clear local
pattern from the most chemically and procedurally similar examples, especially for
```

exact numeric values.
Rules:

  - Work slot by slot, not example by example.

  - Prefer values supported by the most chemically and procedurally similar examples
    for that specific slot.

  - Prefer the nearest local pattern, but do not copy one whole example blindly.

  - Do not average across all examples when a smaller set of closer examples gives a
    clearer signal.

  - Treat categorical value choice, numeric value choice, NO_VALUE choice, and
    formatting choice as separate decisions.

  - Replace [API] with the query API SMILES exactly.

  - Replace [Coformer] with the query coformer SMILES exactly.

  - Replace every other [placeholder] with exactly one concrete value or NO_VALUE, and
    leave no placeholder unreplaced.

  - Use NO_VALUE only when the field is truly not applicable, not when uncertain or
    simply unsupported by some examples.

  - Presence of an optional field requires positive support. Do not add a solvent,
    second solvent, additive, apparatus, mixing frequency, temperature, time, or
    amount just because such fields appear often overall.

  - If a quantity is NO_VALUE, its unit must also be NO_VALUE.

  - If a solvent, second solvent, additive, or apparatus is NO_VALUE, directly
    dependent amount and unit fields must also be NO_VALUE unless the template clearly
    separates them.

  - Keep dependent fields mutually consistent.

Rules for sensitive numeric slots:

  - Choose ratio slots from ratio evidence, not from mass evidence. Do not derive
    [Part of Ratio API] or [Part of Ratio Coformer] from [Amount of API] and [Amount
    of Coformer].

  - Choose amount slots from amount patterns, not from ratio slots. Do not force
    masses to numerically match stoichiometric parts.

  - Keep numbers as plain numbers, units only in unit placeholders, and ratio parts as
    separate numbers unless the template explicitly requires another format.

  - Follow the local numeric scale, precision, and granularity of the nearest relevant
    examples.

  - Do not drift toward convenient anchor values such as round or highly frequent
    numbers unless they are strongly supported by the nearest examples.

  - Keep time, temperature, frequency, amount, and ratio fields separate. Do not
    substitute one for another or infer one from another unless the examples clearly
    establish that relation.

  - Do not invent unusual or highly specific numeric values without strong local
    support.

Rules for solvent-like and optional slots:

  - Solvent identity is highly local. Do not choose a solvent because it is common
    overall or chemically plausible in general.

- Choose the solvent best supported by the nearest relevant examples for this query pair.

- Second solvent or anti-solvent fields require positive local evidence of a real two-solvent pattern. Do not default them to a common solvent, and do not default them to NO_VALUE when the nearest relevant examples show a real second-solvent pattern.

- For optional solvent in grinding-like procedures, use a solvent only when the nearest examples support liquid assistance or there is a clear reason the query pair likely requires it; otherwise use NO_VALUE and keep dependent solvent amount/unit fields consistent.

- For mixing frequency, apparatus, and similar optional operation fields, prefer NO_VALUE unless the nearest examples give positive support that the field is genuinely used for this kind of query/template.

Method-type-specific priorities:

- If method type is grinding: be conservative about adding solvent and mixing frequency. Do not infer liquid-assisted grinding unless the nearest grinding examples support it or there is a clear reason the query pair likely needs it.

- If method type is slurry: do not infer stoichiometric ratio from mg values. Treat solvent identity, mixing time, and mixing temperature as local slurry-pattern choices. Do not collapse to a globally common solvent.

- If method type is solution_crystallization: solvent and second solvent are highly local and should usually be chosen from the nearest solution examples, not from global defaults such as common alcohols. Distinguish mixing time from cooling/evaporation time; do not substitute one for the other.

Formatting rules:

- Use example unit and style conventions whenever possible.

- If a slot behaves like a closed or near-closed vocabulary in the examples, use the canonical spelling/style used in the examples.

- Keep the template text otherwise identical: same wording, order, and punctuation. Only replace bracketed placeholders.

- Every placeholder must be filled exactly once.

- Output exactly one paragraph and nothing except:

  [BEGIN_METHOD]
  <filled template>
  [END_METHOD]

Before finalizing internally, verify:

- The method is intended for the specific query API/coformer pair, not just for a similar example pair.

- [API] and [Coformer] were copied exactly from the query.

- Every placeholder was replaced exactly once.

- NO_VALUE is used only for true non-applicability.

- Dependent fields are consistent with NO_VALUE.

- Ratios were chosen from ratio evidence, not from masses.

- Solvent and second-solvent choices follow the nearest relevant local examples rather than global common values.

```
    - Numeric values stay anchored to local example scale rather than chemistry-driven
      invention.

  Method type:  [Method type]

  Examples:  [Examples]

  Query:
  API SMILES: [API SMILES]
  Coformer SMILES: [Coformer SMILES]

  Template to fill:  [Template]
```

B.3. Solvent Families

A curated synonym dictionary normalizes raw solvent labels to 475 canonical names. Since exact-match accuracy over 475 classes is dominated by long-tail confusions between solvents of similar chemical nature, we additionally group canonical solvents into 13 chemical families. Table 9 lists the families with their descriptions and the number of canonical solvents covered.

Table 9. Solvent families used for categorical solvent aggregation. The 13 chemical families are sorted by the number of canonical solvent names covered.

Family	Num.	Description
Alcohols	97	Protic alcohols, polyols, and phenols
Amines	80	Amines and N-heteroaromatics, basic and coordinating
Hydrocarbons	58	Aliphatic and aromatic hydrocarbons, low polarity
Esters	50	Esters, carbonates, and lactones
Halogenated	48	Halogenated aliphatics and aromatics, polarizable
Carbonyls	44	Ketones and aldehydes, polar aprotic
Ethers	30	Ethers, cyclic ethers, and glymes
Acids	25	Acidic and protic media
Amides	17	Amides, ureas, and lactams, strong solvators
Special	14	Oxidants, gases, and reactive or unusual cases
Nitriles	8	Nitriles, polar aprotic
Sulfur	3	Sulfur-containing solvents such as DMSO and sulfolane
Water	1	Aqueous media, very polar with strong H-bonding

B.4. Numerical tolerance bins

For numerical-parameter evaluation, we additionally used tolerance bins to count near-correct predictions as correct when they fall into the same experimentally meaningful range. The bin definitions were generated from the imputed training data only, using the train split to avoid using test-set information.

For each method, bins were fitted for the numerical columns when the corresponding column was present in the method table. Ratio parameters were treated globally by pooling values across all methods. If a parameter had at most n_{bins} unique values, bins were created from sorted unique values using midpoint boundaries. Otherwise, bins were created using quantile binning and duplicate boundaries removed. All bin boundaries were rounded to three significant figures.

During tolerance-based scoring, a value is assigned to a bin if it satisfies $lo \leq x < hi$; the final bin is open-ended above. A prediction is counted as tolerance-correct when the gold and predicted values fall into the same bin.

B.5. Solvent Confusion Analysis

Figure 8 provides a more detailed view of solvent-prediction errors. The exact-solvent confusion matrix shows that errors are concentrated among high-frequency solvents and the long-tail OTHER class, reflecting the high cardinality and local nature of solvent choice.

The family-level confusion matrix indicates that many mistakes remain chemically structured: predictions often collapse

toward broad, frequently observed families, especially alcohols, rather than arbitrary solvent classes. These results explain why solvent prediction remains challenging despite the use of retrieved few-shot precedents, and support reporting both exact-solvent and solvent-family metrics.

Figure 8. Solvent confusion analysis for solvent prediction. [A] Exact-label confusion matrix for the 12 most frequent solvents, with all remaining labels grouped as OTHER. [B] Confusion matrix after aggregation into solvent families.

B.6. Per-Method Prediction Results

Tables 10–12 report per-parameter Accuracy and weighted F1 for Gemini-2.5-Flash (k=3) across two independent runs. Reported values are mean $\pm$ standard deviation. Fill Rate denotes the proportion of test records containing a valid ground-truth value for a given parameter.

Overall, the results show that the model is consistently more reliable for predicting stoichiometric ratios (Part of Ratio API / Coformer), where performance remains high across all synthesis methods ($F1 > 0.77$). In contrast, absolute quantity parameters (e.g., Amount of API, Amount of Solvent) remain difficult, typically yielding $F1 < 0.30$, likely because these values depend on experiment-specific conditions and are less standardized across publications.

Categorical solvent prediction remains particularly challenging for Slurry and Solution Crystallization. One likely reason is that the model does not have explicit knowledge of the full solvent space and tends to over-rely on solvents observed in the retrieved in-context examples. As a result, predictions become biased toward a narrow subset of frequently occurring solvents, reducing generalization when the correct solvent category is rare or absent from the provided examples.

The low standard deviations across runs indicate that the model behavior is stable, suggesting that these limitations are systematic rather than caused by sampling variability.

C. Agentic System Details

C.1. Domain Tool Groups

The agentic system uses four tool groups, which are shared by the Single-Agent architecture and distributed across specialists in the Multi-Agent architecture.

- **Common Tools** normalize chemical inputs, resolve molecule names, canonicalize structures, identify molecular analogues, and convert structured synthesis parameters into method text.
- **Database Tools** query COSYN-DB for experimental precedents, exact or similar API / coformer pairs, parameter-constrained records, component ratios, and common synthesis conditions.
- **Article-analysis Tools** load scientific articles from DOI, PDF, or TXT inputs, detect whether a synthesis procedure is

Table 10. Per-parameter prediction on the Grinding test set. Fill rate, weighted F1, and accuracy for `Gemini-2.5-Flash` ($k=3$); fill rate is the percentage of test records with an observed gold value.

Type	Parameter	Fill %	F1	Accuracy
Categorical	Solvent	95	0.34 ±0.00	0.35 ±0.00
	Mixing Apparatus	89	0.57 ±0.00	0.57 ±0.00
Numerical	Part of Ratio API	100	0.84 ±0.00	0.82 ±0.01
	Part of Ratio Coformer	100	0.86 ±0.00	0.90 ±0.00
	Amount of API	47	0.20 ±0.00	0.21 ±0.00
	Amount of Coformer	48	0.29 ±0.00	0.27 ±0.00
	Amount of Solvent	63	0.42 ±0.01	0.39 ±0.01
	Mixing Frequency	71	0.46 ±0.01	0.42 ±0.01
	Mixing Time	88	0.31 ±0.00	0.34 ±0.00
Categorical avg.		92.0	0.46 ±0.00	0.46 ±0.01
Numerical avg.		73.9	0.48 ±0.01	0.48 ±0.01
Overall		78.0	**0.48** ±0.01	**0.47** ±0.01

Table 11. Per-parameter prediction on the Solution-crystallization test set. Fill rate, weighted F1, and accuracy for `Gemini-2.5-Flash` ($k=3$); fill rate is the percentage of test records with an observed gold value.

Type	Parameter	Fill %	F1	Accuracy
Categorical	Solvent	98	0.17 ±0.01	0.17 ±0.01
	Second Solvent	100	0.53 ±0.00	0.57 ±0.01
Numerical	Part of Ratio API	100	0.82 ±0.00	0.84 ±0.01
	Part of Ratio Coformer	100	0.77 ±0.00	0.76 ±0.01
	Amount of API	61	0.12 ±0.00	0.11 ±0.00
	Amount of Coformer	61	0.19 ±0.01	0.17 ±0.01
	Amount of Solvent	57	0.20 ±0.00	0.21 ±0.00
	Time of Cooling or Evaporation	44	0.25 ±0.01	0.24 ±0.02
	Mixing Temperature	32	0.46 ±0.00	0.47 ±0.00
	Mixing Time	27	0.22 ±0.00	0.26 ±0.00
Categorical avg.		99.0	0.35 ±0.01	0.37 ±0.01
Numerical avg.		60.3	0.38 ±0.01	0.38 ±0.01
Overall		68.0	**0.37** ±0.01	**0.38** ±0.01

present, classify the synthesis method, and extract method-specific parameters.

- **Prediction Tools** generate candidate synthesis procedures for new molecular pairs, select related examples, extract parameters from generated text, and validate explicit user constraints.

C.2. Agentic System Prompts

This section summarizes the system prompts used for the non-agentic baselines, the Single-Agent system, and the Multi-Agent system. To avoid repeating identical grounding and abstention rules across prompts, we report the prompts as compact templates with explicitly marked variable parts. The prompts differ primarily in the evidence available to the model, the allowed tool groups, and the routing structure.

C.2.1. BASELINE PROMPTS

We evaluated two non-agentic baselines. The LLM-Only baseline receives only the user question and must answer from the model's internal knowledge. The Context-Augmented LLM additionally receives task-relevant static context, such as article text or database-derived evidence, but it cannot iteratively call tools or decompose the task through tool use. The corresponding prompt-field instantiations are summarized in Table 13.

Table 12. Per-parameter prediction on the Slurry test set. Fill rate, weighted F1, and accuracy for `Gemini-2.5-Flash` (k=3); fill rate is the percentage of test records with an observed gold value.

Type	Parameter	Fill %	F1	Accuracy
Categorical	Solvent	100	0.15 ±0.00	0.16 ±0.00
Numerical	Part of Ratio API	100	0.80 ±0.00	0.84 ±0.01
	Part of Ratio Coformer	100	0.88 ±0.00	0.85 ±0.00
	Amount of API	58	0.22 ±0.00	0.21 ±0.00
	Amount of Coformer	59	0.19 ±0.00	0.24 ±0.00
	Amount of Solvent	59	0.16 ±0.00	0.19 ±0.00
	Mixing Time	83	0.25 ±0.01	0.27 ±0.02
	Mixing Temperature	34	0.48 ±0.01	0.41 ±0.00
Categorical avg.		100.0	0.15 ±0.00	0.16 ±0.00
Numerical avg.		70.4	0.43 ±0.01	0.43 ±0.01
Overall		74.1	**0.39** ±0.01	**0.39** ±0.01

Table 13. Baseline prompt configuration. Prompt-field instantiations contrasting the LLM-Only and Context-Augmented non-agentic baselines, which differ in the evidence made available to the model.

Prompt field	LLM-Only	Context-Augmented LLM
Baseline role	An expert assistant for co-crystal synthesis.	A context-grounded assistant for co-crystal synthesis.
Available evidence	Only the user question and the model's own chemistry knowledge.	The user question and static task-relevant context provided with the query.
Evidence-use instruction	Answer from general chemistry knowledge, but do not present the answer as article- or database-supported.	Reason directly from the provided context, but do not infer beyond what the context supports.

```
You are [Baseline role] helping with questions about co-crystal synthesis.
Your job is to answer the user's request using [Available evidence].
Core behavior:

  - Determine which parts of the request can be answered from the available evidence.

  - Answer as fully as possible while clearly marking uncertainty.

  - Do not claim that an article was analyzed, a database was searched, or a
    prediction was validated unless such evidence is explicitly available.

  - Clearly distinguish evidence-based conclusions from general-knowledge reasoning.

Evidence policy:

  - [Evidence-use instruction]

  - If evidence is insufficient for part of the request, state the limitation and
    provide a cautious best-effort answer when appropriate.

  - Do not invent article-specific facts, database-wide statistics, retrieved
    examples, or validated synthesis predictions.

Final answer policy:

  - Provide a concise user-facing answer.

  - For yes/no questions, begin the relevant answer section with an explicit yes or no
    when possible.

  - If the answer is not evidence-backed, make this clear in the wording.
```

C.2.2. SINGLE-AGENT PROMPT

The Single-Agent system uses one controller to decompose the request, select tools, and produce the final answer. It has access to the four domain tool groups defined in Appendix C.1. The detailed tool-selection rules are omitted for brevity and represented as a placeholder in the template below.

```
You are a single-agent assistant helping with questions about co-crystal synthesis.

Your job is to answer the user's request by using the available tools and returning a
grounded final answer.  You may call tools in any order and as many times as needed,
within the step limit.

Core behavior:

  - First determine which subtasks are required by the request.

  - Classify each subtask as article analysis, database retrieval, synthesis
    prediction, or background answering.

  - Keep track of which subtasks are completed, pending, unsupported, or abstained.

  - Do not finalize until every required tool-backed subtask has been completed or
    explicitly marked impossible.

  - If the request combines multiple subtasks, resolve each one separately and then
    merge the results into a single final answer.

  - Avoid redundant tool calls unless new information, a refined constraint, or a
    failed prior attempt justifies them.

                              [Tool policy]

Grounding policy:

  - Use tool outputs as the primary evidence source.

  - Report article facts, database examples, retrieved conditions, and
    prediction-derived details only when they are supported by tool outputs.

  - Do not invent article contents, database matches, retrieved examples, or validated
    predictions.

  - Clearly distinguish tool-grounded conclusions from general-knowledge reasoning.

Failure and abstention behavior:

  - If input is insufficient for a tool, ask for the missing field when appropriate or
    abstain for that subtask.

  - If one part of a multi-part request fails, continue solving the remaining parts
    whenever possible.

  - Do not turn a partial failure into a full failure if other required subtasks can
    still be completed.

  - Clearly mark unsupported or abstained parts in the final response.

Final answer policy:

  - Provide a concise user-facing answer.

  - For yes/no questions, begin the relevant answer section with an explicit yes or
    no.

  - If the user asked for a synthesis method, procedure, or protocol, present it as
    readable method text rather than only as raw parameters unless structured fields
    were explicitly requested.

  - Keep the final answer grounded, complete, and aligned with the resolved subtasks.
```

C.2.3. ORCHESTRATOR PROMPT

The orchestrator controls the hierarchical Multi-Agent workflow. It does not call domain tools directly. Instead, it decomposes the user request, selects the appropriate specialist, and writes the final answer after the required specialist reports are available.

The coordination guardrails require the orchestrator to preserve the user's original constraints, treat multiple constraints as conjunctive, and use only grounded specialist outputs in the final answer. Failed, partial, empty, or unsupported specialist results must not be presented as completed evidence. If a requested item cannot be fully supported, the answer must state the unsupported part rather than fill it with non-matching results.

```
You are the orchestrator for a hierarchical assistant that helps with questions about
co-crystal synthesis.

You do not call domain tools yourself.  Your job is to decompose the user request,
choose the next specialist, and produce the final answer after specialists have
supplied enough evidence.

Available specialists:  article_agent, db_agent, prediction_agent.

Routing policy:

  - Use article_agent for questions about a specific article, PDF, TXT file, DOI,
    extracted synthesis parameters, method presence, method type, or article evidence.

  - Use db_agent for questions asking to find database examples, database conditions,
    common solvents/apparatus/times, exact API/coformer pairs, ratios, or similar
    precedents.

  - Use prediction_agent for questions asking to predict, propose, generate, or
    validate a new synthesis method.

  - If multiple capabilities are needed, call one specialist at a time and then route
    to the next missing specialist.

  - Do not call a specialist again unless the previous output was insufficient,
    failed, or a refined follow-up is needed.

  - Finalize only when every required part is answered or explicitly
    impossible/abstained.

                            [Coordination guardrails]
Return only valid JSON with this schema:
{
  "next_agent": "article_agent" | "db_agent" | "prediction_agent" | "final",
  "specialist_task": "short concrete instruction for the selected specialist,
                      or empty string for final",
  "final_answer": "final user-facing answer if next_agent is final,
                   otherwise empty string",
  "reason": "short routing reason"
}
```

C.2.4. SHARED SPECIALIST PROMPT SKELETON

The Multi-Agent system uses three specialist roles: article_agent, db_agent, and prediction_agent. Their specialist-specific policies follow the tool grouping defined in Appendix C.1.

```
The Multi-Agent system uses three specialist roles:  article_agent, db_agent, and
prediction_agent.  Their specialist-specific policies are derived from the logical
tool grouping defined in Appendix C.1.
```

```
                         [Specialist-specific policy]

Constraint policy:

  • Preserve the original user request and all constraints from the orchestrator task.

  • Use only tools from your assigned capability group, as defined in Appendix C.1.

  • Follow the specialist-specific workflow for the assigned subtask.

  • Do not solve unrelated subtasks assigned to other specialists.

  • If the assigned subtask cannot be completed with the available tools, explicitly
    mark it as unsupported or abstained.

Return a concise report containing:

  • completed steps;

  • extracted, retrieved, or generated facts relevant to the user request;

  • evidence used, such as source path, DOI, database filters, retrieved examples,
    generated method text, or validation result;

  • unsupported or abstained parts, if any.

Assigned specialist subtask:

                              [Assigned subtask]
```

C.3. CoSyn-Bench Examples

Each COSYN-BENCH item consists of a user question, gold answer, required capabilities, and scoring metadata. The first example illustrates a single-capability retrieval task, where the answer can be obtained from COSYN-DB alone.

```
{
  "type": "single_capability",
  "capabilities": ["Retrieval"],
  "question": "Which five solvents are most commonly used in the synthesis of co-crystals
               by the solution crystallization method?",
  "gold_answer": "The five most commonly used solvents in solution crystallization are
                  methanol, ethanol, water, acetonitrile, and acetone.",
  "metadata": {
    "input_artifacts": {
      "method_family": "solution crystallization",
      "target": "most common solvents",
      "k": 5
    },
    "hard_constraints": [
      "return exactly five solvents",
      "count only solution-crystallization records",
      "rank solvents by frequency"
    ],
    "required_items": [
      "five most common solvents used in solution crystallization"
    ]
  }
}
```

The second example combines prediction and retrieval. The system must first generate a grinding-based synthesis method and then compare the predicted apparatus with the most common apparatus for grinding in COSYN-DB.

```
{
  "type": "multi_capability",
  "capabilities": ["Prediction", "Retrieval"],
  "question": "Predict the method for synthesising a co-crystal by grinding, consisting
               of 2-ethoxybenzamide and 2,5-dihydroxybenzoic acid. Does the predicted
               method use the most common mixing apparatus for grinding?",
  "gold_answer": "Yes, the predicted method uses the most common mixing apparatus for
                  grinding. The predicted synthesis method: Mechanochemical synthesis
                  was carried out using a ball mill. The mixture was processed at 20.0
                  Hz for 30.0 min. A blend of 100.0 mg of CCOc1ccccc1C(N)=O and 93.3 mg
                  of O=C(O)c1cc(O)ccc1O was prepared in a ratio of 1:1. Additionally,
                  0.1 mL of methanol was added. The mixture was then ground. The
                  resulting co-crystal was dried. In the predicted method, a ball mill
                  is used as the mixing apparatus. Based on analysis of the database,
                  the ball mill is the most commonly used mixing apparatus for grinding.",
  "metadata": {
    "input_artifacts": {
      "molecular_pair": [
        "2-ethoxybenzamide",
        "2,5-dihydroxybenzoic acid"
      ],
      "method_family": "grinding",
      "db_stat": "most common mixing apparatus for grinding"
    },
    "hard_constraints": [
      "Predict a grinding-based method.",
      "Determine the most common mixing apparatus for grinding.",
      "State whether the predicted method uses that apparatus."
    ],
    "required_items": [
      "Propose the target synthesis procedure.",
      "Identify the mixing apparatus used in the prediction.",
      "Determine the most common apparatus for grinding.",
      "Compare them and answer yes or no."
    ]
  }
}
```

C.4. LLM-as-a-Judge Prompt

We used an LLM-as-a-judge evaluator to score candidate answers against the user question, the gold reference answer, benchmark metadata, and execution context. The judge was instructed to prioritize the gold reference and tool-derived execution context over its own background knowledge, and to penalize unsupported article, database, or prediction claims.

```
You are an expert evaluator for a benchmark on co-crystal synthesis agents.

Evaluate the candidate answer against the user question and the gold reference.
Use the benchmark input artifacts to understand the intended task setup.  Use the
execution input context as the concise record of evidence and tool-derived context
available to the agent.

When scoring groundedness, check whether the candidate answer is supported by
the execution input context and does not invent unsupported article, database,
or prediction details.  Prefer the gold reference and the explicitly provided
requirements over your own background knowledge.

You must score the answer on five criteria:

 1. Task completion.  Measures whether the candidate answer fully addresses the user
    request, including all required subtasks, examples, comparisons, extractions,
    predictions, or checks.

 2. Correctness.  Measures whether the answer is factually and semantically consistent
```

with the gold reference and available context. Minor wording differences are acceptable if the meaning is correct. For prediction tasks, the gold answer should be treated as a reference solution rather than the only valid solution.

3. Constraint adherence. Measures whether the answer follows explicit requirements in the question, such as synthesis method type, number of examples, solvent, temperature range, apparatus, ratio, API, coformer, yes/no decision, or method family.

4. Groundedness. Measures whether the answer is supported by the execution input context, including article evidence, database evidence, retrieved examples, extracted parameters, or prediction context. Unsupported details and hallucinated evidence should be penalized.

5. Answer quality. Measures whether the answer is clear, well-structured, readable, concise, complete, and directly usable.

Special-case scoring rules:

- If a required PDF or article could not be downloaded or accessed, do not penalize the candidate for failing to extract information from that unavailable source. A clear abstention or limitation statement is acceptable, but invented article contents must still be penalized.

- For prediction tasks, treat the gold answer as a reference solution. Chemically plausible and partially correct predictions should receive meaningful partial credit even when they do not match every parameter.

- Placeholders such as NO_VALUE or NOT_DETECTED are acceptable for missing, unavailable, or unsupported parameters. Penalize them only when the evidence clearly supports a concrete value.

- Additional synthesis parameters beyond the gold reference are acceptable if they do not contradict the reference or execution context.

- For yes/no questions, reward answers that state the decision explicitly. If the answer is substantively correct but indirect, reduce task completion or answer quality slightly.

Scoring rubric:

- 5 = fully satisfies the criterion.

- 4 = mostly satisfies it, with only minor issues.

- 3 = partially satisfies it, with meaningful correct content but important omissions, mismatches, or mistakes.

- 2 = limited correctness or coverage, but still contains some non-trivial correct content.

- 1 = completely fails the criterion, is almost entirely incorrect, or is unusable.

Assign only integer scores from 1 to 5 for every criterion. Do not use fractional values. Do not assign 0 under any circumstances.

Return only valid JSON with this schema:

```
{
  "task_completion": 1-5,
  "correctness": 1-5,
  "constraint_adherence": 1-5,
  "groundedness": 1-5,
  "answer_quality": 1-5,
  "rationale": "short explanation"
}
```

C.5. Runtime and Resource Usage

Table 14 reports median runtime, call count, and token usage for each evaluated system. Brackets denote the interquartile range.

Table 14. Runtime and resource usage on CoSyn-Bench. Median time, call count, and token usage per system, with interquartile ranges [Q1, Q3] in brackets.

System	Time (s)	Calls	Tokens
LLM-Only	4.04 [2.31, 5.17]	1.0 [1.0, 1.0]	779 [680, 901]
Context-Augmented LLM	16.52 [11.63, 26.89]	4.0 [2.0, 6.0]	21,590 [12,498, 41,202]
Single-Agent	17.87 [11.42, 25.34]	3.5 [3.0, 4.25]	57,610 [22,246, 105,856]
Multi-Agent	29.24 [21.74, 39.82]	7.0 [5.0, 10.0]	20,130 [14,994, 33,248]