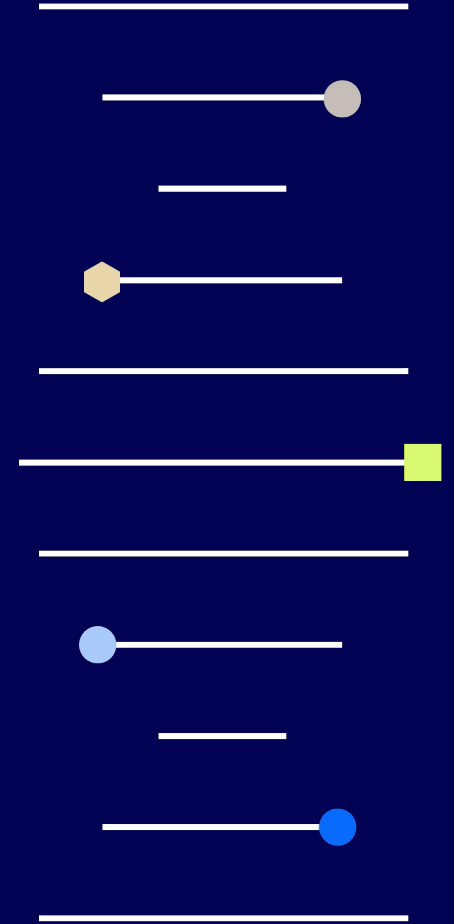


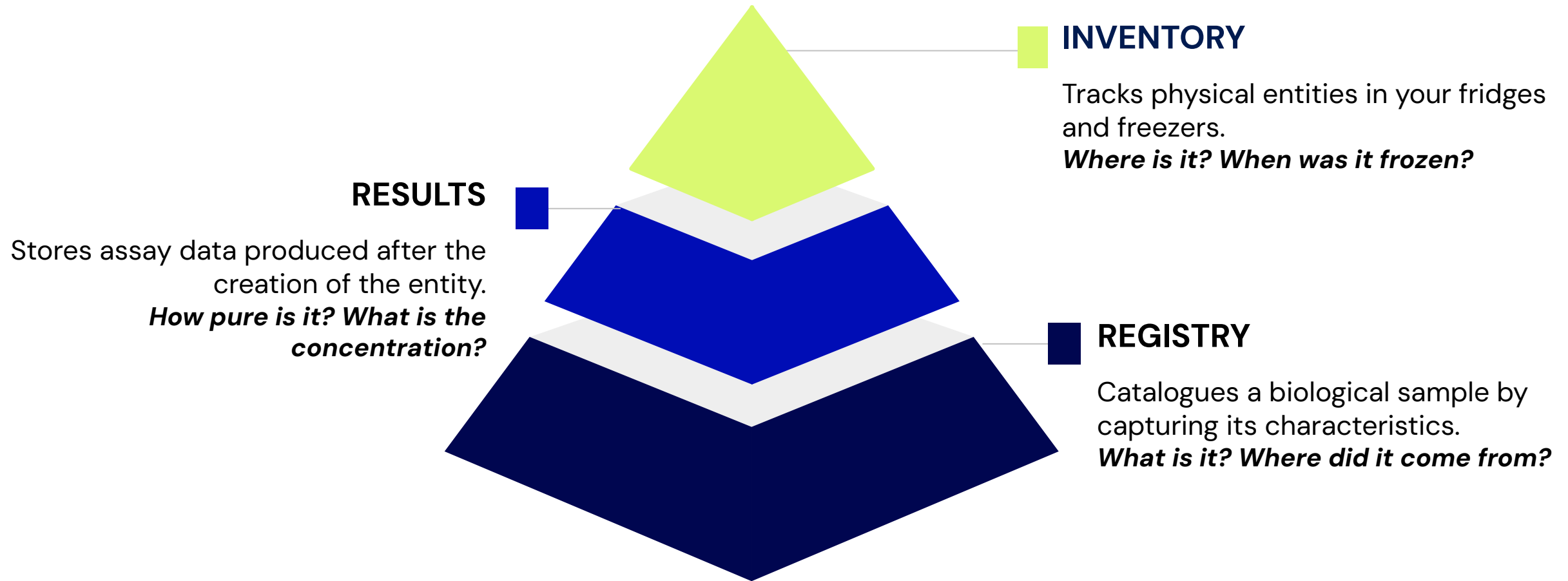
# Results Foundations & Best Practices



# Results Foundations



# Sample Tracking in Benchling





# Benefits of Leveraging Results

## Structured Data Capture

- Standardizing result capture enables easy data comparison
- Templates can be embedded in notebooks and templates to document assays contemporaneously
- Structured data is logged in the data warehouse for downstream analytics

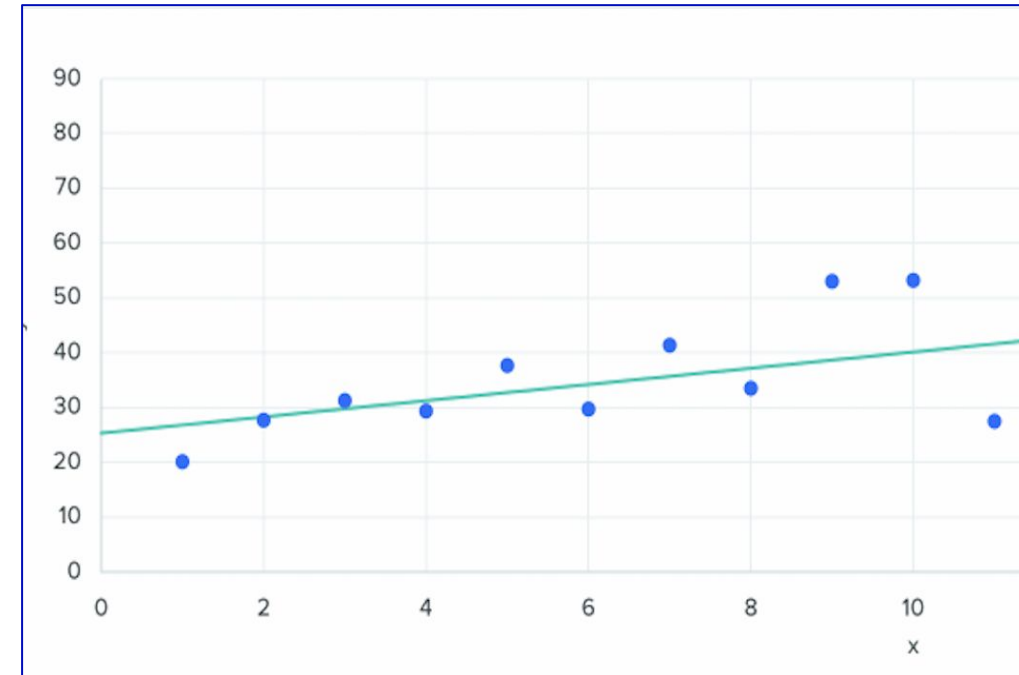
## Compliance

- Validate results alongside notebook entries through the witness and review process
- Eliminate manual transcription of result data via automation

## Analytical

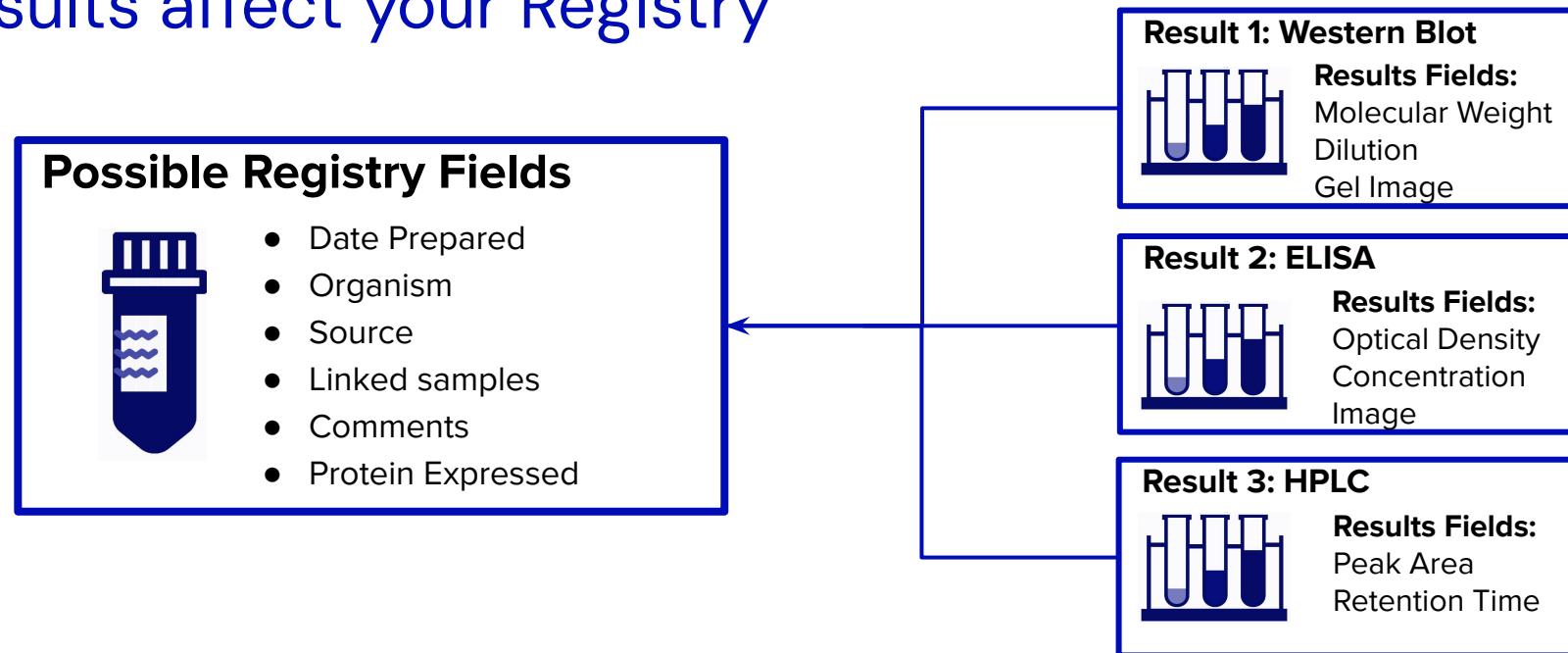
- Identify notebook entries associated with particular results
- Leverage the SQL Warehouse to aggregate results and generate visualizations

Training Plasmid Lot	Concentration	a260/280
TPL025	50	1.2
TPL083	50	1.2
TPL185	48	1.2
TPL332	52	1.2
TPL299	52	1.2





# How Results affect your Registry

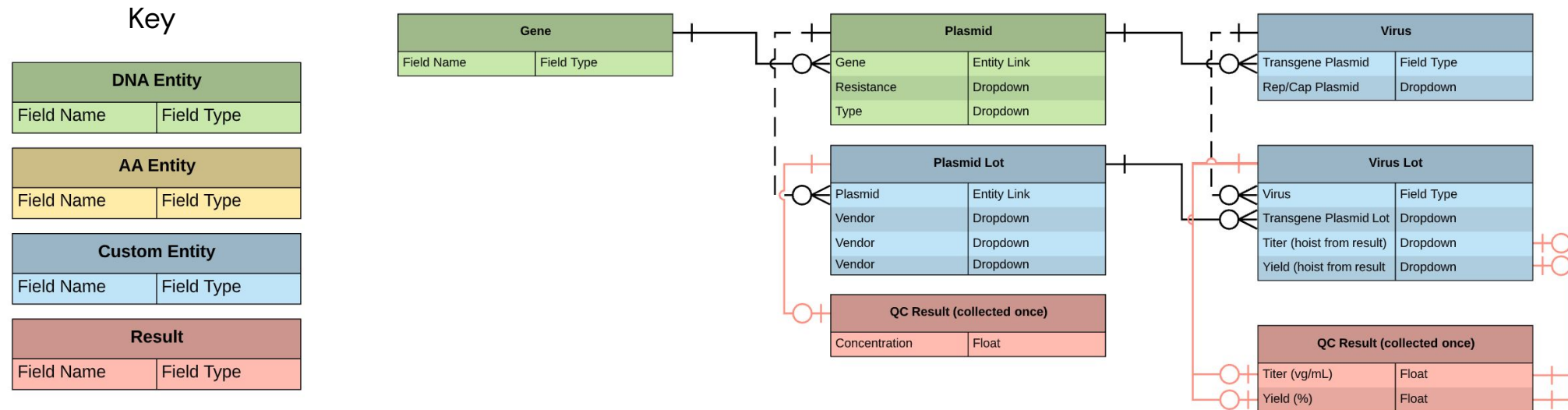


**Results are an equally important part of your data model** as the entity schemas themselves. Results *should* be associated to a registered entity or container:

- Entities can be associated with multiple types of result schemas
- Entities may have multiple results logged against them



# How Results affect your Registry – Downstream Effects

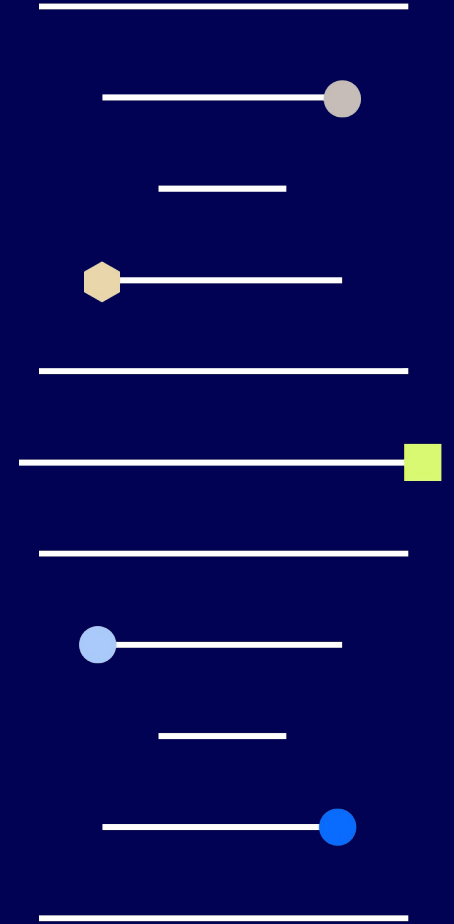


Data scientists (or other data enthusiasts) will be able to pull data against your registered entities and results to:

- Query and compare results from different results schemas across registered entities
- Aggregate large data set from a single result schema from one or many registered entities
- Analyze result data against upstream metadata



# Results Best Practices





## Best Practices: Using Results vs Registry

It can often be complicated deciding on where metadata should be placed. However, to achieve long-term optimization of a data model, consider the following: **Registry fields describe the characteristics of the sample, while Results fields record data produced after registration of an sample.**

### Registry Metadata Guidance:

- Measured/Recorded at time of Creation
- Inherent part of Registry Structured Data
- Best to measure one-time values that will not change often

### Results Metadata Guidance:

- Measured/Recorded at time of assay
- Connected to Registry via entity, not an inherent quality of entity
- Can have multiple measurements against one entity

#### Plasmid Lot (Registry)

Entity fields +	Definition
Parent Plasmid	○ Plasmid
Prep	☰ Prep Type
Date	📅 Date
Resistance $f_x$	☰ Resistance

#### Nanodrop (Result)

Result fields +	Definition
Entity	📄 Any entity
Date Created	📅 Date
Concentration	# Integer
A260/280	# Decimal





# Best Practices: Snapshot Fields

Snapshot fields are computed fields on a Result schema that pull data linked to inventory or registry items, **at a specific point in time**. This will provide a **one-off snapshot** rather than something that updates dynamically.

Snapshot fields, for example, can capture a 'freeze frame' of **what entity is in a container you are testing at the time the Result is captured**

## Registry and Associated Metadata

**METADATA**

DemoAbLot1  
Registry

Authors: Megan

Project Location: Inventory

Registry IDs: AbLot016, Bioprocessing Development

Created: 5/28/2020 09:23 AM

Aliases: +  
This entity has no aliases.

Schema: Antibody Lot (Cannot change schema of registered entities)

FIELD	VALUE
Date Made	5/28/2020
Parent Antibody	Candidate284
Bioreactor Run	AbRunMedia005
Chains $f_x$	DemoHC1, DemoLC1

## Results Tab on Registered Entity

DESCRIPTION STORAGE RESULTS

Results

Antibody Mass Spec (SSF)

	Container	$f_x$ Antibody Lot	$f_x$ Antibody
1	EppTube03 3	DemoAbLot 1	Candidate2 84

Container

Snapshot Fields



# Snapshot Fields vs. Lookup Columns

Use Snapshot Fields to record data in a predefined way for Results:

- Data needs to be **measured at a single point in time** and not dynamically updated
- Configured in **Registry Settings**

Use Lookup Columns to view data from variety of sources:

- Users need **reference information** in notebook tables that **doesn't belong on the schema** the table represents
- Configured in the **Notebook or Templates**

Results QC Testing - Protein Chain 1

Protein Chain Tested	Plasmid that made Protein	#	% Purity	Aq	Pass/Fail
1 Chain 1	New Plasmid 1		99		Pass

Computed field (snapshot)

← *fx* Snapshot Field

🔍 Lookup Column →

Results Antibody Mass Spec (w/out SSF) 1

A - Antibody ...	B - Container	C - Antibody	D - Chains	# E - Molecular weight (kDa)	F - Parent Antibody
DemoAbLot1	EppTube033	=C1			Candidate284

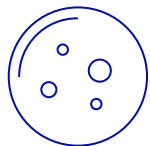
# Best Practices for Results Against Containers

Results against Containers are an example of leveraging [Snapshot Fields](#) to automatically surface information about contents of containers. Results can then be captured against containers, as well as registered entities. Tracking results against containers is useful for:

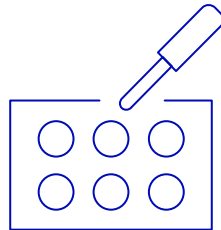
- Improved Sample Tracking and Auditing
- Measuring results against physical objects and their location metadata
- Scanning container labels into result tables

## Real Examples:

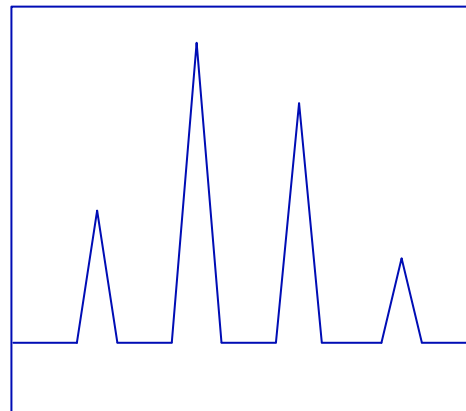
Tumor cells are cultured in multiple wells of a plate



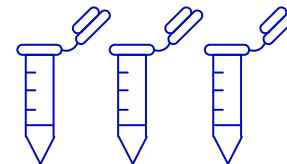
Wells have different chemical treatment applied to test for tumor cell survival



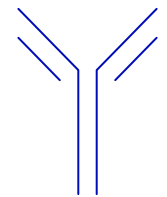
Results against containers allow for tracking of entities and conditions in both of these examples



Containers are stored at different conditions to test stability



Production lot of mAb is aliquoted into multiple containers



**Example 1**

**Example 2**

# Best Practices for Results in Templates

## Standardizing Data Capture

**Templates** ensure your standardized processes are followed. If you are looking to improve your result tracking adoption, adding Results **Structured Tables** to your templates will make it easier for users to have “ready-to-go” results tables in their entries.

	Container*	f(x) Container Sample	# Esimtated Timepoint (hours)	Datetime	# OD600 (dilution)	# Dilution Factor	OD600 (calculated)
1	E_Test Protein-001	Test Protein-001	1	3/20/2023 01:05:19 PM -0700	10	10	100
2	E_Test Protein-002	Test Protein-002	1	3/20/2023 01:05:19 PM -0700	10	10	100
3	E_Test Protein-003	Test Protein-003	1	3/20/2023 01:05:19 PM -0700	10	10	100
4	E_Test Protein-004	Test Protein-004	1	3/20/2023 01:05:19 PM -0700	10	10	100
5	E_Test Protein-005	Test Protein-005	1	3/20/2023 01:05:19 PM -0700	10	10	100
6	E_Test Protein-006	Test Protein-006	1	3/20/2023 01:05:19 PM -0700	10	10	100
7	E_Test Protein-007	Test Protein-007	1	3/20/2023 01:05:19 PM -0700	10	10	100
8	E_Test Protein-008	Test Protein-008	1	3/20/2023 01:05:19 PM -0700	10	10	100
9	E_Test Protein-009	Test Protein-009	1	3/20/2023 01:05:19 PM -0700	10	10	100

**Tips and Tricks:** Try adding a pre-built formula to a result table in a template to save your scientist time from recalculating in their entry

## Best Practices: Required Fields

	Required Fields for Results
<b>Benefits</b>	<ul style="list-style-type: none"><li>+ Standardize the inputs of your registry schema</li><li>+ Define what “complete” metadata is</li><li>+ Commit to data quality up front</li></ul>
<b>Trade-offs</b>	<ul style="list-style-type: none"><li>- Required fields can make results submission difficult, particularly if data is gathered at different times or if some fields only pertain to certain teams</li><li>- Notebook entries cannot be submitted for review if they contain unsubmitted tables</li></ul>

On a Results Schema, you have the option to designate fields as “Required”, meaning you cannot Submit your results without that value being inputted.

When designing a Result Schema, consider this decision carefully. You **cannot** retroactively mark a field on a Result Schema as required if there are already results submitted to that schema.

Result fields +	Warehouse name	Required	Multi-select	Definition
Cell Line	cell_line	✓		Cell Line (bt)
Antigen	antigen			Antigen
Expression Protocol	expression_protocol			Entry
Yield (mg/L)	yield_mgl			Decimal

# Growing with Results

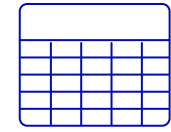
## Best Practices for Different Stages

<b>Basic</b> Focus on Adoption	<ul style="list-style-type: none"><li>Develop Result Schemas for your established assays</li><li>Train users on Result Table usage or add them to Notebook Templates</li><li>Communicate value of adding results into Benchling</li></ul>
<b>Intermediate</b> Focus on Optimization	<ul style="list-style-type: none"><li>Version control your Result Schemas as processes evolve</li><li>Use Lookup Tables to quickly communicate results to downstream stakeholders</li><li>Create a functional group responsible for communicating changes needed for schemas</li></ul>
<b>Advanced</b> Focus on Technical Maturity	<ul style="list-style-type: none"><li>Build Insights dashboards to visualize Result data</li><li>Utilize Developer Platform functionality to import and analyze Result data</li><li>Apply change management for new Result functionality/process implementation</li><li>Create team-based Result Schemas and implement tighter permissions around results</li></ul>

# Resources for Results



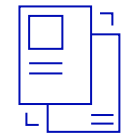
[Creating Result schemas and tables](#)



[Create equations for Results tables in an entry template](#)



[Configuring Snapshot fields](#)



[Using Results tables](#)