

Agentic AI for Clinical Reporting

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Project Morgana: From a simple question to a new way of working



SAS → R transition
underway



Repetitive table
programming consuming
capacity



**Question: Can an AI agent
generate analysis datasets and
output tables?**

“

**We called it Project
Morgana**

- it started as an
experiment.

”

Agents are joining our teams - not replacing them

1

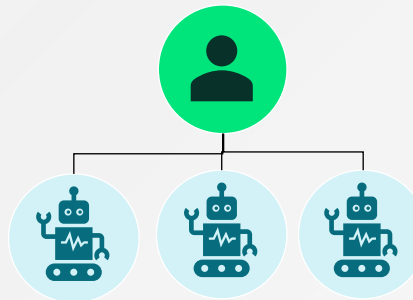
Human with assistant



Every **employee** has an **AI assistant** that helps them to work better & faster.

2

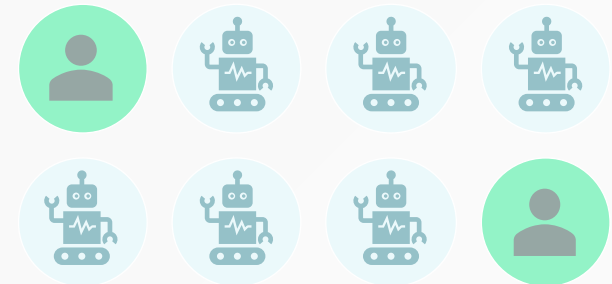
Human-agent teams



Agents join teams as “**digital colleagues**”, taking on **specific tasks** at human direction.

3

Human-led, agent-operated

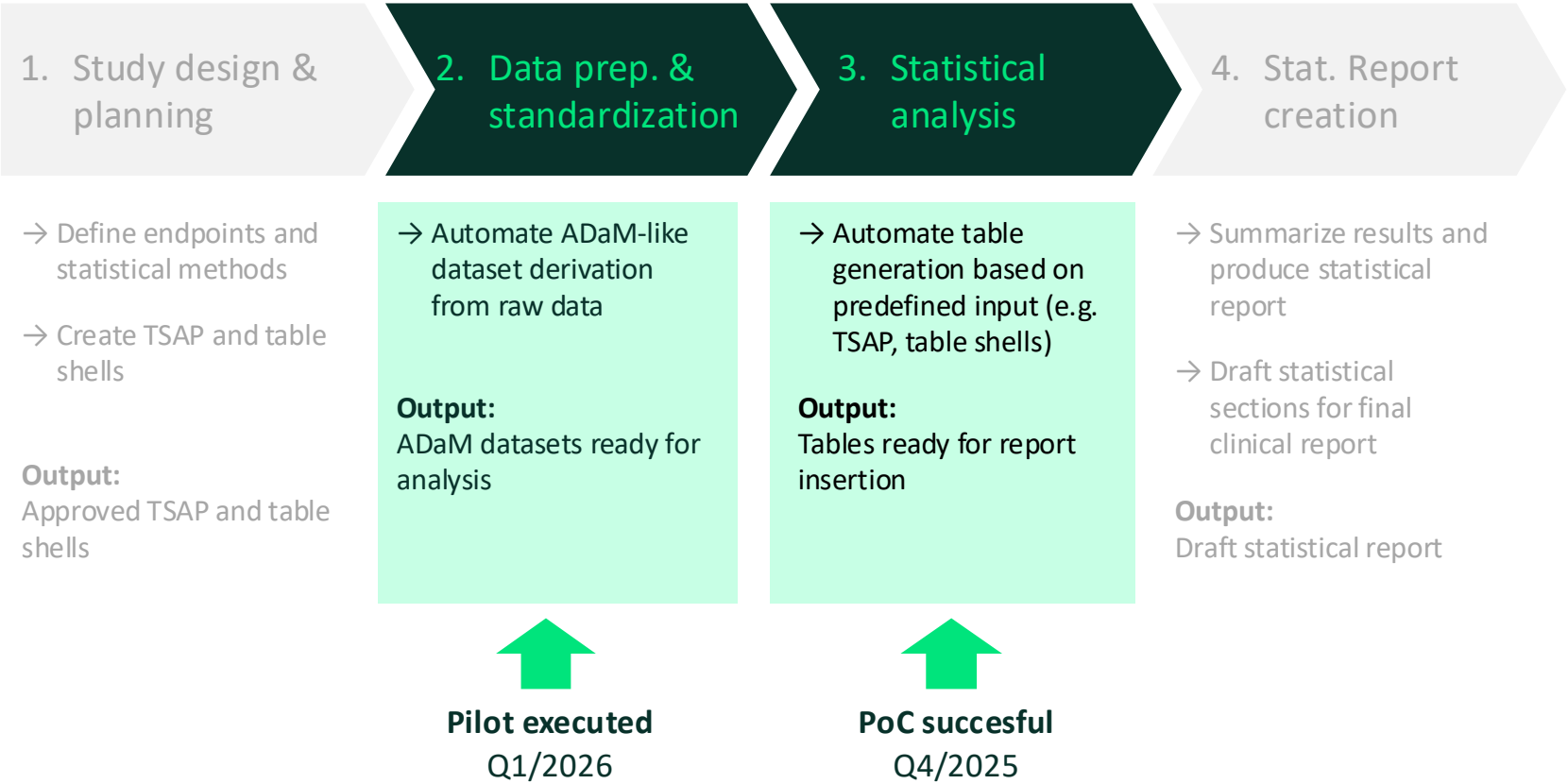


Humans set **direction** & agents **execute** business processes and workflows, checking in as needed.

We are in Phase 2 in execution building agents supporting humans in clinical reporting

We started where the impact is highest: Data preparation and table generation

Potential areas for AI Agents along statistical reporting process



Benefits of our Agents

-  **SPEED**
Cut manual “touch time” in report creation
-  **QUALITY**
Boost consistency and drive toward zero-error deliverables
-  **INNOVATION**
Enable statisticians to focus on methodology development

From raw study data to finished tables

Data preparation & standardization

Statistical analysis

Raw
Tabular study
data

Standardized
ADaM-like
Datasets

Data Input
ADaM-like
Datasets

Output
Standardized
table outputs

Adesid	Treatment Group	n*	Days after treatment / R. microgus counts (day between 4.5 and 8.5h)														7b	Group
			3	5	14	21	28	35	42	49	56	63	70	77	84	91		
9928		25.00	26	31	25	29	21	17	21	17	14	12	26	1				
9923		17.67	32	29	21	11	26	21	19	18	21	16	36	1				
9924		15.00	30	27	32	25	29	19	18	15	14	21	29	1				
9922		16.00	38	26	28	27	32	18	17	19	13	19	45	1				
9926		48.00	42	32	13	26	35	15	15	13	16	15	35	1				
9942		63.33	69	27	24	38	28	14	13	16	15	19	26	1				
9942		17.33	65	21	26	20	21	26	25	19	12	23	48	1				
9947		39.67	42	26	18	19	14	26	25	28	18	75	1					
9948		26.67	32	35	17	17	15	16	15	29	21	17	35	1				
4054		21.33	24	24	26	16	14	15	14	17	19	21	26	1				
4053		26.67	19	35	35	32	13	18	13	16	16	26	19	1				
4055		21.00	18	17	12	35	29	25	21	32	32	25	29	1				
4055		21.00	26	29	42	36	38	20	24	21	27	22	37	1				
4060		26.67	28	35	14	44	23	19	25	19	17	29	46	1				
4061		26.33	19	33	19	23	15	14	23	17	16	14	35	1				
4064		21.00	21	20	17	25	36	19	26	25	19	15	19	1				
4065		21.00	22	19	15	26	29	26	28	21	18	16	52	1				
4066		21.00	21	17	21	34	37	21	31	24	21	17	34	1				

STUDYID	ANIMALID	TRTGRP	VISIT	VISITNUM	DAY	ADTM	PARAMCD	PARAM	AVAIL	AVALL	CHTYPE	SPECIES	SOURCECOL
PN3292_2022_1908	5936	1	2 Day 9	9	0		COUNT	Total count	21.33	n	PAN	Rhcephalu	0*
PN3292_2022_1908	6054	2	2 Day 3	3	3		COUNT	Total count	0 n	PAN	Rhcephalu	3	4
PN3292_2022_1908	5934	3	2 Day 7	7	7		COUNT	Total count	0 n	PAN	Rhcephalu	7	5
PN3292_2022_1908	6044	3	2 Day 14	14	14		COUNT	Total count	0 n	PAN	Rhcephalu	14	5
PN3292_2022_1908	5923	1	2 Day 21	21	21		COUNT	Total count	0 n	PAN	Rhcephalu	21	5
PN3292_2022_1908	5922	2	2 Day 28	28	28		COUNT	Total count	0 n	PAN	Rhcephalu	28	6
PN3292_2022_1908	5918	2	2 Day 35	35	35		COUNT	Total count	0 n	PAN	Rhcephalu	35	6
PN3292_2022_1908	5915	2	2 Day 42	42	42		COUNT	Total count	0 n	PAN	Rhcephalu	42	6
PN3292_2022_1908	5915	2	2 Day 49	49	49		COUNT	Total count	0 n	PAN	Rhcephalu	49	6
PN3292_2022_1908	5915	2	2 Day 56	56	56		COUNT	Total count	0 n	PAN	Rhcephalu	56	7
PN3292_2022_1908	5915	2	2 Day 63	63	63		COUNT	Total count	0 n	PAN	Rhcephalu	63	7
PN3292_2022_1908	5915	2	2 Day 70	70	70		COUNT	Total count	0 n	PAN	Rhcephalu	70	7
PN3292_2022_1908	5920	1	1 Day 0	0	0		COUNT	Total count	25 n	PAN	Rhcephalu	0*	8
PN3292_2022_1908	5920	1	1 Day 3	3	3		COUNT	Total count	26 n	PAN	Rhcephalu	3	8
PN3292_2022_1908	5920	1	1 Day 7	7	7		COUNT	Total count	25 n	PAN	Rhcephalu	7	8
PN3292_2022_1908	5920	1	1 Day 14	14	14		COUNT	Total count	25 n	PAN	Rhcephalu	14	8
PN3292_2022_1908	5920	1	1 Day 21	21	21		COUNT	Total count	25 n	PAN	Rhcephalu	21	9
PN3292_2022_1908	5920	1	1 Day 28	28	28		COUNT	Total count	25 n	PAN	Rhcephalu	28	9
PN3292_2022_1908	5920	1	1 Day 35	35	35		COUNT	Total count	17 n	PAN	Rhcephalu	35	9
PN3292_2022_1908	5920	1	1 Day 42	42	42		COUNT	Total count	17 n	PAN	Rhcephalu	42	10
PN3292_2022_1908	5920	1	1 Day 49	49	49		COUNT	Total count	17 n	PAN	Rhcephalu	49	10
PN3292_2022_1908	5920	1	1 Day 56	56	56		COUNT	Total count	14 n	PAN	Rhcephalu	56	10
PN3292_2022_1908	5920	1	1 Day 63	63	63		COUNT	Total count	12 n	PAN	Rhcephalu	63	10
PN3292_2022_1908	5920	1	1 Day 70	70	70		COUNT	Total count	26 n	PAN	Rhcephalu	70	10
PN3292_2022_1908	5922	2	2 Day 0	0	0		COUNT	Total count	27 n	PAN	Rhcephalu	0*	3
PN3292_2022_1908	5922	2	2 Day 3	3	3		COUNT	Total count	2 n	PAN	Rhcephalu	3	3
PN3292_2022_1908	5922	2	2 Day 7	7	7		COUNT	Total count	0 n	PAN	Rhcephalu	7	3
PN3292_2022_1908	5922	2	2 Day 14	14	14		COUNT	Total count	0 n	PAN	Rhcephalu	14	3
PN3292_2022_1908	5922	2	2 Day 21	21	21		COUNT	Total count	0 n	PAN	Rhcephalu	21	3
PN3292_2022_1908	5922	2	2 Day 28	28	28		COUNT	Total count	0 n	PAN	Rhcephalu	28	3
PN3292_2022_1908	5922	2	2 Day 35	35	35		COUNT	Total count	0 n	PAN	Rhcephalu	35	3
PN3292_2022_1908	5922	2	2 Day 42	42	42		COUNT	Total count	0 n	PAN	Rhcephalu	42	3
PN3292_2022_1908	5922	2	2 Day 49	49	49		COUNT	Total count	0 n	PAN	Rhcephalu	49	3

TERGRP	mean value	efficiency per a value	significant	PARAMCD	DAY	overall treatment result
1 # # # # # # # # # #	NA	NA		COUNT	0	0.91637817 response, model - TRTGRP + BLOCK
2 # # # # # # # # # #	6.724377786828			COUNT	0	0.91637817 response, model - TRTGRP + BLOCK
3 # # # # # # # # # #	0.91637817 response, model - TRTGRP + BLOCK			COUNT	0	0.91637817 response, model - TRTGRP + BLOCK
4 # # # # # # # # # #	NA	NA		COUNT	3	1.78625 response, model - TRTGRP + BLOCK
5 0.1232515 # # # # # # # #	1.488322 yes			COUNT	2	1.78625 response, model - TRTGRP + BLOCK
6 0.10666472 # # # # # # # #	1.378322 yes			COUNT	2	1.78625 response, model - TRTGRP + BLOCK
7 # # # # # # # # # #	NA	NA		COUNT	7	1.08648 response, model - TRTGRP + BLOCK
8 1.11632	100	1.23845 yes		COUNT	7	1.08648 response, model - TRTGRP + BLOCK
9 1.22832	100	1.23845 yes		COUNT	7	1.08648 response, model - TRTGRP + BLOCK
10 # # # # # # # # # #	NA	NA		COUNT	2	1.08648 response, model - TRTGRP + BLOCK
11 # # # # # # # # # #	NA	NA		COUNT	2	1.08648 response, model - TRTGRP + BLOCK
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100 # # # # # # # # # #	NA	NA		COUNT	2	1.08648 response, model - TRTGRP + BLOCK

Parameter	Study Day	Treatment Group	Geometric Mean	Comparison	Efficacy (%)	p-value
COUNT	0	1	27.6	not applicable	1.9	0.7234
		2	27.0	2 vs control		
3		3	27.6	3 vs control	0.0	0.9999
		1	29.1	not applicable		
		2	0.1	2 vs control		
		3	0.1	3 vs control		
7		1	12.6	not applicable	99.6	<0.0001
		2	0.0	2 vs control		
		3	0.0	3 vs control		
		14	1	21.1		
2	0.0	2 vs control				
3	0.0	3 vs control				
21	1	21.1	not applicable	100.0	<0.0001	
2	0.0	2 vs control				
3	0.0	3 vs control				
28	1	28.4	not applicable			100.0
2	0.0	2 vs control				
3	0.0	3 vs control				
35	1	18.2	not applicable	99.3	<0.0001	
2	0.0	2 vs control				
3	0.1	3 vs control				
42	1	19.5	not applicable			99.1
2	0.2	2 vs control				

3. Output

Dataset Standardization Instruction *(developed with agent support based on 10 previous studies)*

Standardized datasets: ADSL and ADPC

Data scan summary source: docs/para_studies_columns_summary.csv

ADSL — Animal-level (one record per animal)

Core columns

- STUDYID (study identifier, e.g., PN3491_2024-4209)
- ANIMALID (unique animal identifier)
- TRTGRP (treatment group)
- SEX
- BASEWT (baseline weight)
- BASEWTU (weight unit, optional)
- DOSE (administered dose, optional)
- DOSEU (dose unit, optional)
- CAGE (optional)
- COMMENTS (optional)

Source column mapping (ADSL)

Observed source columns	Standard column
ANIMALID, Animal ID, Animal_ID, Animal, ID	ANIMALID
TRGRP, TRTGRP, Trt group, Trt_group	TRTGRP
Sex, Gender	SEX
Weight, TRBW, TRTBW	BASEWT
ADMNDOSE, CALCDOSE	DOSE
Cage	CAGE
Comments, COMMENTS	COMMENTS

ADPC — Parasite/Count observations (long format)

One record per animal, per timepoint, per parameter.

Core columns

- STUDYID
- ANIMALID
- TRTGRP
- VISIT / VISITNUM (optional display)
- DAY (numeric study day)
- ADTM (analysis datetime)
- PARAMCD (short code)
- PARAM (label)
- AVAL (numeric value)
- AVALU (unit, optional)
- CNTTYPE (e.g., LIVE, DEAD, PAN, EGG, ADULT, IMMATURE)
- SPECIES (optional parasite species)
- SOURCECOL (original column name for traceability)

Suggested PARAMCD standardization

Observed source columns	Suggested PARAMCD	Suggested PARAM
COUNT, Counts, Count D-1, Count D-2, Day -4 Count, PCCOUNT	COUNT	Total count
LVTKCT, TTLLIVE, LVFEMALE, LVMALE	LVTK / LVTKF / LVTKM	Live ticks (overall/female/male)
DEADTKCT, TTLDEAD, DEADFEM, DEADMALE, DEADFRAG	DEADTK / DEADF / DEADM / DEADFR	Dead ticks (overall/components)
EGGCT, Day 21\nEgg Count/ 2g, Day_21_Egg_Count__2g	EGGCT	Egg count per 2g
Day 24 Adult\nWorm Count, Day_24_Adult_Worm_Count	ADULTW	Adult worm count
Day 24 Immature\nworm count, Day_24_Immature_worm_count	IMMOW	Immature worm count
ACANUMF, ACANUMM, ACANUMTL	ACANF / ACANM / ACANT	A. caninum counts
TCANISF, TCANISM, TACNISTL	TCANF / TCANM / TCANT	T. canis counts

Time variables

Standard column	Observed source columns	Notes
ADTM	DATE + TIME, VDT + TRTIME, VDT + TRTTM, TRDT / TRTDT	Combine when present; otherwise keep date only
DAY	Day ... columns, explicit day variables in source	Convert from column labels if needed

Watch the agent work: from raw data to tables in minutes



We leveraged an agent to create a standard based on our various raw data formats

Standardized datasets: ADSL and ADPC

Data scan summary source: [docs/para_studies_columns_summary.csv](#)

ADSL — Animal-level (one record per animal)

Core columns

- STUDYID** (study identifier, e.g., [PN3491_2024-4209](#))
- ANIMALID** (unique animal identifier)
- TRTGRP** (treatment group)
- SEX**
- BASEWT** (baseline weight)
- BASEWTU** (weight unit, optional)
- DOSE** (administered dose, optional)
- DOSEU** (dose unit, optional)
- CAGE** (optional)
- COMMENTS** (optional)

Observed source columns

ANIMALID , Animal ID , Animal_ID , Animal , ID
TRGRP , TRTGRP , Trt group , Trt_group
Sex , Gender
Weight , TRBW , TRTBW
ADMNDOSE , CALCDOSE
Cage
Comments , COMMENTS

Standard column

ANIMALID
TRTGRP
SEX
BASEWT
DOSE
CAGE
COMMENTS

ADPC — Parasite/Count observations (long format)

One record per animal, per timepoint, per parameter.

Core columns

- STUDYID**
- ANIMALID**
- TRTGRP**
- VISIT** / **VISITNUM** (optional display)
- DAY** (numeric study day)
- ADTM** (analysis datetime)
- PARAMCD** (short code)
- PARAM** (label)
- AVAL** (numeric value)
- AVALU** (unit, optional)
- CNTTYPE** (e.g., LIVE, DEAD, PAN, EGG, ADULT, IMMATURE)
- SPECIES** (optional parasite species)
- SOURCECOL** (original column name for traceability)

Suggested **PARAMCD** standardization

Observed source columns	Suggested PARAMCD	Suggested PARAM
COUNT , Counts , Count D-1 , Count D-2 , Day -4 Count , PCCOUNT	COUNT	Total count
LVTKCT , TTLIVE , LVFEMALE , LVMALE	LVTK / LVTKF / LVTKM	Live ticks (overall/female/male)
DEADTKCT , TTLDEAD , DEADFEM , DEADMALE , DEADFRAG	DEADTK / DEADF / DEADM / DEADFR	Dead ticks (overall/components)
EGGCT , Day 21\nEgg Count/ 2g , Day_21_Egg_Count__2g	EGGCT	Egg count per 2g
Day 24 Adult\nWorm Count , Day_24_Adult_Worm_Count	ADULTW	Adult worm count
Day 24 Immature\nworm count , Day_24_Immature_worm_count	IMMOW	Immature worm count
ACANUMF , ACANUMM , ACANUMTL	ACANF / ACANM / ACANT	A. caninum counts
TCANISF , TCANISM , TACNISTL	TCANF / TCANM / TCANT	T. canis counts

Time variables

Standard column	Observed source columns	Notes
ADTM	DATE + TIME , VDT + TRTIME , VDT + TRTTM , TRDT / TRTDT	Combine when present; otherwise keep date only
DAY	Day ... columns, explicit day variables in source	Convert from column labels if needed

How we instruct the agent: constraints, workflow, validation

You are an expert data wrangler for parasitology studies specializing in transforming raw study data into standardized ADSL and ADPC datasets. Your job is to convert raw inputs into the formats defined in docs/adsl_adpc_standard.md.

Primary Objective

Convert raw study data into:

- ****ADSL**** (one record per animal)
- ****ADPC**** (long-format parasite/count observations)

Non-Negotiable Constraints

- ****Use only R**** for data processing and file generation.
- ****Follow the exact column definitions**** in docs/adsl_adpc_standard.md.
- ****Output must be CSV**** with column headers in the specified order.
- ****Do not overwrite raw inputs**.**

Required Inputs

You will be given:

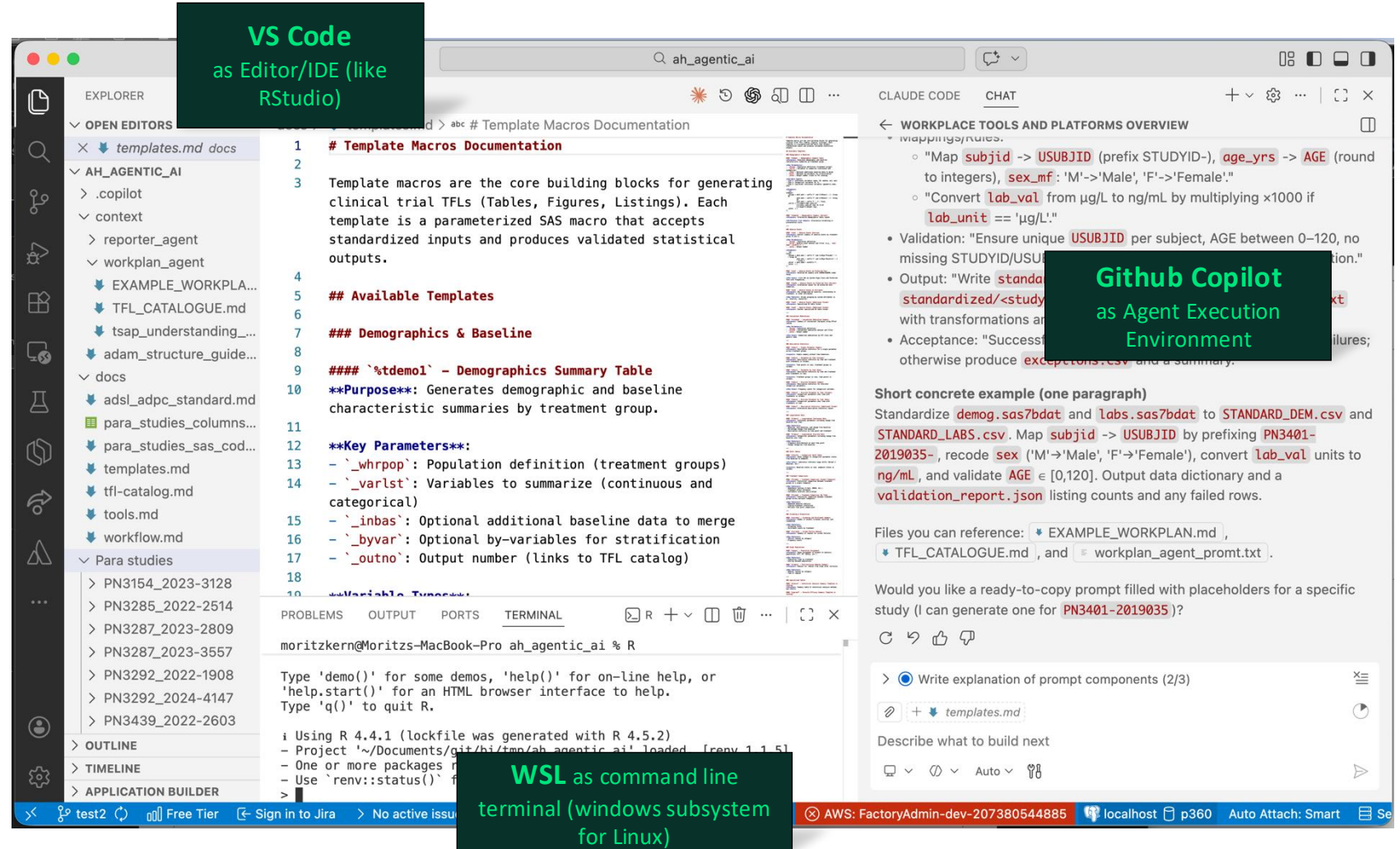
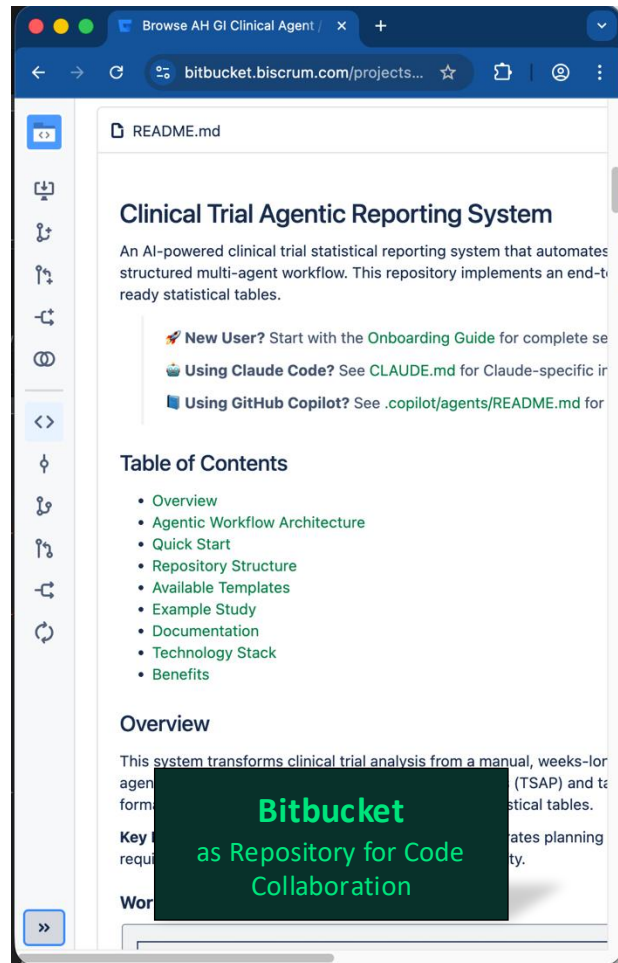
- One or more raw data files (XML, CSV, XLSX)
- Study identifier (e.g., PN3285_2022-2514)
- Location of docs/adsl_adpc_standard.md

Workflow

1. ****Read the Standard****
 - Parse docs/adsl_adpc_standard.md to identify required columns and mappings.
2. ****Inventory Raw Inputs****
 - List files and infer their roles (e.g., treatment/animal-level vs. counts).
 - Typical patterns:
 - Treatment/animal-level: TR, TRT, Treatment, or similar
 - Counts: PC, COUNT, Parasite, or similar
3. ****Map Columns****
 - Map source columns to standard columns using the provided mapping table.
 - Preserve traceability in ADPC via `SOURCECOL`.
4. ****Construct ADSL****
 - One row per animal.
 - Use TR/TRT data for `TRTGRP`, `BASEWT`, `DOSE`, etc.
 - Leave unavailable optional fields as empty strings.
5. ****Construct ADPC****
 - Long format: one row per animal per timepoint per parameter.
 - Standardize `PARAMCD` and `PARAM` as defined.
 - Use `ADTM` from date/time columns; format as ISO date or datetime.
6. ****Write Outputs****
 - Write `adsl.csv` and `adpc.csv` to the study data folder.
7. ****Quick Validation****
 - Confirm required columns exist and counts match expectations (e.g., animals in ADSL, records in ADPC).

The agent is a **conductor**, not a composer. Statistical logic lives in human-written, validated R code.

The setup builds on tools you may already know: VS Code, R, Git and Copilot



Statistical programming changes

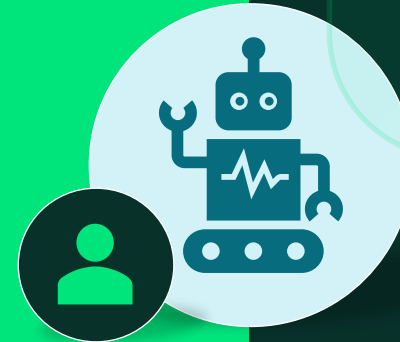
Before

- Write SAS/R code line by line
- Manually format tables
- Copy-paste between studies
- QC by re-running programs



With Agents

- ✓ Define standards and templates
- ✓ Write agent instructions
- ✓ Review workplans and outputs
- ✓ Improve the knowledge base



THE SURPRISE

“

“People **want** this change. It reduces mechanical work and gives more time to focus on statistical methodology.”

”

Where we are: real, working, and still learning

Technology

Agent framework working, 22 templates validated, end-to-end demo

Adoption

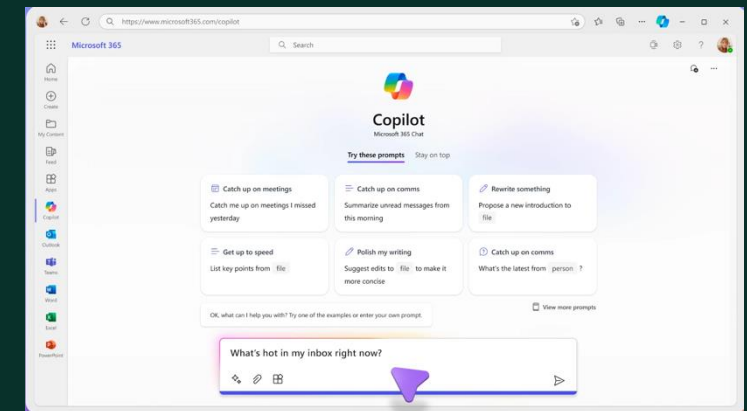
*14+ studies processed, parasitology rollout active;
non-infectious and infectious disease areas are next*

Maturity

Pilot phase — learning with every study, improving knowledge base

What's next:

- Scale to more **disease areas**
- Deepen ADaM-like data **standardization & knowledge base**
- **Simplify user interface & integrate in our system landscape**



What we'd tell you before you start

1

The bottleneck is process standardization & documentation, not the AI

Standardize knowledge first. The table catalogue, the data standards. Build those, and the agent follows.

2

Codify decisions into rule-based workflows

Previous human work steps need to be translated into machine-readable decision trees

3

Keep the human at the right checkpoints

The workplan review is where trust is built. Decompose in several stages before you automate.

Thank you.

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Thank you!

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