Council on Dairy Cattle Breeding

Data Ingestion System

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CDCB data ingestions systems

- **All** records that you submit are processed by a group of programs that perform the “ETL” process: extract, transform, load.
- The 2 main programs are named:
  - **EDCLOB** (genomic data ingestion system)
  - **EDITS** (pretty much everything else you submit, including nominations)
    - The *true* data ingestion system
- Runs and writes locally to the database.
- Active 24/7

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CDCB data ingestions systems – What we're changing

- CDCB is investing resources in modernizing its infrastructure.
  - Is there something wrong with EDITS?
  - Create more opportunities for the US industry
  - Use more modern programming infrastructure and languages
  - Full documentation of business rules
  - Succession plan
- WebConnect was the first of such projects.
- **CDCB data ingestion systems are next.**
  - **EDITS is phase 1.**

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CDCB data ingestions system project

- The project concept was developed way before it was even conceived.
- First **key** step: full documentation of business rules
- Thousands of lines of documentation.. inside the code.
  - … and in the head of a CDCB employee.

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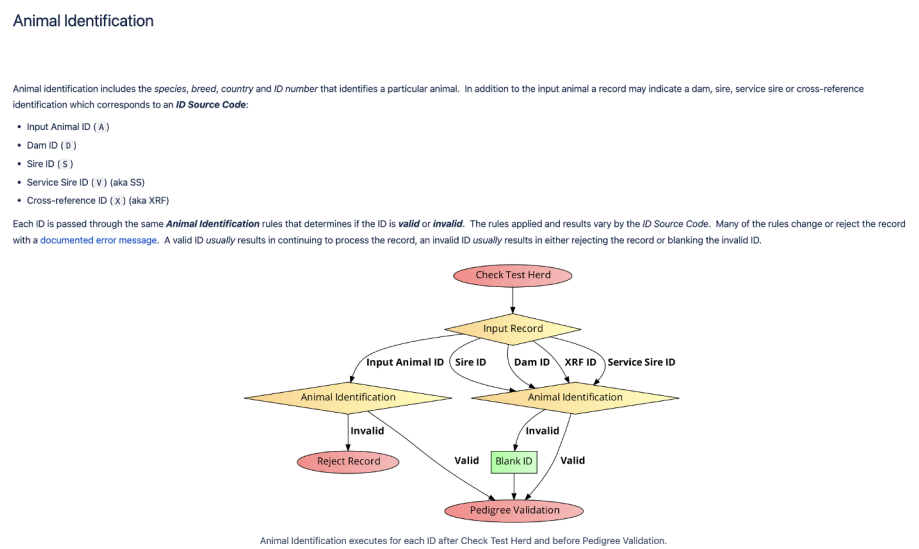
Documentation of business rules

- Formal documentation in Confluence
- Collaboration with iYotah solutions
- ~ 1 year for one highly skilled programmer.
- Independent review of business rules
  - very little # of discarded rules
- Reality check.



Animal Identification

Animal identification includes the *species*, *breed*, *country* and *ID number* that identifies a particular animal. In addition to the input animal a record may indicate a dam, sire, service sire or cross-reference identification which corresponds to an ***ID Source Code***:

- Input Animal ID (`A`)
- Dam ID (`D`)
- Sire ID (`S`)
- Service Sire ID (`V`) (aka SS)
- Cross-reference ID (`X`) (aka XRF)

Each ID is passed through the same ***Animal Identification*** rules that determines if the ID is ***valid*** or ***invalid***. The rules applied and results vary by the *ID Source Code*. Many of the rules change or reject the record with a documented error message. A valid ID *usually* results in continuing to process the record, an invalid ID *usually* results in either rejecting the record or blanking the invalid ID.

Animal Identification executes for each ID after Check Test Herd and before Pedigree Validation.

The next phase: planning the future

- There is nothing wrong with the current system
- But there are limitations:
  - Local server (pros and cons!)
  - Physical storage space
  - A single CDCB staff member in control of development
    - Very difficult skills to pass along
  - Tightly linked to structured DHI data
  - Not possible to integrate to new tools developed or in development

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Planning the future

- CDCB is expected to:
  - Provide solutions at a faster rate than in the past
  - Accept a wider range of data formats and providers
  - Deal with (and store) bigger data (e.g. sensors, MIR)
  - Provide better feedback to users (and help them solve issues)
  - Store large amounts of raw data for researchers to make sense of
  - Etc…
- The original Data Ingestion System was not designed to tackle these new needs

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Designing the future …

- Re-write the Data Ingestion System (~ no change of rules) to enhance its functionalities:
  - ID database
  - File queuing system
  - New universal format for data submission
  - API capabilities
  - Cloud-based system
  - Unlimited storage
  - New reporting capabilities & direct integration with WebConnect

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… without disrupting operations

- Biggest challenge of all
- Interaction with our current (local) database
- Minor / no changes to user interaction necessary
- Mixed system: local + cloud interaction needs to be seamless

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Developing the future

- A year and a half project signed with iYotah solutions
- Exclusive team of 6 people dedicated to this project
- 5 deliverables
- Objective: reproduce EDITS on a cloud environment successfully interacting with CDCB system (full integration).
  - Barely any change for users of the *current* system in the delivery of v.1.0
  - Further enhancements already spec'd, planned, and documented

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What “barely any change” means

- No change for the user, completely new system “under the hood”
- Maintaining rules and functionalities will allow thorough testing and guarantee seamless continuity of services
- Some new features:
  - possibility of submitting files via API
  - some basic statistics shown in WebConnect
  - “visible” queuing system (where is my file in the queue?)

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Where are we in the project

- **Deliverable 1**: “Cloud + Data lake + infrastructure setup” (Q2 2023)
- **Deliverable 2**: “Universal format, API” (Q3 2023)
- **Deliverable 3**: “Testing framework, ID database” (Q4 2023)
- **Deliverable 4**: “EDITS code, integration testing” (Q2 2024) - Expected Q3 2024
- **Deliverable 5**: “UAT” (Q3 2024) – Expected Q4 2024

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How testing will work

- The project includes 4 phases of testing:
  - iYotah's testing platform (cucumber): automated set of rules continuously maintained to ensure future changes to the code do not affect other parts of the system.
  - CDCB targeted testing: CDCB staff is testing features as they are released
  - CDCB integrated testing: CDCB staff will test integration of the new system vs the old by running both in parallel and compare output
  - UAT: Selected industry partners will test the system before launch

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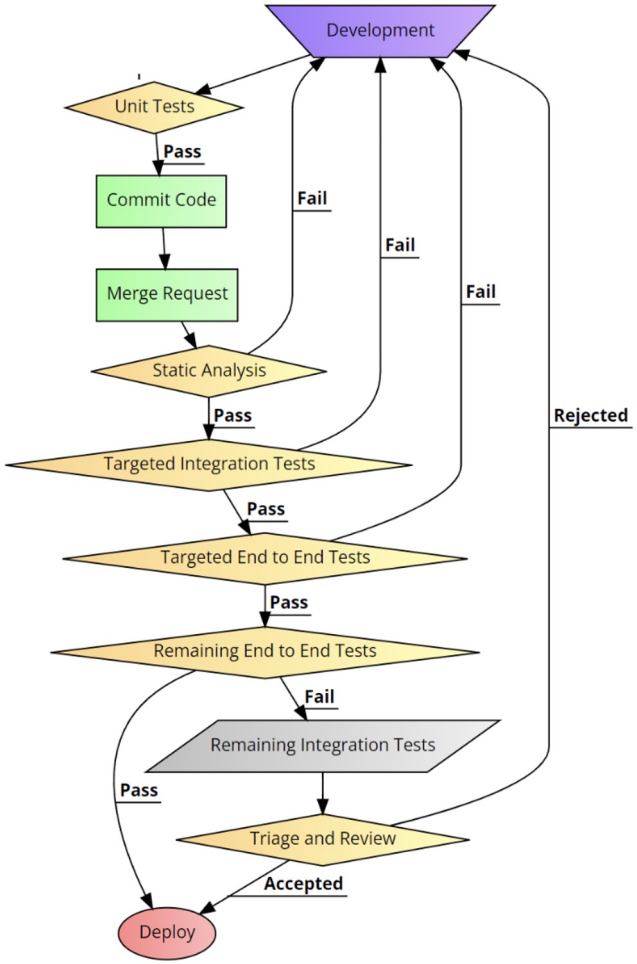
And after the project?

- Similar process to development, but faster
- *Continuous testing*
- Human (CDCB staff) approval still req'd!



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Setting clear expectations

- Release 1.0 will be a slightly enhanced version of the current system.
  - Maximum success: nothing changes for you
  - We *must* ensure the system is stable and robust before building new features
  - New features will be easier to implement
    - Much more flexibility
    - Automated testing infrastructure assistance
    - Can work in “islands” and deploy as many testing environments as we need
- Future releases:
  - Harvesting ideas internally first, will interact with industry next.
  - Nearly perfect alignment with feedback from 2023 CDCB Industry Workshop
  - Need to shape the future

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SOME “VISUAL” RESULTS

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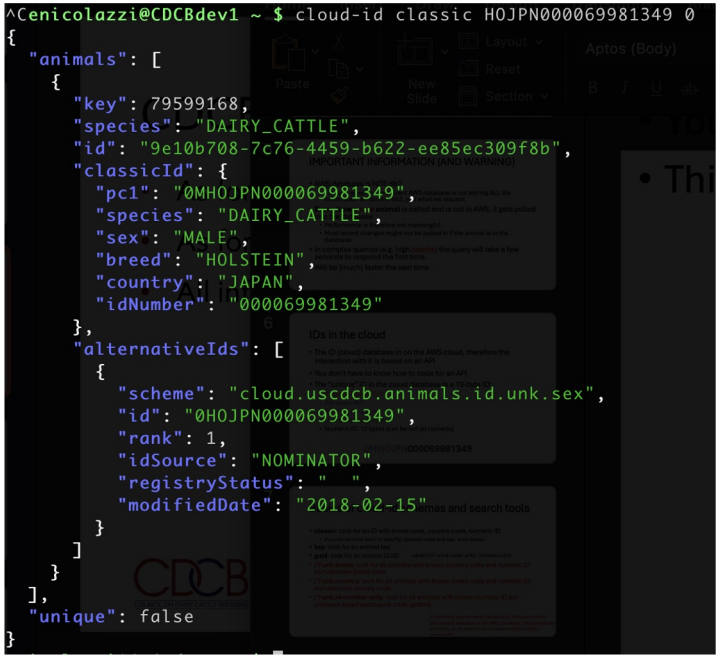
CDCB ID database

- As for the user, there is no change to current practices.
- As for CDCB, this is planned to be the core of animal identification.
- All information linked to an animal, gathered centrally in a database.



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Queue system

- Replaces bash automations handling files from your “in” folders to EDITS and back to your “out” folders
  - (CDCB only) Processing a file from.. anywhere
  - (CDCB only) Processing a file from a cloud location
  - (all) Automated processing of a file in a folder (e.g. SFTP)
  - (all) API submission to the queue

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Queue system – Automated processing

```
enicolazzi@CDCBdev1 ~ $ cloud-notify -l
[
  {
    "path": "cdcb/enicolazzi/SFTP/in",
    "output": "cdcb/enicolazzi/SFTP/out",
    "source": "BREED_ASSOCIATION",
    "user": "HO",
    "requester": "HO",
    "center": "HO",
    "enabled": true,
    "autoQueue": true
  },
  {
    "path": "iyotah/breedho/in",
    "output": "iyotah/breedho/out",
    "source": "BREED_ASSOCIATION",
    "user": "HO",
    "requester": "HO",
    "center": "HO",
    "enabled": true,
    "autoQueue": false
  }
]
```

```
enicolazzi@CDCBdev1 ~ $ ll /S3/cdcb/cdcb/enicolazzi/SFTP/in
total 4
drwxrwxr-x 1 enicolazzi enicolazziGr   0 Jun 11 04:10 .
drwxrwxr-x 1 enicolazzi enicolazziGr   0 Jun 11 04:10 ..
-rwxrwxr-x 1 enicolazzi enicolazziGr 124 Jun 11 04:24 misanim1fmt1.1
-rwxrwxr-x 1 enicolazzi enicolazziGr 124 Jun 11 04:33 misanim3fmt1.1
-rwxrwxr-x 1 enicolazzi enicolazziGr 124 Jun 11 04:33 misdam2fmt1.1
-rwxrwxr-x 1 enicolazzi enicolazziGr 124 Jun 11 04:12 misdamfmt1.1
-rwxrwxr-x 1 enicolazzi enicolazziGr 124 Jun 11 04:22 missexfmt1.1
-rwxrwxr-x 1 enicolazzi enicolazziGr 125 Jun 11 04:33 shiftfmt1.1
-rwxrwxr-x 1 enicolazzi enicolazziGr 124 Jun 11 04:33 wrongspecfmt1.1
enicolazzi@CDCBdev1 ~ $ ll /S3/cdcb/cdcb/enicolazzi/SFTP/out
total 3
drwxrwxr-x 1 enicolazzi enicolazziGr   0 Jun 11 04:10 .
drwxrwxr-x 1 enicolazzi enicolazziGr   0 Jun 11 04:10 ..
-rwxrwxr-x 1 enicolazzi root         363 Jun 11 04:24 misanim1fmt1.1e
-rwxrwxr-x 1 enicolazzi root         363 Jun 11 04:33 misanim3fmt1.1e
-rwxrwxr-x 1 enicolazzi root         363 Jun 11 04:33 misdam2fmt1.1e
-rwxrwxr-x 1 enicolazzi root         363 Jun 11 04:12 misdamfmt1.1e
-rwxrwxr-x 1 enicolazzi root         363 Jun 11 04:22 missexfmt1.1e
-rwxrwxr-x 1 enicolazzi root           0 Jun 11 04:33 shiftfmt1.1e
-rwxrwxr-x 1 enicolazzi root         363 Jun 11 04:33 wrongspecfmt1.1e
```

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Queue system – CLOUD / API submission

Cloud File Submission (cloud-submit)

The `cloud-submit` tool is used to submit files that are already uploaded to the data lake, or to submit the contents of a file directly to the API without the need to upload it to the data lake first. To submit a file automatically when it is uploaded to a data lake folder see the `cloud-notify` tool to configure automatic submission of files instead. The advantage of using `cloud-submit` is that you have control over the parameters the file is processed with such as *source*, *requester*, *center* whereas with `cloud-notify` those options are configured per folder.

Additionally, for testing purposes only, `cloud-submit` allows processing the file immediately, bypassing the data lake and normal file processing system entirely to return the results and additional debugging information right away.

```
Usage: cloud-submit [options]

Options:
  -p [file]      Add a file that is already in the data lake by path and filename, pa
  -f [file]      Directly submit the contents of the specified file
  -s [source]    Source to use when processing the file
  -r [requester] Requester to use when processing the file
  -c [center]    Center to use when processing the file
  -a [aiplCode]  AIPL code to use when processing the file
  -u [user]      User to associate with the file, defaults to your login username
  -Q             Do NOT automatically queue the file for processing, you will have to
  -i             Process the records immediately and return the result, bypasses the

Examples:
  cloud-submit -f myfolder/myfile.1 -s B -r HO
    Submit the contents of myfile.1 and process as a BREED_ASSOCIATION with a request

  cloud-submit -p cdcb/breedho/in/20200401.4 -s D -r WI -c WI
    Submit the file already present on the data lake at breedho/in with filename 2020
    source of DRPC and a requester of WI and a center of WI.
```

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Initial processing

```
cloud-queue -id f7042330-934a-4351-ad43-9632758c037a
```


```
{
  "data": {
    "id": "f7042330-934a-4351-ad43-9632758c037a",
    "path": "cdcb/iyotah/breedho/in",
    "name": "pedigree4.fmt1",
    "source": "BREED_ASSOCIATION",
    "user": "HO",
    "requester": "HO",
    "center": "UT",
    "aiplCode": " ",
    "results": {
      "totalRecords": "9",
      "pedigreeRecords": "4",
      "genomicRecords": "0",
      "0Eb": "5",
      "unknownRecords": "5",
      "1Aa": "2",
      "4Aa": "1",
      "healthRecords": "0",
      "reproductiveRecords": "0",
      "changedRecords": "0",
      "1Fp": "1",
      "lactationRecords": "0",
      "rejectedRecords": "8",
      "notifyRecords": "1"
    },
    "filePathAndName": "cdcb/iyotah/breedho/in/pedigree4.fmt1"
  },
  "system": {
    "id": "f7042330-934a-4351-ad43-9632758c037a",
    "version": 4,
    "status": "PROCESSED",
    "createdTime": "2024-05-15T15:29:52.447993787Z",
    "modifiedTime": "2024-05-15T15:33:26.764828611Z",
    "queued": false,
    "processedTime": "2024-05-15T15:33:21.088679723Z"
  }
}
```


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Take home messages

- CDCB is investing heavily in the future of dairy industry
- Modernizing infrastructure (both hardware and software)
- Special attention to continuity and seamless integration
- Documenting all processes
- Future enhancements easier to implement
- **Work is in progress!**

*Disclaimer: Picture **not** approved by CDCB communications and outreach specialists*

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THANK YOU FOR YOUR ATTENTION

ACKNOWLEDGMENTS

U.S. dairy producers
Member sectors and collaborators
USDA AGIL
CDCB staff

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