WebConnect

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WebConnect

- **WebConnect** is a user interface introduced in August, 2022

- For **public** and **members**

 (accessibility depends on configuration of the user)

- **WebConnect** made it easy for users to query, update, edit, add, remove, move and view data

- Some **batch submissions** and **API** interactions became possible

- Animal centric

- Interactive query system

Tabs

Welcome, KaoTok

Queries Data Exchange ▾ Special Section ▾ Top Animal Listing Summary Stats ▾ National Performance Metrics

QUERIES

ID + Sex Code (18 bytes)

- **Queries**

  ID/pedigree, evaluation, Genotype, Error tabs for given Animal ID (s) (Max. 50 records)

- **Data Exchange**

  ID/pedigree, Genotype, Special Section, Reports, and Create report (for list of animals **requested in a file)**

- **Special Section**

  Includes tools for specific tasks or information handled differently from regular queries

- **Top Animal Listing, Summary Stats, National Performance Metrics**

  General information on the National Cooperator Database

- Please **see slide 33 and after** for detailed info

WebConnect

Web Queries	Data Exchange User Interface	Data Exchange 3rd Party
• Graphical User Interface • Animal-centric • Limitation of a result set of 50 records or less	• Graphical User Interface • Intermediate between Web Queries and Data Exchange 3rd party • Allows exchange information in files (input and output)	• Data exchange done through API* (not GUI) • Requires Advanced (skilled) API usage • Ezequiel will talk about it tomorrow

application programming interface (**API***) is a way for two or more computer programs or components to communicate with each other. (wikipedia)

WebConnect : Jan-June 2024

	Web Queries	Data- Exchange UI	Data-Exchange 3rd Party
Total Users	96 users (18 Organizations)	54 Users (8 Organizations + 'unmapped')	3 Users (3 Organizations)
Total Submissions	158,569	11,936	406,140
Most used Queries	FIX_FMT1, Bulk Nomination, edit_ped	Genotype Query, Get Animal Formatted Pedigree, Genotype Nomination Status	Genotype Query, Get Animal Formatted Pedigree, Get Animal ID by 12 Character ID Number

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Nomination using WebConnect - **login**

Login

Queries   Special Section ▾   Top Animal Listing   Summary Stats ▾   National Performance Metrics

COLLABORATOR LOGIN

The CDCB database is currently locked. Any submissions made will be processed automatically as soon as the database is unlocked.

Message if database is locked

Session timed out. Please log-in again.

Username or Email

kaori.tokuhisa@uscdcb.com

Password

••••••••••••

Credentials

Account help

Forgot Password?   Register New Account?

LOGIN

WARNING: This system is for the use of authorized users only. Individuals using this computer system without authority, or in excess of their authority, are subject to having all of their activities on this system monitored and recorded by system personnel.
In the course of monitoring individuals improperly using this system, or in the course of system maintenance, the activities of authorized users may also be monitored.
Anyone using this system expressly consents to such monitoring and is advised that if such monitoring reveals possible evidence of criminal activity, system personnel may provide the evidence of such monitoring to law enforcement officials.

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https://webconnect.uscdcb.com/#/account/log-in

Queries Tab – **ID entry**

Choose a format of the ID that you are querying from the dropdown menu

Queries Tab – **ID entry**

1. Enter Animal ID in a format you chose
2. Click "Run Query"

Queries Tab - **ID/Pedigree**

Available information:
- General profile of the animal
- Three-generation family tree
- Cross reference information
- Clonal family information
- Edit button available if you have privilege to edit the info

Queries Tab - **ID/Pedigree Query (Edit)**

1. Edit/add information in the box
2. Click "Next" to proceed with submission

Queries Tab – **Evaluation Query**

Available information
-Latest triannual evaluation result if available

Queries Tab – **Genotype Query**

Available information:
-General profile of the animal
-Nomination status
-Billing status
-Sample status
-Genotype confirmations
-Genotype conflicts
-Other close relatives

Queries Tab – **Genotype Query (Nomination)**

Click on Edit button in Nomination table

Queries Tab– **Genotype (Nomination)**

Name:
DOB: 2024-05-14
Sex: F
Multi-Birth Code: 1
ID Source Code: N
Pedigree Source Code: N
Blend Code: S

Genotyped:

Dam

ID: HO840003219051524
Name:
DOB: 2021-06-29
Source Code: D
Genotyped: No

ID: BS840003137435947
Name: MEADO-BROOK SAGES SHINE ET
DOB: 2017-07-26
Source Code: B
Genotyped: Yes

Does the animal qualify for a genomic evaluation?

Nomination Status

ANIMAL: BS840003272897071

① Update Information   ② Review Changes and Submit

Nomination Status

Nom. Date	Requester	Group/Herd	Herd Code Difference Reason Code	Fee	Fee Assigned Date	Mod Date	
2024-07-12	Zoetis	35631032		1	2024-07-12	2024-07-12 16:44	✏

[+]

Cancel   **Next**

Sample ID	Barcode & Position	Chip Name	Fee	Usability Ind.	Pending		...ival Date	Conflicts
Z004648446	208160920079 R23C04	ZOC	1	N			07-25	YES

Genotype Confirmations, Conflicts, and Other Close Relatives

MGS Info   PGS Info   Show All

Sample ID: Z004648446 – Barcode & Position: 208160920079 R23C04 – Usability Indicator: N

Confirmation Type	Genotype Confirmations		Pedigree Chang...	USE
	Confirmed Genotype			Y
	Barcode & Position	Confirmed ID		
No records available.				

1. Click on the edit button (right) if you want to edit existing nomination
2. Click on + if you are entering new nomination
3. Click on "Next"(middle)to proceed

Queries Tab– Genotype (Entering Nomination)

1. Enter nomination information in boxes
2. Click on ✓ to save

3. Click on "Next" to proceed

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Queries Tab– Genotype (submitting Nomination)

Sample Status

ANIMAL: BS840003272897071

① Update Information

② Review Changes and Submit

Nomination Status

Nom. Date	Requester	Group/Herd	Herd Code Difference Reason Code	Fee	Fee Assigned Date	Mod Date	Note
2024-07-12	Zoetis	35631032		1	2024-07-12	2024-07-12 16:44	
2024-07-26	AIPL	35631032		1			New

Back   Submit

1. Review and confirm entered information
2. Click on "Submit"

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Queries Tab– Errors

HORUS003213001120

| ID/Pedigree | Evaluation | Genotype | Progeny | **Errors** | Lactations |

Available information:
- General profile of the animal
- Available error format

Sire

ID:	HO840003149798696
Name:	MAPLEHURST DELROY TORRO-ET
DOB:	2018-08-11
Source Code:	B
Genotyped:	Yes

Animal

Preferred ID:	HORUS003213001120
Name:	
DOB:	2020-03-16
Sex:	M
Multi-Birth Code:	1
ID Source Code:	N
Pedigree Source Code:	N

Dam

ID:	
Name:	
DOB:	
Source Code:	
Genotyped:	

Error Information

☐ Select All ☑ Pedigree (fmt-1) ☐ Lactation (fmt-4) ☐ Reproduction (fmt-5) ☐ Health (fmt-6)   **Apply**   **Clear**

1

2

1. Select error format(s) you would like to receive
2. Click Apply

Queries Tab– Errors

Available information:
-Error from the latest submission

Data Exchange Tab – Create Format

Both input and output are files in Data Exchange!!

Data Exchange Tab – Create Format1

1. Select format of IDs you are entering to query the information at **SEARCH FOR** box
2. Choose the **SEX** of the animals you are querying for by clicking Male or Female
3. Choose what kind of record you want to generate by clicking on a format(s) in the **API REQUEST** box
4. Drop the file in the drop box
5. Click on "Create Format"

Data Exchange Tab – Create Format1

1. Just be patient while file is being generated...
2. You will see complete when the file is ready
3. Click on ⬇ to download the output file

Data Exchange Tab – Create Format1
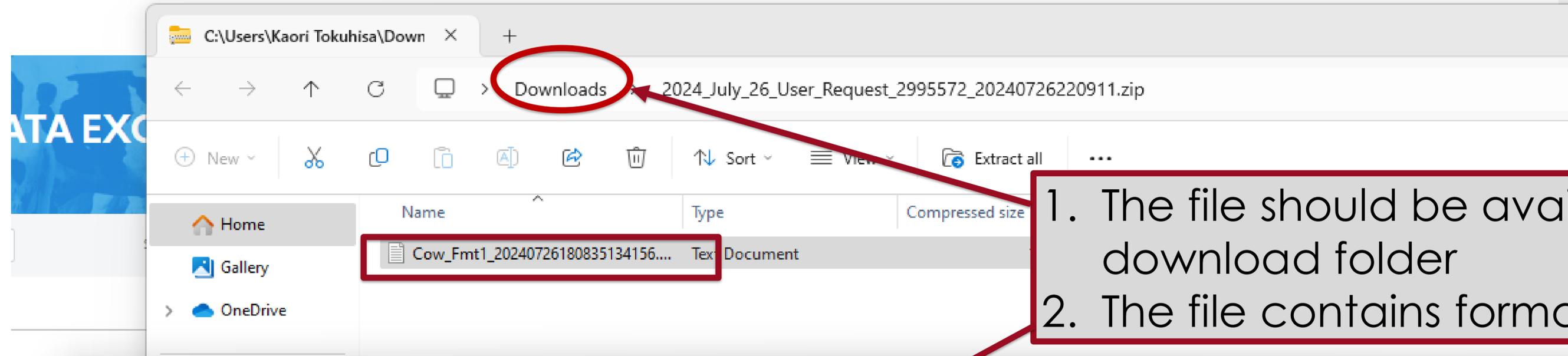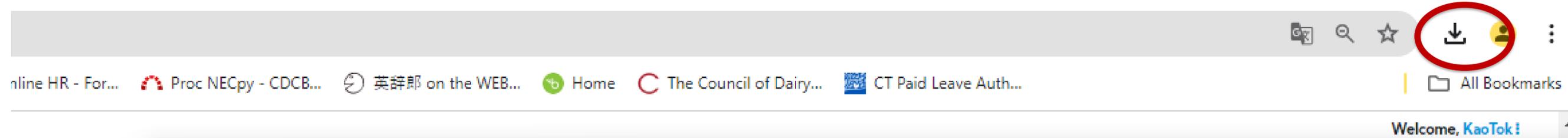
1. The file should be available in your download folder
2. The file contains format1 records
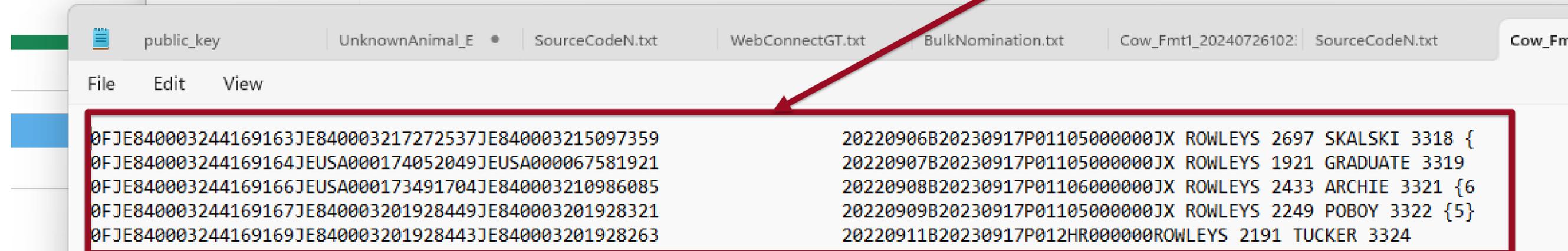
C:\Users\Kaori Tokuhisa\Down

Downloads ... 2024_July_26_User_Request_2995572_20240726220911.zip

Name | Type | Compressed size
Cow_Fmt1_20240726180835134156.... | Text Document

public_key   UnknownAnimal_E   SourceCodeN.txt   WebConnectGT.txt   BulkNomination.txt   Cow_Fmt1_20240726102   SourceCodeN.txt   Cow_Fmt1

File   Edit   View

```
0FJE840003244169163JE840003217272537JE840003215097359            20220906B20230917P011050000000JX ROWLEYS 2697 SKALSKI 3318 {
0FJE840003244169164JEUSA000174052049JEUSA000067581921            20220907B20230917P011050000000JX ROWLEYS 1921 GRADUATE 3319
0FJE840003244169166JEUSA000173491704JE840003210986085            20220908B20230917P011060000000JX ROWLEYS 2433 ARCHIE 3321 {6
0FJE840003244169167JE840003201928449JE840003201928321           20220909B20230917P011050000000JX ROWLEYS 2249 POBOY 3322 {5}
0FJE840003244169169JE840003201928443JE840003201928263           20220911B20230917P012HR000000ROWLEYS 2191 TUCKER 3324
```

Special Section - Reports

Affiliate specific reports (conflicts, fee, PVR etc) can be queried

Special Section – Reports

Queries Data Exchange ▾ Special Section ▾ Top Animal Listing Summary St...

REPORTS

REPORT Lists Genotypes with Fee Code=N Loaded in the past 6 Months ▾

AFFILIATE AIPL

Lists Genotypes with Fee Code=N Loaded in the past 6 Months

Fee Codes for Genotypes Loaded Since the Last Invoice

Lists Parentage only Genotypes Since the Previous Genomic Run

Reports Missing Animal ID for a Requester

Conflicts for Genotypes Loaded in the Past 45 Days

Clear

1. Find and choose a report that you would like to retrieve

Queries Data Exchange ▾ Special Section ▾ Top Animal Listing Summary Stats ▾ National...

REPORTS

REPORT Lists Genotypes with Fee Code=N Loaded in the past 6 Months ▾

AFFILIATE AIPL

Run Query Clear

2. Click on "Run Query" after choosing a report

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Special Section – Reports (Results)

1. Result(s) will be displayed in the table
2. CSV file is also available

Special Section – Bulk Nomination

Useful when you nominate many animals with similar nomination information (same non-US country, herd ID, etc)

Special Section – Bulk Nomination

1. Enter a list of animal IDs in the input box (17 or 18-bytes animal ID only)
2. Select nomination information (requesterID, Group/herd ID, and fee code are required)
3. Click on "Run Query"

Special Section – Bulk Nomination

	Animal	Sex	Sire	Dam	DOB	MBC	SRC	
▾	JE840003244169163	F	JE840003217272537	JE840003215097359	2022-09-06	1	B	⊗

	Requester	Fee	Group/Herd	HRC
	AIPL ⌄	CDCB Determined ⌄	USA00000	⌄

	Animal	Sex	Sire	Dam	DOB	MBC	SRC	
▾	JE840003244169164	F	JEUSA000174052049	JEUSA000067581921	2022-09-07	1	B	⊗

	Requester	Fee	Group/Herd	HRC
	AIPL ⌄	CDCB Determined ⌄	USA00000	⌄

	Animal	Sex	Sire	Dam	DOB
▾	JE840003244169166	F	JEUSA000173491704	JE840003210986085	2022-09-08

	Requester	Fee	Group/Herd	HRC
	AIPL ⌄	CDCB Determined ⌄	USA00000	⌄

	Animal	Sex	Sire	Dam	DOB
▾	JE840003244169167	F	JE840003201928449	JE840003201928321	2022-09-09

	Requester	Fee	Group/Herd	HRC
	AIPL ⌄	CDCB Determined ⌄	USA00000	⌄

	Animal	Sex	Sire	Dam	DOB	MBC	SRC	
▾	JE840003244169169	F	JE840003201928443	JE840003201928263	2022-09-11	2	B	⊗

	Requester	Fee	Group/Herd	HRC
	AIPL ⌄	CDCB Determined ⌄	USA00000	⌄

1. Update information (Requester ID, fee code, group ID/herd ID or HRC)if necessary

2. Click "Next" to proceed

Next

Special Section – Bulk Nomination

Animal	Nom. Date	Requester	Group/Herd	HRC	Fee	Fee Assigned Date	Mod. Date	Note
JE840003244169163	2024-07-26	AIPL	USA00000		2	2022-10-08	2022-10-08 03:50	Update Requester, Group/Herd
JE840003244169164	2024-07-26	AIPL	USA00000		2	2022-10-08	2022-10-08 03:50	Update Requester, Group/Herd
JE840003244169166	2024-07-26	AIPL	USA00000		2	2022-10-08	2022-10-08 03:50	Update Requester, Group/Herd
JE840003244169167	2024-07-26	AIPL	USA00000		2	2022-10-08	2022-10-08 03:50	Update Requester, Group/Herd
JE840003244169169	2024-07-26	AIPL	USA00000		2	2022-10-08	2022-10-08 03:50	Update Requester, Group/Herd

Animal ID (17 bytes)/ Animal ID + Sex Code (18 bytes)   Found 5/5 result sets   REQUESTER*   AIPL

1. Input Information
2. Review Changes and Submit

Back   Submit

1. Confirm entered information
2. Click submit

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Special Section – Nominate Animal

Only query in WebConnect that allows you to nominate animals unknown to CDCB!

1. Select "Nominate" Animal
2. Enter a (list of) animal ID(s) in the input box (17 or 18-bytes animal ID only)
3. Click on "Run Query"

Special Section – Nominate Animal

Special Section – Nominate Animal

CREATE ANIMAL AND NOMINATE

1. Navigated to input information page

Sex	Animal	Sire	Dam	DOB	MBC	Name	Group/Herd	Fee	HRC	Requester
⌄	HOBEL000566137100			📅	⌄			⌄	⌄	AIPL ⌄	✖

Cancel   Next

Sex	Animal	Sire	Dam	DOB	MBC	Name	Group/Herd	Fee	HRC	Requester
F ⌄	HOBEL000566137100	HODNK000000257840	HOBEL000163563018	2024-07-26 📅	3 ⌄	EXAMPLE	BEL00000	7 ⌄	⌄	AIPL ⌄	✖

2. Enter nomination information
3. Click "Next" button

Cancel   Next

4. Review entered information and click on "Submit" button

Sex	Animal	Sire	Dam	DOB	MBC	Name	Group/Herd	Fee	HRC	Requester
F	HOBEL000566137100	HODNK000000257840	HOBEL000163563018	2024-07-26	3	EXAMPLE	BEL00000	7		AIPL

Back   Submit

Documentation

- https://redmine.uscdcb.com/projects/webconnect-production/wiki

Thank you for your attention!

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Description of Each Query
(supplemental info)

Queries Tab

Types of information	Contents
ID/pedigree	• Basic animal's pedigree information • Mostly for viewing info, but **edit function is available**
Evaluation	• Displays the **latest tri-annual evaluation** if available
Genotype	• Nomination status, Sample status, Genotype Confirmations, Conflicts, and Other Close Relatives are included • Adding and updating nomination, and fixing genomic conflicts can be done
Error	• Error information can be queried

Data Exchange Tab

Types of information	Contents
ID/Pedigree	• Variety of pedigree info can be queried (ex. 5gen ped, Aliases, Clonal members, PVR etc)
Genotype	• Nomination Status • Genotype info
Special Section	• Herd code+ fee for specific ID, GT history, SampleID lookup, suggested Dam • GT history is useful to see if the GT was ever moved from one ID to another
Reports	• **Affiliate specific** reports (conflicts,fee, etc) can be queried
Create Format	• **Format1** (M/F per query) can be generated

Special Section(1)

Types of information	Contents
Get Fee	• Query for **fee code and amount** based on animal ID or Herd ID • Useful to find out the fee code **before making nomination**
Nominate Animal	• **This is the only place in WebConnect where unknown ID to CDCB can be nominated**
Bulk Nomination	• Useful when animals are from same herd or from same non-US country
Bulk Transfer	• **Up to 50** animal's requester ID can be transferred to another requester ID

Special Section(2)

Types of information	Contents
Move Genotype	• Genotypes should be reassigned to another animal by using this app • **Sample ID** should be used for Negative/zero key Genotypes
Check Fmt1 Record	• Checks format1 to see input is consistent with existing pedigree. Validity of the contents is not checked
Suggested Dam	• Suggests dam based on group name, birth date and discovered MGS
Find Grandsire	• Finds possible grandsires based on SNP % conflict

Special Section(3)

Types of information	Contents
History of Genotypes	• useful to see if the Genotype was ever moved from one ID to another
Sample ID Look Up	• Finds sample information based on animal ID/sample ID • ID, sex, NAAB code, Name, DOB, Sample ID, status, Usability, and requester ID
ID Range	• lists all the IDs that are in the CDCB database for a specified range of IDs • Useful to see if a range of IDs has already been used.
Reports	• **Affiliate specific** reports (conflicts, fee, etc) can be queried
Performance Metrics	• Reports monthly performance on each metric for past 13 months • This report is reviewed during annual review