Default values

Pedigree named fields:

- 1. PedID: Pedigree ID Animal primary unique identifier
- 2. ISUBPID:
- 3. PreRegID: temporary pesudo "intl_ID" by a temp serial together with 'ISU' after essential parts of an Intl_ID; To be replaced by Intl_ID.
- 4. HealthID: 5. DummyID:
- 6. Data ownership: Data "ownership" is represented by BOTH `labID' and `UserID' because (1) responsibility will be traceable to individuals; (2) personel changes - eventual owner is `lab'. For this reason when a user works for both labs, s/he needs to have 2 separate accounts.
- 7. DamPedID/SirPedID: Added in 2024, replacing (old)Dam_ID/Sir_ID which use Intnl_IDs. The current DamPedID/SirPedID are looked up and filled by PedID composed following above rules matching their Intl_IDs.
- 8. SpecificDates:
- Weaning date
- Yearling date
- Slaughter date
- Castration date
- 9. Flag Controlled vocabulary

Basic structural types:

- Styles of table box borders:
- 1. Thick lines: key info tables.
- 2. Solid lines: existing tables 3. Dotted lines: concepturally proposed (not yet implemented)
- 4. Gray lines: existing tables 5. Black lines: existing and potentially to
- improve 6. light cyan lines: newly created tables since year 2000.

Controlled vocabulary

vocID

vocabulary

Record_Date

meaning

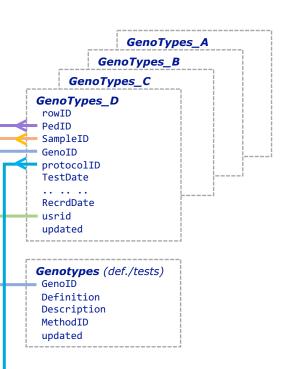
Fit data flows:

Pedigree pID (primary) AnimalRegIDs Location_track **Breed_associations** PedID • rID Asso_ID FarmID PedID ' PedID Name updated PreRegID AssoID Address Intl_ID RegNum Phone ISUBPID RegDate Emails HealthID usrid Website Farms DummyID updated Notes FarmID Sex (M/F)Created Name BirthDate Address WeaningDate Phone YearlingDate Email **CastrOvrmDate** Samples Created LeaveDate SampID **SlaughterDate** PedID **DamPedID** New; use PedID; SampleName SirePedID New; use PedID Tissue_Type Dam_ID Old; Use Intl_ID; Descrition Sire_ID Old; Use Intl_ID SamplProtocol BreedID SampleDate TWorET ProcessProtocol AIorNA ProcessDate isSteer (yes/NULL) ProcessedProduct Season (S|F)Storage_PlateLoc Presentation usrid Remarks updated **Breeds** Genotyped Breed_ID Lab Breed_name cuid LBO_ID created Notes updated Created Labs labID Name Piname Piemail Piphone PhenoMeasure_A Address PhenoMeasure_B Notes PhenoMeasure_C Created **PhenoMeasures** rowID AnimalPedID SampleID (Lab storage) Users Repeats (eg. 3 milkings a day, etc) uid PhenoID (Phenotype definition) UserName MeasuredValue fname MeasurmtUnit lname MeasurDate labID SDID • roles Protocol email israt. Password usrid access_ipv4 Record_Date randomstr TotalLogins RecentLogin created **SpecificDate** SizeMeasurements Ped_ID SpecificDate AnimalID (intl_ID) created Date User_login_track MeasurementID tkID SDID UserName Value Units access_ipv4 UnitsID UnitsID access dir created UnitsType access_time created Measurement_names MeasurementID Past routine data MeasurementType Multiple (Y|N) (??) collections from farms created scope (pheno/geno/farm/etc) Phenotypes (traits) phenoID 3181. Name Definition Descriptin vt_ID (mappings)

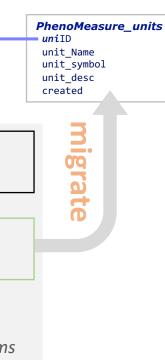
lpt_ID (mappings) cmo_ID (mappings) UBR_ID (mappings)

updated

A Conceptual ER for ISU Cattle DataBase



Protocols ProtocolID
Name category (geno/pheno/sampl) Protocol Description usrid
created updated



Development history

2023.01.21	introduced " Birth_ID " as the unique animal identifier.
2023.01.25	Created "Breeds" table
2023.01.26	Created "Labs " and "Users " tables. Created table "Farms " to record animal
2024.01.19:	locations (go with birth, sampling data
2024 01 10	etc).
2024.01.19	Created a "AnimalRegIDs" table, to
	accommodate multiple breed association register IDs.
2024.01.15	Created "Breed_associations" table.
2024.01.26	Added ` Phenotypes ` table to hold phenotype definitions
2024.01.26	Added Protocols table for protocols
	used in phenotype/genotype and other procedures in the lab.
2024.02.05	Created table ` Location_track ` to record animal locations and movements
	(eg. not just birth time location);
2024.02.21	Added 'AIorNA' field to Pedigree table to
2024.02.21	indicate it's from artificial inseminations or
2024.03.19	natural pregnancy. added ` Phenomeasures `table along
	with `Samples ` table. These work with
	<i>`Phenotypes'</i> to record phnotype
	measurements data. (experimenting in
	DairyDB)
2024.04.26	Changed primary animal ID from Birth_ID
	to PedID (for 'pedigree ID') to avoide
2024 05 06	confusings.
2024.05.06	added DamPedID and SirePedID (to
	replace old <i>Dam_ID/Sire_ID</i> which uses Int'l IDs)
2024.05.09	Subsequence changes from AnimalBirID
	to PedID (<i>AnimalPedID</i>) for all tables.
2024.06.29	to PedID (<i>AnimalPedID</i>) for all tables. Determined the table structure of old trait
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