

**File 711**

for the upload of genotypes sharing permission in GenoEx-GDE database.

**711 format**

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| **Field Name** | Format | Example |
| Record Type 1 | alphanumeric 3 | 711 |
| Animal ID 2 - Breed Code 3 | character 3 | BSW |
| Animal ID - Nation Code 4 | character 3 (with the exception of 840) | AUS |
| Animal ID - Sex Code | character 1 | M |
| Animal ID - Registration | alphanumeric 12 | 000000A12345 |
| UUID 5 | alphanumeric 36 | assigned automatically by gxprep.py program |
| Shareable with organization(s) 7 | character, repeatable | BFRO,IBC |

1. Record type is always 711 for this File Format
2. Please see Interbull Bulletin 28. Each file can only contain any given animal in one row.
3. Breed of animal (3 characters, capital letters)
4. ISO 3166-1 alpha-3 codes (3 characters)
5. UUID, used as reference to every uploaded genotype sequence in the 706 file. Additional information about generation of UUID can be found at <https://wiki.interbull.org/public/uuid?action=print>
6. Comma-separated list of zero or more organizations that should be allowed to download the associated genotype

**711 example**

711;BSW;ITA;M;000000A12345;09c98b1e-6af8-4254-9768-58d7cd1ddafd;BFRO,IBC

**711 upload**

In order to upload sharing permissions to GenoEx-GDE database, the file in 711 format has to be zipped. Zip format is the only one accepted by GenoEx-GDE database for the upload. Zip file can be created using the support program gxprep.py as described here: <https://wiki.interbull.org/public/GDE_gxprep_manual#zip_example> If the User creates the zip file without using the support program gxprep.py, please make sure that the file containing sharing permissions in 711 format and compressed in the zip archive has name **711\_file.csv**.