



To be returned by e-mail to: DNA@icar.org

Annex VII

## Renewal form for the ICAR accreditation for Parentage Testing in Cattle by STR and SNP

### 1. ADDRESS DETAILS (fill out)

Country: .....

Laboratory name: .....

Contact person: .....

Address: .....

Telephone: .....

E-mail: .....

### 2. BILLING INFORMATION (fill out)

Name : .....

Address.....

VAT Number .....

Contact person and email.....

### 3. RENEWAL APPLICATION (check box)

Microsatellite-based parentage  YES  NO

Year of previous accreditation .....

SNP-based parentage  YES  NO

Year of previous accreditation .....

### 4. RELEVANT CHANGES IN PROCEDURES & MANAGEMENT

- a. Change in head of laboratory  YES  NO  
If Yes, please explain: .....
- b. Changes in equipment  YES  NO  
If Yes, please explain: .....



**5. Dates of last revisions of equipment:**

Type of equipment	Date of purchase	Date of last revision

**6. CERTIFICATION AND NUMBER OF SAMPLES TESTED**

a. Certification (tick the box and send a copy of the certification, and English translation if necessary, to ICAR Secretariat. Please note that an ISO certification is a minimum requirement!):

ISO17025 certification  
.....

ISO9001 certification  
.....

Other or no certification (No need to continue application in this case)

**7. PARTICIPATION AND PERFORMANCE IN ISAG RING TESTS**

- No ring test participation
- ISAG ring test participation

Year of the last ISAG ring test \_\_\_\_\_

In the last ISAG ring test (2018), the laboratory achieved the following result for the Typing Comparison Test:

- Absolute genotyping accuracy rank 1
- Absolute genotyping accuracy rank 2

**Please notice:**

1. provide a copy of your ISAG certificate, when available and describe only the results obtained with the compulsory ISAG recommended markers ;
2. to clarify the meaning of “number of genotypes,” a ring test on 20 individuals analysed with 12 microsatellites produces 240 genotypes, for example.

- \_\_\_\_\_ Number of samples
- \_\_\_\_\_ Number of markers
- \_\_\_\_\_ Number of correct genotypes
- \_\_\_\_\_ Number of missing genotypes
- \_\_\_\_\_ Number of incorrect genotypes



**8. MARKER SET AND NOMENCLATURE**

Changes in marker sets  YES  NO

If Yes, please explain: .....

Changes in nomenclature  YES  NO

If Yes, please explain: .....

**Number of animals typed with markers**

Last year: .....

This year (expected): .....