

ACCESSION FORM FOR PUBLIC DEPOSIT OF MICROORGANISM

Microorganism type

☐ bacteria ☐ yeast ☐ filamentous fungi

To be filled by CIM

Collection accession number:

Reception date:

Deposit date:

Microorganism designation

Existing or proposed scientific name:

Additional classification info (subspecies, biovar, serotype, mutant etc):

Strain designation used by the depositor:

Microorganism characteristics

Risk Assessment

Risk group (CIM accepts only microorganisms from group 1 and 2): ☐ group 1 ☐ group 2

According to which regulations has the assessment to a risk group been carried out?:

☐ Directive 2000/54/EC

☐ Other (please, specify):

Microorganism is pathogenic for: ☐ humans ☐ animals ☐ plants
☐ unknown ☐ other specify:

Growth and maintenance

Temperature [°C]:

Incubation time:

Metabolism type: ☐ Strict anaerobe ☐ Facultative anaerobe ☐ Microaerophilic ☐ Aerobic

Special growth requirements (light, gas phases, etc.):

Gram staining:

Is the microorganism genetically modified? ☐ Yes (Fill in annex I) ☐ No

Is the microorganism capable of surviving freezing process? ☐ Yes ☐ No ☐ Unknown

Recommended long-term preservation methods and/or conditions:

Culture medium composition:

Genetic material

Please provide below GenBank accession number or attach file with microorganism's DNA sequence, if available:

Strain properties.

Please describe below as thoroughly as possible (production, antimicrobial resistance, sensitivity, auxotrophy, biological indicator, etc.):

Strain sampling and isolation:

Sample source:

Sampling localization:

Sampling date (YYYY-MM-DD):

Sampled by (person/institution):

Isolation localization:

Isolation date (YYYY-MM-DD):

Isolated by (person/institution)*:

*If you did not isolate the strain, please indicate all scientists or laboratories which maintained it before you.

Reasons for deposit

Please list below (biotechnological applications, quality control, etc.):

Methods used for strain identification

Identified by (person/institution):

Attach document or describe employed methods below:

Nagoya Protocol (www.cbd.int/abs) related information

Geographical location of sampling:

Internationally Recognized Certificate of Compliance (IRCC):

☐ Applicable (attach document) ☐ Not applicable

Sampling permit/Prior Informed Consent (PIC): ☐ Applicable (attach document) ☐ Not applicable

Details of Mutually Agreed Terms – MAT: ☐ Applicable (attach document) ☐ Not applicable

AGREEMENT FOR DEPOSIT IN THE OPEN COLLECTION:

1. Information regarding public deposits that are pending of taxonomic description will be kept confidential until description is published or until the depositor authorizes the publication in the open catalogue. If the information is not published after 4 years from the deposit, CIM reserves the right to release the strain to the public catalogue.
2. According to the CBD (Convention on Biological Diversity), is the depositor responsibility to ensure that the samples from which the strains were isolated, were collected according to the legislation of the country of origin and that the deposit of the strains in an open collection does not infringe any national regulation.
3. The undersigned declares that information provided in this form is true and accepts that CIM distributes subcultures of the deposited strain according to the terms specified in the CIM MTA.

Depositor's data

First name:

Family name:

Institution name:

Postal Address:

Postal code and city name:

Country:

Telephone:

Fax:

E-mail:

Date:

Signature:

ANNEX I

Fill only if the microorganism is genetically manipulated

Host organism

Scientific name:

Risk group: ☐ group 1 ☐ group 2

Pathogenic for: ☐ humans ☐ animals ☐ plants specify:

Sensitivities:

Resistances:

Auxotrophies:

Special properties (genetic recombination, restriction/modification system, etc.):

Donor organism

Scientific name:

Risk group: ☐ group 1 ☐ group 2

Pathogenic for: ☐ humans ☐ animals ☐ plants specify:

Size of the cloned DNA fragment (in bp):

☐ complete genome ☐ subgenomic ☐ subgenic

Additional description of the DNA cloned fragment (attach relevant documents or describe below):