

# SNIF: GMOB-2024-30970

**Domain:**

GMO

**Authorisation type:**

Deliberate release of GMO or a combination of GMOs for any other purpose than for placing on the market

**Application type:**

Summary notification for the release of genetically modified higher organisms other than higher plants in line with Decision 2002/813/EC Annex, Part 1.

**Recipient Member State:**

Norway

**Competent Authority:**

Section for Invasive Species and International Trade

# **A- General information**

## **Details of notification**

### **Details of notification**

Study YTB323

### **Member State of notification**

NOT PROVIDED

### **Title of the project**

CD19-directed CAR-T cell therapy (YTB323) in patients with B-cell malignancies or autoimmune diseases.

### **Proposed period of release**

#### **Starting date**

2024-10-31

#### **Finishing date**

2034-10-31

## **Notifier**

### **Name of institute or company**

Novartis Pharma AG, Postfach, 4002 Basel, Switzerland

### **Email**

Not provided

### **Phone number**

Not provided

### **Website**

Not provided

### **Address**

Not provided

### **Post code**

Not provided

### **Country**

Not provided

## **GMO characterisation**

### **(a) Indicate whether the GMO is a:**

#### **Viroid**

No

#### **RNA virus**

No

#### **DNA virus**

No

#### **Bacterium**

No

#### **Fungus**

No

#### **Animal**

Yes

**Select from following options:**

**Mammal**

Yes

**Insect**

No

**Fish**

No

**Other animal**

No

**Other**

No

**(b) Identity of the GMO (genus and species)**

human

**(c) Genetic stability - according to Annex IIIa, II, A(10)**

yes

**Is the same GMO release planned elsewhere in the Community (in conformity with Article 6(1)), by the same notifier?**

Yes

**Country**

Austria

France

Germany

Spain

Italy

Netherlands

**Has the same GMO been notified for release elsewhere in the Community by the same notifier?**

Yes

**Country**

AT

**Notification number**

Not provided

**Country**

DE

**Notification number**

Not provided

**Country**

IT

**Notification number**

Not provided

**Country**

FR

**Notification number**

Not provided

**Country**

ES

**Notification number**

Not provided

**Has the same GMO been notified for release or placing on the market outside the Community by the same or other notifier?**

Yes

**Country**

United States

**Notification number**

Not provided

**Country**

Australia

**Notification number**

Not provided

**Country**

Japan

**Notification number**

Not provided

**Summary of the potential environmental impact of the release of the GMOs.**

An environmental impact is not expected as the release of YTB323 (transduced autologous T cells) is limited to patient administration in hospital settings. According to the environmental risk assessment YTB323 will not reach the environment at large.

## **B. Information relating to the recipient or parental organisms from which the GMO is derived**

### **1. Recipient or parental organism characterisation**

**Indicate whether the recipient or parental organism is a:**

**Viroid**

No

**RNA virus**

No

**DNA virus**

No

**Bacterium**

No

**Fungus**

No

**Animal**

Yes

**Select from following options:**

**Mammal**

Yes

**Insect**

No

**Fish**

No

**Other animal**

No

**Other**

No

### **2. Name**

**(i) Order and/or higher taxon (for animals)**

Primates

**(ii) Genus**

Homo

**(iii) Species**

Homo sapiens

**(iv) Subspecies**

NA

**(v) Strain**

NA

**(vi) Pathovar (biotype, ecotype, race, etc.)**

NA

### **3. Geographical distribution of the organism**

**(a) Indigenous to, or otherwise established in, the country where the notification is made:**

yes

**(b) Indigenous to, or otherwise established in, other EC countries:**

notKnown

**(c) Is it frequently used in the country where the notification is made?**

Yes

**(d) Is it frequently kept in the country where the notification is made?**

Yes

#### **4. Natural habitat of the organism**

**(a) Is the organism a microorganism ?**

No

**Specify**

**(b) Is the organism an animal?**

Yes

**Natural habitat or usual agroecosystem**

NA, Parental organism and recipient are human

#### **5(a) Detection Techniques**

**Detection Techniques**

Common techniques of blood cell analysis

#### **5(b) Identification Techniques**

**Identification Techniques**

Common techniques of blood cell analysis

**6. Is the recipient organism classified under existing Community rules relating to the protection of human health and/or the environment?**

No

**7. Is the recipient organism significantly pathogenic or harmful in any other way (including its extracellular products), either living or dead?**

no

#### **8. Information concerning reproduction**

**(a) Generation time in natural ecosystems:**

NA

**(b) Generation time in the ecosystem where the release will take place:**

NA

**(c) Way of reproduction**

Sexual

**(d) Factors affecting reproduction:**

NA

#### **9. Survivability**

**(a) Ability to form structures enhancing survival or dormancy:**

**(i) endospores**

No

**(ii) cysts**

No

**(iii) sclerotia**

No

**(iv) asexual spores (fungi)**

No

**(v) sexual spores (fungi)**

No

**(vi) eggs**

No

**(vii) pupae**

No

**(viii) larvae**

No

**Other**

Yes

**Specify**

Not applicable for human T cells as they cannot survive outside the human body

**(b) Relevant factors affecting survivability**

The survival of human blood cells requires a complex combination of special media, temperature and CO<sub>2</sub>. The environmental conditions outside the host are substantially different and not appropriate for its survival (temperature, pH, UV, and a change in the biophysical and biochemical conditions).

**10(a) Ways of dissemination**

Blood cells can only be transmitted between individuals through injection. No dissemination in the environment is possible due to fast inactivation.

**10(b) Factors affecting dissemination**

The immune system of people other than the donor will eliminate the blood cells.

**11. Previous genetic modifications of the recipient or parental organism already notified for release in the country where the notification is made (give notification numbers)**

NA

## **C. Information relating to the genetic modification**

### **1. Type of the genetic modification**

**Insertion of genetic material**

Yes

**Deletion of genetic material**

No

**Base substitution**

No

**Cell fusion**

No

**Other**

No

### **2. Intended outcome of the genetic modification**

YTB323 is a novel, investigational, adoptive cancer immunotherapy whereby autologous T cells are genetically modified to express a transmembrane chimeric antigen receptor (CAR) to target CD19 on the cell surface of (malignant) B cells. YTB323 is also being explored in autoimmune diseases involving pathogenic B cells such as severe refractory Systemic Lupus Erythematosus (srSLE).

### **3(a) Has a vector been used in the process of modification?**

Yes

### **3(b) If yes, is the vector wholly or partially present in the modified organism?**

Yes

Yes

### **4. If the answer to 3(b) is yes, supply the following information**

#### **(a) Type of vector**

**Plasmid**

No

**Bacteriophage**

No

**Virus**

Yes

**Cosmid**

No

**Transposable element**

No

**Other**

No

#### **(b) Identity of the vector**

VSV-G pseudotyped replication-deficient HIV-1-derived viral vector of the 3rd generation.

#### **(c) Host range of the vector**

The viral vector is pseudotyped, since it has the Vesicular Stomatitis Virus (VSV) envelope VSV-G, which provides the capacity to transduce many different non-dividing human and animal cells.



**(d) Presence in the vector of sequences giving a selectable or identifiable phenotype**

No

**(e) Constituent fragments of the vector**

Self-inactivating replication deficient lentiviral vector including minimal HIV-1 derived sequences required for vector packaging, reverse transcription and integration of the vector genome into the host cell genome (LTRs, packaging signal, RRE and CPPT) in addition the expression cassette for the expression of an anti-CD19 directed chimeric antigen receptor.

**(f) Method for introducing the vector into the recipient organism**

**(i) transformation**

No

**(ii) electroporation**

No

**(iii) macroinjection**

No

**(iv) microinjection**

No

**(v) infection**

No

**Other**

Yes

**Specify**

transduction

**6. Composition of the insert**

**(a) Composition of the insert**

The vector sequence integrated into the CTL019 cell genome consist of minimal HIV-1 derived self-inactivating lentiviral sequences required for vector packaging, reverse transcription and integration of the vector genome into the host cell genome (LTRs, packaging signal, RRE and cPPT) in addition to the transgene expression cassette. The transgene expression cassette contains the human elongation factor 1 (EF-1) promoter controlling transgene expression, the transgene and a modified woodchuck hepatitis virus posttranscriptional regulatory element (WPRE), wherein the promoter and X-protein start codon have been mutated to prevent expression, for improved RNA translation and hence increased expression. The transgene is a chimeric antigen receptor targeted against the CD19 antigen (CAR-19). It consists of a murine anti-CD19 scFv, a human CD8 hinge and transmembrane domain, and human 4-1BB (CD137) and CD3 (T-cell receptor ) intracellular signalling domains.

**(b) Source of each constituent part of the insert**

Lentiviral sequences from HIV; WPRE from the Woodchuck HBV virus; anti-CD19 scFV from mouse; and other parts of the CAR from human, as indicated above in (a).

**(c) Intended function each constituent part of the insert in the GMO**

See above.

**(d) Location of the insert in the host organism**

**On a free plasmid**

No

**Integrated in the chromosome**

Yes

**Other**

No

**(e) Does the insert contain parts whose product or function are not known?**

No

## **D. Information on the organism(s) from which the insert is derived**

### **1. Indicate whether it is a:**

**Viroid**

No

**RNA virus**

Yes

**DNA virus**

No

**Bacterium**

No

**Fungus**

No

**Animal**

Yes

### **Select from following options:**

**Mammal**

Yes

**Insect**

No

**Fish**

No

**Other animal**

No

**Other**

No

### **2. Complete name**

**(i) Order and/or higher taxon (for animals)**

NA

**(ii) Family name (for plants)**

NA

**(iii) Genus**

Retrovirus

**(iv) Species**

Human Immunodeficiency virus

**(v) Subspecies**

NA

**(vi) Strain**

HIV-1

**(vii) Cultivar/Breeding line**

NA

**(viii) Pathovar**

NA

**(ix) Common name**

NA

**3. Is the organism significantly pathogenic or harmful in any other way (including its extracellular products), either living or dead?**

yes

**If yes, specify the following:**

**(a) to which of the following organisms?**

**humans**

Yes

**animals**

No

**plants**

No

**Other**

No

**(b) are the donated sequences involved in any way to the pathogenic or harmful properties of the organism?**

yes

**Give the relevant information specified under Annex III A, point II, (A)(11)(d) of Directive 2001/18/EC**  
Wild type HIV is classified as group 3 organism. However, the replication-defective lentiviral vector used for transduction of T cells is not pathogenic anymore as no infectious viral particles can be produced after transduction.

**4. Is the donor organism classified under existing Community rules relating to the protection of human health and the environment, such as Directive 90/679/EEC on the protection of workers from risks related to exposure to biological agents at work?**

No

**5. Do the donor and recipient organism exchange genetic material naturally?**

no

## **E. Information relating to the genetically modified organism**

### **1. Genetic traits and phenotypic characteristics of the recipient or parental organism which have been changed as a result of the genetic modification**

**(a) Is the GMO different from the recipient as far as survivability is concerned?**

no

**Specify**

Not provided

**(b) Is the GMO in any way different from the recipient as far as mode and/or rate of reproduction is concerned?**

no

**Specify**

Not provided

**(c) Is the GMO in any way different from the recipient as far as dissemination is concerned?**

no

**Specify**

Not provided

**(d) Is the GMO in any way different from the recipient as far as pathogenicity is concerned?**

no

**Specify**

Not provided

### **2. Genetic stability of the genetically modified organism**

The chimeric antigen receptor is introduced in the T cells via lentiviral gene transfer and after integration of the SIN vector the gene modified autologous T cells are genetically stable and an integral part of the host DNA.

### **3. Is the GMO significantly pathogenic or harmful in any way (including its extracellular products), either living or dead?**

no

**(b) Give the relevant information under Annex III A, point II (A)(11)(d) and II( C)(2)(i):**

The replication-deficient lentiviral vector genome is integrated as provirus in the T cell genome. No new viral particles can be assembled in the final host cell since the gag gene is not present. In addition, all accessory elements are absent from this viral vector. The transgenes inserted in the lentiviral vector do not code for pathogenicity factors, cytokine-coding sequences, oncogenes, antibiotic resistance genes or otherwise hazardous inserts.

### **4. Description of identification and detection methods**

**(a) Techniques used to detect the GMO in the environment**

Post-administration monitoring of patients for persistence of YTB323 is done using qPCR of the transgene.

**(b) Techniques used to identify the GMO**

Identity of YTB323 is determined by qPCR in transduced cells.

## **F. Information relating to the release**

### **1. Purpose of the release (including any significant potential environmental benefits that may be expected)**

Treatment of B cell malignancies and autoimmune diseases as well as neuroscience involving pathogenic B cells, in which B cells play a key role in pathogenesis through the secretion of pathogenic autoantibodies. YTB323 treatment is not expected to have any effects on the environment, at large, neither negative nor positive.

### **2. Is the site of the release different from the natural habitat or from the ecosystem in which the recipient or parental organism is regularly used, kept or found?**

No

### **3. Information concerning the release and the surrounding area**

#### **(a) Geographical location (administrative region and where appropriate grid reference):**

Hospitals across Europe

#### **(b) Size of the site (m<sup>2</sup>)**

##### **(i) actual release site (m<sup>2</sup>)**

NA, hospital

##### **(ii) wider release area (m<sup>2</sup>)**

NA, hospital

#### **(c) Proximity to internationally recognised biotopes or protected areas (including drinking water reservoirs), which could be affected:**

NA

#### **(d) Flora and fauna including crops, livestock and migratory species which may potentially interact with the GMO**

NA

### **4. Method and amount of release**

#### **(a) Quantities of GMOs to be released:**

YTB323 is a single infusion treatment. The maximum target dose a patient might receive is  $12.5 \times 10^6$  transduced viable T cells per dose.

#### **(b) Duration of the operation:**

The administration will take up to 60 minutes.

**(c ) Methods and procedures to avoid and/or minimise the spread of the GMOs beyond the site of the release**

Standard Hospital Hygiene Measures.

**5. Short description of average environmental conditions (weather, temperature etc.)**

Short description of average environmental conditions (weather, temperature, etc.) Hospital rooms have to fulfill hygiene conditions required for the treatment of immune- compromised patients.

**6. Relevant data regarding previous releases carried out with the same GMO, if any, specially related to the potential environmental and human health impacts from the release**

Various clinical studies in ALL, CLL, and NHL have been carried out and are ongoing with Kymriah product which uses the same lentiviral vector. A long term follow-up study, required for patients exposed to gene therapy products, is ongoing. The GMO has already been released to the environment as part of completed or ongoing clinical trials of Kymriah product without evidence of environmental or human health impacts.

## **G. Interactions of the GMO with the environment and potential impact on the environment, if significantly different from the recipient or parent organism**

### **1. Name of target organisms (if applicable)**

**Applicable?**

Yes

**(i) Order and/or higher taxon (for animals)**

Primates

**(ii) Family name (for plants)**

NA

**(iii) Genus**

Homo

**(iv) Species**

Homo sapiens

**(v) Subspecies**

NA

**(vi) Strain**

NA

**(vii) Cultivar/Breeding line**

NA

**(viii) Pathovar**

NA

**(ix) Common name**

human

### **2. Anticipated mechanism and result of interaction between the released GMOs and the target organism (if applicable)**

**Applicable?**

No

### **3. Any other potentially significant interactions with other organisms in the environment**

None expected.

### **4. Is post-release selection such as increased competitiveness, increased invasiveness for the GMO likely to occur?**

no

### **5. Types of ecosystems to which the GMO could be disseminated from the site of release and in which it could become established**

None, except the dedicated patients who receive the product. Exposure requires direct injection of YTB323. Immune-repressed individuals other than the patients will not participate in the administration of YTB323.



Persons with a functional immune system would eliminate YTB323 upon accidental injection. Simple contact exposure to blood from treated patients will not result in transmission of YTB323 as cells are quickly inactivated under environmental conditions.

**6. Complete name of non-target organisms which (taking into account the nature of the receiving environment) may be unintentionally significantly harmed by the release of the GMO**

**(i) Order and/or higher taxon (for animals)**

NA

**(ii) Family name (for plants)**

NA

**(iii) Genus**

NA

**(iv) Species**

NA

**(v) Subspecies**

NA

**(vi) Strain**

NA

**(vii) Cultivar/Breeding line**

NA

**(viii) Pathovar**

NA

**(ix) Common name**

NA

**7. Likelihood of genetic exchange in vivo**

**(a) from the GO to other organisms in the release ecosystem**

Highly unlikely

**(b) from other organisms to the GMO**

Highly unlikely

**(c) likely consequences of gene transfer**

Highly unlikely

**8. Give references to relevant results (if available) from studies of the behaviour and characteristics of the GMO and its ecological impact carried out in simulated natural environments (e.g. microcosms etc.):**

Not applicable. No such studies have been conducted.

**9. Possible environmentally significant interactions with biogeochemical processes (if different from the recipient or parental organism)**

None

## **H. Information relating to monitoring**

### **H. Information relating to monitoring**

#### **1. Methods for monitoring the GMOs**

No specific GMO monitoring is proposed.

#### **2. Methods for monitoring ecosystem effects**

NA

#### **3. Methods for detecting transfer of the donated genetic material from the GMO to other organisms**

NA

#### **4. Size of the monitoring area (m<sup>2</sup>)**

NA

#### **5. Duration of the monitoring**

NA

#### **6. Frequency of the monitoring**

NA

# **I. Information on post release and waste treatment**

## **I. Information on post release and waste treatment**

### **1. Post-release treatment of the site**

Standard Hospital Hygienic Procedures

### **2. Post-release treatment of the GMOs**

None

### **3(a) Type and amount of waste generated**

Contaminated material used for the administration of YTB323 is composed of disposables.

### **3(b) Treatment of waste**

All disposable waste that has been in contact with the genetically modified human cells during preparation and administration will be disposed of as specific hospital waste or as waste containing GMOs.

## **J. Information on emergency response plans**

### **J. Information on emergency response plans**

#### **1. Methods and procedures for controlling the dissemination of the GMO(s) in case of unexpected spread**

No spread of the GMO is expected. In case of spills decontamination with disinfectants and cleaning according to standard hospital procedures.

#### **2. Methods for removal of the GMOs of the areas potentially affected**

Decontamination with disinfectants and cleaning according to standard hospital procedures.

#### **3. Methods for disposal or sanitation of plants, animals, soils etc. that could be exposed during or after the spread**

NA

#### **4. Plans for protecting human health and the environment in the event of an undesirable effect**

NA