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(57) **Abrégé/Abstract:**

The invention relates to an infectious arenavirus particle that is engineered to contain a genome with the ability to amplify and express its genetic information in infected cells but unable to produce further infectious progeny particles in normal, not genetically engineered cells. One or more of the four arenavirus open reading frames glycoprotein (GP), nucleoprotein (NP), matrix protein Z and RNA-dependent RNA polymerase L are removed or mutated to prevent replication in normal cells but still allowing gene expression in arenavirus vector-infected cells, and foreign genes coding for an antigen or other protein of interest or nucleic acids modulating host gene expression are expressed under control of the arenavirus promoters, internal ribosome entry sites or under control of regulatory elements that can be read by the viral RNA- dependent RNA polymerase, cellular RNA polymerase I, RNA polymerase II or RNA polymerase III. The modified arenaviruses are useful as vaccines and therapeutic agents for a variety of diseases.



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(57) **Abstract:** The invention relates to an infectious arenavirus particle that is engineered to contain a genome with the ability to amplify and express its genetic information in infected cells but unable to produce further infectious progeny particles in normal, not genetically engineered cells. One or more of the four arenavirus open reading frames glycoprotein (GP), nucleoprotein (NP), matrix protein Z and RNA-dependent RNA polymerase L are removed or mutated to prevent replication in normal cells but still allowing gene expression in arenavirus vector-infected cells, and foreign genes coding for an antigen or other protein of interest or nucleic acids modulating host gene expression are expressed under control of the arenavirus promoters, internal ribosome entry sites or under control of regulatory elements that can be read by the viral RNA-dependent RNA polymerase, cellular RNA polymerase I, RNA polymerase II or RNA polymerase III. The modified arenaviruses are useful as vaccines and therapeutic agents for a variety of diseases.

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Replication-defective arenavirus vectors

Field of the invention

- 5 The invention relates to genetically modified arenaviruses suitable as vaccines or gene therapy vectors, and to methods of using these in vaccination and treatment of diseases.

Background of the invention

- 10 Preventive vaccines represent one of the most successful chapters of modern medicine, having led to the worldwide eradication of smallpox and to the control of polio, measles and many other devastating infectious diseases. More recently, vaccines have become available that prevent cancer, and strong efforts are ongoing to exploit "vaccines" in a therapeutic fashion, raising hope for both infection and malignancy. Historically,
- 15 vaccination strategies have comprised a variety of approaches: Starting with the use of wild type infectious agents and the auto-(re)-inoculation of tumor cells, followed by live-attenuated agents and killed tumor tissues, clinical medicine has over time moved more and more to the use of (inert) proteins and/or other extracts (commonly referred to as "antigen") derived from infectious agents or tumors, respectively. This gradual process
- 20 represents the search for safer vaccine formulation, often accompanied, however, by a relative loss in efficacy. In recent years the advancement of biological engineering has made possible yet an additional approach that currently is widely considered among the most promising ones: infectious agents serving as a "ferry" (called "vector") are equipped with an antigen from the pathogen or tumor of choice. Thereby, the immune response of
- 25 the vaccine recipient recognizes the antigen of interest in the context of a strongly immune-enhancing ("immunogenic") context conferred by the vector.

- The "vector approach" has also made possible the directed introduction of foreign genes into living cells at the level of tissue culture but also in multicellular organisms including
- 30 man, and vectors can therefore also be exploited for the expression of genes in cultured cells or in gene therapy.

- A variety of vectors are currently in experimental use, both for vaccination and gene therapy, with the ultimate goal of optimizing efficacy and safety for clinical application
- 35 (vaccinology and gene therapy) or for biotechnology (gene transfer in cell culture).

As a common observation, vectors tend to share general traits of the organism, e.g. virus, they are derived from. The exploitation of a novel family of viruses for vector design promises therefore a novel combination of traits that may confer this new type of vector with unprecedented capabilities and corresponding applications in biomedical application.

- 5 Vector design needs, however, to take into account the safety profile of the organism used, and must come up with a strategy of how to eliminate the organism's pathogenic potential in a manner that does not interfere with desirable traits such as immunogenicity for administration as a vaccine.
- 10 Arenaviruses in general and lymphocytic choriomeningitis virus (LCMV) in particular have been known for more than seventy years to elicit extraordinarily strong and long-lasting humoral and cell-mediated immune responses. Of note, though, protective neutralizing antibody immunity against the viral envelope glycoprotein (GP) is minimal, meaning that infection results in minimal antibody-mediated protection against re-infection if any. Also it
- 15 has been firmly established for decades that owing to their non-cytolytic (not cell-destroying) nature, arenaviruses can, under certain conditions, maintain long-term antigen expression in animals without eliciting disease. Recently, reverse genetic systems for the manipulation of the infectious arenavirus genome (L. Flatz, A. Bergthaler, J.C. de la Torre, and D.D. Pinschewer, Proc Natl Acad Sci U S A 103:4663-4668, 2006; A.B. Sanchez and
- 20 J.C. de la Torre, Virology 350:370, 2006) have been described, but arenaviruses have not so far been exploited as vaccine vectors. Two major obstacles are mainly responsible: i) Arenaviruses can cause overwhelming infection which then can result in serious disease and immunosuppression. ii) The incorporation of foreign antigens of choice has not been possible.

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Summary of the invention

The invention relates to an infectious arenavirus particle that is engineered to contain a genome with the ability to amplify and express its genetic information in infected cells but

30 unable to produce further infectious progeny particles in normal, not genetically engineered cells.

More specifically the invention relates to such arenavirus particles comprising additional ribonucleic acids coding for proteins of interest or modulating host gene expression.

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An arenavirus of the invention comprises a modified genome, wherein

- i) one or more of the four arenavirus open reading frames glycoprotein (GP), nucleoprotein (NP), matrix protein Z and RNA-dependent RNA polymerase L are removed or mutated to prevent propagation of infectivity in normal cells but still allowing gene expression in such cells;
- 5 ii) foreign ribonucleic acids coding for one or more proteins or modulating host gene expression are introduced and are transcribed from one or more of the four arenavirus promoters 5' UTR and 3' UTR of the S segment, and 5' UTR and 3' UTR of the L segment, or from additionally introduced promoters that can be read by the viral RNA-dependent RNA polymerase, by cellular RNA polymerase I, RNA polymerase II or RNA polymerase
- 10 III, respectively, and wherein ribonucleic acids coding for proteins or modulating host gene expression are transcribed either by themselves or as read-through by fusion to arenavirus protein open reading frames; and optionally
- iii) one or more internal ribosome entry sites are introduced in the viral transcript sequence to enhance expression of proteins in the arenavirus-infected cell.

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The invention furthermore relates to vaccines and pharmaceutical preparations comprising such genetically engineered arenaviruses, and to methods of vaccination and gene therapy using these genetically engineered arenaviruses.

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The invention furthermore relates to expression of a protein of interest in a cell culture or to modulation of gene expression in cell culture wherein the cell culture is infected with genetically engineered arenaviruses.

Brief description of the Figures

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Figure 1

Owing to the changes made to the wild type arenavirus genome, the derived arenavirus vectors replicate only in complementing cells.

30

A: Arenavirus vectors (1) can infect both normal cells (2) or complementing cells (C-cells, 3). Upon infection of C-cells further infectious vector progeny are formed whereas infection of normal cells yields either no vector particles or non-infectious ones.

B: C-cells (1) and normal cells (2) were infected with an LCMV-based vector

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(rLCMV/GFP) expressing green fluorescent protein (GFP) instead of LCMV-GP, and

samples of supernatant collected at various time points (3, given in hours). Infectivity in the supernatant (4, given as PFU/ml) was determined in a focus forming assay.

C: The wild type arenavirus genome consists of a large (L; 1) and a small (S; 2) segment. The L segment expresses the L (3) and Z (4) genes, whereas the S segment carries the NP (5) and GP (6) genes. One strategy to generate a replication-deficient arenavirus vector can be to substitute the GP gene for genes of interest, for example GFP (7) or ovalbumin (OVA; 8).

Figure 2

10

Schematic organization of complementing plasmid (C-plasmid), of plasmids for intracellular expression of transacting factors (TF-plasmids) and of plasmids for intracellular expression of arenavirus vector genome segments (GS-plasmids).

15 A: example of a C-plasmid.

B: examples of TF-plasmids expressing the viral NP and L protein, respectively.

C: examples of GS-plasmids expressing the arenavirus vector S and L segments, respectively.

1: polymerase II promoter; 2: viral gene to be expressed for complementation; 3: internal
20 ribosome entry site; 4: mammalian selection marker, such as puromycin resistance gene;
5: polyadenylation signal; 6: ampicillin resistance cassette; 7: origin of replication; 8: viral
trans-acting factor, e.g. NP ORF; 9: viral trans-acting factor, e.g. L ORF; 10: promoter
driving expression of arenavirus genome segment in C-cells, e.g. polymerase I promoter;
11: 5' UTR of the S segment; 12: antigen of interest; 13: IGR of the S segment; 14: NP
25 gene; 15: 3' UTR of the S segment; 16: polymerase I terminator; 17: 5'UTR of the L
segment; 18: Z gene; 19: IGR of the L segment; 20: L gene; 21: 3'UTR of the L segment.

Figure 3

30 Arenavirus vectors are cleared within days after inoculation, and accordingly do not cause immunosuppression in vaccine recipients.

A: On day 0, mice were immunized intravenously with rLCMV/GFP (1). At different time points thereafter (2; given in days), viral genome copies (3, given in log10) were
35 measured in spleen.

B: As primary immunization/infection (1°), mice were either immunized with rLCMV/OVA (5) or were left uninfected (6) or were infected with LCMV wild type (7). On day 20, all mice were given an intraperitoneal infection (2°) with vesicular stomatitis virus (VSV, 8). Subsequently, blood was collected for measuring antiviral and anti-vector T cell responses (3) and antiviral antibody responses (4). On day 28, H-2K^b-SIINFEKL (ovalbumin-derived CD8+ T cell epitope) specific CD8+ T cells (9) and H-2K^b-VSV-NP52-29 (VSV-NP-derived CD8+ T cell epitope) specific CD8+ T cells (10) were measured in peripheral blood by intracellular staining for interferon gamma upon peptide restimulation (values indicate the percentage of specific cells amongst CD8+ T cells). On day 27 (indicated as "d7" referring to the time point after VSV infection), day 29 (indicated as "d9") and day 61 (indicated as "d41"), serum was tested for total VSV neutralizing antibodies (11) and for beta-mercapto-ethanol-resistant IgG (12) in a 50% plaque reduction assay (values given as -log2 of 40-fold prediluted serum).

15 *Figure 4*

Arenavirus vectors elicit high frequencies of long-lived memory CD8+ T cells and long-lived antibody memory at high titers.

20 A: Mice were immunized with rLCMV/OVA and blood samples were collected over time (2; given in days) for measuring the frequency of H2K^b-OVA/SIINFEKL specific CD8+ T cells using MHC class I tetramers (3, values indicate the frequency of tetramer-positive CD8+ T cells within the CD8+ T cell compartment).

25 B: Mice were immunized with rLCMV/OVA at the indicated dose (1), either via the subcutaneous (s.c.) or intravenous (i.v.) route, and OVA-specific IgG in the serum was measured by ELISA on day 14 (5) and day 58 (6). Values are expressed as the dilution of serum yielding twice background optical density (OD) measurements.

Figure 5

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Arenavirus vectors do not cause central nervous system disease

35 Mice were inoculated intracerebrally with rLCMV/OVA (open squares) or with wild type LCMV (closed circles), and were monitored at the indicated time points (1, indicated in days) for clinical signs of terminal choriomeningitis. For each time point, the number of healthy animals per number of animals tested (2) is displayed.

Figure 6

5 Arenavirus vectors confer T cell and antibody-mediated protection against infectious challenge.

10 A) On day 0 of the experiment, mice were vaccinated using either rLCMV/OVA (group AA) or using a rLCMV control vector expressing the irrelevant Cre recombinase antigen as negative control (group BB). Intravenous challenge with recombinant *Listeria monocytogenes* expressing OVA was performed after an interval of either 16 or 58 days (d16, d58). Four days after challenge, bacterial titers (1) were measured in the spleen of the animals (displayed in log₁₀ colony forming units per organ). Black circles indicate values of individual mice. Vertical lines indicate the mean value of each group.

15 B) Type I interferon receptor-deficient mice were vaccinated (closed squares) with a LCMV vector that expresses an antigenic but non-functional variant of the vesicular stomatitis virus envelope protein G (modified by insertion of a foreign peptide sequence in its ectodomain) or were left without vaccination (open circles). One month later, all animals were challenged intravenously with 2x10⁶ PFU vesicular stomatitis virus. At the indicated time points after challenge (2, indicated in days), the animals were monitored for clinical signs of terminal myeloencephalitis. For each time point and group, healthy survival is indicated as the number of healthy animals per number of animals tested (3).

Detailed description of the invention

25 The invention relates to infectious arenavirus particles, referred to as arenavirus vectors, that are engineered to contain a genome with the ability to amplify and express its genetic information in infected cells but unable to produce further infectious progeny particles in normal, not genetically engineered cells. This principle is shown schematically in Figure 1A. Example data are presented in Figure 1B.

30

Replication of arenavirus vectors requires genetically engineered cells complementing the replication-deficient vector. Upon infection of a cell, the arenavirus vector genome expresses not only arenavirus proteins but also additional proteins of interest, for example antigens of interest. Arenavirus vectors are produced by standard reverse genetic techniques as described for LCMV (L. Flatz, A. Bergthaler, J.C. de la Torre, and D.D. Pinschewer, Proc Natl Acad Sci U S A 103:4663-4668, 2006; A.B. Sanchez and J.C. de la

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Torre, Virology 350:370, 2006) , but their genome is modified in one or more of the following ways, resulting in the above-mentioned characteristics:

- i) One or more, e.g. two, three or four, of the four arenavirus open reading frames (glycoprotein (GP); nucleoprotein (NP); the matrix protein Z; the RNA-dependent RNA polymerase L) are removed or mutated to prevent formation of infectious particles in normal cells albeit still allowing gene expression in arenavirus vector-infected cells.
- ii) Foreign nucleic acids coding for one or more proteins can be introduced. Alternatively or in addition, foreign nucleic acids may be incorporated for modulating host gene expression. These include but are not limited to short hairpin RNAs (shRNA), small interfering RNA (siRNA), micro RNAs (miRNA), and precursors thereof. These foreign nucleic acids are transcribed from one or more, e.g. two or three of the four arenavirus promoters 5' UTR and 3' UTR of the S segment, and 5' UTR and 3' UTR of the L segment, or from additionally introduced promoter sequences that can be read by the viral RNA-dependent RNA polymerase, by cellular RNA polymerase I, RNA polymerase II or RNA polymerase III, such as duplications of viral promoter sequences that are naturally found in the viral UTRs, the 28S ribosomal RNA promoter, the beta-actin promoter or the 5S ribosomal RNA promoter, respectively. The ribonucleic acids coding for proteins or modulating host gene expression are transcribed and translated either by themselves or as read-through by fusion to arenavirus protein open reading frames, and expression of proteins in the host cell may be enhanced by introducing in the viral transcript sequence at the appropriate place(s) one or more, e.g. two, three or four, internal ribosome entry sites.

"Modulating host gene expression" as understood herein refers to reduction of expression of host genes or the enhancement thereof, either in all vector-targeted cells or in a cell type-specific manner. These desirable features can be achieved by adapting the nucleic acid sequence incorporated into vectors.

Arenavirus vectors can be used to improve life and health in general, and to immunize (in a preventive manner) or treat (in an immunotherapeutic manner) animals including men in a variety of contexts including but not limited to

- i) infections including but not limited to viruses such as human immunodeficiency virus (HIV), hepatitis B virus (HBV), hepatitis C virus (HCV), influenza viruses, and respiratory syncytial virus (RSV), bacteria such as mycobacteria, haemophilus spp., and pneumococcus spp., and parasites such as plasmodia, amebiasis, and filaria, and prions such as the infectious agents causing classical and variant Creutzfeldt-Jakob disease and mad cow disease;

- ii) autoimmune diseases including but not limited to type I diabetes, multiple sclerosis, rheumatoid arthritis, lupus erythematosus, and psoriasis;
- iii) neoplastic diseases including but not limited to melanoma, prostate carcinoma, breast carcinoma, lung carcinoma and neuroblastoma;
- 5 iv) metabolic diseases including but not limited to type II diabetes, obesity, and gout;
- v) degenerative diseases including but not limited to Alzheimer's disease, and Parkinson's disease;
- vi) inherited diseases including but not limited to Huntington's disease, severe combined immunodeficiency, and lipid storage diseases;
- 10 vii) substance dependences including but not limited to tobacco and alcohol abuse; and
- viii) allergic diseases including but not limited to seasonal or perennial rhinoconjunctivitis, asthma and eczema.

With the same intention, arenavirus vectors can be used to introduce a gene of interest, e.g. foreign nucleic acids, into cells of living animals including men, i.e. as gene therapy, or they can be used to introduce and express a gene product of interest in biotechnological applications. Abolishing replication of arenavirus vectors by deleting from their genome e.g. the Z gene which is required for particle release, or the GP gene which is required for infection of target cells (compare also Figure 3), the total number of infected cells is limited by the inoculum administered, e.g. to a vaccinee or to a recipient of gene therapy, or accidentally transmitted to personnel involved in medical or biotechnological applications or to animals. Arenavirus disease and immunosuppression in wild type arenavirus infection are both known to result from unchecked viral replication. Therefore, abolishing replication of arenavirus vectors prevents pathogenesis as a result of intentional or accidental transmission of vector particles. In this invention, one important aspect consists in exploiting the above necessity of abolishment of replication in a beneficial way for the purpose of expressing one or more foreign proteins, e.g. antigens of interest: Removal, e.g. structurally by deletion or functionally by mutagenesis, of one or more of the arenavirus genes frees the respective viral promoters for expression of the proteins of choice.

A number of combined advantages characterize the present invention on arenavirus vector strategy: Of note, the retained exquisite immunogenicity of arenavirus vectors – retained despite the inability of arenavirus vectors to spread – comes as a great surprise to immunologists working in the field of arenavirus immunology. A substantial virus and antigen load over a critical period of time is generally considered essential for the

unmatched immunogenic properties of arenaviruses. With regard to safety, the virus' (and the vector's) non-cytolytic behavior is a major advantage over most available vector systems, and the same applies to the lack of oncogenic potential of arenaviruses in general. Also, the inability of arenavirus vectors to replicate is of much importance with regard to safety. Very advantageous, particularly for the application as vaccines, is also the high level of resistance of arenavirus vectors to antibody neutralization. This property is inherent to many arenavirus envelopes and allows repeated immunization with the same arenavirus vector resulting in repeated boosting of the immune response. Similarly, pre-existing immunity against arenaviruses is very low or negligible in the human population.

Arenaviruses considered are Old World viruses, for example Lassa virus, Lymphocytic choriomeningitis virus (LCMV), Mobala virus, Mopeia virus, or Ippy virus, or New World viruses, for example Amapari virus, Flexal virus, Guanarito virus, Junin virus, Latino virus, Machupo virus, Oliveros virus, Paraná virus, Pichinde virus, Pirital virus, Sabiá virus, Tacaribe virus, Tamiami virus, Bear Canyon virus, or Whitewater Arroyo virus. Preferred are members of the Old World viruses, e.g. Lassa virus or LCMV, in particular LCMV.

Foreign nucleic acids coding for one or more proteins of interest are e.g. messenger RNA-derived sequences or RNA corresponding to a primary gene transcript, leading to expression of the protein of interest when arenavirus particles of the invention carrying this RNA infect a cell. Further foreign nucleic acids considered are those modifying gene expression in cells infected with the arenavirus vector particle, e.g. by RNA interference.

Ribonucleic acids of interest considered to be introduced in engineered arenaviruses of the invention are any sequences coding for protein or modulating host gene expression that can be introduced in an arenavirus vector genome by replacement or fusion to the open reading frame of glycoprotein GP, the matrix protein Z, the nucleoprotein NP, or the polymerase protein L, i.e. that can be transcribed and/or expressed under control of the four arenavirus promoters (5' UTR and 3' UTR of the S segment, and 5' UTR and 3' UTR of the L segment), as well as ribonucleic acids that can be inserted with regulatory elements that can be read by the viral RNA-dependent RNA polymerase, cellular RNA polymerase I, RNA polymerase II or RNA polymerase III, such as duplications of viral promoter sequences that are naturally found in the viral UTRs, the 28S ribosomal RNA promoter, the beta-actin promoter or the 5S ribosomal RNA promoter, respectively. The proteins or nucleic acids are transcribed and/or expressed either by themselves or as

read-through by fusion to arenavirus open reading frames and genes, respectively, and/or in combination with one or more, e.g. two, three or four, internal ribosome entry sites. As is demonstrated with genes for GFP and ovalbumin replacing GP, the length of the gene inserted and the properties of the expressed protein are not critical and open the
5 possibility for expression of a large variety of proteins of interest.

Preferred proteins of interest are peptidic or proteinaceous antigens. Peptidic or proteinaceous antigens of the invention may, for example, be selected from the group consisting of (a) proteins or peptides suited to induce or modulate an immune response
10 against infectious diseases; (b) proteins or peptides suited to induce or modulate an immune response against neoplastic diseases, i.e. cancer cells; and (c) proteins or peptides suited to induce or modulate an immune response against allergens. Combinations of antigens, e.g. of antigens derived from one or more infectious organisms or tumors or allergens may be combined to elicit or modulate an immune response
15 protecting or curing more than one infection, type of tumor or allergic disease, respectively.

"Modulating an immune response" as used herein means i) improving, either in quality or quantity, a beneficial immune response of a patient. This is desirable, for example, when
20 enhancing HIV-specific T cell and antibody responses in the context of immunotherapy of an infected individual. The term "modulating an immune response" also refers to ii) the process generally known as desensitization, e.g. desensitization against allergens by suppressing an allergic type of immune response such as the one of the immunoglobulin E isotype, with the attempt of substituting or superimposing a protective immune response
25 or of dampening the pathogenic immune response.

In one specific embodiment of the invention, the antigen is one that is useful for the prevention of infectious disease. Particular examples of antigens or antigenic determinants include the HIV antigens gp 41, gp 120, gag, and pol, non-structural (NS)
30 proteins of Hepatitis C virus, the influenza antigens hemagglutinin and neuraminidase, hepatitis B surface antigen, and circumsporozoite protein of malaria.

Preferably the antigen is selected from respiratory syncytial virus antigens, human immunodeficiency virus antigens, hepatitis C virus antigens, varicella zoster virus
35 antigens, herpes simplex virus antigens, cytomegalovirus antigens and antigens derived from mycobacterium tuberculosis.

The selection of antigens for the composition and method of treatment for cancer would be known to those skilled in the medical art treating such disorders. Representative examples of this type of antigen include the following: HER2/neu (breast cancer), GD2
5 (neuroblastoma), EGF-R (malignant glioblastoma), CEA (medullary thyroid cancer), CD52 (leukemia), MUC1 (expressed in hematological malignancies), gp100 protein, MELAN-A/MART1 or the product of the tumor suppressor gene WT1.

The selection of antigens for the composition and method of treatment for allergy would
10 be known to those skilled in the medical art treating such disorders. Representative examples of this type of antigen include but are not limited to birch pollen antigen Bet v 1 and cat allergen Fel d 1.

The selection of antigens for the composition and method of treatment for obesity would
15 be known to those skilled in the medical art treating such disorders. Representative examples of this type of antigen include but are not limited to ghrelin and gastric inhibitory peptide (GIP).

Design of arenavirus vector genome

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Starting out from a wild type arenavirus genome (Figure 1C), the arenavirus vector genome is designed to retain at least the essential regulatory elements on the 5' and 3' untranslated regions (UTRs) of both segments, and preferentially also the intergenic regions (IGRs). The minimal transacting factors necessary for gene expression in infected
25 cells remain in the vector genome as open reading frames that can be expressed, yet they can be placed differently in the genome and can be placed under control of a different promoter than naturally, or can be expressed from internal ribosome entry sites. At least one of the four viral genes (NP, L, GP, Z) is removed or is functionally inactivated. One or more additional genes or stretches of nucleic acids of interest are inserted into the
30 arenavirus vector genome, placed and oriented to allow their expression in infected cells, either under control of one of the four viral promoters (5' and 3' UTR of the S segment, 5' and 3' UTR of the L segment) or from an internal ribosome entry site or from promoters that can be read by the viral RNA-dependent RNA polymerase, cellular RNA polymerase I, RNA polymerase II or RNA polymerase III. Figure 1C shows one example where the
35 arenavirus GP open reading frame (ORF) is replaced by either an ovalbumin (OVA) or green fluorescent protein (GFP) ORF.

Generation of a complementing cell line

Owing to the "deletion" (referring to either removal or functional inactivation) of one or more of the viral genes in arenavirus vectors (here deletion of the glycoprotein, GP, will be taken as an example), arenavirus vectors must be generated and expanded on cells providing in trans the deleted viral gene(s), e.g. the GP in the present example. Such a complementing cell line, henceforth referred to as C-cells, is generated by transfecting a mammalian cell line such as BHK-21, HEK293, VERO or other (here BHK-21 will be taken as an example) with one or more plasmid(s) for expression of the viral gene(s) of interest (complementation plasmid, referred to as C-plasmid). The C-plasmid(s) (for an example see Figure 2A) express the viral gene(s) deleted in the arenavirus vector to be generated under control of one or more expression cassettes suitable for expression in mammalian cells, e.g. a mammalian polymerase II promoter such as the CMV or EF1alpha promoter with a polyadenylation signal. In addition, the complementation plasmid features a mammalian selection marker, e.g. puromycin resistance, under control of an expression cassette suitable for gene expression in mammalian cells, e.g. polymerase II expression cassette as above, or the viral gene transcript(s) are followed by an internal ribosome entry site, such as the one of encephalomyocarditis virus, followed by the mammalian resistance marker. For production in *E. coli*, the plasmid additionally features a bacterial selection marker, such as an ampicillin resistance cassette.

The cells to be used, e.g. BHK-21, HEK293, MC57G or other, are kept in culture and are transfected with the complementation plasmid(s) using any of the commonly used strategies such as calcium-phosphate-, liposome-based protocols or electroporation. A few days later the suitable selection agent, e.g. puromycin, is added in titrated concentrations. Surviving clones are isolated and subcloned following standard procedures, and high-expressing C-cell clones are identified using Western blot or flow cytometry procedures with antibodies directed against the viral protein(s) of interest. As an alternative to the use of stably transfected C-cells transient transfection of normal cells can complement the missing viral gene(s) in each of the steps where C-cells will be used below.

Plasmids for the recovery of arenavirus vectors

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Plasmids needed are of two types:

i) Two plasmids, referred to as TF-plasmids (for an example see Fig. 2B), for expressing intracellularly in C-cells the minimal transacting factors of the arenavirus the vector is derived from e.g. NP and L proteins of LCMV in the present example.

5 ii) Plasmids, referred to as GS-plasmids (for an example see Fig. 2C), for expressing intracellularly in C-cells the arenavirus vector genome segments, e.g. the segments with designed modifications as described in Figure 1C. TF-plasmids express the NP and L proteins of the respective arenavirus vector under control of an expression cassette suitable for protein expression in mammalian cells, typically e.g. a mammalian polymerase II promoter such as the CMV or EF1alpha promoter, either one of them preferentially in combination with a polyadenylation signal (Fig. 2B). GS-plasmids express the small (S) and the large (L) genome segments of the vector. Typically, polymerase I-driven expression cassettes (Fig. 2C) or T7 bacteriophage RNA polymerase (T7-) driven expression cassettes can be used, the latter preferentially with a 3'-terminal ribozyme for processing of the primary transcript to yield the correct end. In the case of using a T7-
10 based system, expression of T7 in C-cells must be provided by either including in the recovery process an additional expression plasmid, constructed analogously to TF-plasmids, providing T7, or C-cells are constructed to additionally express T7 in a stable manner.

20 *Recovery of the arenavirus vector*

First day: C-cells, typically 80% confluent in M6-well plates, are transfected with a mixture of the two TF-plasmids plus the two GS-plasmids. For this one can exploit any of the commonly used strategies such as calcium-phosphate-, liposome-based protocols or
25 electroporation.

3-5 days later: The culture supernatant (arenavirus vector preparation) is harvested, aliquoted and stored at 4°C, -20°C or -80°C depending on how long the arenavirus vector should be stored prior to use. Then the arenavirus vector preparation's infectious titer is
30 assessed by an immunofocus assay on C-cells.

Titration of arenavirus vector infectivity

For measuring the infectivity of an arenavirus vector preparation C-cells are used for a
35 typical immunofocus assay following commonly used principles in virology as outlined hereinafter:

C-cell monolayers, typically in M24 well plates, 80% confluent, are infected with 10-fold dilutions of the arenavirus vector preparation for 90 min. Subsequently, the cell layer is overlaid with suitable cell culture medium supplemented with 1% methylcellulose. Two
5 to three days later, depending on the permissiveness of the C-cell line used, the culture supernatant is removed, the cell layer is fixed, typically with ethanol/acetone or with formalin 4%, followed by permeabilization of the cell layer using mild detergents. Subsequently, arenavirus-vector-infected cell foci are identified using mono- or polyclonal antibody preparation(s) against one of the proteins in the arenavirus vector to be tested or
10 against the antigen introduced. Bound antibody is detected using appropriate reagents, such as anti-isotype or anti-species antibodies that are conjugated to a system for visualization such as horse radish peroxidase, followed by a color reaction with suitable chromogens such as o-phenylenediamine. The resulting spots on the plate are counted to calculate the number of infectious focus forming units (FFU) per volume of arenavirus
15 vector preparation.

Vaccines and pharmaceutical preparations

The invention furthermore relates to vaccines and pharmaceutical preparations
20 comprising the genetically engineered arenaviruses as described hereinbefore. Vaccines and pharmaceutical preparations for other uses are prepared according to standard procedures in the art.

Compositions for enteral administration, such as nasal, buccal, rectal or oral
25 administration, and for parenteral administration, such as intravenous, intramuscular, intradermal or subcutaneous administration, to warm-blooded animals, especially humans, are preferred. Particularly preferred are compositions for parenteral administration. The compositions comprise the genetically engineered arenaviruses alone or, preferably, together with a pharmaceutically acceptable carrier. The dosage of the
30 active ingredient depends upon the type of vaccination and the disease to be treated and upon the species, its age, weight, and individual condition, the individual pharmacokinetic data, and the mode of administration.

The pharmaceutical compositions comprise from about 10^3 to about 10^{11} focus forming
35 units of the genetically engineered arenaviruses. Unit dose forms for parenteral administration are, for example, ampoules or vials, e.g. vials containing from about 10^3 to

10^{10} focus forming units or 10^5 to 10^{15} physical particles of genetically engineered arenaviruses.

5 Preference is given to the use of suspensions or dispersions of genetically engineered arenaviruses, especially isotonic aqueous dispersions or suspensions. The pharmaceutical compositions may be sterilized and/or may comprise excipients, e.g. preservatives, stabilizers, wetting agents and/or emulsifiers, solubilizers, salts for regulating osmotic pressure and/or buffers and are prepared in a manner known per se, for example by means of conventional dispersing and suspending processes. The said
10 dispersions or suspensions may comprise viscosity-regulating agents. The suspensions or dispersions are kept at temperatures around 2-4°C, or preferentially for longer storage may be frozen and then thawed shortly before use.

The invention relates also to processes and to the use of genetically engineered
15 arenaviruses for the manufacture of vaccines in the form of pharmaceutical preparations, which comprise genetically engineered arenaviruses as active ingredient. The pharmaceutical compositions of the present invention are prepared in a manner known per se, for example by means of conventional mixing and/or dispersing processes.

20 *Administration to vaccinee and to gene therapy recipient*

The invention furthermore relates to methods of vaccination and gene therapy using the genetically engineered arenaviruses as described hereinbefore.

25 Arenavirus vectors are administered for improving the quality of live, including but not limited to vaccination, immunotherapy and gene therapy in order to prevent, treat or improve

i) infections including but not limited to those caused by viruses such as human immunodeficiency virus (HIV), hepatitis B virus (HBV), hepatitis C virus (HCV), influenza
30 viruses, and respiratory syncytial virus (RSV), bacteria such as mycobacteria, haemophilus spp., and pneumococcus spp., parasites such as plasmodia, amebia, and filaria, and prions such as the infectious agents causing classical and variant Creutzfeldt-Jakob disease and mad cow disease;

ii) autoimmune diseases including but not limited to type I diabetes, multiple sclerosis,
35 rheumatoid arthritis, lupus erythematosus, and psoriasis;

- iii) neoplastic diseases including but not limited to melanoma, prostate carcinoma, breast carcinoma, lung carcinoma, and neuroblastoma;
- iv) metabolic diseases including but not limited to type II diabetes, obesity, and gout;
- v) degenerative diseases including but not limited to Alzheimer's disease and Parkinson's disease;
- vi) inherited diseases including but not limited to Huntington's disease, severe combined immunodeficiency, and lipid storage diseases; and
- vii) substance dependence including but not limited to tobacco and alcohol abuse.

10 In particular the invention relates to a method of preventing infections by viruses, bacteria, parasites and prions comprising administering a vaccine comprising genetically engineered arenaviruses to a patient in need thereof, and likewise to a method of preventing neoplastic diseases and degenerative diseases as listed hereinbefore.

15 Furthermore the invention relates to a method of treating infections by viruses, bacteria, parasites and prions, autoimmune diseases, neoplastic diseases, metabolic diseases, degenerative diseases, inherited diseases, or substance dependence comprising administering a pharmaceutical preparation comprising genetically engineered arenaviruses to a patient in need thereof.

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Arenavirus vectors are administered to a vaccinee either by one or by multiple ones of the available routes including but not limited to intramuscular, intradermal, subcutaneous, peroral, intranasal, or intravenous routes, e.g. as in the experiment outlined in Figure 3A. This results in infection of cells, amplification of the viral genome segments in these very same initially infected cells, e.g. after intravenous inoculation. This comprises dendritic cells of the spleen that can trigger T cell responses. Owing to the inability of the arenavirus vector to replicate in cells of a vaccinee, lacking the complementing viral protein present in C-cells, the levels of arenavirus vector RNA decline rapidly over time and the viral genome approaches its extinction within days after inoculation of the arenavirus vector (Figure 3A). Owing to the lack of arenavirus vector replication and persistence, and in contrast to infection with the same dose of wild type virus, arenavirus vector immunization does not cause immunosuppression (Figure 3B) or disease (Figure 5). This is tested in mice infected either with wild type LCMV or with an LCMV-based vector expressing OVA instead of LCMV-GP (rLCMV/OVA; compare Fig. 1C).

35 Subsequent infection with vesicular stomatitis virus elicited a normal CD8 T cell and antibody response in animals previously immunized with rLCMV/OVA, but was

suppressed in animals previously infected with wild type LCMV. Similarly, wild type LCMV elicited lethal choriomeningitis in mice when administered intracranially, but rLCMV/OVA did not elicit any clinically detectable signs of illness (Figure 5).

5 Despite its transient nature, expression of the antigen of interest does, however, evoke a strong and long-lasting T cell response (Figure 4A) and evokes high titers of specific antibodies (Figure 4B). This response is dose-dependent but even small doses are efficient (Figure 4B). The protective capacity of T cell responses elicited by LCMV-based vaccine vectors is tested in mice. Immunization with rLCMV/OVA protected against
10 infectious challenge with recombinant *Listeria monocytogenes* expressing OVA (rLM/OVA). This was evident in strongly reduced or undetectable rLM/OVA titers in the spleen of vaccinated animals (Figure 6A). Induction of antibody-mediated protection by LCMV vector is tested in mice lacking the type I interferon receptor. These mice are highly susceptible to vesicular stomatitis virus (VSV), with a 50% lethal dose (LD₅₀) in the range
15 of 50 PFU. For immunization against VSV, a LCMV vector was used that expresses an antigenic but non-functional variant of the vesicular stomatitis virus envelope protein G (modified by insertion of a foreign peptide sequence in its ectodomain). Immunized mice survived a challenge infection with 2x10⁶ PFU VSV (i.e. >10'000-fold LD₅₀), whereas unimmunized control mice developed terminal myeloencephalitis within two to three days
20 after VSV challenge (Figure 6). Of note, though, inactivation of the arenavirus vector genome by UV irradiation abrogated its immunogenicity, demonstrating that replication and gene expression of the viral vector in infected cells is essential for its efficacy as a vaccine. Additionally, T cell and antibody responses can be enhanced by repeated applications of the same (homologous) or different (heterologous) arenavirus vectors, i.e.
25 in the form of booster immunization. In homologous prime-boost regimens, the virtual absence of neutralizing antibody induction renders booster immunizations particularly efficient.

When used for gene therapy, arenavirus vectors can be applied systemically, e.g.
30 intravenously, or topically, e.g. by stereotactic injection using appropriate equipment, for targeting and delivery to specific tissues where the antigen of interest should be expressed. Owing to its non-cytolytic nature, the arenavirus vector does not harm the cell it infects and can functionally substitute for a gene of interest.

35 As an alternative way of exploiting arenavirus vectors for treatment of multicellular organisms, complementing (C-cells) or non-complementing (normal) cells are implanted

into a recipient's body encapsulated by biocompatible materials preventing immune rejection by the recipient yet allowing for the constant release of infectious (implantation of infectious C-cells) or non-infectious (implantation of infected normal cells) particles or of proteins and/or ribonucleic acids from the encapsulated cells across the capsule into the recipient's tissues.

Expression of a protein of interest in a cell culture

The invention furthermore relates to expression of a protein of interest in a cell culture wherein the cell culture is infected with genetically engineered arenaviruses. When used for expression of a protein or stretch of nucleic acids of interest, e.g. an antigen of interest, in cultured cells, the following two procedures are envisaged:

- i) The cell type of interest is infected with the arenavirus vector preparation at a multiplicity of infection (MOI) of one or more, e.g. two, three or four, resulting in production of the protein of interest in all cells already shortly after infection.
- ii) Alternatively, a lower MOI can be used and individual cell clones can be selected for their level of virally driven protein expression. Subsequently individual clones can be expanded infinitely owing to the non-cytolytic nature of arenavirus vectors. Irrespective of the approach, the protein(s) of interest can subsequently be collected (and purified) either from the culture supernatant or from the cells themselves, depending on the properties of the protein(s) produced.

However, the invention is not limited to these two strategies, and other ways of driving expression of proteins or nucleic acids of interest using genetically engineered arenaviruses as vectors may be considered.

Claims

1. An arenavirus particle comprising a genome wherein at least one of the four arenavirus open reading frames of the genome coding for a glycoprotein (GP), a nucleoprotein (NP), a matrix protein Z or a RNA-dependent RNA polymerase L is removed or functionally inactivated, and wherein the arenavirus particle is infectious, but unable to produce further infectious progeny particles in non-complementing cells.
2. The arenavirus particle according to claim 1, wherein the genome comprises a foreign ribonucleic acid sequence coding for one or more protein or peptide of interest.
3. The arenavirus particle according to claim 1, wherein the genome comprises a foreign ribonucleic acid sequence for increasing or decreasing host gene expression.
4. The arenavirus particle according to claim 2 or 3, wherein the foreign ribonucleic acid sequence is under the control of:
 - i) one or more of the arenavirus promoters within a 5' UTR or a 3' UTR of an arenavirus S segment or L segment, and/or
 - ii) a regulatory element that can be read by the viral RNA-dependent RNA polymerase, cellular RNA polymerase I, RNA polymerase II or RNA polymerase III, andwherein the foreign ribonucleic acid sequence is expressed either by itself or as read-through by fusion to an arenavirus protein open reading frame.
5. The arenavirus particle according to any one of claims 1 to 4, wherein the arenavirus open reading frame coding for the glycoprotein (GP) is removed or functionally inactivated.
6. The arenavirus particle according to claim 5 wherein the arenavirus open reading frame coding for the glycoprotein (GP) is removed and replaced by a foreign ribonucleic acid sequence

coding for one or more protein or peptide of interest or a foreign ribonucleic acid sequence for increasing or decreasing host gene expression.

7. The arenavirus particle according to claim 5 wherein the arenavirus open reading frame coding for the glycoprotein (GP) is removed and replaced by a foreign ribonucleic acid sequence coding for an antigen of an infectious organism, tumor or allergen.

8. The arenavirus particle according to claim 5 wherein the arenavirus open reading frame coding for the glycoprotein (GP) is removed and replaced by a foreign ribonucleic acid sequence selected from the group consisting of a short hairpin RNA (shRNA), a small interfering RNA (siRNA) and a micro RNA (miRNA).

9. The arenavirus particle according to any one of claims 1 to 8 wherein the arenavirus is lymphocytic choriomeningitis virus (LCMV).

10. The arenavirus particle according to claim 2 or 6, wherein the one or more protein or peptide of interest is an antigen selected from the group consisting of a respiratory syncytial virus antigen, a human immunodeficiency virus antigen, a hepatitis B virus antigen, a hepatitis B surface antigen, a hepatitis C virus antigen, a varizella zoster virus antigen, a herpes simplex virus antigen, a cytomegalovirus antigen, a mycobacterium tuberculosis antigen, and an antigen suitable to induce an immune response against a neoplastic disease or a tumor.

11. The arenavirus particle according to claim 10, wherein the neoplastic disease is selected from the group consisting of melanoma, prostate carcinoma, breast carcinoma, lung carcinoma, and neuroblastoma.

12. The arenavirus particle according to claim 10, wherein the antigen suitable to induce an immune response against a neoplastic disease or a tumor is selected from the group consisting of HER2/neu, GD2, EGF-R, CEA, CD52, MUC1, gp100 protein, MELAN-A/MART1, and the product of the tumor suppressor gene WT1.

13. The arenavirus particle of any one of claims 1 to 12, wherein the open reading frames coding for the nucleoprotein (NP) and the RNA-dependent RNA polymerase L of the arenavirus are intact.

14. An arenavirus particle comprising a genome, wherein at least one of the two arenavirus open reading frames of the genome coding for a glycoprotein (GP) or a matrix protein Z is removed or functionally inactivated, wherein the open reading frames of the genome coding for a nucleoprotein (NP) and a RNA-dependent RNA polymerase L are intact, and wherein the arenavirus particle is infectious, but unable to produce further infectious progeny particles in non-complementing cells.

15. The arenavirus particle of claim 14, wherein the arenavirus open reading frame coding for the glycoprotein (GP) is removed or functionally inactivated, and wherein the open reading frame coding for the matrix protein Z is intact.

16. A pharmaceutical preparation comprising an arenavirus particle according to any one of claims 2 to 15.

17. Use of the arenavirus particle of any one of claims 2 to 15 for the prevention or treatment of infections caused by viruses, bacteria, parasites or prions, or the prevention or treatment of autoimmune diseases, neoplastic diseases, metabolic diseases, degenerative diseases inherited diseases, allergic diseases, or substance dependence in a patient, wherein the foreign ribonucleic acid sequence coding for one or more protein or peptide of interest is a foreign ribonucleic acid sequence coding for one or more therapeutic protein or peptide of interest or the foreign ribonucleic acid sequence for increasing or decreasing host gene expression is therapeutic.

18. Use of the arenavirus particle of any one of claims 2 to 15 in the manufacture of a medicament for the prevention or treatment of infections caused by viruses, bacteria, parasites or prions, or the prevention or treatment of autoimmune diseases, neoplastic diseases, metabolic diseases, degenerative diseases, inherited diseases, allergic diseases, or substance dependence in

a patient, wherein the foreign ribonucleic acid sequence coding for one or more protein or peptide of interest is a foreign ribonucleic acid sequence coding for one or more therapeutic protein or peptide of interest or the foreign ribonucleic acid sequence for increasing or decreasing host gene expression is therapeutic.

19. A method of expressing a protein or peptide of interest comprising infecting cells in a cell culture with the arenavirus particle according to any one of claims 2 and 4 to 13.

20. An arenavirus genome wherein at least one of the four arenavirus open reading frames of the genome coding for a glycoprotein (GP), a nucleoprotein (NP), a protein Z, or a RNA dependent RNA polymerase L is removed or functionally inactivated, and wherein an arenavirus particle comprising the genome is infectious, but unable to produce further infectious progeny particles in non-complementing cells.

21. The arenavirus genome of claim 20, further comprising a foreign ribonucleic acid sequence coding for an antigen selected from the group consisting of a respiratory syncytial virus antigen, a human immunodeficiency virus antigen, a hepatitis B virus antigen, a hepatitis B surface antigen, a hepatitis C virus antigen, a varicella zoster virus antigen, a herpes simplex virus antigen, a cytomegalovirus antigen, a mycobacterium tuberculosis antigen, and an antigen suitable to induce an immune response against a neoplastic disease or a tumor.

22. The arenavirus genome of claim 21, wherein the neoplastic disease is selected from the group consisting of melanoma, prostate carcinoma, breast carcinoma, lung carcinoma, and neuroblastoma.

23. The arenavirus genome of claim 21, wherein the antigen suitable to induce an immune response against a neoplastic disease or a tumor is selected from the group consisting of HER2/neu, GD2, EGF-R, CEA, CD52, MUC1, gp100 protein, MELAN-A/MART1, and the product of the tumor suppressor gene WT1.

24. The arenavirus genome of claim 20, further comprising a foreign ribonucleic acid sequence coding for HIV gp41, gp120, gag, or pol, a hepatitis C virus non-structural (NS) protein, influenza virus hemagglutinin or neuroaminidase, or a malarial circumsporozoite protein.

25. The arenavirus genome of any one of claims 20-24, wherein the arenavirus open reading frame coding for the glycoprotein (GP) is removed and replaced by a foreign ribonucleic acid sequence coding for one or more protein or peptide of interest or a foreign ribonucleic acid sequence for increasing or decreasing host gene expression.

26. The arenavirus genome of any one of claims 20-24, wherein the arenavirus open reading frame coding for the glycoprotein (GP) is removed and replaced by a foreign ribonucleic acid sequence coding for an antigen of an infectious organisms, tumor or allergens.

27. The arenavirus genome of any one of claims 20-24, wherein the arenavirus open reading frame coding for the glycoprotein (GP) is removed and replaced by a foreign ribonucleic acid sequence selected from the group consisting of a short hairpin RNA (shRNA), a small interfering RNA (siRNA) and a micro RNA (miRNA).

28. The arenavirus genome of any one of claims 20-27, wherein the arenavirus genome is a lymphocytic choriomeningitis virus (LCMV) genome.

29. The arenavirus genome of claim 20, wherein the genome comprises a foreign ribonucleic acid sequence for increasing or decreasing host gene expression.

30. The arenavirus genome of any one of claims 21-29, wherein the foreign ribonucleic acid sequence is under the control of:

- i) one or more of the arenavirus promoters within a 5' UTR or a 3' UTR of an arenavirus S segment or L segment, and/or
- ii) a regulatory elements that can be read by the viral RNA dependent RNA polymerase, cellular RNA polymerase I, RNA polymerase II or RNA polymerase III, and

wherein the foreign ribonucleic acid sequence is expressed either by itself or as read-through by fusion to an arenavirus protein open reading frame.

31. A vector comprising the arenavirus genome of any one of claims 20-30.

32. A complementing cell for the production of infectious arenavirus particles, wherein the complementing cell

(i) comprises the arenavirus genome of any one of claims 20-30, and

(ii) expresses the open reading frame that has been removed or functionally inactivated from the arenavirus genome of any one of claims 20-30.

33. A method for producing infectious arenavirus particles comprising:

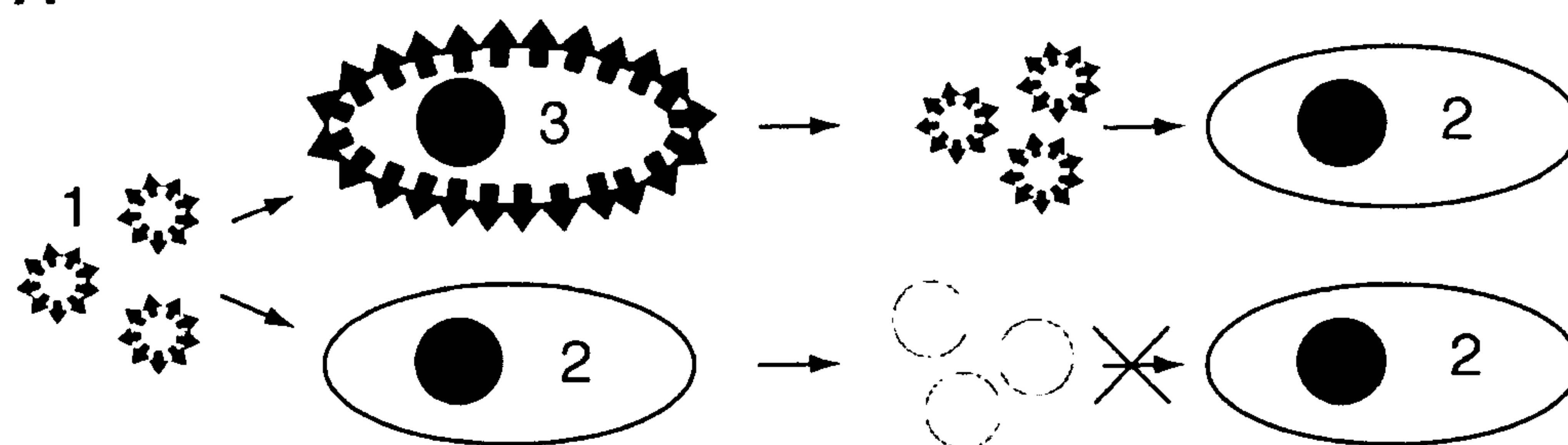
(a) culturing the complementing cell of claim 32; and

(b) harvesting the culture supernatant comprising arenavirus particles.

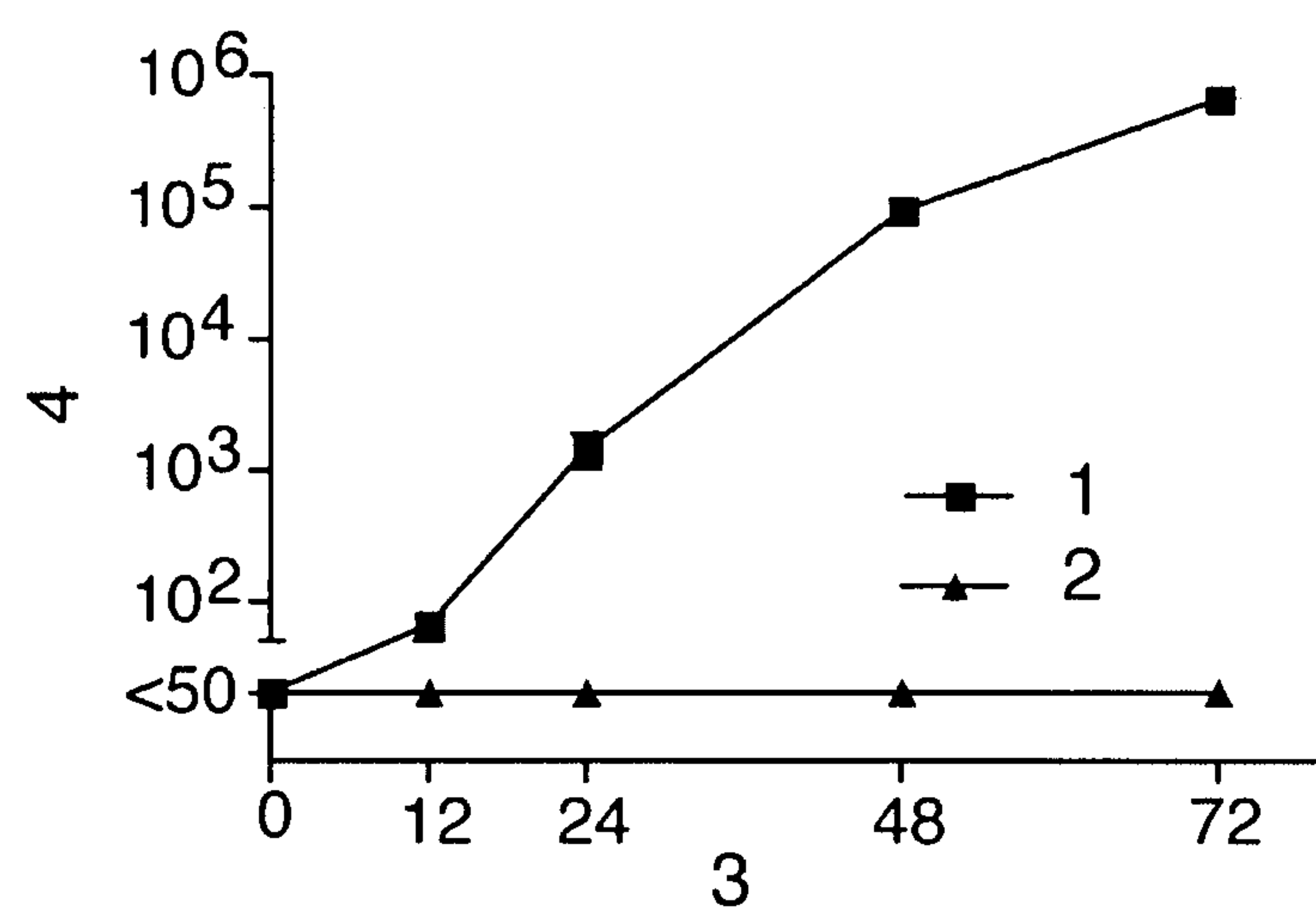
34. A vaccine comprising an arenavirus particle according to any one of claims 2 and 4 to 12.

Fig. 1

A



B



C

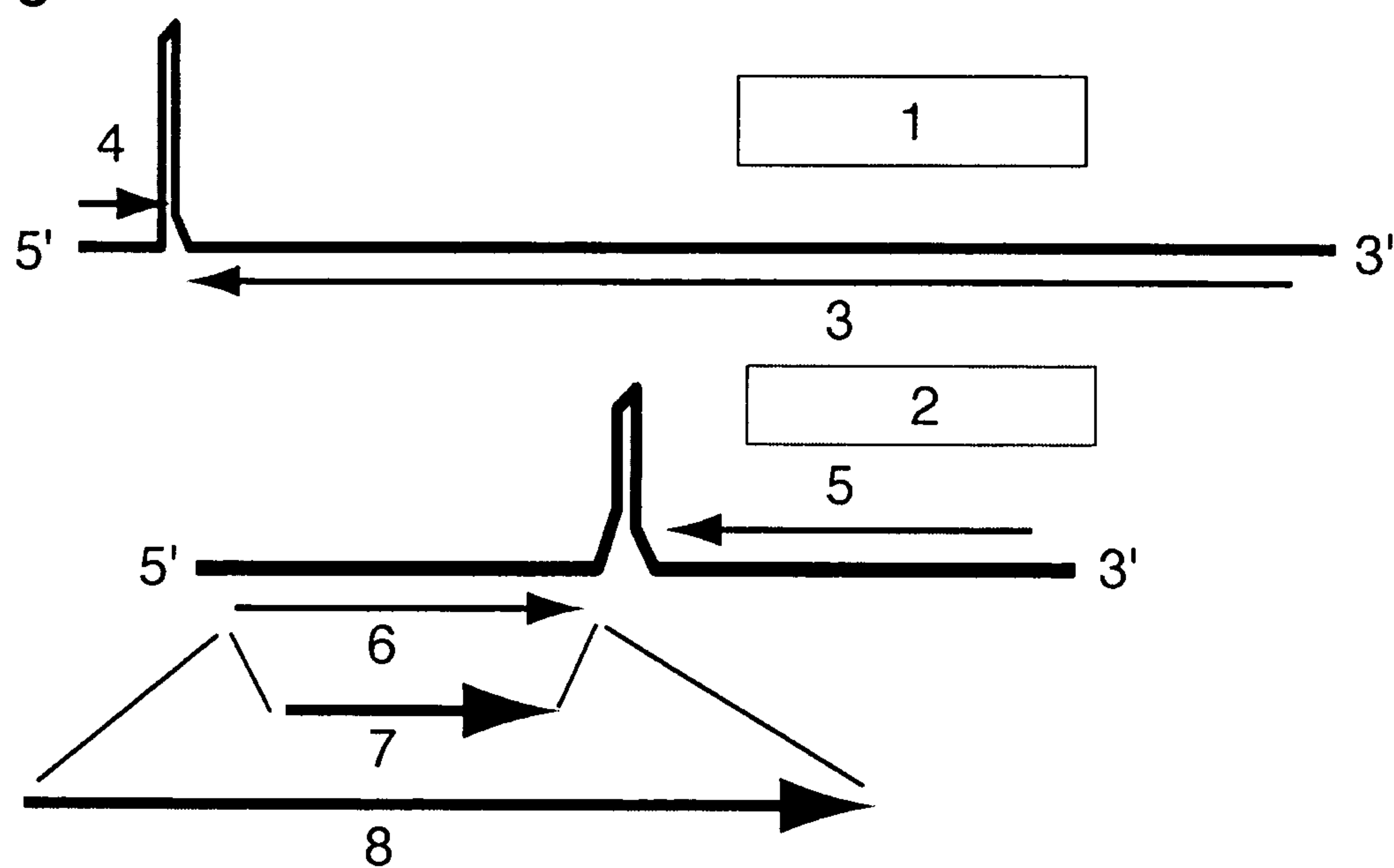


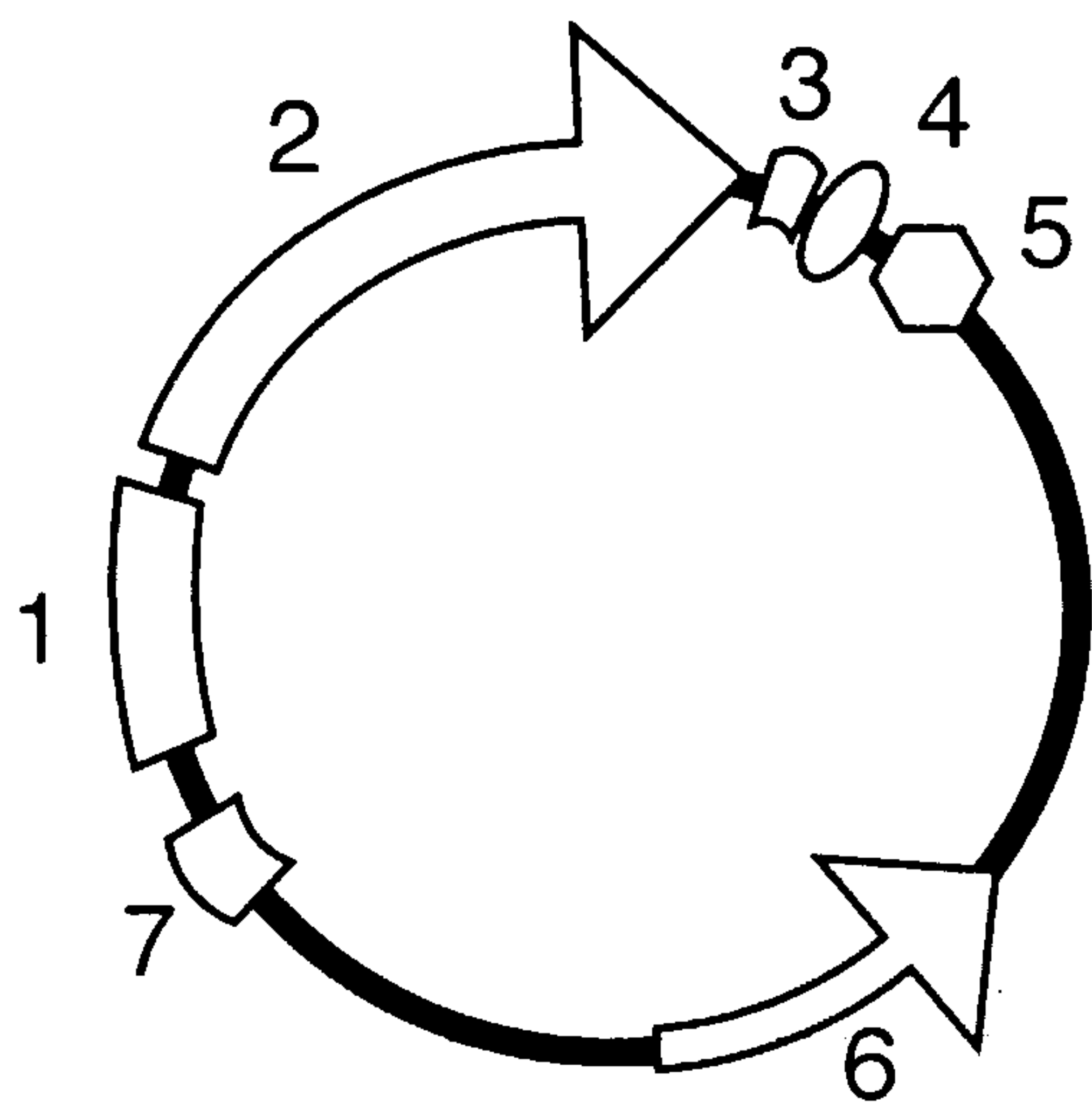
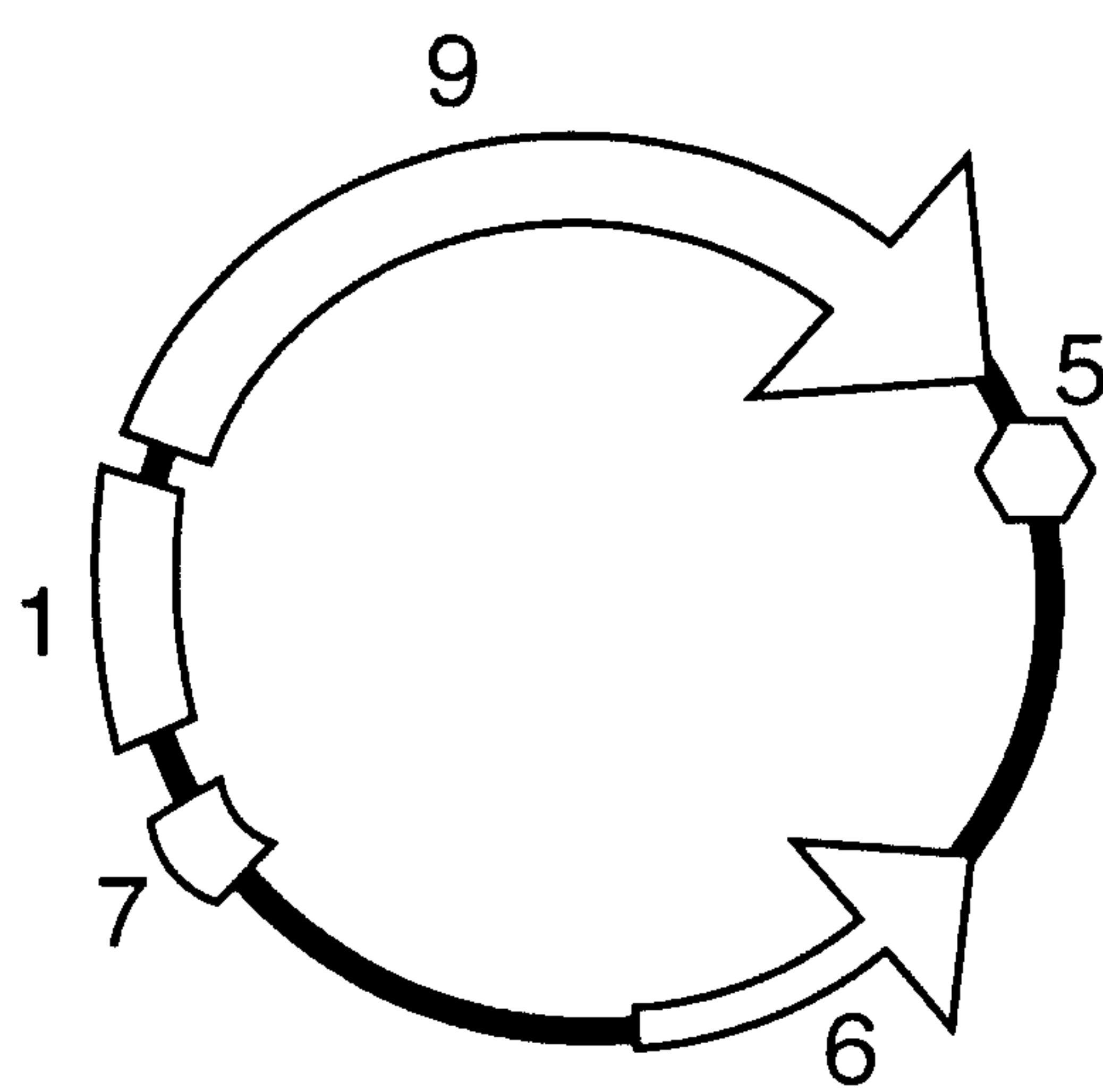
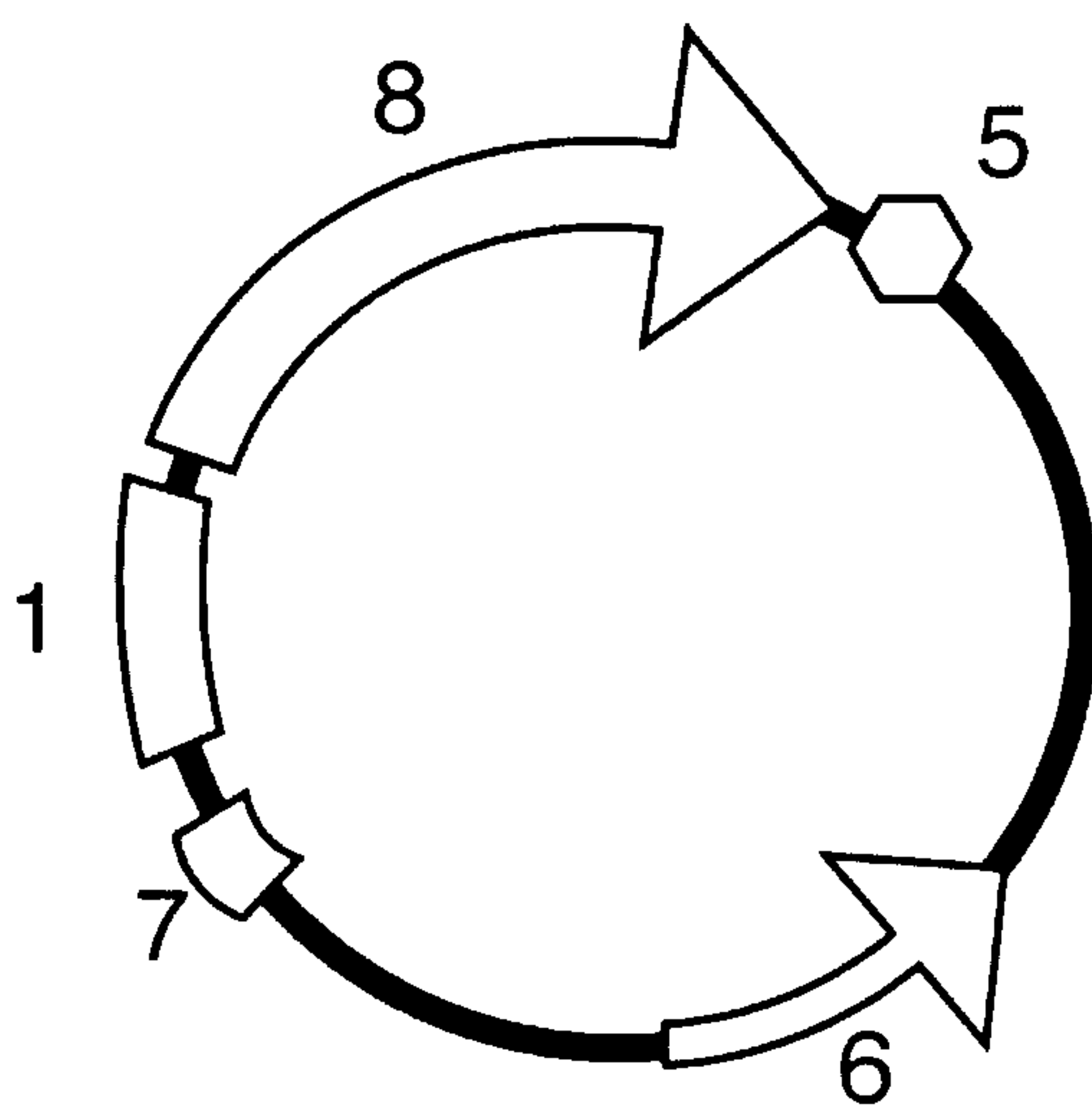
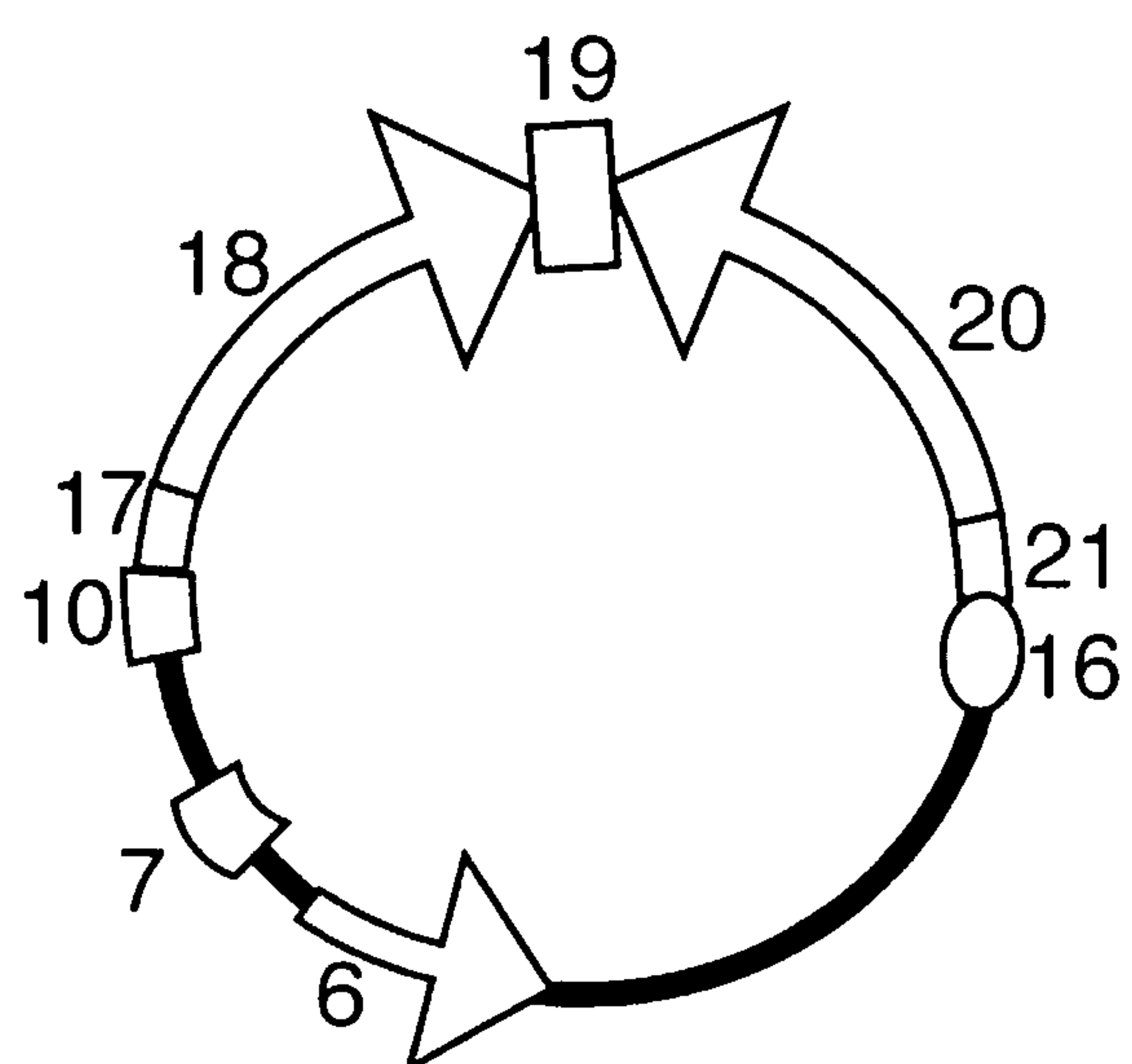
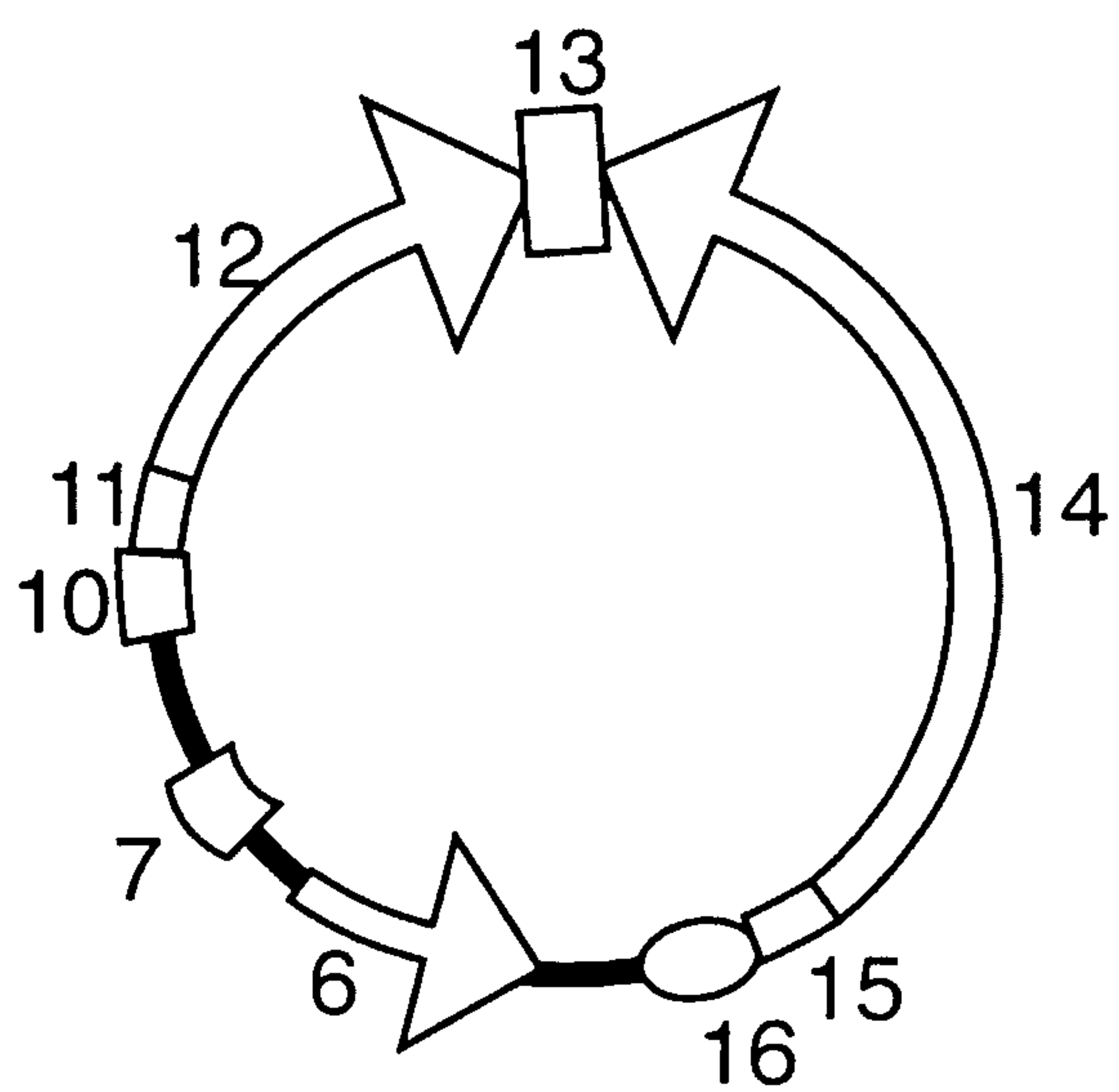
Fig. 2**A****B****C**

Figure 1 consists of a schematic and a graph. The schematic shows a mouse receiving an injection (1), followed by a 2-day period (2), and then a 3-day period (3) before being sacrificed. The graph shows the number of plaques (Y-axis, 0 to 8) versus the number of days post-injection (X-axis, 2, 7, 21, 63). The number of plaques decreases significantly over time, with a horizontal line indicating the limit of detection at <2.

Days post-injection	Number of plaques (individual data points)	Mean ± SD
2	7.5, 7.5, 7.5, 7.5	7.5 ± 0
7	4.5, 4.5, 3.5, 4.5	4.1 ± 0.5
21	2.5, 2.5, 1.5	2.2 ± 0.2
63	<2, <2, <2, <2, <2, <2, <2	<2

The diagram illustrates the experimental protocol in four steps:

1. A mouse is injected with a syringe.
2. A mouse is injected with a syringe.
3. A mouse is shown with a bowl of food.
4. A mouse is shown with a bowl of food.

The diagram also includes a grid with the number 3 and a chemical structure of a branched alkene.

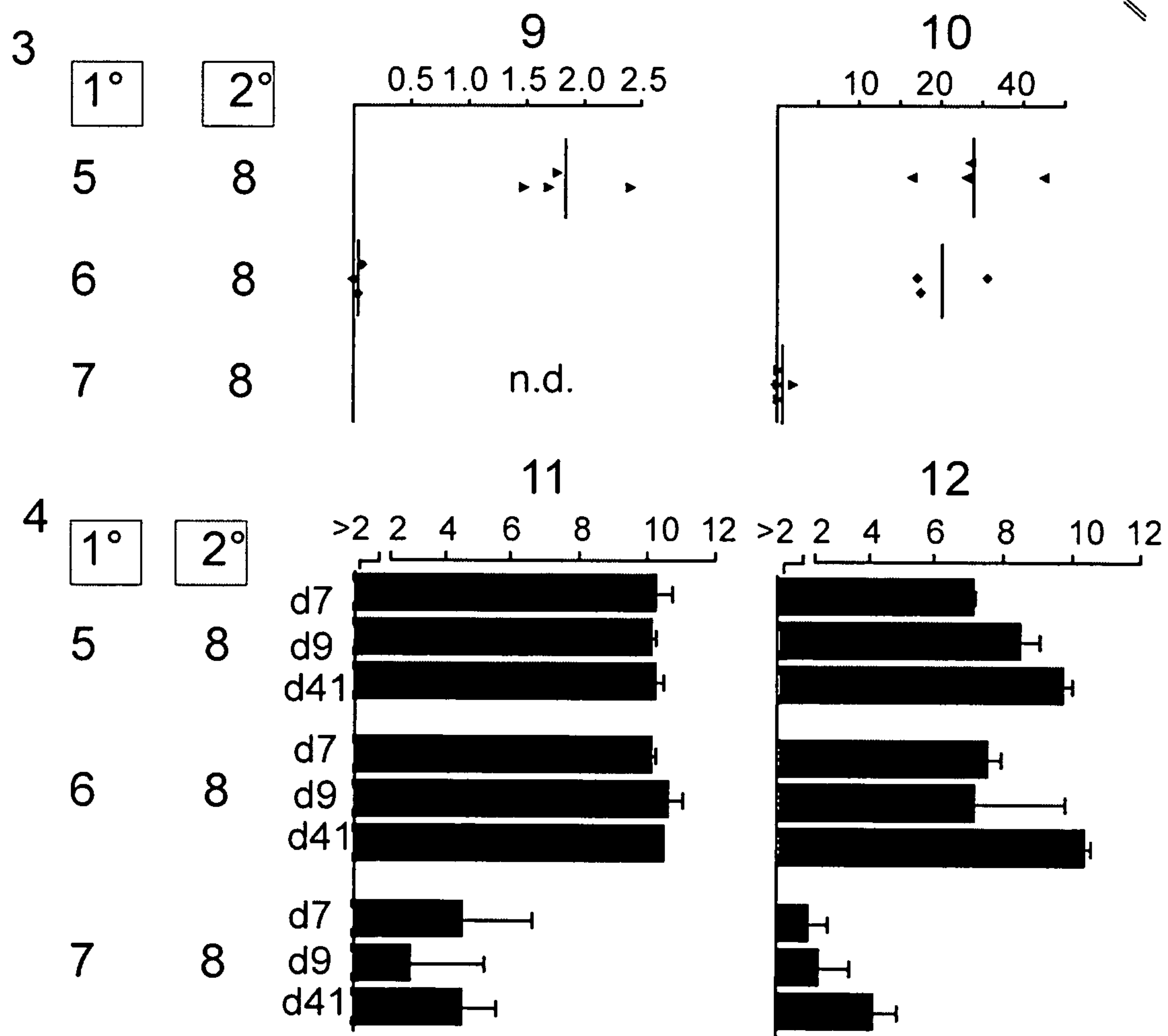
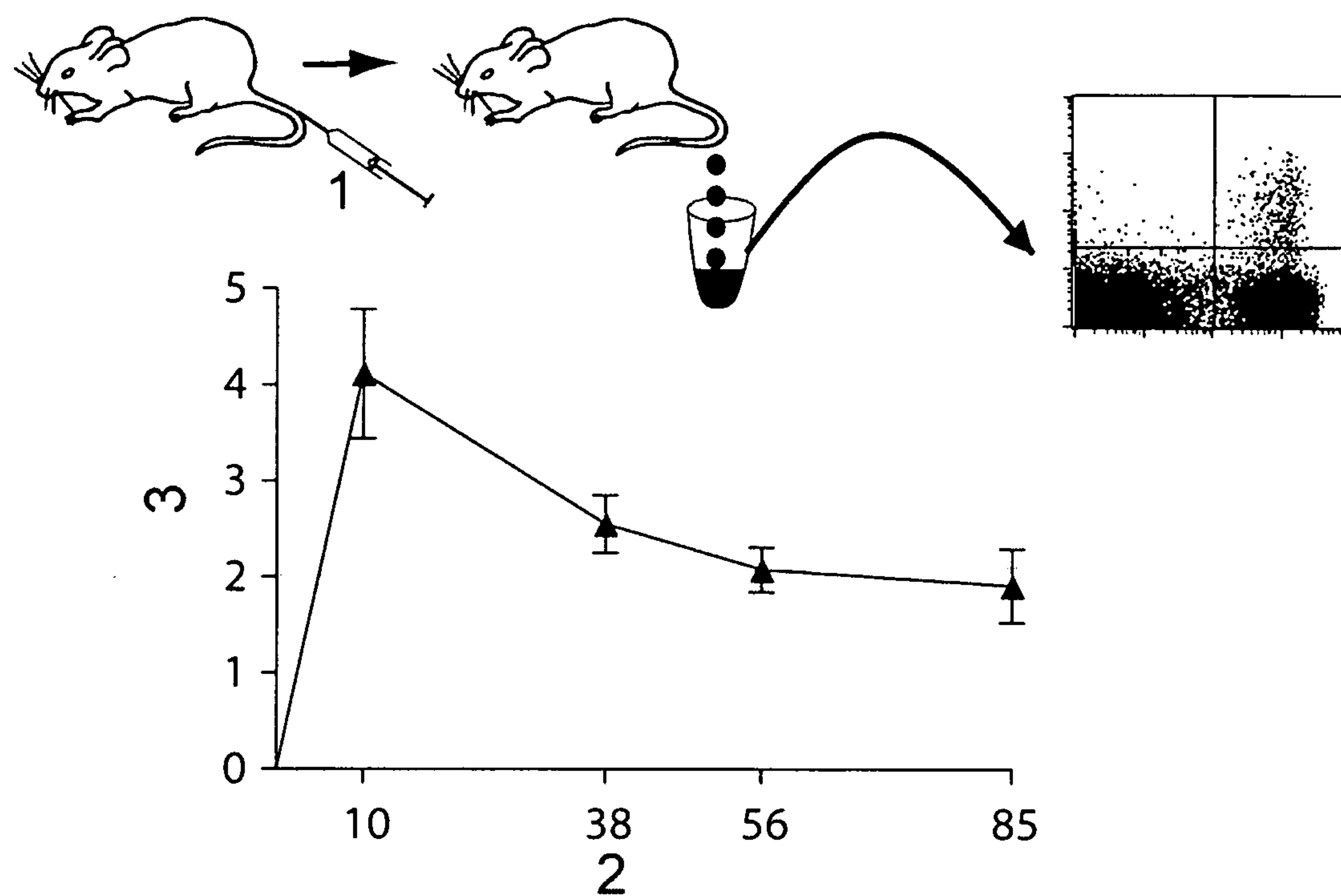


Fig. 4

A



B

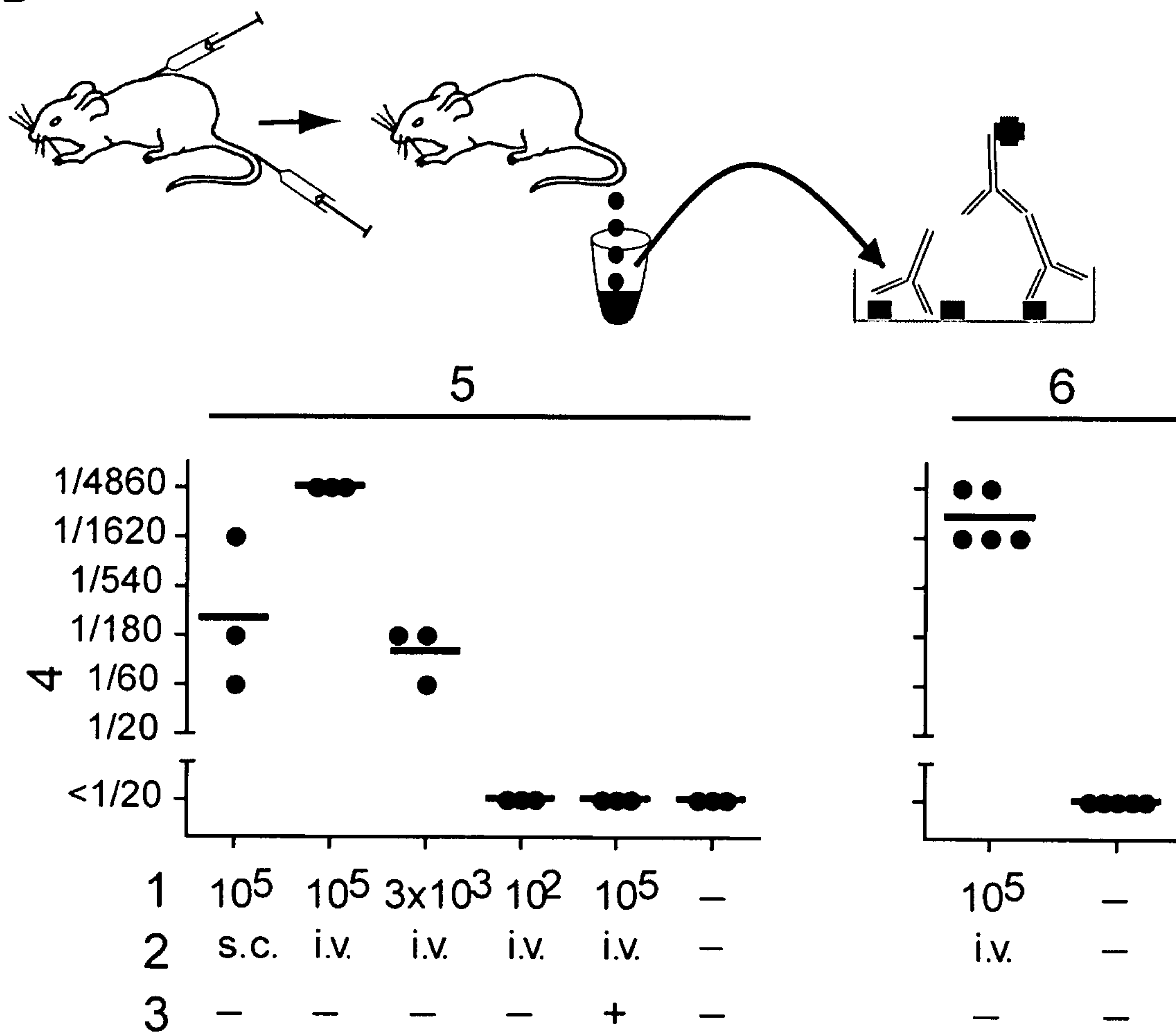


Fig. 5

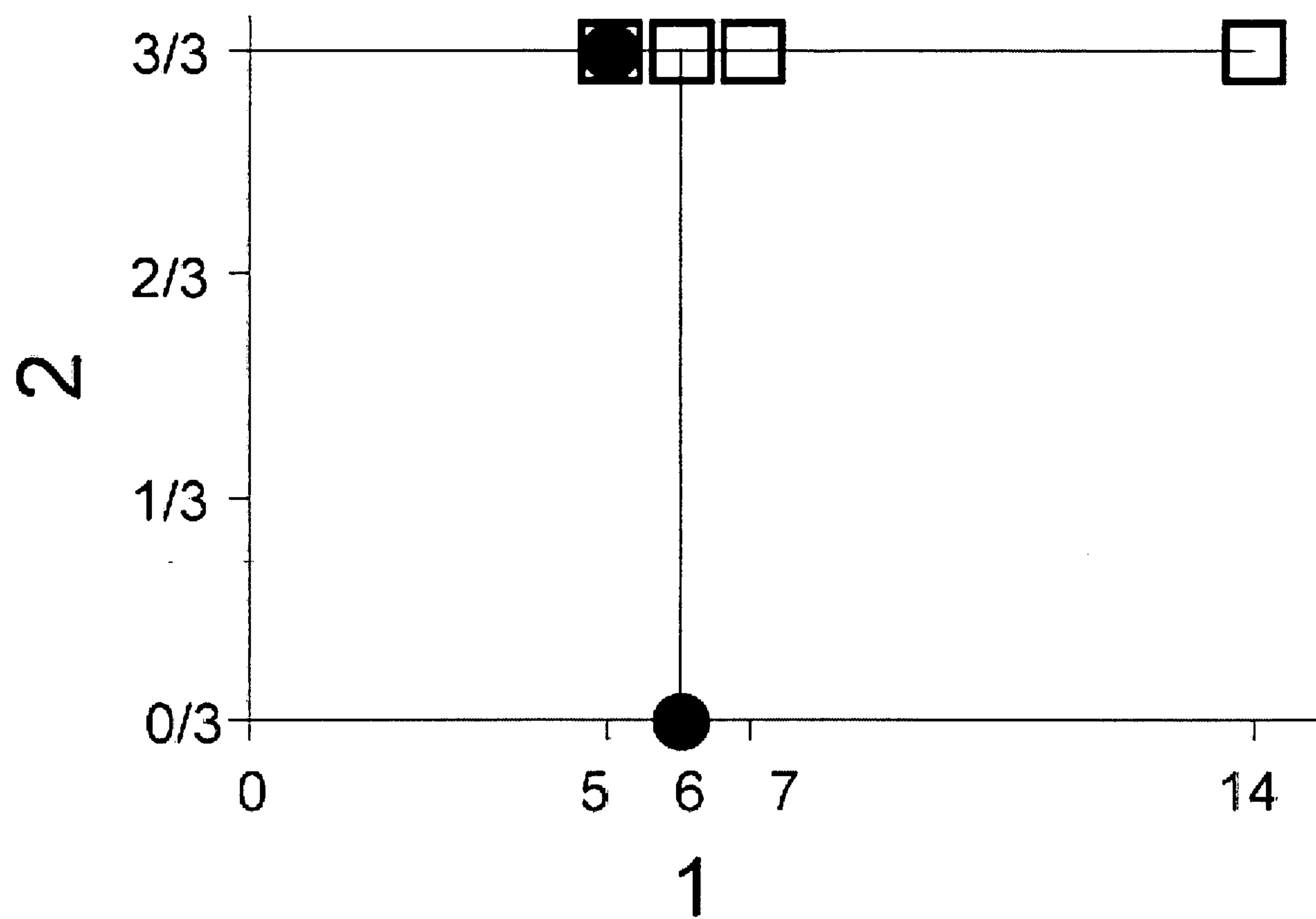


Fig. 6

