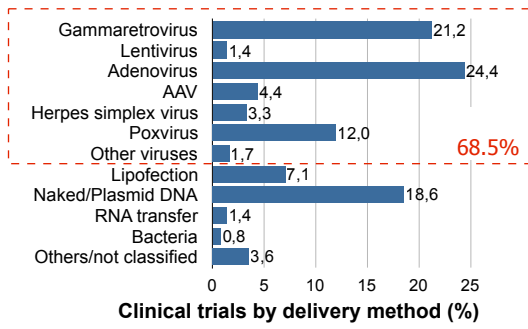
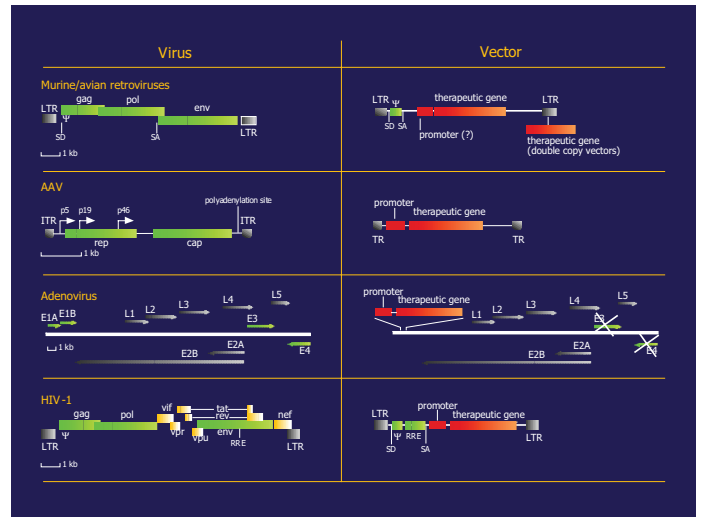


Gene therapy clinical trials by delivery method

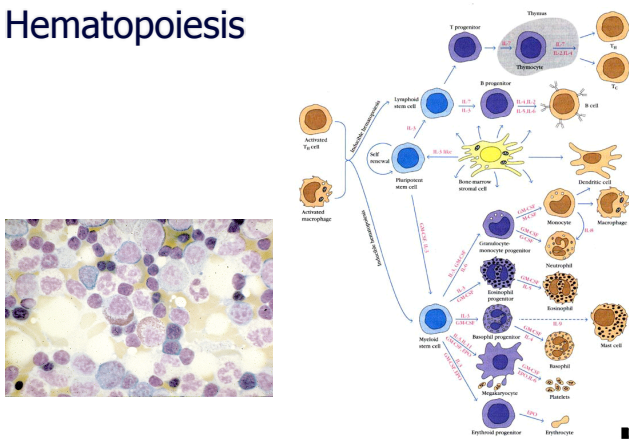


Clinical trials by delivery method (%)

Giacca, M. 2010. Gene Therapy. Springer

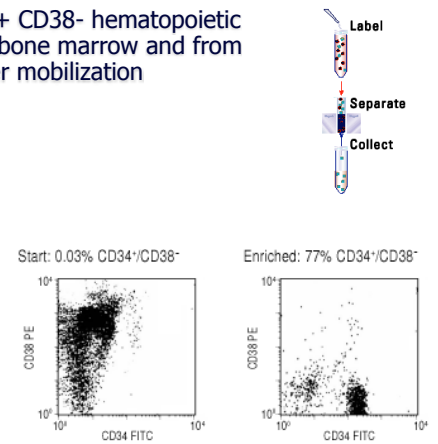
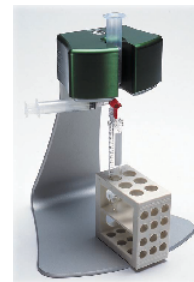


Hematopoiesis



The whole hematopoietic system can be reconstituted by a single HSC

Enrichment of CD34+ CD38- hematopoietic precursors from the bone marrow and from peripheral blood after mobilization

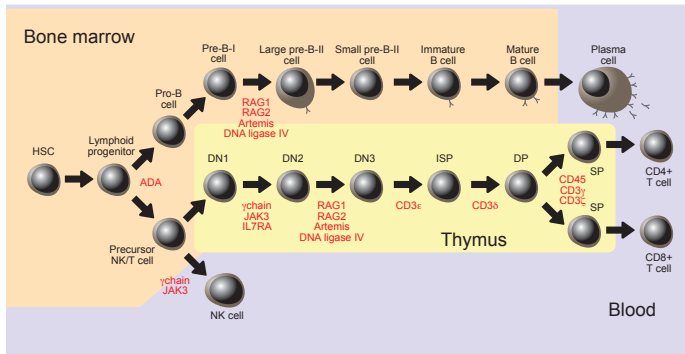


Principali difetti molecolare che portano allo sviluppo di SCID. AR: autosomica recessiva; X-L: legata al cromosoma X

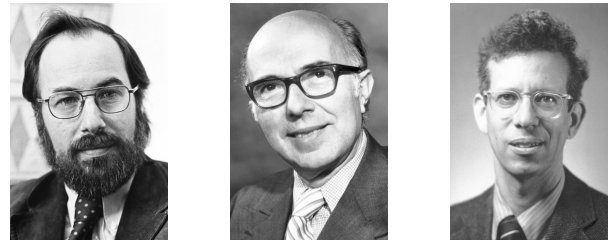
| Meccanismo | Gene mutato | Ereditarietà | Cellule affette |
|--|-------------------------------------|--------------|-----------------|
| Morte prematura delle cellule | ADA | AR | T, B, NK |
| Difetto nella sopravvivenza dovuto alla mancanza di segnali attivatori da parte di citochine | catena comune γ (cy) | X-L | T, NK |
| | JAK-3 | AR | T, NK |
| Difetto nel riarrangiamento V(D)J | IL7RA | AR | T |
| | RAG1 o RAG2 | AR | T, B |
| Difetto nella segnalazione da parte del pre-TCR o del TCR | Artemis | AR | T, B |
| | CD3 δ , ζ , ϵ | AR | T |
| | CD45 | AR | T |

- La convenzione classica della terapia genica non corregge ma aggiunge una copia sana del gene mutato
- Principali successi ad oggi ottenuti per le malattie AR
- Nuove prospettive con gene editing

Defects leading to the development of severe combined immunodeficiency (SCID)

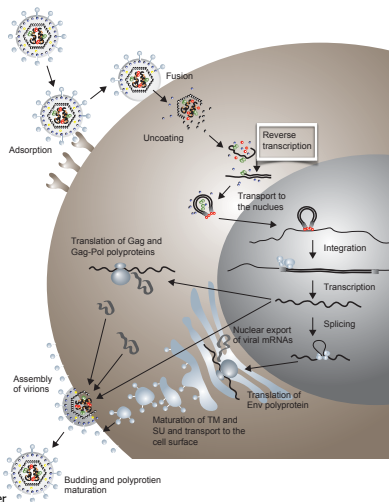


The Nobel Prize in Physiology or Medicine 1975



The Nobel Prize in Physiology or Medicine 1975 was awarded jointly to David Baltimore, Renato Dulbecco and Howard Martin Temin "for their discoveries concerning the interaction between tumour viruses and the genetic material of the cell."

Retrovirus life cycle

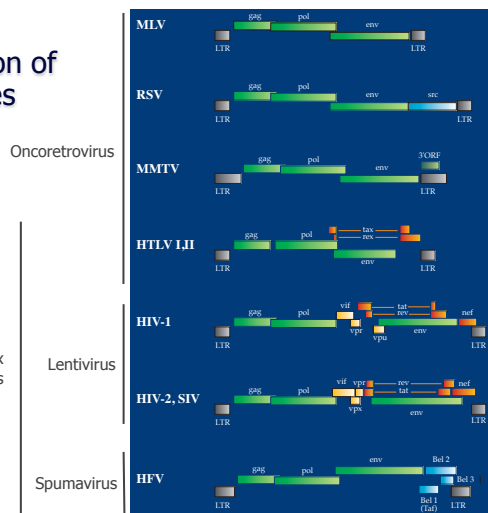


Giacca, M. 2010. Gene Therapy. Springer

Taxonomy of the *Retroviridae* family

| Subfamily | Genus | Former classifications | Main species | Prototype viruses | |
|-------------------|-----------------|--|--|---|--|
| Alpharetrovirus | Alpharetrovirus | Avian type C retroviruses; Avian sarcoma/leukosis viruses (ASLV) | Avian leukosis virus Rous sarcoma virus | ALV RSV | |
| | | Betaretrovirus | Mouse mammary tumor virus Mason-Pfizer monkey virus | MMTV MPMV | |
| Gammaretrovirus | Gammaretrovirus | Mammalian type B retroviruses; Type D retroviruses | Murine leukemia virus Feline leukemia virus Gibbon ape leukemia virus Harvey murine sarcoma virus Moloney murine sarcoma virus | Abelson-MLV, Friend-MLV, Moloney-MLV FeLV GaLV Ha-MSV Mo-MSV | |
| | | Mammalian type C retroviruses | Simian sarcoma virus Reticuloendotheliosis virus Bovine leukemia virus | SSV REV-A, REV-T BLV | |
| | | Deltaretrovirus | BLV-HLV group retroviruses | Primate T-lymphotropic viruses (human and simian) | HTLV-1, STLV-1, HTLV-2, STLV-2, STLV-3 |
| | | Epilontoretrovirus | Fish retroviruses | Walleye dermal sarcoma virus Bovine immunodeficiency virus | WDSV BIV |
| | | Lentivirus | Lentivirus | Equine infectious anemia virus Feline immunodeficiency virus Caprine arthritis encephalitis virus Visna/Maedi virus Human immunodeficiency virus 1 and 2 Simian immunodeficiency virus | EIAV FIV-O, FIV-P CAEV VISNA HIV-1, HIV-2 SIV-agm, SIV, SIV-zpz, SIVmac |
| Spumaretrovirinae | Spumavirus | Spumavirus | Simian foamy virus Bovine foamy virus Equine foamy virus Feline foamy virus Human foamy virus | SFVmac (SFV-1 and SFV-2), SFVagm (SFV-3), SFVcpz and SFVcpz(hu) BFV EFV FFV HFV or HSRV | |

Genetic organization of retroviruses

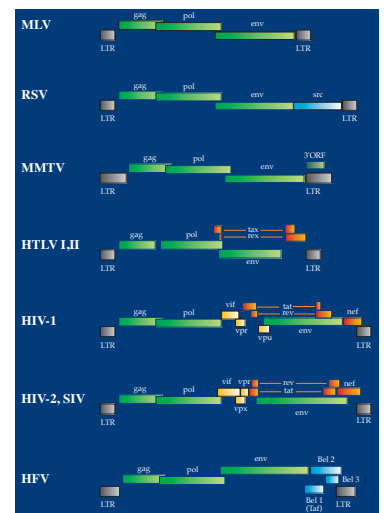


Common features of retroviruses

They all contain LTRs (400-700 nt), which form in the integrated provirus

Viral particles contain mRNA

They all contain gag, pol and env genes



RNA Tumor Viruses - The Rous Sarcoma Virus Story

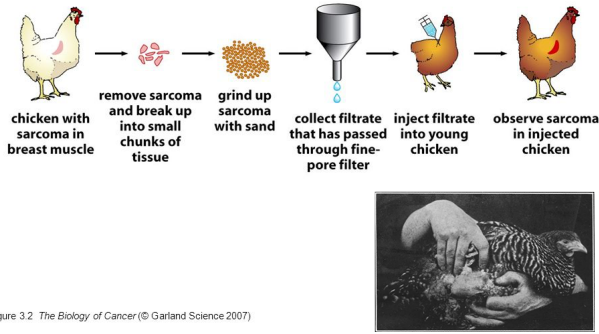


Figure 3.2. The Biology of Cancer (© Garland Science 2007)

The discovery of proto-oncogenes: a version of the src gene carried by RSV is also present in uninfected cells

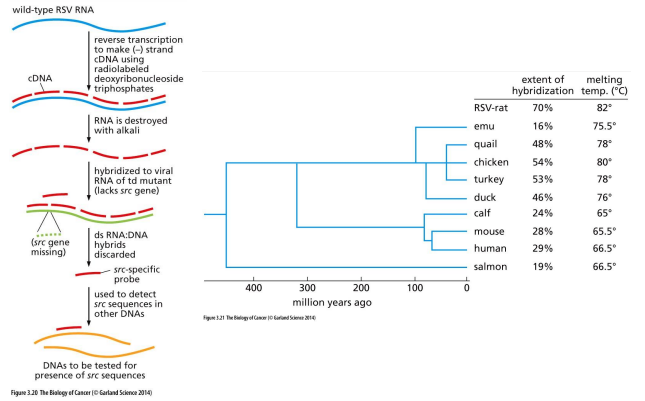


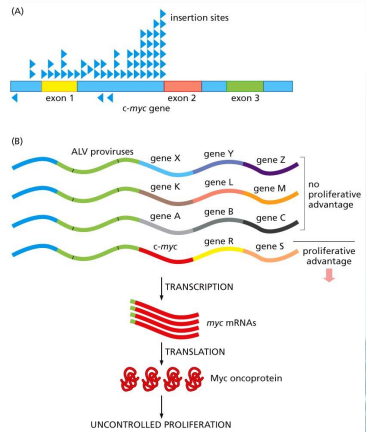
Figure 3.2b. The Biology of Cancer (© Garland Science 2016)

Examples of retroviruses carrying viral oncogenes (v-onc)

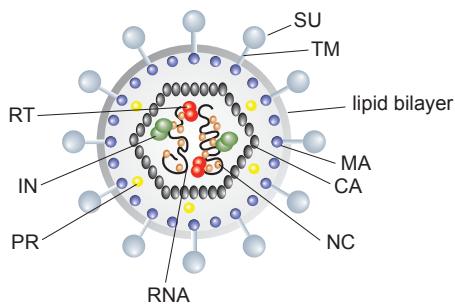
| Parental /helper virus | Retrovirus | Acronym | v-onc |
|--|--|----------|---------|
| | Rous sarcoma virus | RSV | src |
| Avian leukosis virus (ALV) | Avian myeloblastosis virus | AMV | myb |
| | Avian erythroblastosis virus | AEV | erbA, B |
| | Avian myelocytomatosis virus 29 | AMCV-29 | myc |
| | Y73 sarcoma virus | Y73SV | yes |
| | Avian sarcoma virus 17 | ASV-17 | jun |
| Moloney-Murine leukemia virus (Mo-MLV) | Abelson murine leukemia virus | Ab-MLV | abl |
| | Harvey murine sarcoma virus | Ha-MSV | ras |
| | Moloney murine sarcoma virus | Mo-MSV | mos |
| | Finkel-Biskis-Jenkins murine sarcoma virus | FBJ-MSV | fos |
| Feline leukemia virus (FeLV) | Snyder-Theilen feline sarcoma virus | ST-FeSV | fes |
| | Gardner-Arnstein feline sarcoma virus | GA-FeSV | |
| | Susan McDonough feline sarcoma virus | SM-FeSV | fms |
| | Hardy-Zuckerman 4 feline sarcoma virus | HZ4-FeSV | kit |
| Simian sarcoma virus (SSV) | Woolly monkey sarcoma virus | WMSV | sis |

Slowly transforming retroviruses activate protooncogenes by inserting their genomes adjacent to these cellular genes

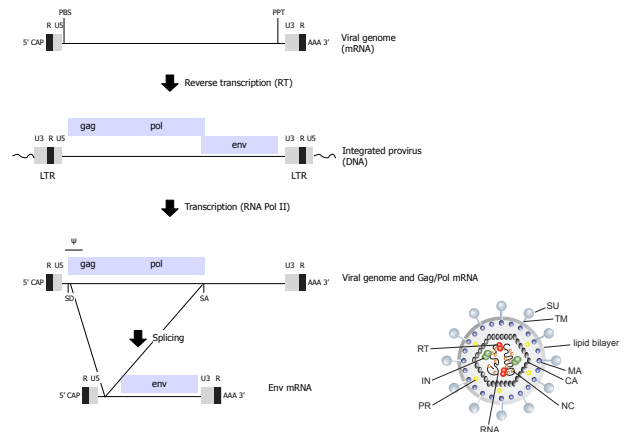
Insertional mutagenesis



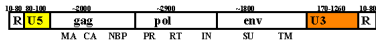
Retrovirus virion



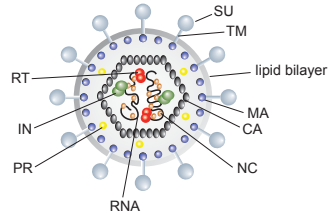
Genetic organization of generalized retrovirus



Genetic organization of generalized retrovirus

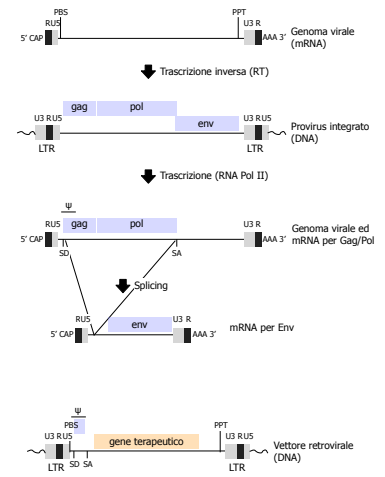


MA Matrix
CA Capsid
NBP Nuclear Binding Protein
PR Protease
RT Reverse Transcriptase
IN Integrase
SU Surface protein
TM Transmembrane Protein



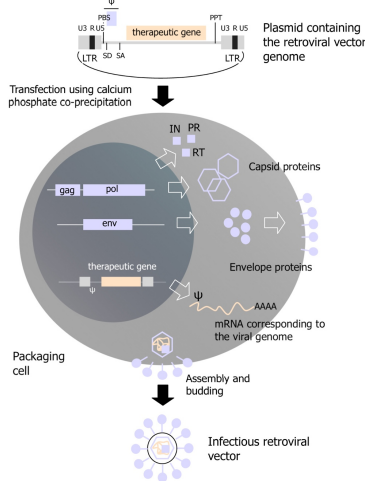
Elementi genetici dei Retrovirus

La trascrizione parte da LTR del virus integrato

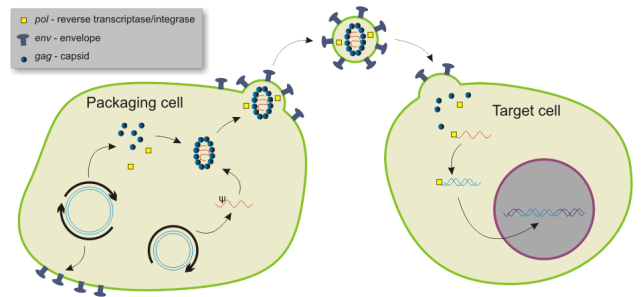


Nel vettore virale rimane:
- LTR che serve per la trascrizione,
- una regione di gag che serve per incapsidamento
- una porzione che produce un tRNA che funziona da primer per la trascrittasi inversa

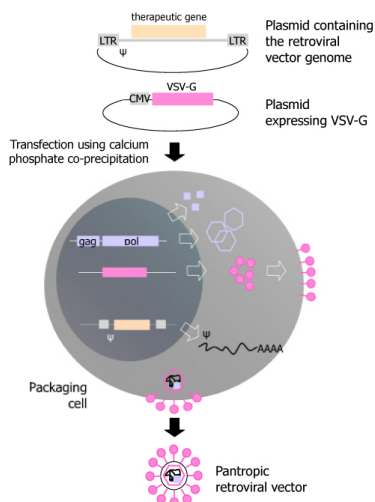
Packaging of gammaretroviral vectors



Retroviral vector integration results in transgene transcription (no additional particles produced)



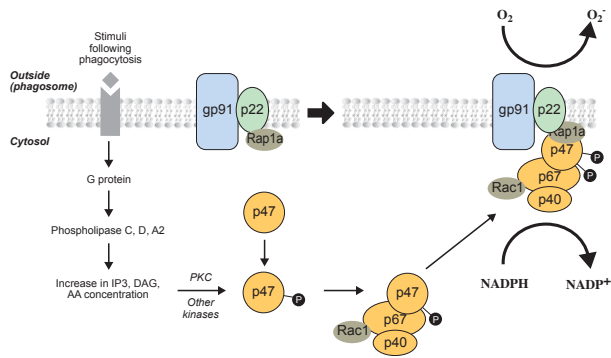
Pseudotyping of gammaretroviral vectors



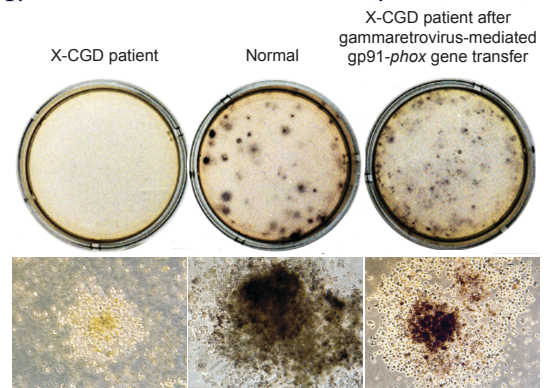
Monogenic hereditary disorders for which gene therapy clinical trials were conducted by gene transfer into HSCs

| Disease group | Disease | Defective gene |
|---|-------------------------------------|---|
| Severe combined immunodeficiency syndromes (SCID) | SCID-X1 | Gamma common (γ c) chain of interleukin receptors |
| | ADA-SCID | Adenosine deaminase |
| | | JAK-3 |
| | PNP-SCID | Purine-nucleoside phosphorylase (PNP) |
| Lysosomal storage disorders | Hurler's disease (MPS I) | α -L-iduronidase |
| | Hunter's disease (MPS II) | Iduronate-2-sulfatase |
| | Gaucher's disease | Glucocerebrosidase (β -glucosidase) |
| | Fabry's disease | α -galactosidase A |
| | Sly syndrome (MPS VII) | β -glucuronidase |
| Defects of phagocytes | Chronic granulomatous disease (CGD) | gp91 ^{phox} , p47 ^{phox} |
| | Leukocyte adhesion disorder | CD18 (β 2-integrin) |
| Other diseases | Fanconi anemia, group C | FANCC |

Activation of phagocyte NADPH oxidase



Functional correction of NADPH activity in myeloid colonies from an X-CGD patient after gene transfer of the gp91^{phox} cDNA into CD34⁺ hematopoietic stem cells



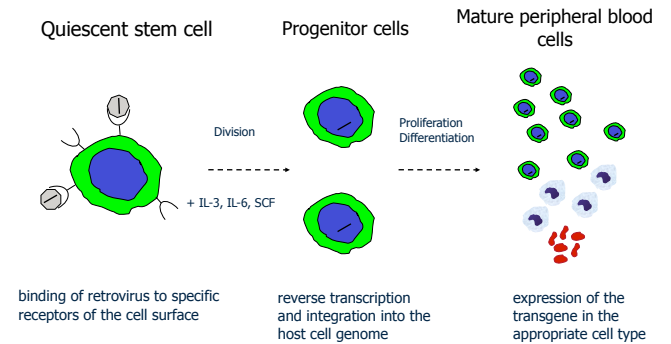
Zentilin, L. et al. 1996. Exp. Cell. Res. 225, 257.

Gene therapy of hematopoietic stem cells: Conclusions from clinical trials

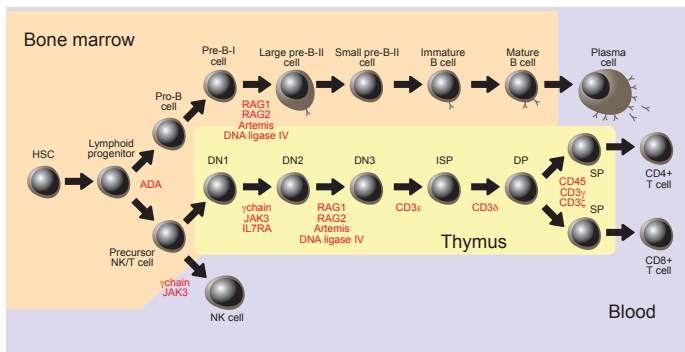
- Virus-positive cells are detectable in peripheral blood after several years from treatment
- Only a very small fraction (0.01-0.1%) of reconstituting HSCs are transduced with the currently available protocols



Gene therapy of hematopoietic stem cells



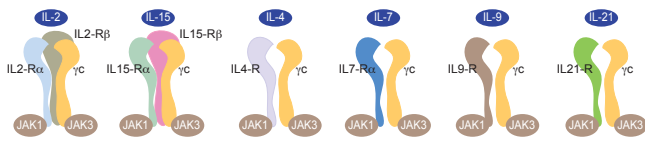
Defects leading to the development of severe combined immunodeficiency (SCID)



SCID-X1, the bubble boy disease



Molecular structure of the interleukin receptors



Gene Therapy of Human Severe Combined Immunodeficiency (SCID)-X1 Disease

Marina Cavazzana-Calvo,^{*1,2,3} Salima Hacein-Bey,^{*1,2,3}
 Geneviève de Saint Basile,¹ Fabian Gross,² Eric Yvon,³
 Patrick Nusbaum,² Françoise Selz,¹ Christophe Hue,^{1,2}
 Stéphanie Certain,¹ Jean-Laurent Casanova,^{1,4} Philippe Bousso,⁵
 Françoise Le Deist,¹ Alain Fischer^{1,2,4†}

Severe combined immunodeficiency-X1 (SCID-X1) is an X-linked inherited disorder characterized by an early block in T and natural killer (NK) lymphocyte differentiation. This block is caused by mutations of the gene encoding the γ c cytokine receptor subunit of interleukin-2, -4, -7, -9, and -15 receptors, which participates in the delivery of growth, survival, and differentiation signals to early lymphoid progenitors. After preclinical studies, a gene therapy trial for SCID-X1 was initiated, based on the use of complementary DNA containing a defective γ c Moloney retrovirus-derived vector and ex vivo infection of CD34⁺ cells. After a 10-month follow-up period, γ c transgene-expressing T and NK cells were detected in two patients. T, B, and NK cell counts and function, including antigen-specific responses, were comparable to those of age-matched controls. Thus, gene therapy was able to provide full correction of disease phenotype and, hence, clinical benefit.

Science. 2000. Vol. 288, pp. 669-672

Leukemia case triggers tighter gene-therapy controls

Trials of gene therapy for SCID were halted in the United States and France following the report that a three-year-old patient treated by Alain Fischer in Paris had developed leukemia after being treated with a retroviral vector (ex vivo transduction of bone marrow stem cells).

In October 2002 an advisory committee to the FDA ruled that gene therapy trials of that kind should now continue. However, there must be increased monitoring for adverse events (abnormal activity of certain cells, integration sites), and patients must receive modified informed consent forms to explain the chances of this side effect occurring. "One adverse event, as serious as it is, in the context of the whole field is not enough to put all programs on hold".



A baby cured of SCID by gene therapy

Using a PCR-based technique, it was discovered that the retroviral vector had inserted into more than 40 sites in the genome of different repopulating cells. In the T-cell clone that grew abnormally, it had inserted in the *LMO-2* oncogene, causing increased expression of the gene. Increased activity of the T-cell clone carrying the *LMO-2* integration was detected in blood samples taken from the boy as early as 13 months after treatment, well before he showed any clinical symptoms. However, this event was probably not sufficient for leukemia, but a second event was required for cancer to ensue.

Another question is the possibility that the boy had a genetic predisposition to leukemia, as there have been two childhood cancers in the family.

γ c gene therapy trial

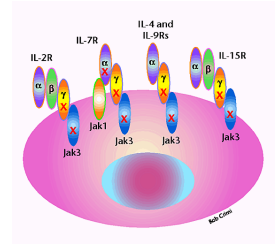
A. Fisher, Paris 2000

Eligibility

SCID-X1 (proven γ c gene mutation)
 Lack of an HLA identical donor

Protocol

Bone marrow harvesting (30-100 ml)
 CD34⁺ cell separation (immunomagnetic micro beads)
 One day pre-activation with SCF, FLT 3L, IL-3 and MGDF
 Three rounds of infection with the MFG γ c vector-containing supernatants in CH-296 fibronectin fragment-coated bags
 I.V. infusion

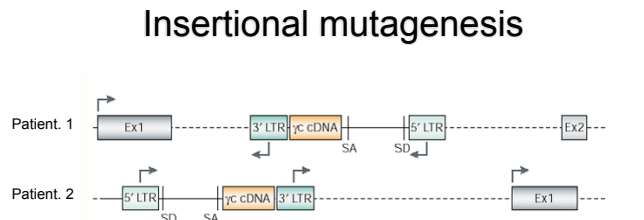


SCID-X1 has been a suitable and attractive setting for the clinical translation of targeted gene correction strategies and adoptive transfer of gene-corrected cells, as cells bearing a functional gamma chain show a positive selective advantage in vivo in the affected patients

December 2002

Second Child in French Trial Is Found to Have Leukemia

| Gene Therapy Studies Under Review | | | |
|-----------------------------------|------------------------------------|------------------|---------|
| Lead investigator | Institution | Disease | Status |
| A. Fischer | Necker Hospital, Paris | X-SCID | on hold |
| H. Malech and J. Puck | NIH | X-SCID | on hold |
| B. Sorrentino and R. Buckley | St. Jude Children's, Memphis; Duke | JAK-3 deficiency | on hold |
| A. Thrasher | Inst. of Child Health, London | X-SCID | on hold |
| K. Weinberg and D. Kohn | NIH and Children's Hospital, LA | X-SCID | on hold |
| F. Condotti and D. Kohn | NIH and Children's Hospital, LA | ADA-SCID | on hold |
| C. Bordignon | San Raffaele Institute, Milan | ADA-SCID | on hold |



- LMO2 encodes a LIM domain protein that binds to transcription factors SCL/TAL1, GATA1, GATA2
- Expressed by haematopoietic progenitors and cells of myeloid lineage, but not in post-thymic T cells
- LMO2 is activated in childhood ALL and in other spontaneous human T cell leukaemias
- LMO2 is leukaemogenic when overexpressed in transgenic mice

Good news for gene therapy

Gene Therapy Insertional Mutagenesis Insights

Utpal P. Davé, Nancy A. Jenkins, Neal G. Copeland*

SCIENCE VOL 303 16 JANUARY 2004

The finding that a retrovirally induced mouse leukaemia contains integrations at both *Lmo2* and γ C loci provided genetic evidence for cooperativity between LMO2 and γ C

In most gene therapy trials, the transplanted gene is unlikely to be oncogenic and occurrences of insertional mutagenesis will be low, as has been seen in trials conducted during the past several years

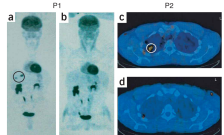


Figure 4 Fixed PEIT scans of P1 (a,b) and fixed PEIT scans of P2 (c,d) before (a,c) and 50 (b) or 53 (d) after gene therapy. Circle in a denotes the active site from the GFP gene transfer.

Correction of X-linked chronic granulomatous disease by gene therapy, augmented by insertional activation of *MDS1-EV11*, *PRDM16* or *SETBP1*

Marion G Ott^{1,16}, Manfred Schmidt^{2-4,16}, Kerstin Schwarzwaldler^{3-5,16}, Stefan Stein^{3,16}, Ulrich Siler^{7,16}, Ulrike Koehl⁸, Hanno Glimm^{9,3}, Klaus Kühlike⁸, Andrea Schilz⁹, Hana Kunke⁸, Sonja Naundorff⁸, Andrea Brinkmann⁸, Annette Deichmann⁸, Marlene Fischer^{3,5}, Claudia Ball^{3,5}, Ingo Pilz^{3,5}, Cynthia Dunbar¹⁰, Yang Du¹¹, Nancy A Jenkins¹¹, Neal G Copeland¹¹, Ursula Lüthi¹², Moustapha Hassan¹³, Adrian J Thrasher¹⁴, Dieter Hoelzer¹, Christof von Kalle^{2-3,15,16}, Reinhard Seger^{7,16} & Manuel Grez¹⁶

Gene transfer into hematopoietic stem cells has been used successfully for correcting lymphoid but not myeloid immunodeficiencies. Here we report on two adults who received gene therapy after nonmyeloablative bone marrow conditioning for the treatment of X-linked chronic granulomatous disease (X-CGD), a primary immunodeficiency caused by a defect in the oxidative antimicrobial activity of phagocytes resulting from mutations in *gp91^{pho}*. We detected substantial gene transfer in both individuals' neutrophils that lead to a large number of functionally corrected phagocytes and notable clinical improvement. Large-scale retroviral integration site-distribution analysis showed activating insertions in *MDS1-EV11*, *PRDM16* or *SETBP1* that had influenced regulation of long-term hematopoiesis by expanding gene-corrected myelopoiesis three- to four-fold in both individuals. Although insertional influences have probably reinforced the therapeutic efficacy in this trial, our results suggest that gene therapy in combination with bone marrow conditioning can be successfully used to treat inherited diseases affecting the myeloid compartment such as CGD.

Correction of ADA-SCID by Stem Cell Gene Therapy Combined with Nonmyeloablative Conditioning

Alessandro Aiuti,¹ Shimon Slavin,² Memet Aker,² Francesca Ficari,¹ Sara Deola,¹ Alessandra Mortellaro,¹ Shoshana Morecki,² Grazia Andolfi,¹ Antonella Tabucchi,³ Filippo Carlucci,³ Enrico Marinello,³ Federica Cattaneo,¹ Sergio Vai,¹ Paolo Servida,⁴ Roberto Miniero,⁵ Maria Grazia Roncarolo,^{1,6 *} & Claudio Bordignon^{1,6,*} †

Science, 2002

The first report of immune restoration in 2 patients with ADA-deficient SCID

These subjects were not treated with PEG-ADA enzyme-replacement therapy, thought to reduce selective advantage of the genetically corrected cells

These subjects received BM cytoablation with a moderate dosage of busulphan, which could promote engraftment of the genetically modified cells

Gene Therapy for Immunodeficiency Due to Adenosine Deaminase Deficiency

Alessandro Aiuti, M.D., Ph.D., Federica Cattaneo, M.D., Stefania Galimberti, Ph.D., Ulrike Benninghoff, M.D., Barbara Cassani, Ph.D., Luciano Callegaro, R.N., Semra Scarlata, Ph.D., Grazia Andolfi, M.D., Massimo Miano, B.Sc., Innocenzo Briganti, B.Sc., Antonella Turchetti, Ph.D., Filippo Cellucci, Ph.D., Martha Esai, M.D., Mirella Mili, M.D., Shimon Shavit, M.D., Haroud Al-Mousa, M.D., Abdulaziz Al-Chrom, M.D., Alina Foster, M.D., Andrea Duppenthaler, M.D., Luigi Neselegro, M.D., Lutz Wittgenberg, M.D., Rebecca H. Buckley, M.D., Marco Bregni, M.D., Sarah Marktel, M.D., Maria Grazia Valsecchi, Ph.D., Paolo Rossi, M.D., Fabio Cirot, M.D., Roberto Miano, M.D., Claudio Bordignon, M.D., and Maria Grazia Roncarolo, M.D.

ABSTRACT

BACKGROUND: We investigated the long-term outcome of gene therapy for severe combined immunodeficiency (SCID) due to the lack of adenosine deaminase (ADA), a fatal disorder of purine metabolism and immunodeficiency.

DESIGN: We infused autologous CD34+ bone marrow cells transduced with a retroviral vector containing the ADA gene into 20 children with SCID due to ADA deficiency who had an HLA-identical sibling donor, after nonmyeloablative conditioning with busulfan. Enzyme-replacement therapy was not given after infusion of the cells.

RESULTS: All patients are alive after a median follow-up of 4.0 years (range, 1.8 to 8.6). Transduced hematopoietic stem cells have engrafted and differentiated into myeloid cells containing ADA (mean range at 1 year in bone marrow [range, 3.5 to 8.9%] and lymphoid cells [mean range in peripheral blood, 5.2 to 88.0%]). Eight patients do not require enzyme-replacement therapy; their blood cells continue to express ADA, and they have no signs of defective classification of purine metabolites. Nine patients had immune reconstitution with increases in T-cell counts (median count at 3 years, 1.0/10⁶ per liter) and normalization of T-cell function. In the five patients in whom intravenous immune globulin replacement was discontinued, antigen-specific antibody responses were detected after exposure to vaccines or viral antigens. Effective protection against infectious and improvement in physical development made a normal life expectancy possible. Serious adverse events included prolonged neutropenia (in two patients), hypertension (in one), central-nervous-system-related infections (in one), Epstein-Barr virus reactivation (in one), and autoimmune hepatitis (in one).

CONCLUSIONS: Gene therapy, combined with reduced-intensity conditioning, is a safe and effective treatment for SCID in patients with ADA deficiency. (ClinicalTrials.gov numbers, NCT00984841 and NCT00999781.)

Table 3. Long-Term Immune Reconstitution after Gene Therapy.*

| Variable | Patients with Normal Value no./total no. |
|--|--|
| Cell count | |
| CD3+ T cells | 5/9 |
| CD4+ T cells | 4/9 |
| Natural killer cells | 3/9 |
| B cells | 4/9 |
| In vitro proliferative responses | |
| PHA mitogen | 9/9 |
| Anti-CD3 mitogen | 9/9 |
| Candida albicans | 7/9 |
| Allergens | 8/9 |
| Serum immunoglobulins | 1/5 |
| IgG | 5/9 |
| IgM | 7/9 |
| IgA | 5/9 |
| Antibodies to specific antigens | |
| Vaccine including TT, DT, BPT, and Hib | 5/5 |
| Pneumococcus (IgM) | 4/5 |
| MMR vaccine or other viral antigens† | 5/5 |

*Results are from the most recent time point at which the patient was not receiving polyethylene glycol-modified bovine adenosine deaminase (PEG-ADA). Data for Patient 8 are not included here; this patient was evaluated for safety only because PEG-ADA was reintroduced 0.4 year after gene therapy. Some results are listed only for the five patients whose serum IgG levels were within the normal range after discontinuation of intravenous immune globulin supplementation. The normal values of cell counts, serum immunoglobulin levels, and in vitro proliferative responses are those reported for age-matched subjects (Elli et al.¹⁷; Cantoni-Bitter et al.¹⁸ and laboratory controls). BPT denotes Bordetella pertussis toxin; DT, diphtheria toxin; Hib, Haemophilus influenzae type b; MMR, measles, mumps, and rubella; PHA, phytohemagglutinin; and TT, tetanus toxoid.

†Viral antigens consisted of antibodies against varicella, Epstein-Barr virus, or cytomegalovirus.

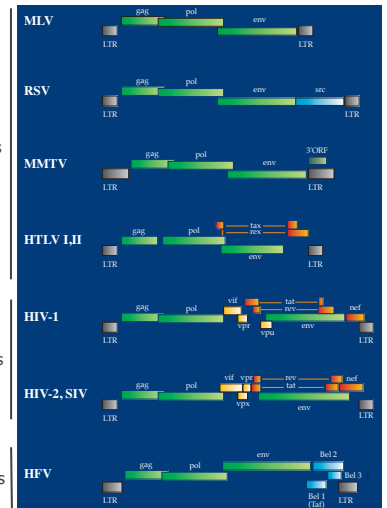
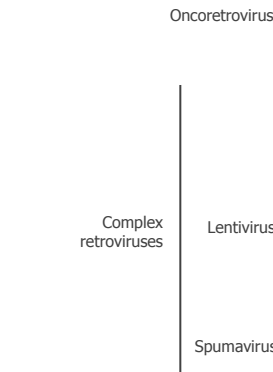
Retroviruses: historical introduction

- 1908 Chicken leukemia is caused by a virus (Ellerman and Bang)
- 1911 Cell-free transmission of sarcoma in chickens (Rous)
- 1936 Mammary carcinoma in mice caused by a filterable agent (Bittner)
- 1958 Development of the focus assay for RSV (Temin and Rubin)
- 1964 Provirus hypothesis (generation of viral DNA copy and integration in cellular genome) (Temin)
- 1970 Reverse Transcriptase (Temin and Mizutani; Baltimore)
- 1976 Probe for src oncogene hybridizes with cellular DNA (Steinlin)
- 1980/82: First human retrovirus (HTLV-I)
- 1983: HIV-1

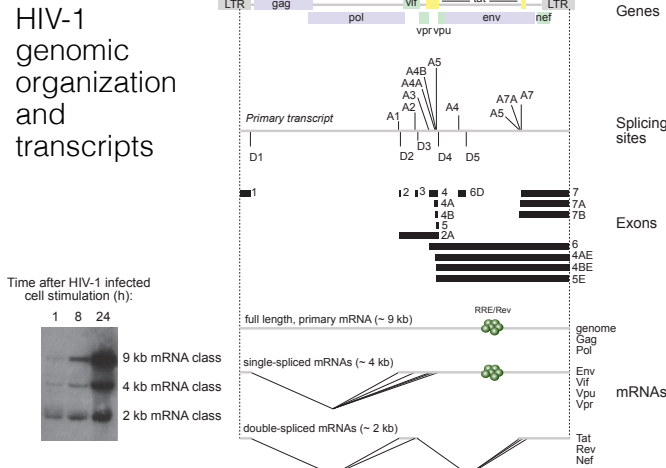
Modern taxonomy of the Retroviridae family

| Subfamily | Genus | Former classifications | Main species | Prototype viruses | | | |
|--------------------|-----------------|--|--------------------------------------|--------------------------------------|--|--------------------|--|
| Alpharetrovirus | | Avian type C retroviruses; Avian sarcoma/leukosis viruses (ASLV) | Avian leukosis virus | ALV | | | |
| | | | Rous sarcoma virus | RSV | | | |
| Betaretrovirus | | Mammalian type B retroviruses; Type D retroviruses | Mouse mammary tumor virus | MMTV | | | |
| | | | Mason-Pfizer monkey virus | MPMV | | | |
| Gammaretrovirus | | Mammalian type C retroviruses | Murine leukemia virus | Abelson-MLV, Friend-MLV, Moloney-MLV | | | |
| | | | Feline leukemia virus | FeLV | | | |
| | | | Gibbon ape leukemia virus | GalV | | | |
| | | | Harvey murine sarcoma virus | Ha-MSV | | | |
| | | | Moloney murine sarcoma virus | Mo-MSV | | | |
| Orthoretrovirinae | Deltaretrovirus | SLV-HTLV group retroviruses | Simian sarcoma virus | SSV | | | |
| | | | Reticuloendotheliosis virus | REV-A, REV-T | | | |
| | | | Bovine leukemia virus | BLV | | | |
| Epsilonretrovirus | | Fish retroviruses | Walleye dermal sarcoma virus | WDSV | | | |
| | | | Bovine immunodeficiency virus | BIV | | | |
| | | | Equine infectious anemia virus | EIAV | | | |
| | | | Feline immunodeficiency virus | FIV-O, FIV-P | | | |
| | | | Caprine arthritis encephalitis virus | CAEV | | | |
| | | | Viral immunodeficiency virus | VISNA | | | |
| | | | Human immunodeficiency virus 1 and 2 | HIV-1, HIV-2 | | | |
| | | | Simian immunodeficiency virus | SIV-agm, SIV, SIV-cpx, SIVmac | | | |
| | | | Spumaretrovirinae | Spumavirus | | Simian foamy virus | SFVmac (SFV-1 and SFV-2), SFVagm (SFV-3), SFVcpz and SFVcpz(h) |
| | | | | | | Bovine foamy virus | BFV |
| Equine foamy virus | EFV | | | | | | |
| Feline foamy virus | FFV | | | | | | |
| Human foamy virus | HFV or HSRV | | | | | | |

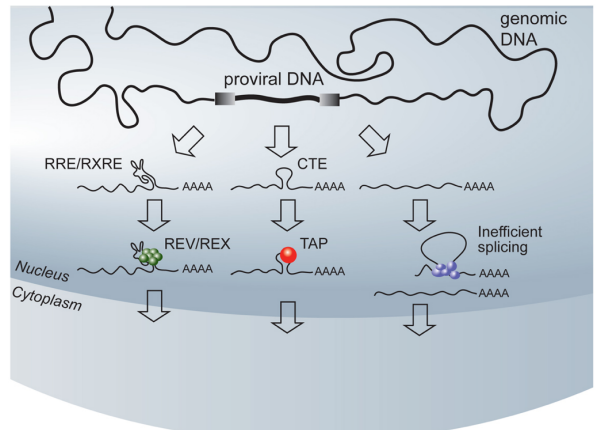
Genetic organization of retroviruses



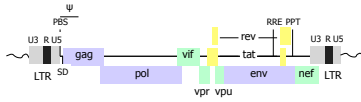
HIV-1 genomic organization and transcripts



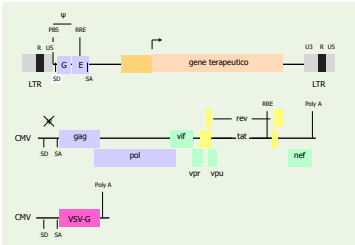
Export of unspliced viral mRNAs outside the nucleus



Vettori lentivirali



Vettori Lentivirali di prima generazione



3 plasmidi

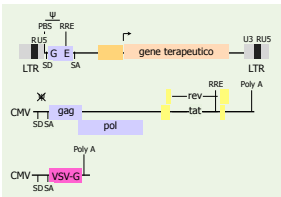
1. segnali regolatori, sito di legame per REV (RRE), promotore e gene terapeutico

2. gag, pol e 6 geni accessori

3. VSV-G (env di HIV lega CD4, espresso essenzialmente in linfociti e macrofagi)

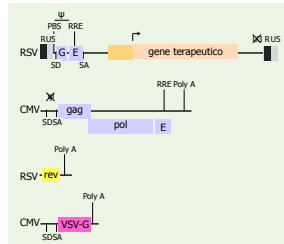
Vettori lentivirali

Vettori lentivirali di seconda generazione



assenza dei fattori di virulenza vif, vpr, vpu, nef

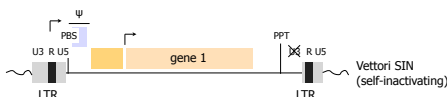
Vettori lentivirali di terza generazione



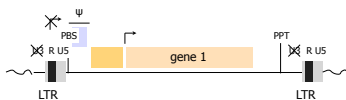
rev in un plasmide separato: ridotta probabilità di ricombinazione

tat non necessaria se trascrizione attivata da CMV

Variazioni nella costruzione dei vettori gammaretrovirali



Trascrizione inversa

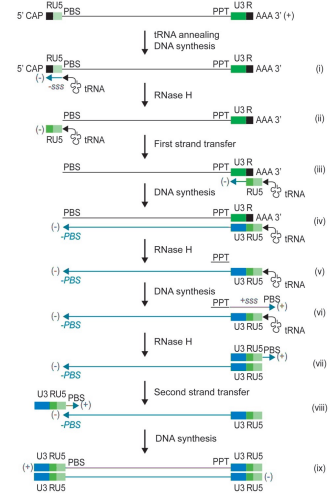


la delezione in U3 viene mantenuta quando il genoma viene retrotrascritto, il che distrugge l'attività di promotore/enhancer del LTR

Safety Concerns Specific to Lentiviral vectors

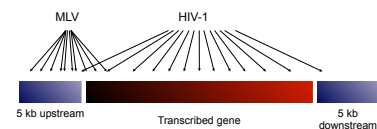
- Recombination during **manufacture** may generate a replication-competent lentivirus (RCL)
 - HIV a known human pathogen
 - vesicular stomatitis virus (VSV-G) envelope broadens tropism
- Recombination with wild type virus in **HIV+ subjects**
- Lentiviral vector **mobilization** by wild type virus

Reverse transcription



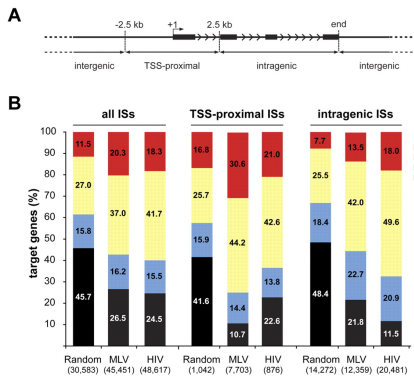
Retroviruses integrate near transcriptionally active regions of DNA

- Acceptor sites for retroviral integrations map near DNase I hypersensitive sites in chromatin (S. Vijaya et al. J. Virol. 1986)
- Retrovirus integration and chromatin structure: Moloney murine leukemia proviral integration sites map near DNase I-hypersensitive sites (H. Rohdewohld et al. J. Virol. 1987)
- Chromosome structure and human immunodeficiency type 1 cDNA integration: centromeric alphoid repeats are a disfavored target (S. Carteau et al. J. Virol. 1998)
- HIV-1 integration in the human genome favors active genes and local hotspots (A.R.W. Schroder et al. Cell 2002)



Integration is not random

MLV: transcriptional start site
HIV: transcriptional units



Lentiviral Hematopoietic Stem Cell Gene Therapy Benefits Metachromatic Leukodystrophy

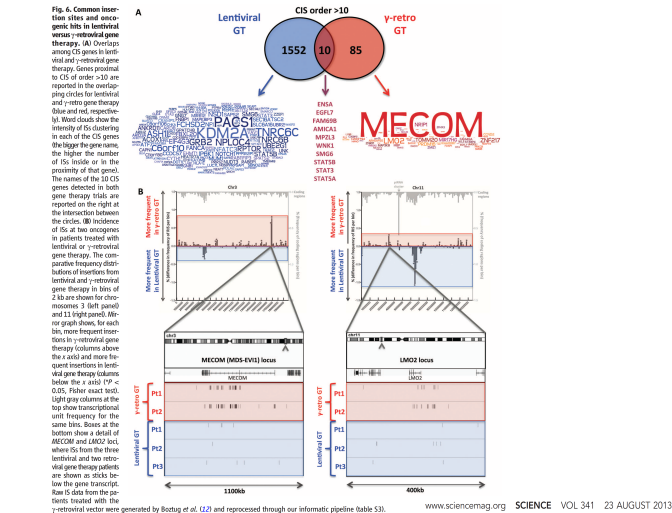
Alessandra Biffi,^{1,2,3,4} Eugenio Montini,^{1,2,3,4} Laura Loric,^{1,2,3,4} Martina Cesani,¹ Francesca Fumagalli,^{1,2,3,4} Tiziana Plat,¹ Cristina Baldoni,¹ Sabata Martino,¹ Andrea Calabria,¹ Sabrina Canale,¹ Fabrizio Benedicenti,¹ Giuliana Vallanti,¹ Luca Bisio,¹ Simone Leo,¹ Nabli Kabbara,¹ Gianrico Zanetti,¹ William B. Rizzo,¹ Nelli A. L. Mehta,¹ Maria Pia Cicalese,^{1,2} Miriam Casiraghi,¹ Jaap J. Roeters,¹ Ubaldo Del Carlo,¹ David J. Dow,¹ Manfred Schmidt,^{1,2} Andrea Asanelli,^{1,2} Victor Medina,¹ Cecilia Di Serio,¹ Ella Stajka,¹ Jason Gardner,¹ Christof von Kalle,^{1,2} Claudio Bordignon,^{1,2} Fabio Cicci,^{1,2} Attilio Rovelli,^{1,2} Maria Grazia Roncarolo,^{1,2,3,4} Alessandro Aiuti,^{1,2,3,4} Maria Serra,^{1,2,3,4} Luigi Naldini^{1,2}

Metachromatic leukodystrophy (MLD) is an inherited lysosomal storage disease caused by arylsulphatase A (ARSA) deficiency. Patients with MLD exhibit progressive motor and cognitive impairment and die within a few years of symptom onset. We used a lentiviral vector to transfer a functional ARSA gene into hematopoietic stem cells (HSCs) from three presymptomatic patients who showed genetic, biochemical, and neurophysiological evidence of late infantile MLD. After reinfusion of the gene-corrected HSCs, the patients showed extensive and stable ARSA gene replacement, which led to high enzyme expression throughout hematopoietic lineages and in cerebrospinal fluid. Analyses of vector integrations revealed no evidence of aberrant clonal behavior. The disease did not manifest or progress in the three patients 7 to 21 months beyond the predicted age of symptom onset. These findings indicate that extensive genetic engineering of human hematopoiesis can be achieved with lentiviral vectors and that this approach may

Lentiviral Hematopoietic Stem Cell Gene Therapy in Patients with Wiskott-Aldrich Syndrome

Alessandro Aiuti,^{1,2,3,4} Luca Bianco,¹ Samantha Scaramuzza,¹ Francesca Ferrua,^{2,3,5} Maria Pia Cicalese,^{1,2} Cristina Baricordi,¹ Francesca Diomiso,¹ Andrea Calabria,¹ Stefania Gianelli,¹ Maria Carmela Caroleola,¹ Maria Bortolotto,¹ Costanza Evangelini,^{1,3} Andrea Asanelli,^{1,2} Miriam Casiraghi,¹ Sara Di Nunzio,¹ Luciano Callegaro,¹ Claudia Benati,¹ Paolo Rizzardi,¹ Danilo Pellico,¹ Celia Di Serio,¹ Manfred Schmidt,¹ Christof von Kalle,¹ Jason Gardner,¹ Nelli Mehta,¹ Victor Medina,¹ David J. Dow,¹ Anne Galy,^{1,2} Roberto Miniero,^{1,2} Andrea Fioschi,¹ Ayele Mettu,¹ Pinaki P. Banerjee,¹ Jordan S. Orange,^{1,3} Stefania Giallombardo,¹ Maria Grazia Roncarolo,^{1,2,3,4} Alessandra Biffi,^{1,2,3,4} Eugenio Montini,^{1,2} Anna Vella,^{1,2} Fabio Cicci,^{1,2} Maria Grazia Roncarolo,^{1,2,3,4} Luigi Naldini^{1,2}

Wiskott-Aldrich syndrome (WAS) is an inherited immunodeficiency caused by mutations in the gene encoding WASP, a protein regulating the cytoskeleton. Hematopoietic stem/progenitor cell (HSPC) transplants can be curative, but, when matched donors are unavailable, infusion of autologous HSPCs modified *in vivo* by gene therapy is an alternative approach. We used a lentiviral vector encoding functional WASP to genetically correct HSPCs from three WAS patients and reinfused the cells after a reduced-intensity conditioning regimen. All three patients showed stable engraftment of WASP-expressing cells and improvements in platelet counts, immune functions, and clinical scores. Vector integration analyses revealed highly polyclonal and multilineage hematopoiesis resulting from the gene-corrected HSPCs. Lentiviral gene therapy did not induce selection of integrations near oncogenes, and no aberrant clonal expansion was observed after 20 to 32 months. Although extended clinical observation is required to establish long-term safety, lentiviral gene therapy represents a promising treatment for WAS.



REVIEW ARTICLE

Acute lymphoblastic leukaemia
Clinical use of lentiviral vectors
Michael C. Milone^{1,2}, Una O'Doherty³

Table 1 Ongoing clinical trials using lentiviral vectors to modify hematopoietic stem cells

| Condition | Phase | NCT number |
|---|-------|-------------|
| Transfusion-dependent β -thalassaemia | 1/2 | NCT02453477 |
| | 3 | NCT02906202 |
| Cerebral adrenoleukodystrophy | 2/3 | NCT01896102 |
| Sickle cell disease | 1 | NCT02140554 |
| Metachromatic leukodystrophy and adrenoleukodystrophy | 1 | NCT02193191 |
| | 1/2 | NCT02559830 |
| Wiskott-Aldrich syndrome | 1/2 | NCT01347346 |
| | 1/2 | NCT01347242 |
| X-SCID | 1/2 | NCT02333760 |
| | 1/2 | NCT01306019 |
| ADA-SCID | 1/2 | NCT01512888 |
| | 1/2 | NCT02999984 |
| Fanconi anemia | 1/2 | NCT01380990 |
| | 1/2 | NCT02931071 |
| X-linked chronic granulomatous disease | 1/2 | NCT02249334 |

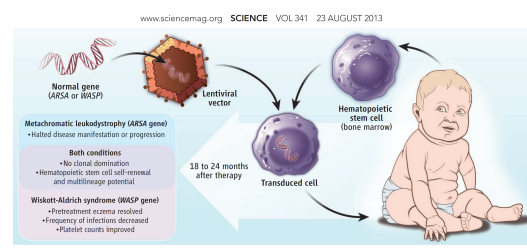
ADA adenosine deaminase, SCID severe combined immunodeficiency

Gene Therapy That Works

Indar M. Verma

The concept of gene therapy is disarmingly simple: introduce a healthy gene in a patient and its product should alleviate the defect caused by a faulty gene or slow the progression of disease (1). Why then, over the past three decades, have there been so few clinical successes in treating patients with this approach? A major obstacle has been the delivery of genes to the appropriate cell, tissue, and organ. How does one introduce a gene into the brain with trillions of cells, or the liver with billions of cells, or the rare hematopoietic adult stem cell that has the

Gene therapy trials show a beneficial effect in children suffering from a neurodegenerative disorder or an immunodeficiency disease.



Targeting β -thalassaemia

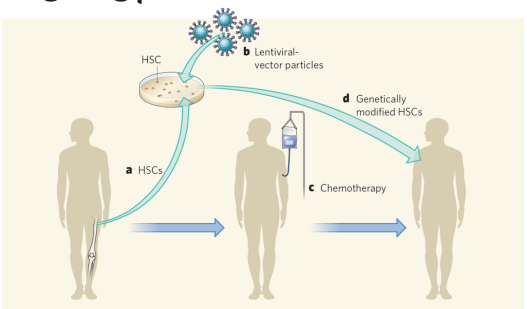


Figure 1 Gene-therapy procedure. **a**, Cavazzana-Calvo *et al.*² collected haematopoietic stem cells (HSCs) from the bone marrow of a patient with β -thalassaemia and maintained them in culture. **b**, The authors then introduced lentiviral-vector particles containing a functional β -globin gene into the cells and allowed them to expand further in culture. **c**, To eradicate the patient's remaining HSCs and make room for the genetically modified cells, the patient underwent chemotherapy. **d**, The genetically modified HSCs were then transplanted into the patient.

LETTERS

Transfusion independence and *HMG2* activation after gene therapy of human β -thalassaemia

Maria Casazza-Cala^{1,2}, Emanuel Payer^{1,2,3}, Oliver Nage^{1,2,3}, Gary Wang^{1,2}, Kathleen Haber^{1,2}, Susanne Hoff^{1,2}, Julian Dool^{1,2}, Maria Drouot^{1,2}, Tracy Brady^{1,2}, Kevin Watters^{1,2}, Rex Cavazos^{1,2}, Brian Lee-Sargent^{1,2}, Laura Gonzalez^{1,2}, Nicolas Spina^{1,2}, Luis Mancera-Ortiz^{1,2}, Francisco Bernaudin^{1,2}, Basim Ghali^{1,2}, Abdul-Djalil, Leif-Jan Nader^{1,2}, Adnan Hachem^{1,2}, Amir Bial^{1,2}, Jean-Sébastien Archin^{1,2}, Nabil Kabbaj^{1,2}, Bruno Datta^{1,2}, Bernard Goumenf^{1,2}, Gerard Sock^{1,2}, Slony Christian^{1,2}, Nathanael Carter^{1,2}, Patrick Kulkarni^{1,2}, Adam Fisher^{1,2}, Adam Cornish^{1,2}, Frédéric Gachon^{1,2}, Yves Seisard^{1,2}, Elvira Guzman^{1,2}, Frederick Bushman^{1,2}, Salim Hasnain^{1,2}, Philippe Leboucq^{1,2,3}

The β -haemoglobinopathies are the most prevalent inherited disorders worldwide. Gene therapy of β -thalassaemia is particularly challenging given the requirement for massive haemoglobin production in a lineage-specific manner and the lack of selective advantage for corrected haematopoietic stem cells. Compound β^0/β^0 thalassaemia is the most common form of severe thalassaemia in southeast Asian countries and their diaspora^{1,2}. The β^0 -globin allele bears a point mutation that causes alternative splicing. The abnormally spliced form is non-coding, whereas the correctly spliced messenger RNA expresses a mutated β^0 -globin with partial instability^{3,4}. When this is compounded with a non-functional β^0 allele, a profound decrease in β -globin synthesis results, and approximately half of β^0/β^0 -thalassaemia patients are transfusion-dependent^{5,6}. The only available curative therapy is allogeneic haematopoietic stem cell transplantation, although most patients do not have a human-leukocyte-antigen-matched, gene-identical donor, and those who do still risk rejection or graft-versus-host disease. Here we show that, 35 months after lentiviral β -globin gene transfer, an adult patient with severe β^0/β^0 -thalassaemia dependent on monthly transfusions since early childhood has become transfusion independent for the past 21 months. Blood haemoglobin is maintained between 9 and 10 g dL⁻¹, of which one-third contains vector-encoded β -globin. Most of the therapeutic benefit results from a dominant, myeloid-biased cell clone, in which the integrated vector causes transcriptional activation of *HMG2* in erythroid cells with further increased expression of a truncated *HMG2* mRNA insensitive to degradation by let-7 microRNAs. The clonal dominance that accompanies therapeutic efficacy may be coincidental and stochastic or result from a hiterto benign cell expansion caused by dysregulation of the *HMG2* gene in stem/progenitor cells.

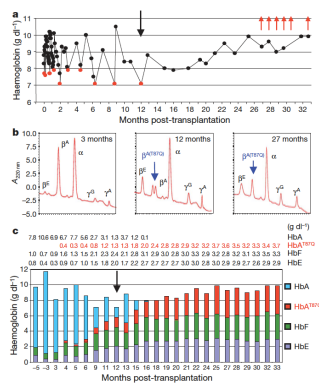


Figure 1 Conversion to transfusion independence. **a**, Total Hb concentrations in whole blood. Red dots, transfusion time points; black vertical arrow, the last time the patient was transfused; red arrows, phlebotomies (200 ml each) to remove excess iron. **b**, HPLC blood globin chain profiles. Note that β^0 only derives from blood transfusions. **c**, Contribution of each Hb species, quantified by HPLC, to total blood Hb concentrations (in g dL⁻¹). Actual numbers for each Hb species are indicated above the chart.

Lentiviral Vector Generations Summary Table

| | First Generation | Second Generation | Third Generation |
|---|-------------------------------|-------------------------------|--|
| Plasmids | 3 | 3 | 4 |
| Deletion in 3' LTR - SIN | No | No | Yes |
| Packaging plasmids with HIV genes | 1 | 1 | 2 |
| Accessory genes: vif, vpr, vpu, nef | All absent | All absent | All absent |
| tat and rev genes | On a single packaging plasmid | On a single packaging plasmid | tat is absent; rev on a separate plasmid |
| gag and pol genes | Same plasmid | Same plasmid | Same plasmid |
| Recombination events needed to generate Replication Competent Lentiviruses (RCL)* | 2 recombinations | 3 recombinations | 4 recombinations between plasmids without homology & must pick a promoter to complement SIN deletion |

Articles

Pro and cons of lentiviral vectors

- Can carry large transgenes (up to 8 Kb)
- Efficient gene transfer
- Infects dividing and non-dividing cells
- No immunogenic proteins generated
- Stable integration into the host genome and stable expression of the transgene

- Potential for generation of RCL
- Potential for insertional mutagenesis: Even replication-incompetent lentiviruses with human tropism are able to infect human cells and integrate their genome into the host cells → risk in case of accidental exposure
- In vivo inactivation by the complement system
- Do not work in all tissues (muscle, heart, vessels)
- No packaging cells for scaling up

Real applications for ex vivo gene therapy (HSC, epithelia)

Lentiviral haemopoietic stem/progenitor cell gene therapy for treatment of Wiskott-Aldrich syndrome: interim results of a non-randomised, open-label, phase 1/2 clinical study

Francesca Fenu¹, Maria Pia Cicales², Stefania Gallimberti, Stefania Giannelli, Francesca Dionisio, Federica Barzaghi, Maddalena Migliaiocco, Maria Ester Bernardi, Valeria Calbi, Andrea Angelo Assanelli, Marcello Facchini, Claudia Fossati, Elena Albertazzi, Samantha Scaramuzza, Immacolata Brigida, Serena Scalo, Luca Basso-Ricci, Roberta Pugno, Miriam Casirogli, Daniele Canarutto, Federica Andreatta, Michael H Albert, Antonella Bartoli, Hermann M Wolf, Rossana Fiori, Paolo Silvani, Salvatore Gattillo, Anna Villa, Luca Biasco, Christopher Dott, Emilij Culme-Seymour, Koenraad van Rossem, Gillian Atkinson, Maria Grazia Valceschi, Maria Grazia Roncarolo, Fabio Cicci, Luigi Naldini, Alessandro Aiuti

Interpretation Data from this study show that gene therapy provides a valuable treatment option for patients with severe Wiskott-Aldrich syndrome, particularly for those who do not have a suitable HSPC donor available.

Funding Italian Telethon Foundation, GlaxoSmithKline, and Orchard Therapeutics.