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# GATEWAY AND TOPO CLONING

#### Introduction to Gateway Cloning

Gateway Cloning Technology is Invitrogen's universal cloning system that provides a rapid and highly efficient route to protein expression, functional analysis and cloning/subcloning of DNA segments. This system is based on the well-characterized bacteriophage lambda-based site-specific recombination system (atrL x atrR ↔ atrB x atrP). In overview, Gateway is a 2-step cloning process; in the first step a sequence of interest is inserted into an Entry Clone. The second step transfers the sequence into a variety of atrB-containing Expression Clones that can be propagated and expressed in a range of host cells for a given experiment.

TOPO Technology is a fast, efficient way to clone. The key to TOPO Cloning is the enzyme DNA topoisomerase I, whose biological role is to cleave and rejoin DNA during replication. To harness this activity, vectors are linearized and each end is conjugated with topoisomerase on the 3 phosphate. This enables fast ligation of DNA sequences with compatible ends. After 5 minutes at room temperature, enzyme is released, the ligation is complete and the recombinant molecule is ready for transformation into E. coli. Many Invitrogen expression vectors are adapted for one-step TOPO Cloning of PCR products in both directional and non-directional formats. Other vectors contain att recombination sequences exterior to the TOPO cloning sites so that cloned inserts are ready for entry into the Gateway system.

Vector NTI Advance 10 offers an interlocking set of computational tools for planning Gateway or TOPO cloning experiments from beginning to end. These allow you to simulate recombinant strategy, as well as to validate design of your reagents before proceeding to primer ordering, bench-top operations, etc. Steps necessary to identify and screen transformants, expedite purification, fuse an insert with vector tags, etc. can be identified and modifications made before

# **Topo Gateway Cloning Manual**

Milos Pawlowski, Yavor Mendel, John Kaisermann

# **Topo Gateway Cloning Manual:**

**Human Stem Cell Manual** Suzanne Peterson, Jeanne F. Loring, 2012-08-27 This reader friendly manual provides a practical hands on guide to the culture of human embryonic and somatic stem cells By presenting methods for embryonic and adult lines side by side the authors lay out an elegant and unique path to understanding the science of stem cell practice

Autophagy in plants and algae Diane C Bassham, Jose L Crespo, 2015-05-15 Autophagy also known as macroautophagy is an evolutionarily conserved process by which cytoplasmic components are nonselectively enclosed within a double membrane vesicle known as the autophagosome and delivered to the vacuole for degradation of toxic components and recycling of needed nutrients This catabolic process is required for the adequate adaptation and response of the cell and correspondingly the whole organism to different types of stress including nutrient starvation or oxidative damage Autophagy has been extensively investigated in yeasts and mammals but the identification of autophagy related ATG genes in plant and algal genomes together with the characterization of autophagy deficient mutants in plants have revealed that this process is structurally and functionally conserved in photosynthetic eukaryotes Recent studies have demonstrated that autophagy is active at a basal level under normal growth in plants and is upregulated during senescence and in response to nutrient limitation oxidative stress salt and drought conditions and pathogen attack Autophagy was initially considered as a non selective pathway but numerous observations mainly obtained in yeasts revealed that autophagy can also selectively eliminate specific proteins protein complexes and organelles Interestingly several types of selective autophagy appear to be also conserved in plants and the degradation of protein aggregates through specific adaptors or the delivery of chloroplast material to the vacuole via autophagy has been reported This research topic aims to gather recent progress on different aspects of autophagy in plants and algae We welcome all types of articles including original research methods opinions and reviews that provide new insights about the autophagy process and its regulation **Characterization of Arabidopsis Basic Science Methods for** Thaliana Defense Responses to the Silverleaf Whitefly Louisa Anne Kempema, 2007 Clinical Researchers Morteza Jalali, Francesca Yvonne Louise Saldanha, Mehdi Jalali, 2017-03-31 Basic Science Methods for Clinical Researchers addresses the specific challenges faced by clinicians without a conventional science background The aim of the book is to introduce the reader to core experimental methods commonly used to answer questions in basic science research and to outline their relative strengths and limitations in generating conclusive data This book will be a vital companion for clinicians undertaking laboratory based science It will support clinicians in the pursuit of their academic interests and in making an original contribution to their chosen field In doing so it will facilitate the development of tomorrow's clinician scientists and future leaders in discovery science Serves as a helpful guide for clinical researchers who lack a conventional science background Organized around research themes pertaining to key biological molecules from genes to proteins cells and model organisms Features protocols techniques for troubleshooting common problems and an

explanation of the advantages and limitations of a technique in generating conclusive data Appendices provide resources for practical research methodology including legal frameworks for using stem cells and animals in the laboratory ethical considerations and good laboratory practice GLP Structure-based Drug Discovery Harren Jhoti, Andrew R. Leach, 2007-05-24 Structure based drug discovery methods have been transformed in the last 5 10 years and are now having a major impact on the discovery of new drugs Some of the most exciting developments in the field such as Fragment based methods are described in this book The book describes the latest developments in technologies that can be used to obtain the 3 D structures including the high profile structural genomics approaches being utilised worldwide The use of 3 D protein structures in new Fragment based approaches to drug discovery are described in some detail This book includes experimental approaches using X ray crystallography and NMR for Fragment based screening as well as other biophysical methods for studying protein ligand interactions In addition developments in computational chemistry methodology are covered together with an assessment of practical applications חחחחחחח חחחחחחח חחחחחחחח II Milos Pawlowski, Yavor Mendel, John Kaisermann, 1960 DNA RNA CRISPR CRISPR Prime Anti CRISPR knock in knockout GeneTalk Haplarithm Haplarithmisis Helicase dependent amplification Immunoprecipitation Isopeptag Jumping library Knockout moss Kodecyte Kodevirion Ligase transfection MassTag PCR Maxam Microsatellite enrichment Minusheet MNase seg plasmon Northern Nuclease Oligomer Oligotyping Oligotyping Paired end tag pBLU pBR322 Peak calling Perturb seg Photoaffinity hybridization Plasmidome Promoter bashing pUC19 Recombinase northern blot transfection Ribosome RNase H PCR Sanger DNA SMILE Seg snRNA seg Sono Seg Southern Southwestern blot Strep tag Streptamer Subcloning TA cloning TBST TCP seg Toeprinting assay DNA Univec VectorDB ViroCap Western blot Western blot **Plant Secondary Metabolism Engineering** Arthur Germano Fett-Neto, 2010-05-06 This book presents detailed practical information on important methods used in the engineering of plant secondary metabolism pathways and the acquisition of essential knowledge in performing this activity including important advances and emerging strategies **Primary and Stem Cells** Uma Lakshmipathy, Bhaskar Thyagarajan, 2011-10-31 This book describes basic cell engineering methods emphasizing stem cell applications and use of the genetically modified stem cells in cell therapy and drug discovery Together the chapters introduce and offer insights on new techniques for engineering of stem cells and the delivery of transgenes into stem cells via various viral and non viral systems. The book offers a guide to the types of manipulations currently available to create genetically engineered stem cells that suit any investigator's purpose whether it's basic science investigation creation of disease models and screens or cells for therapeutic applications Teknik Biologi Molekul II Yavor Mendel, John Kaisermann, Milos Pawlowski, Sejak sekitar tahun 1960 ahli biologi molekul telah mengembangkan kaedah untuk mengenal pasti mengasingkan dan memanipulasi komponen molekul dalam sel termasuk DNA RNA dan protein Kandungan buku ini CRISPR penyuntingan gen CRISPR Prime penyuntingan Anti CRISPR Transfeksi Gen knock in Gen knockout GeneTalk

Haplarithm Haplarithmisis Helicase dependent amplification Immunoprecipitation Pemfokusan isoelektrik Isopeptag Jumping library Knockout moss Kodecyte Kodevirion Tindak balas rantai ligase Ligasi biologi molekul Magnet dibantu transfection MassTag PCR Penjujukan Maxam Gilbert Kaedah untuk menyiasat interaksi protein protein Bahan gelap mikroba Microsatellite enrichment Sistem tanaman perfusi Minusheet MNase seg Resonans plasmon permukaan multi parametrik Mutagenesis teknik biologi molekul Northern blot Northwestern blot Nuklease protection assay Nucleasing acid assiction Oligomer sekatan Oligotyping sequencing Oligotyping taksonomi Overlap extension polymerase chain tindak balas Paired end tag pBLU pBR322 Peak calling Perturb seg Pelabelan fotoaffinity Pemetaan fizikal Vektor transformasi tumbuhan Plak hybridization Plasmid Plasmidome tindak balas rantai Polimerase PRIME PRobe Incorporation Mediated by Enzymes Promoter bashing pUC19 Sentrifugasi kadar zon Penguatan polimerase recombinase Reverse northern blot Reverse transfection Analisis spacer intergenik Ribosom profil Ribosome RNase H bergantung PCR Run off transcription Sanger sequencing Selection and amplification binding assay Single cell sequencing Single penjujukan helai templat sel DNA transkripomik sel tunggal SMiLE Seg snRNA seg Sono Seg Southern blot Southwestern blot Probe isotop stabil Process Staggered extension Strep tag Streptamer Subcloning Surround optik fiber immunoassay Teknologi susunan penggantungan tanaman segerak TA cloning TBST TCP seg Toeprinting assay Inferensi DNA lintasan Urutan mikroskopi elektron DNA penjujukan Univec VectorDB Ujian daya maju ViroCap Western blot Western blot normalisasi **Two-Component** Signaling Systems, Part B, 2007-07-03 Multicellular organisms must be able to adapt to cellular events to accommodate prevailing conditions Sensory response circuits operate by making use of a phosphorylation control mechanism known as the two component system Sections in Two Component Signaling Systems Part B include Structural Approaches Reconstitution of Heterogeneous Systems Intracellular Methods and Assays Genome Wide Analyses of Two Component Systems Presents detailed protocols Includes troubleshooting tips Almindelige molekylærbiologiteknikker II John Kaisermann, Milos Pawlowski, Yavor Mendel, Siden omkring 1960 har molekyl rbiologer udviklet metoder til at identificere isolere og manipulere molekyl re komponenter i celler herunder DNA RNA og proteiner Indholdet af denne bog CRISPR genredigering CRISPR Prime redigering Anti CRISPR Transfektion Gen knock in Gen knockout GeneTalk Haplarithm Haplarithmisis Helicase dependent amplification Immunoprecipitation Isoelektrisk fokusering Isopeptag Jumping library Knockout moss Kodecyte Kodevirion Ligasek dereaktion Ligation molekyl rbiologi transfection Magnetassisteret MassTag PCR Maxam Gilbert sekventering Metoder til unders gelse af protein protein interaktioner Mikrobiel m rk stof Microsatellite enrichment Minusheet perfusionsafgr desystem MNase seg Multi parametrisk overfladeplasmonresonans Mutagenese molekyl rbiologisk teknik Northern blot Northwestern blot Nuclease beskyttelsesassay Nukleinsyrestrukturbestemmelse Oligomer restriktion Oligotyping sekventering Oligotyping taxonomi Overlap ekstensionspolymerasek de reaktion Paired end tag pBLU pBR322 Peak calling Perturb seg Fotoaffinitetsm rkning Fysisk kortl gning Plantetransformationsvektor Plak hybridization Plasmid

Plasmidom Polymerase k dereaktion PRIME PRobe inkorporering medieret af enzymer Promoter bashing pUC19 rate zonal centrifugering Rekombinase polymerase amplifikation Reverse northern blot Reverse transfection Ribosomal intergenisk spacer analyse Ribosome profilering RNase H afh ngig PCR Sanger afk rselstransskription Sanger sekventering selektion og amplificeringsbindingsassay Enkeltcelle sekventering Enkelt celle DNA skabelonstrengsekvensering enkeltcelle transkriptomik SMiLE Seg snRNA seg Sono Seg Southern blot Southwestern blot Stabil isotop sondering forskudt forl ngelsesproces Strep tag Streptamer Subcloning Surround optisk fiberimmunoanalyse Suspension array teknologi Synkron afgr de TA cloning TBST TCP seg Toeprinting assay baneinferens transmissionselektronmikroskopi DNA sekventering Univec VectorDB levedygtighedsassay ViroCap Western blot Western blot normalisering □□□□□□□□□**IV** John Kaisermann, CRISPR CRISPR CRISPR Prime Anti CRISPR transfection RNA transfection knock in knockout knockout GeneTalk Haplarithm Haplarithmisis Helicase dependent amplification Immunoprecipitation Isopeptag Jumping library Knockout moss Kodecyte Kodevirion DNA transfection MassTag PCR Maxam Gilbert Техники на молекулярна биология II Yavor Mendel, John Kaisermann, Milos Pawlowski, 1960 DNA RNA CRISPR CRISPR Prime Anti CRISPR knock in knockout GeneTalk Haplarithm Haplarithmisis Helicase dependent amplification Immunoprecipitation Isopeptag Jumping library Knockout moss Kodecyte Kodevirion transfection MassTag PCR Microsatellite enrichment Minusheet MNase seg Northern Paired end tag pBLU pBR322 Peak calling Perturb seg hybridization PRIME PRobe Incorporation Mediated by Enzymes Promoter bashing pUC19 pUC19 northern blot transfection Ribosomal Ribosome RNase H PCR PCR Sanger DNA SMiLE Seq snRNA seg Sono Seg Southern Southwestern blot Strep tag Strep tag Streptamer Subcloning TA cloning TBST TCP seg Toeprinting assay Toeprinting assay DNA Univec VectorDB ViroCap Western blot Western blot Técnicas de biología molecular II John Kaisermann, Milos Pawlowski, Yavor Mendel, Desde alrededor de 1960 los bi logos moleculares han desarrollado m todos para identificar aislar y manipular componentes moleculares en c lulas que incluyen DNA RNA y prote nas Contenido de este libro CRISPR edici n de genes CRISPR Prime edici n Anti CRISPR Transfecci n Gen knock in Gene knockout GeneTalk Haplarithm Haplarithmisis Helicase dependent amplification Immunoprecipitation Enfoque isoel ctrico Isopeptag Jumping library Knockout moss Kodecyte Kodevirion Reacci n en cadena de la ligasa ligadura biolog a molecular asistida por im n transfection MassTag PCR secuenciaci n de Maxam Gilbert m todos para investigar las interacciones prote na prote na materia oscura microbiana Microsatellite enrichment sistema de cultivo de perfusi n Minusheet MNase seg Resonancia de plasmones de superficie multiparam trica mutag nesis t cnica de biolog a molecular Northern blot Northwestern blot ensayo de protecci n de nucleasa determinaci n de estructura de cido nucleico restricci n de olig meros oligotipado secuenciaci n oligotipado taxonom a cadena de polimerasa de extensi n de solapamiento reacci n Paired end tag pBLU pBR322 Peak calling Perturb seg Etiquetado de fotoafinidad mapeo f sico vector de transformaci n de plantas placa hybridization pl smido plasmidoma reacci n en cadena de la polimerasa PRIME incorporaci n de PRobe mediada por enzimas

Promoter bashing pUC19 centrifugaci n de velocidad de zona amplificaci n de la polibrasa recombinante Inverso northern blot Inverso transfection an lisis de espaciador interg nico ribos mico perfil Ribosome dependiente de RNasa HPCR transcripci n de escorrent a secuenciaci n Sanger ensayo de selecci n y uni n de amplificaci n secuenciaci n de c lulas individuales Single secuenciaci n de cadena de plantilla de c lula DNA transcript mica de c lula nica SMiLE Seg snRNA seg Sono Seg Southern blot Southwestern blot sondeo de is topos estables proceso de extensi n escalonada Strep tag Streptamer Subcloning inmunoensayo de fibra ptica envolvente tecnolog a de matriz de suspensi n cultivo sincr nico TA cloning TBST TCP seg Toeprinting assay inferencia de trayectoria microscop a electr nica de transmisi n DNA secuenciaci n Univec VectorDB ensayo de viabilidad ViroCap Western blot Western blot normalizaci n **Molecular and Genetic** Characterization of Acp29AB and Identification of Acp Interactors in Drosophila Melanogaster Shannon Noelle Albright,2004 Molekuláris biológiai technikák II Yavor Mendel, John Kaisermann, Milos Pawlowski, 1960 k r l a molekul ris biol gusok m dszereket dolgoztak ki a sejtek molekul ris komponenseinek azonos t s ra izol l s ra s manipul l s ra ide rtve a DNA RNA s feh rj ket A k nyv tartalma CRISPR g nszerkeszt s CRISPR Prime szerkeszt s Anti CRISPR transzfekci g n knock in q n knockout GeneTalk Haplarithm Haplarithmisis Helicase dependent amplification Immunoprecipitation izoelektromos f kusz l s Isopeptag Jumping library Knockout moss Kodecyte Kodevirion Lig z l ncreakci Lig ci molekul ris biol gia M gneses asszisztens transfection MassTag PCR Maxam Gilbert szekven l s M dszerek protein protein k lcs nhat sok vizsg lat ra Mikrobi lis s t t anyag Microsatellite enrichment Minusheet perf zi s n v nyi rendszer MNase seg Multiparam teres fel leti plazmonrezonancia mutagenezis molekul ris biol giai technika Northern blot szaknyugati blot nukle z v delmi vizsg lat nukleinsav szerkezet meghat roz sa oligomer restrikci oligot pus meghat roz s szekven l s oligotip l s taxon mia tfed ses kiterjeszt s polimer z l nc reakci Paired end tag pBLU pBR322 Peak calling Perturb seg Fotoaffinit si jel l s Fizikai felt rk pez s N v nytranszform ci s vektor Plakk hybridization Plazmid Plasmidom Polimer z l ncreakci PRIME PRobe be p t s az enzimek ltal k zvet tve Promoter bashing pUC19 Speed zonal centrifug l s Rekombin z polimer z amplifik ci Ford tott northern blot Ford tott transfection Riboszom lis intergenikus spacer anal zis Ribosome profiloz s RN z H f gg PCR Run off transzkripci Sanger szekven l s Szelekci s amplifik ci s k t d si vizsg lat Egysejt szekven l s Egy sejt DNA templ t sz l szekven l s egysejt transzkriptika SMiLE Seg snRNA seg Sono Seg Southern blot Southwestern blot stabil izot p pr ba szakaszos kiterjeszt si folyamat Strep tag Streptamer Subcloning optikai sz las t rbeli immunoassay szuszpenzi s t mb technol gia szinkron v g s TA cloning TBST TCP seq Toeprinting assay p lya k vetkeztet s transzmisszi s elektronmikroszk pia DNA szekven l s Univec VectorDB letk pess gi vizsg lat ViroCap Western blot Western blot normaliz l sa Методы молекулярной биологии II Yavor Mendel, John Kaisermann, Milos Pawlowski, 1960 DNA RNA CRISPR CRISPR Prime Anti CRISPR knock in knockout GeneTalk Haplarithm Haplarithmisis Helicase dependent amplification Immunoprecipitation Isopeptag Jumping library Knockout moss Kodecyte Kodevirion transfection MassTag PCR Microsatellite enrichment Minusheet MNase seg Northern

Paired end tag pBLU pBR322 Peak calling Perturb seg hybridization PRIME PRobe Incorporation Promoter bashing pUC19 northern blot transfection Ribosome H PCR Sanger DNA SMiLE Seg snRNA seg Sono Seg Southern Southwestern blot Strep tag Streptamer Subcloning TA cloning TBST TCP seg Toeprinting assay Univec DNA Univec VectorDB ViroCap Western blot Techniki biologii molekularnej II Milos Pawlowski, Yavor Mendel, John Kaisermann, Od oko o 1960 roku biolodzy molekularni opracowali metody identyfikacji izolacji i manipulowania komponentami molekularnymi w kom rkach w tym DNA RNA i bia kami Zawarto tej ksi ki CRISPR edycja gen w CRISPR Prime edycja Anti CRISPR Transfekcja Gene knock in Gene knockout GeneTalk Haplarithm Haplarithmisis Helicase dependent amplification Immunoprecipitation ogniskowanie izoelektryczne Isopeptag Jumping library Knockout moss Kodecyte Kodevirion Reakcja a cuchowa ligazy ligacja biologia molekularna wspomaganie magnesem transfection MassTag PCR sekwencjonowanie Maxama Gilberta metody badania interakcji bia ko bia ko ciemna materia drobnoustroj w Microsatellite enrichment system perfuzji minusheet MNase seg Wieloparametryczny rezonans plazmon w powierzchniowych Mutageneza technika biologii molekularnej Northern blot Northwestern blot Test ochrony przed nukleaz Okre lanie struktury kwasu nukleinowego Ograniczenie oligomer w Oligotyping sekwencjonowanie Oligotyping taksonomia Overlap extension chain reakcja Paired end tag pBLU pBR322 Peak calling Perturb seg Znakowanie fotopowinowactwa mapowanie fizyczne wektor do transformacji ro lin p ytka hybridization plazmid plazmidom reakcja a cuchowa polimerazy PRIME PRobe Incorporation Mediated by Enzymes Promoter bashing pUC19 wirowanie w strefach szybko ci amplifikacja polimerazy rekombinazowej Odwrotna northern blot odwrotna transfection mi dzygenowa analiza przerywnika rybosomalnego profilowanie Ribosome zale ne od RNazy H PCR transkrypcja Run off sekwencjonowanie Sanger test wi zania selekcji i amplifikacji sekwencjonowanie pojedvnozych kom rek sekwencjonowanie nici matrycowej kom rki DNA Transkryptomika pojedynczych kom rek SMiLE Seg snRNA seg Sono Seg Southern blot Southwestern blot stabilne sondowanie izotop w proces Strep tag roz o enia naprzemiennego Strep tag Streptamer Subcloning test immunologiczny na wiat owodzie surround technologia macierzy zawieszenia uprawa synchroniczna TA cloning TBST TCP seq Toeprinting assay wnioskowanie o trajektorii transmisyjna mikroskopia elektronowa DNA sekwencjonowanie Univec VectorDB test ywotno ci ViroCap Western blot Western blot normalizacja Milos Pawlowski, Yavor Mendel, John Kaisermann, 1960 DNA RNA CRISPR CRISPR Prime Anti CRISPR knock in knockout GeneTalk Haplarithm Haplarithmisis Helicase dependent amplification Immunoprecipitation Isopeptag Jumping library Knockout moss Kodecyte Kodevirion transfection MassTag PCR Maxam Gilbert Microsatellite enrichment Minusheet MNase seq Northern Paired end tag pBLU pBR322 Peak calling Perturb seq hybridization PRIME PRobe Promoter bashing pUC19 northern blot transfection Ribosome RNase H PCR Sanger DNA SMiLE Seg snRNA seg Sono Seg Southern Southwestern blot Strep tag Streptamer Subcloning TA cloning TBST TCP seg Toeprinting assay DNA Univec VectorDB ViroCap Western blot Western blot **Técnicas usuais de biologia molecular II** Milos Pawlowski, Yavor Mendel, John Kaisermann, Desde por

volta de 1960 os bi logos moleculares desenvolveram m todos para identificar isolar e manipular componentes moleculares em c lulas incluindo DNA RNA e prote nas Conte do deste livro CRISPR edi o gen tica CRISPR Prime edi o Anti CRISPR Transfec o gene knock in gene knockout GeneTalk Haplarithm Haplarithmisis Helicase dependent amplification Immunoprecipitation Foco Isopeptag isoel trico Isopeptag Jumping library Knockout moss Kodecyte Kodevirion Rea o em cadeia da ligase Liga o biologia molecular transfection MassTag PCR Sequencia o de Maxam Gilbert M todos para investigar intera es prote na prote na Mat ria escura microbiana Microsatellite enrichment Sistema de cultivo de perfus o em miniplacas MNase seg Resson ncia Northern multiparam trica de plasm nio de superf cie Mutag nese t cnica de biologia molecular Northern blot Northwestern blot Ensaio de prote o de nucleases Determina o da estrutura de cidos nucleicos Restri o de olig meros Oligotipagem seguenciamento Oligotipagem taxonomia Cadeia de polimerase de extens o de sobreposi o rea o Paired end tag pBLU pBR322 Peak calling Perturb seg hybridization Marca o de fotoafinidade Mapeamento f sico Vetor de transforma o vegetal Placa hybridization Plasm deo Plasmidoma rea o em cadeia da polimerase PRIME PRobe Incorporation Mediada por Enzimas Promoter bashing pUC19 Centrifuga o por pUC19 zonas de taxa Amplifica o de polimerase de recombina o northern blot An lise reversa northern blot reversa transfection espa ador interg nico riboss mico Ribosome perfil dependente de RNase H PCR transcri o de Sanger escoamento superficial sequencia o Sanger ensaio de liga o para sele o e amplifica o sequenciamento de c lulas nicas seg enciamento de cadeia de modelo de c lula DNA transcript mica de c lula nica SMiLE Seg snRNA seg Sono Seg Southern blot Southwestern blot Southwestern blot Sondagem de is topos est veis Processo de extens o escalonado Strep tag Streptamer Subcloning Imunoensaio de fibra ptica surround Tecnologia de matriz de suspens o Colheita s ncrona TA cloning TBST TCP seq Toeprinting assay infer ncia Traject ria microscopia Electr nica de Transmiss o DNA sequencia o Univec VectorDB ensaio de viabilidade ViroCap Western blot Western blot normaliza o

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