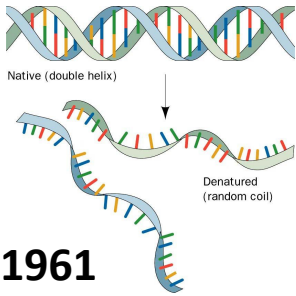


Métodos de Análise do DNA, de Genes e Genoma I

Genética Molecular
RIB-0102

Wilson Araújo Silva Jr
FMRP/USP
wilsonjr@usp.br

DNA Recombinante ...

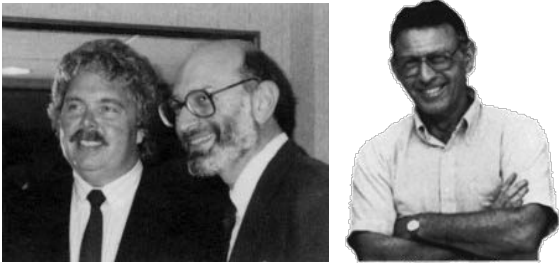


1961
Marmur e Doty
Renaturação de DNA



1966
Nirenberg, Ochoa e Khorana
Elucidaram o código genético

	T			C			A			G		
T	TTT	Ph	F	TC	TCC	S	TAT	Tyr	Y	TGT	Cys	C
	TTG	Leu	L	TCA	TCG		TAA	STOP		TGC		
	TTC			TCT			TAG			TGG	Trp	W
C	CTT	Leu	L	CCT	Pro	P	CAT	His	H	CGT	Arg	R
	CTC			CCC			CAC			CCG		
	CTA			CCA			CAG	Gln	Q	CCG		
	CTG			CCG			CAG			CCG		
A	ATT	Ile	I	ACT	Thr	T	ATG	Met	M	AGT	Ser	S
	ATC			ACC			AAC			AGC		
	ATA			ACA			AAA	Lys	K	AGA	Arg	R
	ATG	Met	M	ACG			AGG			AGG		
G	GTT	Val	V	GCT	Ala	A	GAT	Asp	D	GGT	Gly	G
	GTG			GCC			GAC			GGC		
	GTA			GCA			GAA	Glu	E	GGA		
	GTG			GCG			GAG			GGG		



1972
Boyer, Cohen e Berg
Clonagem gênica

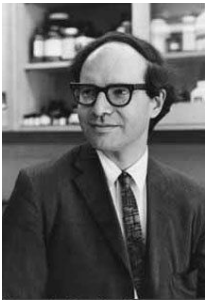
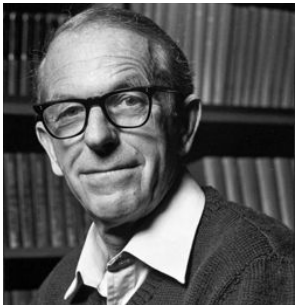
1962
Wener Arber
Enzimas de restrição

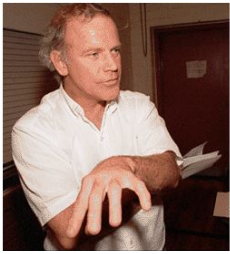


1967
Martin Gellert
DNA ligase



1975-1977
Frederick Sanger e Walter Gilbert
Sequenciamento do DNA





1987

Kary Mullis

Reação em cadeia da polimerase (PCR)

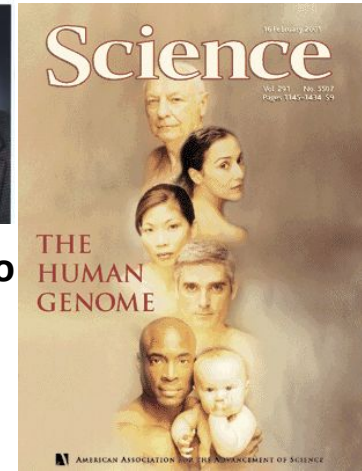
Francis Collins
Diretor do "National Human Genome Research Institute"



2001

Seqüenciamento completo do Genoma Humano

Craig Venter
Co-fundador da "Celera Genomics Corporation"

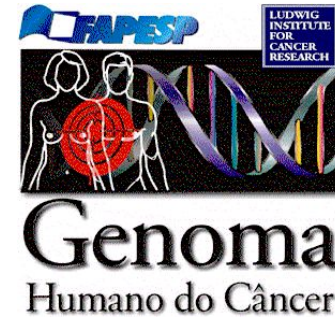


1990

Lançado o Projeto Genoma Humano (HGP)

Esforço de 15 anos coordenado pelo "Department of Energy – DOE" e pelo "National Institutes of Health – NIH" dos USA para o seqüenciamento completo do genoma humano.

1999



Tecnologia do DNA recombinante

Conceito:

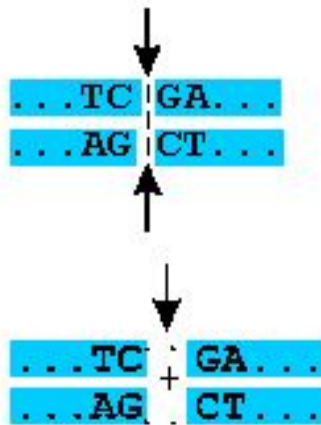
É um conjunto de técnicas de manipulação de DNA em um mesmo organismo ou entre espécies diferentes.

Engenharia genética
Biotecnologia

Tecnologia do DNA recombinante

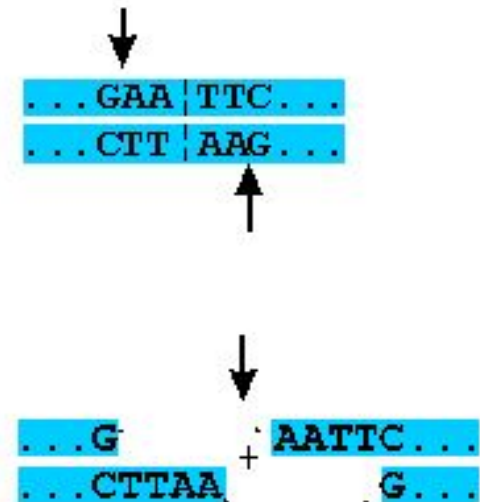
Enzima de restrição:

a) Clivagem no eixo de simetria



Moléculas com extremidades abruptas

b) Clivagem simetricamente situada ao redor do eixo de simetria

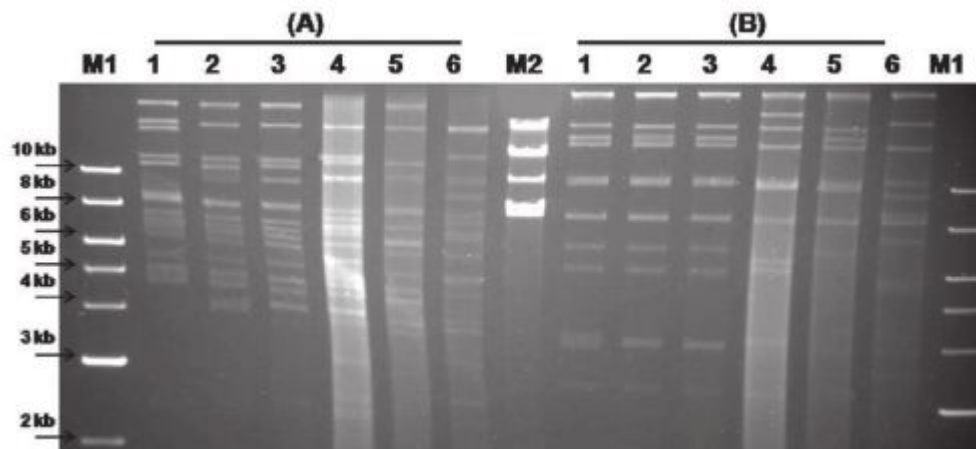
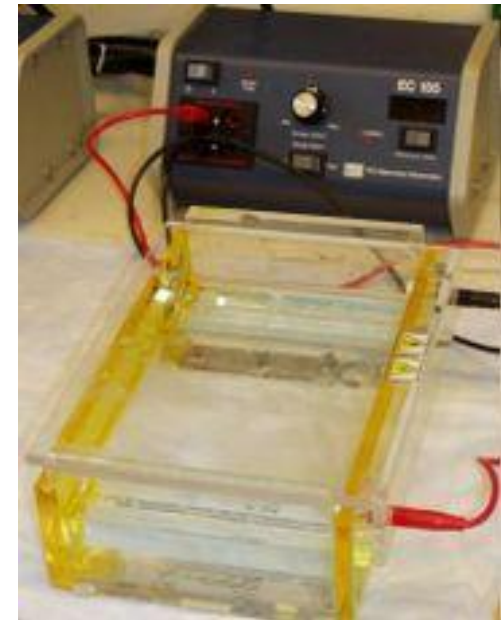
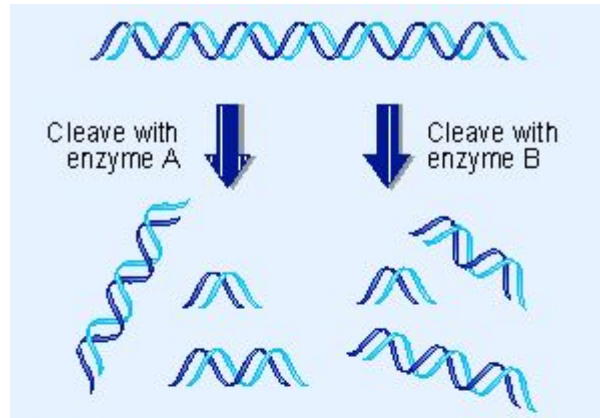


Moléculas com extremidades coesivas

Enzima de restrição:

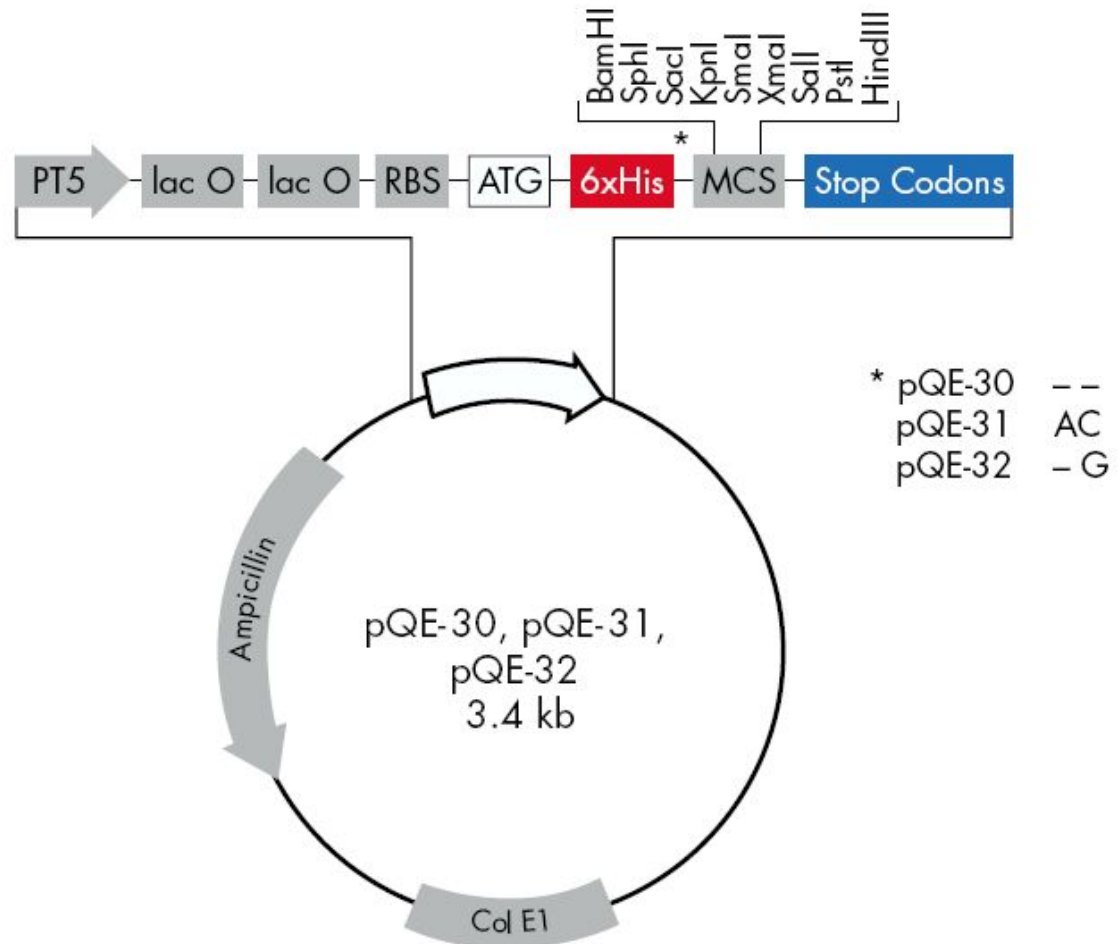
MICROORGANISM	ENZYME ABBREVIATION	SEQUENCE
<i>Haemophilus aegyptius</i>	<i>HaeIII</i>	5' ... G G C C ... 3' 3' ... C C G G ... 5'
<i>Thermus aquaticus</i>	<i>TaqI</i>	5' ... T C G A ... 3' 3' ... A G C T ... 5'
<i>Haemophilus haemolyticus</i>	<i>HbaI</i>	5' ... G C G C ... 3' 3' ... C G C G ... 5'
<i>Desulfovibrio desulfuricans</i>	<i>DdeI</i>	5' ... C T N A G ... 3' 3' ... G A N T C ... 5'
<i>Moraxella bovis</i>	<i>MboII</i>	5' ... G A A G A (N) ₈ ... 3' 3' ... C T T C T (N) ₇ ... 5'
<i>Escherichia coli</i>	<i>EcoRV</i>	5' ... G A T A T C ... 3' 3' ... C T A T A G ... 5'
	<i>EcoRI</i>	5' ... G A A T T C ... 3' 3' ... C T T A A G ... 5'
<i>Providencia stuarti</i>	<i>PstI</i>	5' ... C T G C A G ... 3' 3' ... G A C G T C ... 5'
<i>Microcoleus</i>	<i>MstII</i>	5' ... C C T N A G G ... 3' 3' ... G G A N T C C ... 5'
<i>Nocardia otitidis-caviarum</i>	<i>NotI</i>	5' ... G C G G C C G C ... 3' 3' ... C G C C G G C G ... 5'

Tecnologia do DNA recombinante



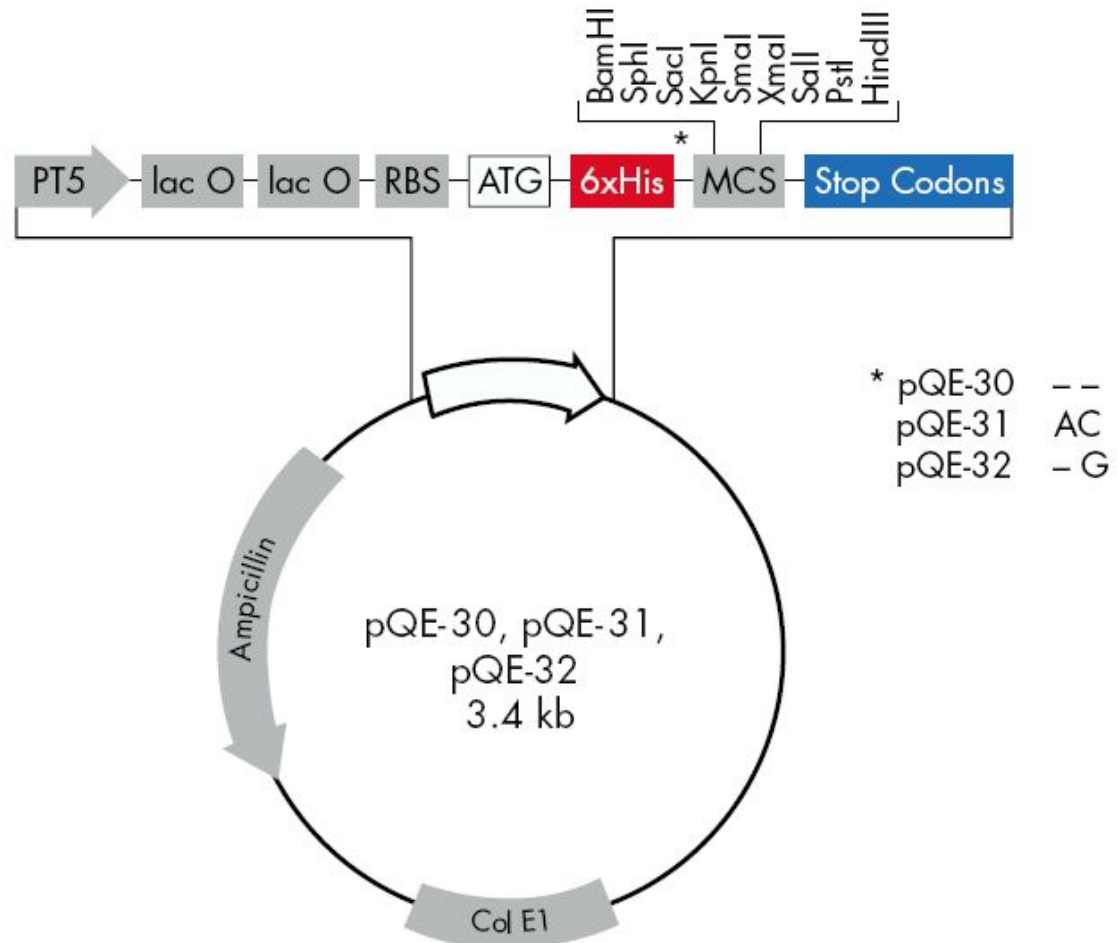
Tecnologia do DNA recombinante

Vetores:



Tecnologia do DNA recombinante

Vetores:



Outros vetores:

Cosmídeos

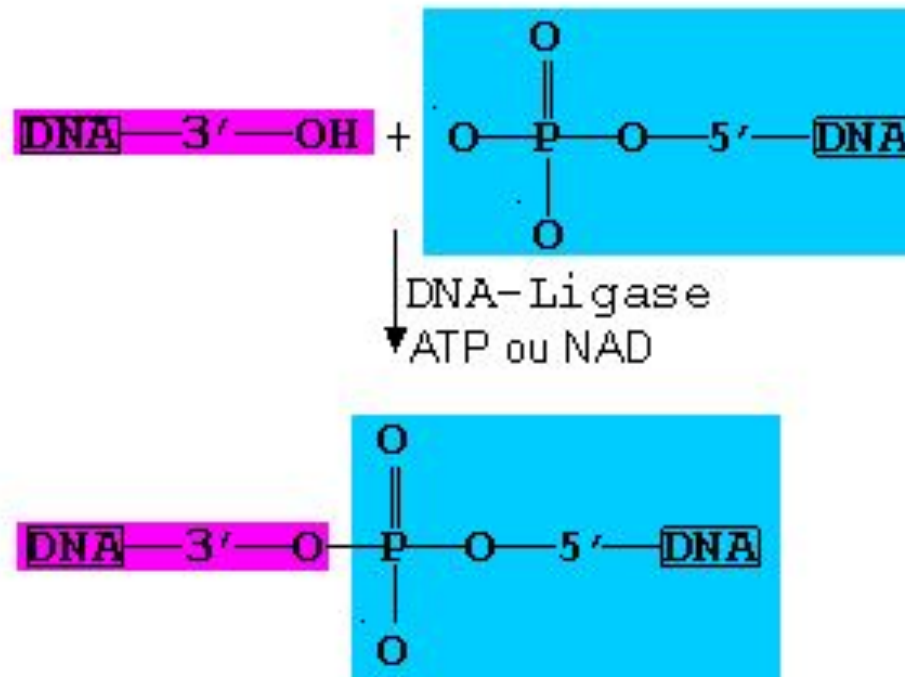
BACs

YACs

Bacteriófagos

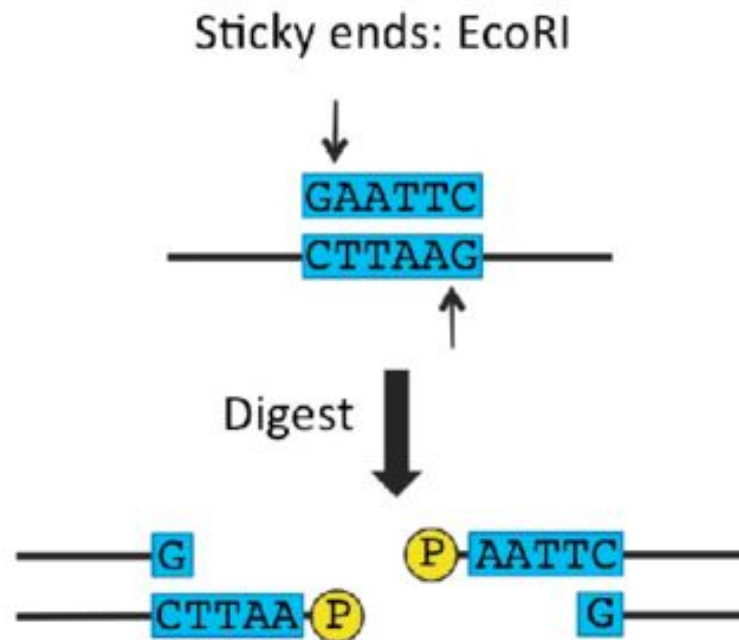
Tecnologia do DNA recombinante

DNA ligase:



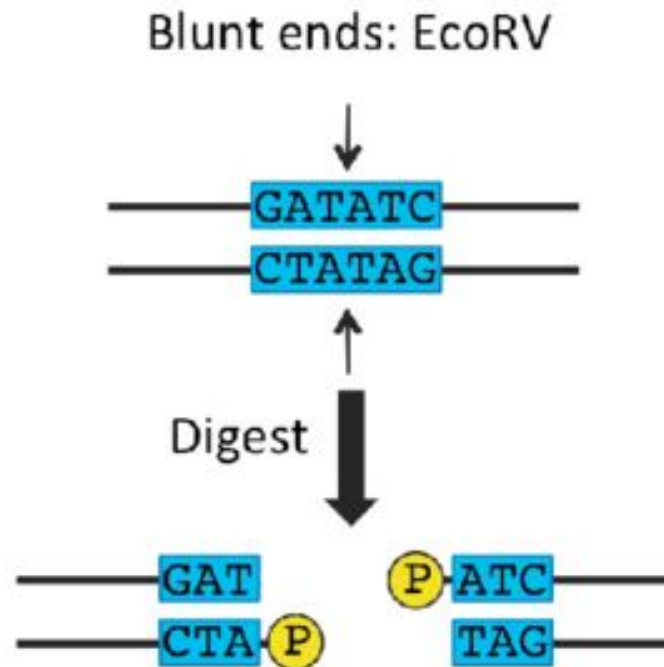
Tecnologia do DNA recombinante

Digestão com extremidades coesivas:



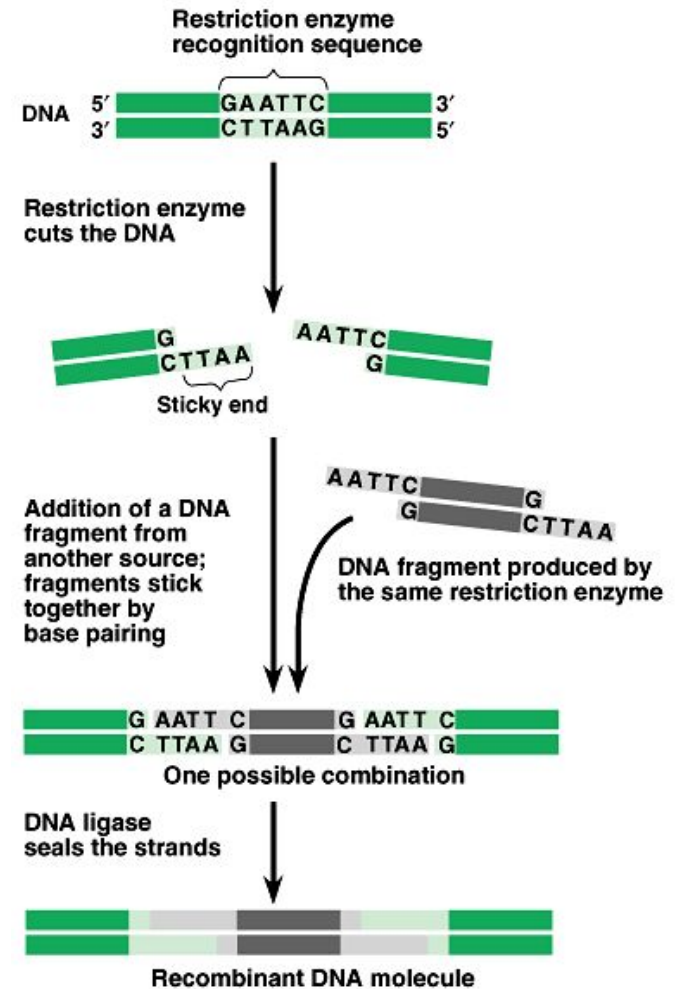
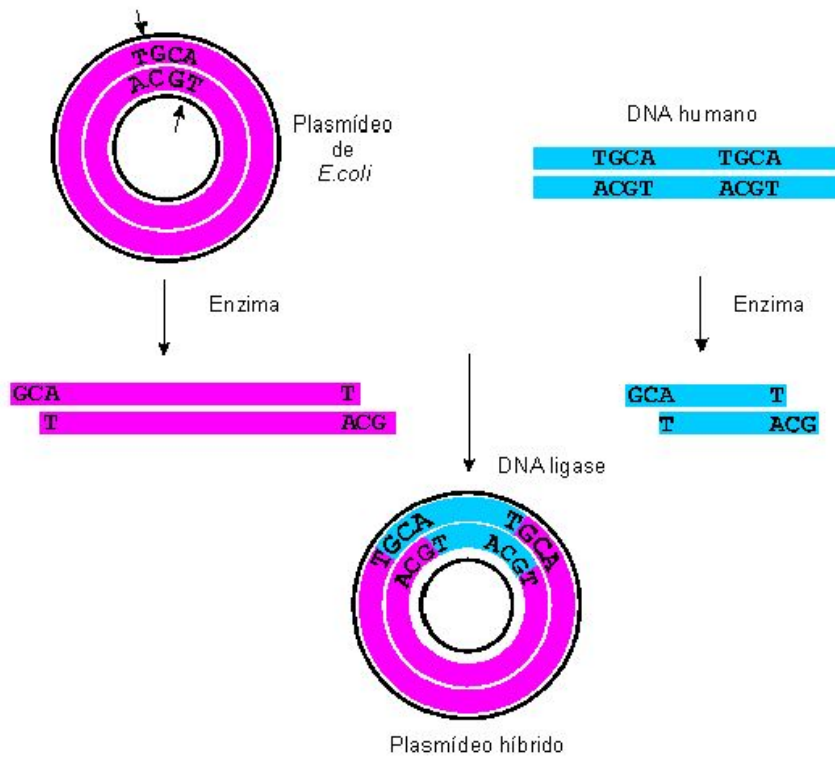
Tecnologia do DNA recombinante

Digestão com extremidades não coesivas:



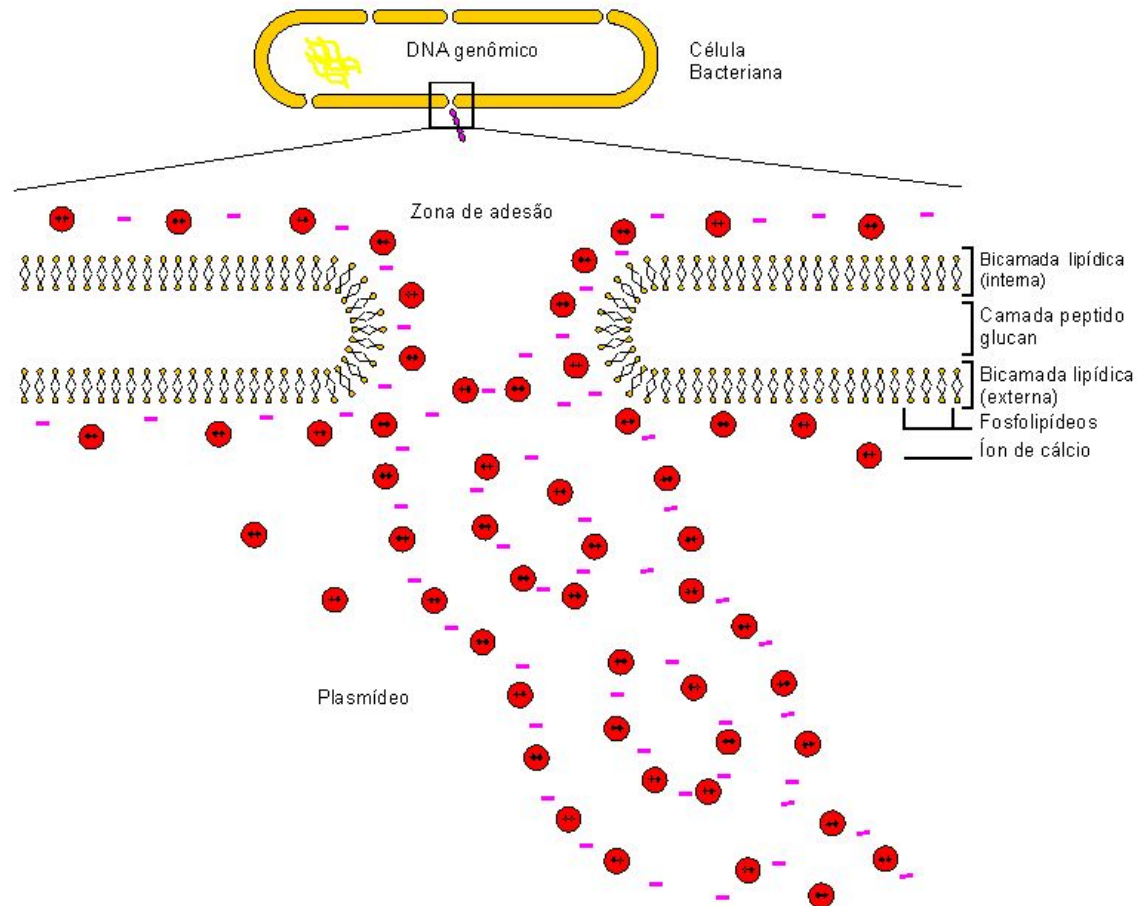
Tecnologia do DNA recombinante

Clonagem:



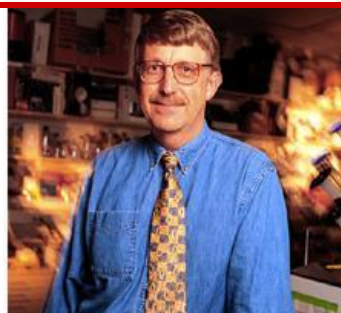
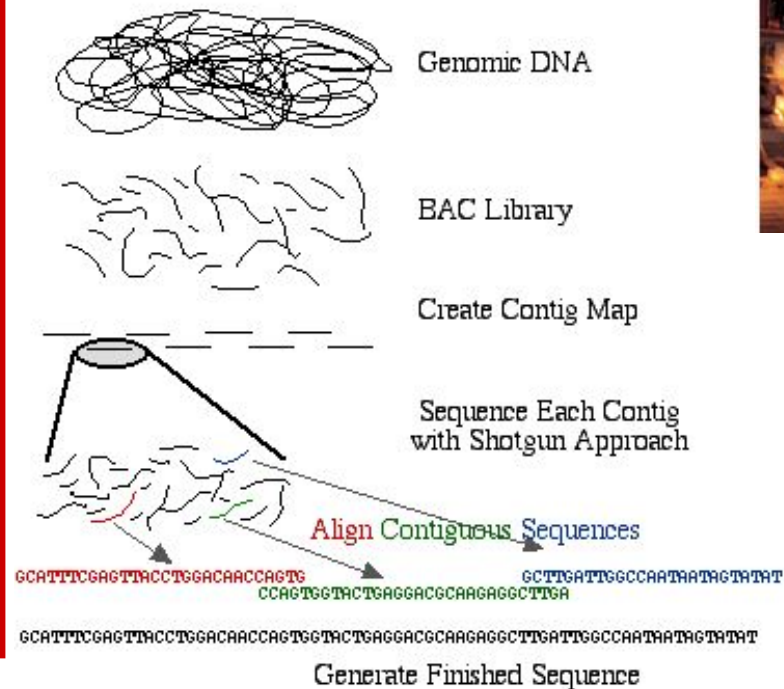
Tecnologia do DNA recombinante

Transformação:



Biblioteca Genômica

Hierarchical Shotgun Sequencing Method



Francis Collins

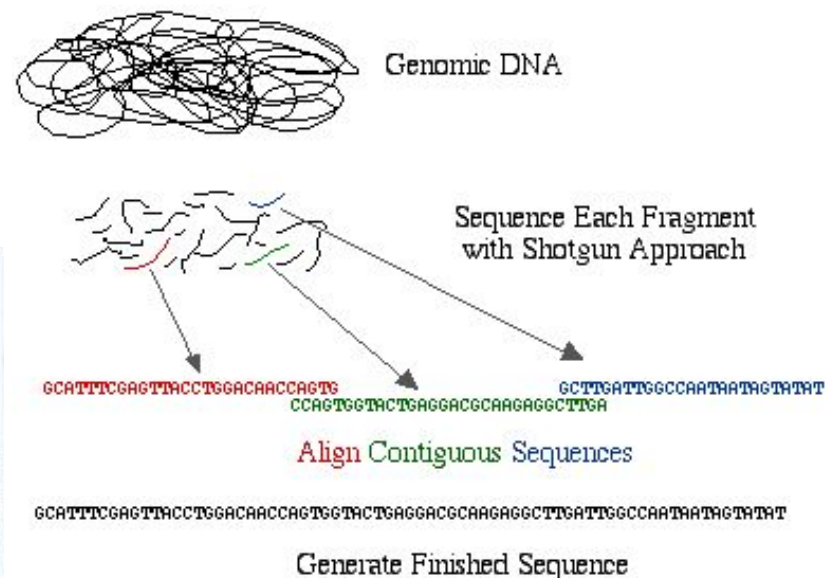
Diretor do "National Human Genome Research Institute"

Craig Venter

Co-fundador da "Celera Genomics Corporation"



Whole Genome Shotgun Sequencing Method



Biblioteca Genômica

Hierarchical Shotgun Sequencing Method



Genomic DNA



BAC Library

Create Contig Map



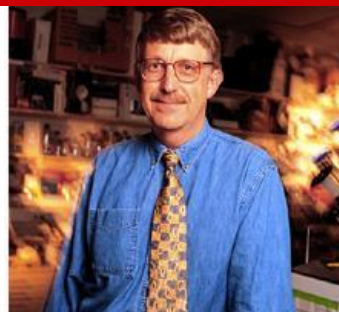
Sequence Each Contig with Shotgun Approach

Align Contiguous Sequences

GCATTTCGAGTTACCTGGACACCAAGTG GCTTGATTGGCCATATAGTATAT
CCAGTGGTACTGAGGACGCAAGAGGCTTGA

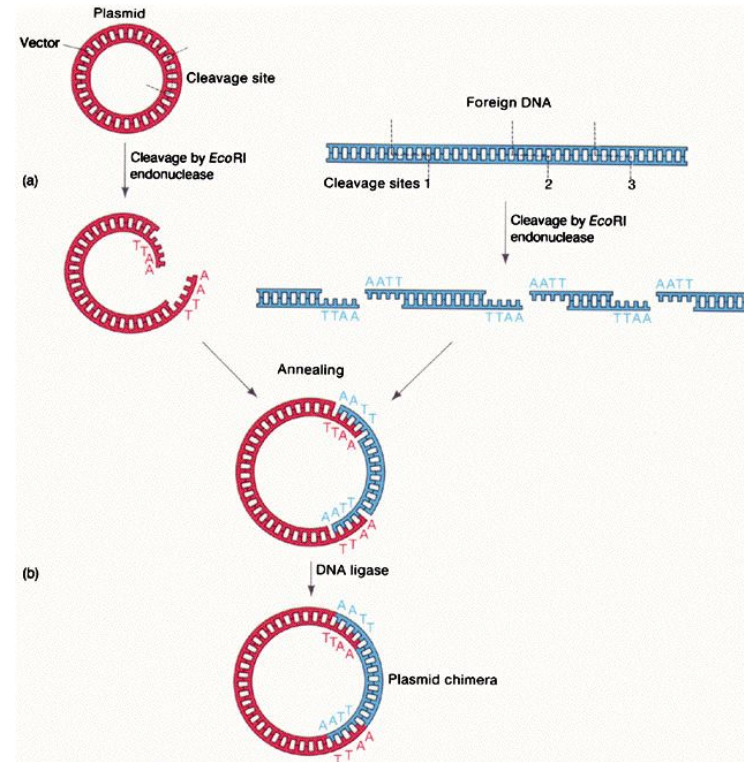
GCATTTCGAGTTACCTGGACACCAAGTGGTACTGAGGACGCAAGAGGCTTGATTGGCCATATAGTATAT

Generate Finished Sequence

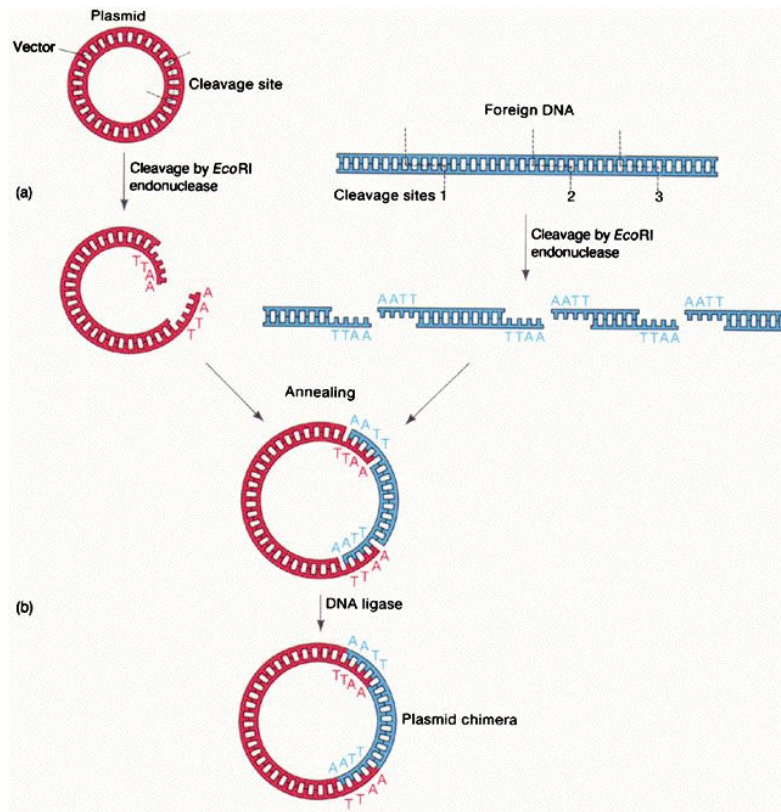


Francis Collins

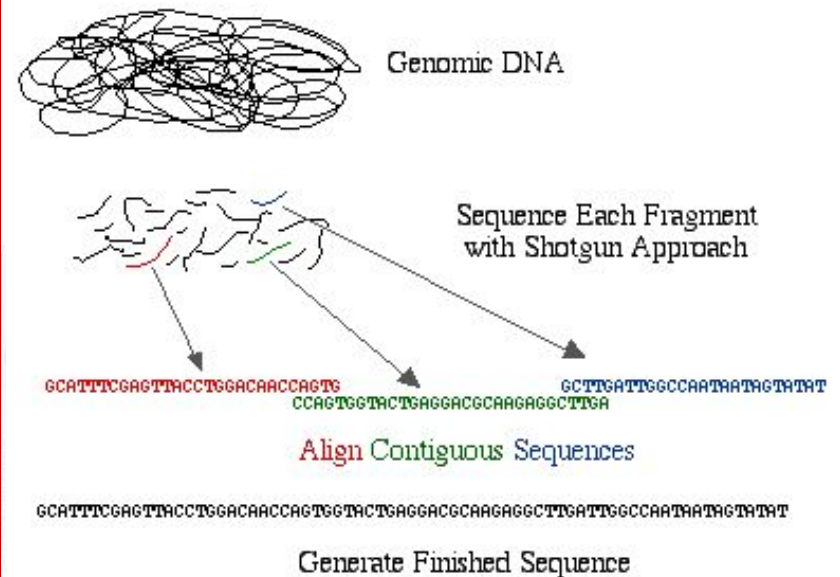
Diretor do “National Human Genome Research Institute”



Biblioteca Genômica



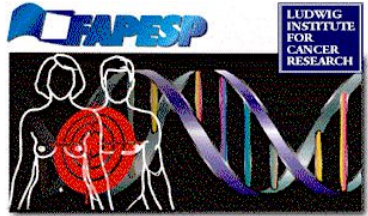
Whole Genome Shotgun Sequencing Method



Craig Venter
Co-fundador da "Celera Genomics Corporation"



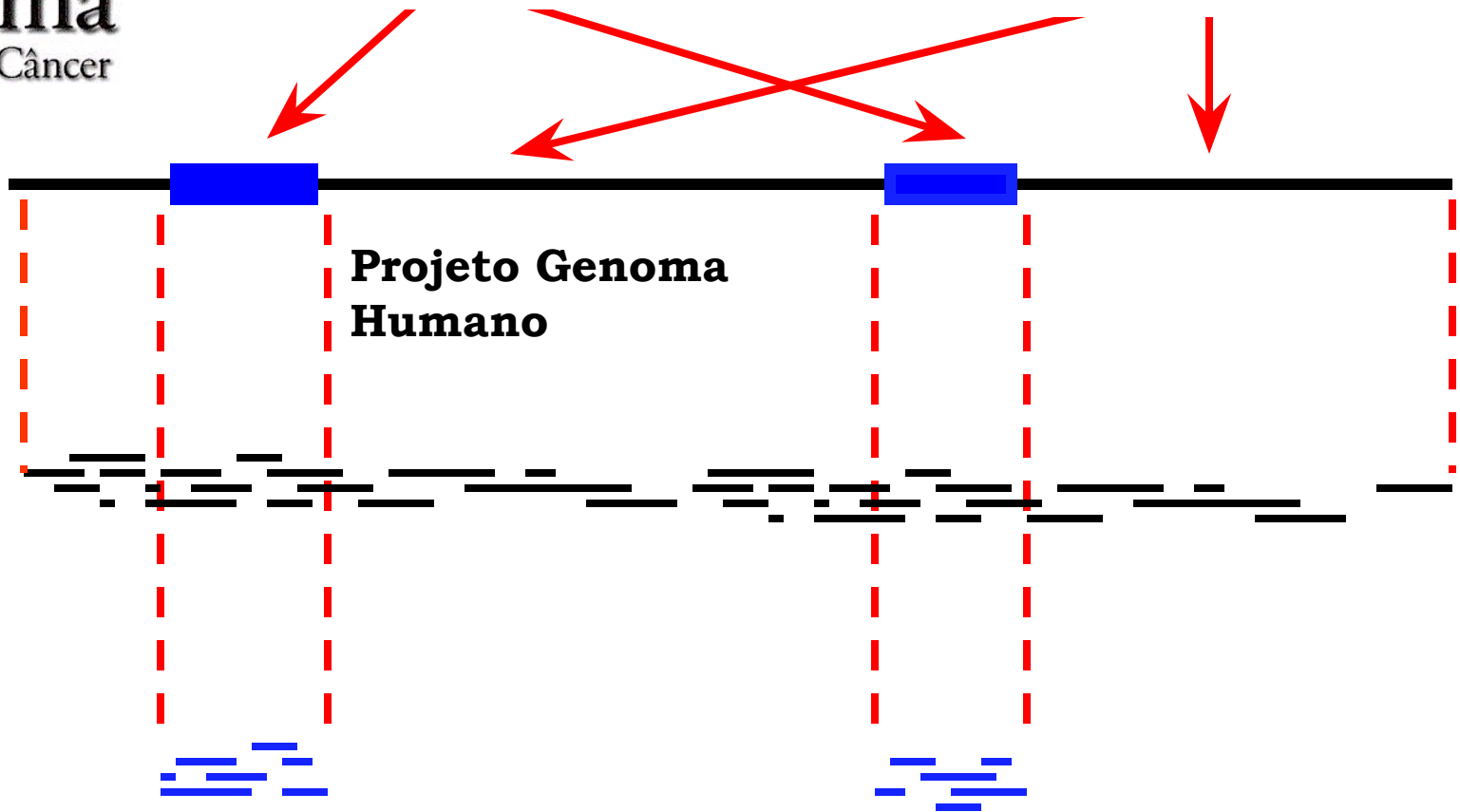
Biblioteca de cDNA



Genoma
Humano do Câncer

Genes = 3%

Não-codificante = 97%



Orestes FAPESP-Ludwig