

Lecture 9: Mapping a gene defined by the mutation

I. Classical mapping

II. Molecular mapping

III. Positional cloning

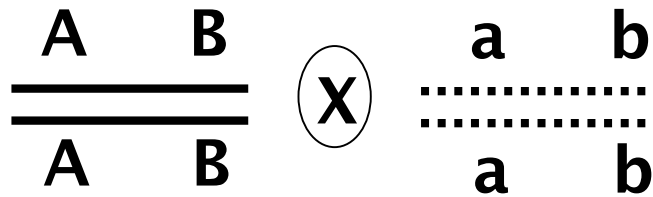
Read 387-398

Fig. 11.17; 11.19; 11.20; 11.21; 11.22

Self-reviewing of classical genetics

Read: 114-117; 123-127

Fig. 5.2; 5.3; 5.4; 5.5; 5.10; 5.12



F1:



F2:

A	B/a b		90
a	b/a b		90
A	b/a b		10
a	B/a b		10

$$\frac{Ab + aB}{\text{total}} = \text{rec, freq.}$$

$$\frac{10 + 10}{200} = 0.1 = 10\%$$

Fig. 5.2

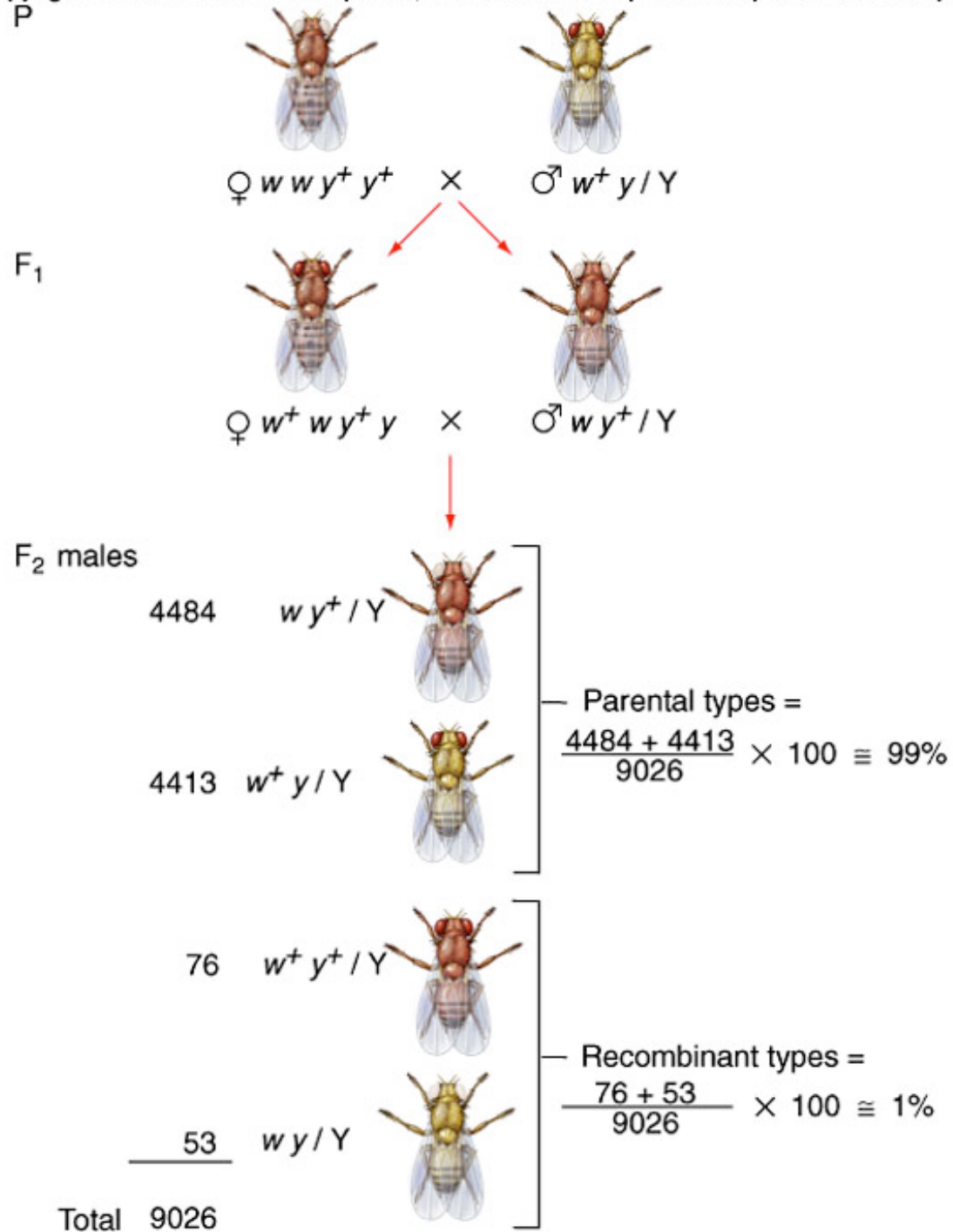


Fig. 5.4

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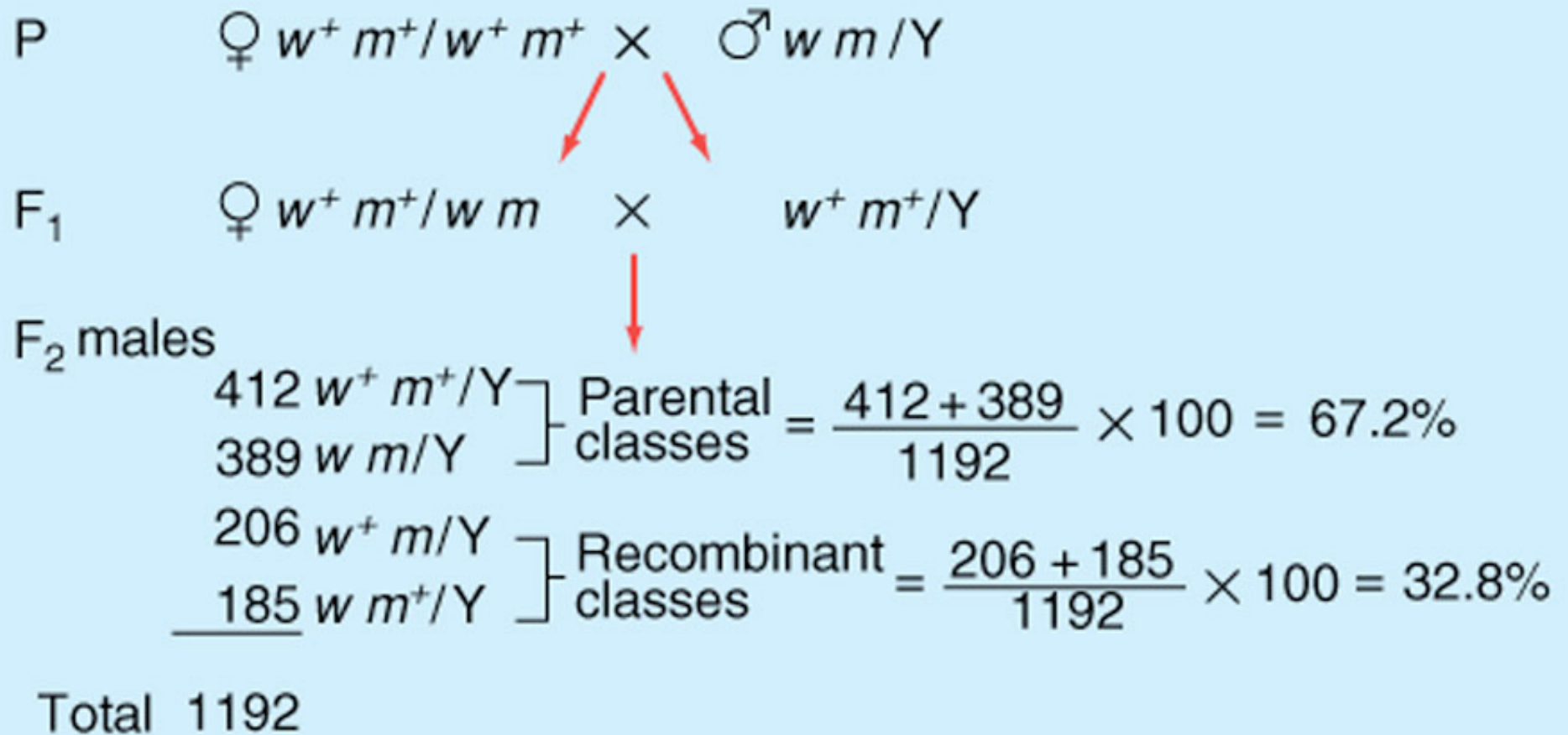


Fig. 5.5

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P $\text{♀ } b c^+ / b c^+ \times \text{♂ } b^+ c / b^+ c$

F₁ (all identical) $b c^+ / b^+ c$

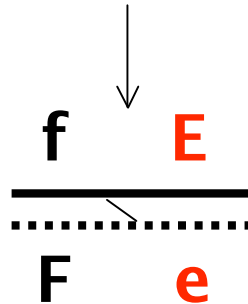
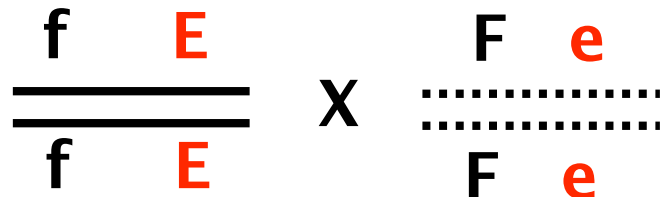
Testcross $\text{♀ } b c^+ / b^+ c \times \text{♂ } b c / b c$

Testcross progeny $\begin{array}{l} 2934 \text{ } b c^+ / b c \\ 2768 \text{ } b^+ c / b c \end{array} \left. \vphantom{\begin{array}{l} 2934 \\ 2768 \end{array}} \right\} \text{Parental classes} = \frac{2934 + 2768}{7419} \times 100 = 77\%$

$\begin{array}{l} 871 \text{ } b c / b c \\ \underline{846 \text{ } b^+ c^+ / b c} \end{array} \left. \vphantom{\begin{array}{l} 871 \\ 846 \end{array}} \right\} \text{Recombinant classes} = \frac{871 + 846}{7419} \times 100 = 23\%$

Total 7419

Mapping with a CAPS marker



E/e: CAPS marker
F/f: a mutation causing abnormal flowers
Ler: solid line
Col: dotted line

self

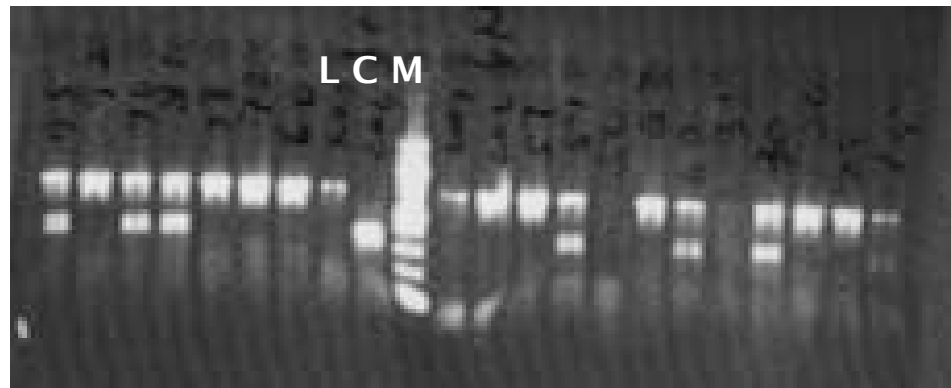
No. of plants

No. of
recombinant Chr.

11 $\hline \hline$ 0

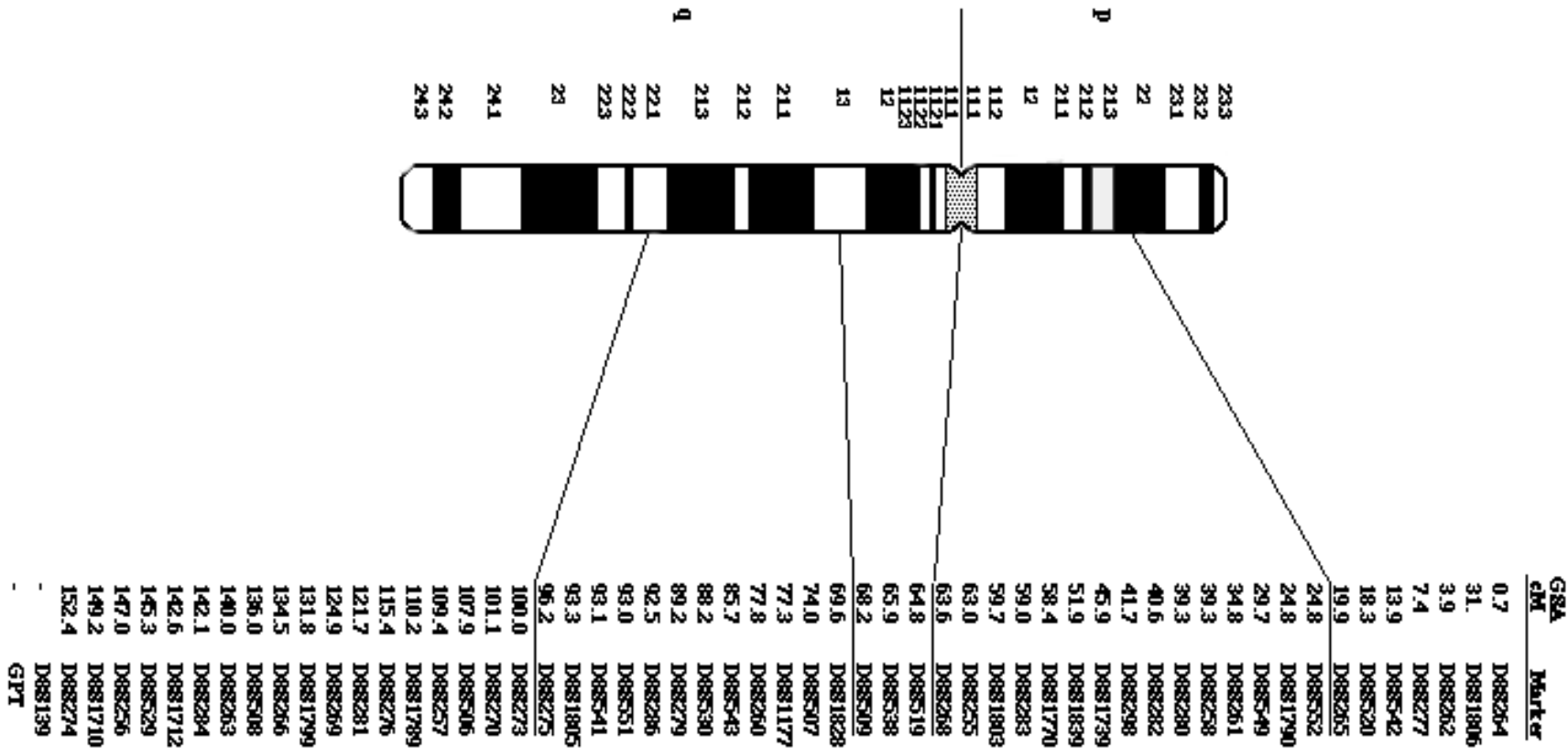
6 $\hline \hline \cdots$ 1

0 $\hline \hline \cdots$ 2

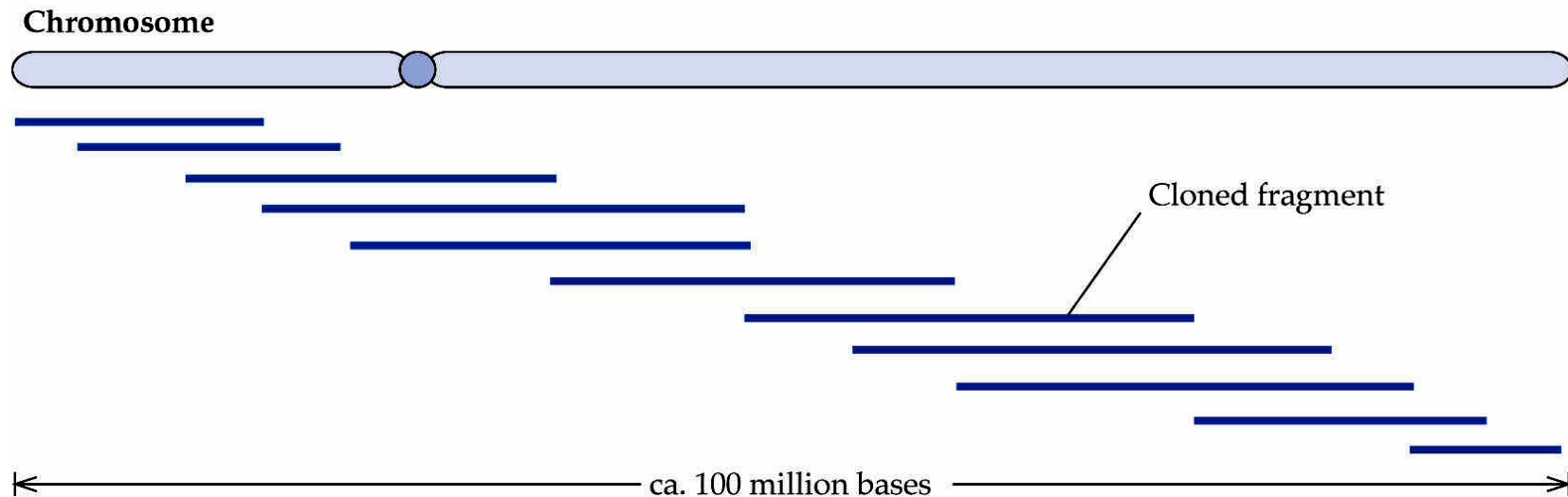


$$\frac{6}{17 \times 2} = 0.17$$

A genetic map of DNA markers

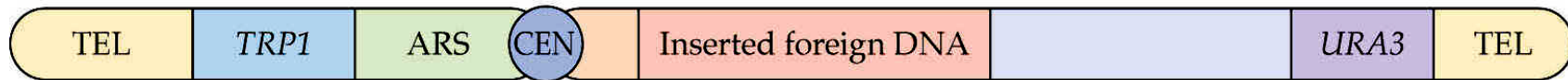


Physical map



In Arabidopsis: 1 cM = about 200 kb (50 genes)
In human 1cM = 1000 kb (~17 genes)

Yeast Artificial Chromosomes YAC



<u>Vector</u>	<u>insert size</u>	<u>Host</u>
YAC:	100-1000kb	yeast
BAC:	80-300kb	bacterium
Cosmid:	20-50 kb	bacterium
Lamda:	10-20kb	bacterium
Plasmid:	0.2-15kb	bacterium

A PHYSICAL MAP OF HUMAN CHROMOSOME 11

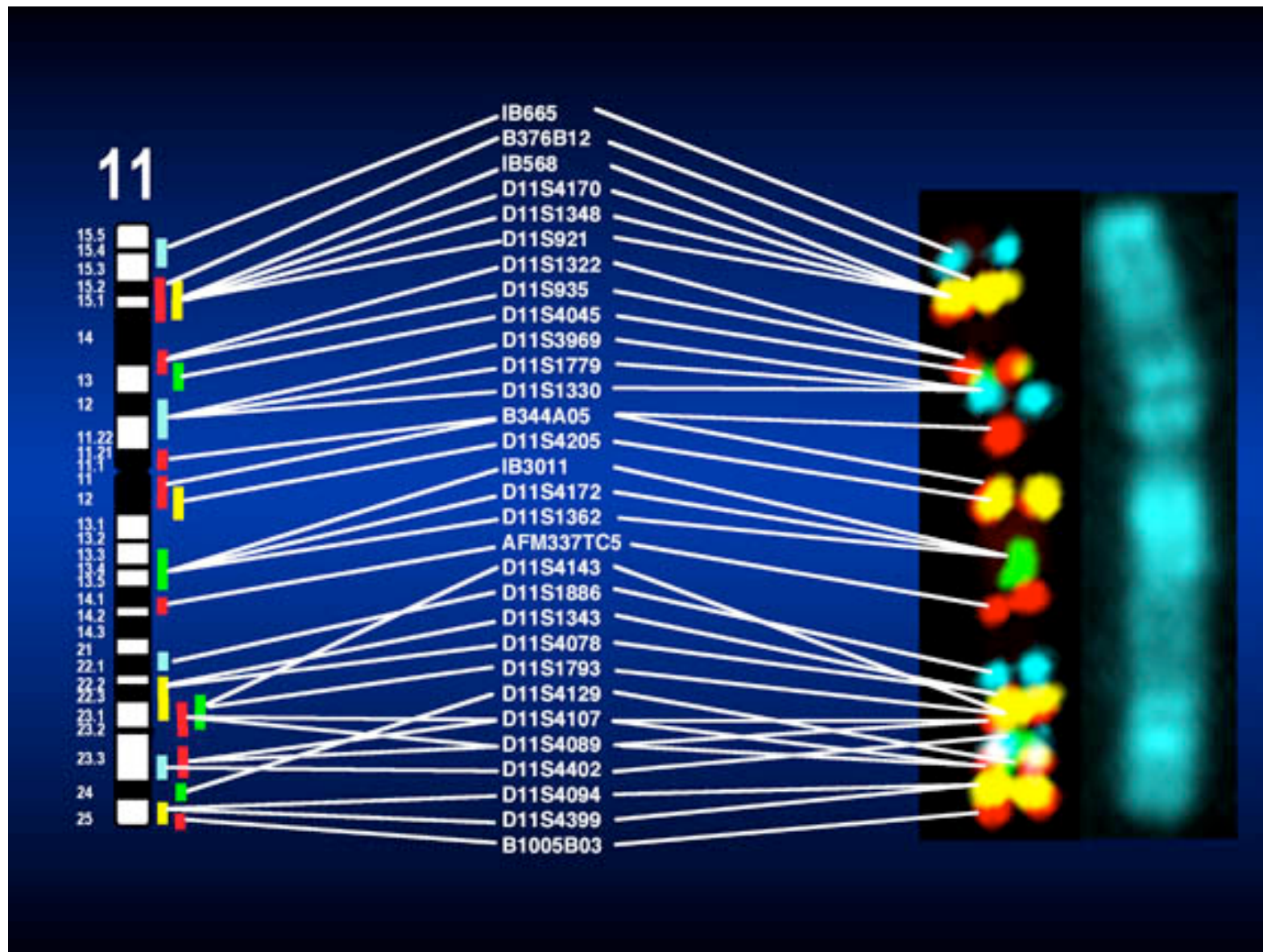
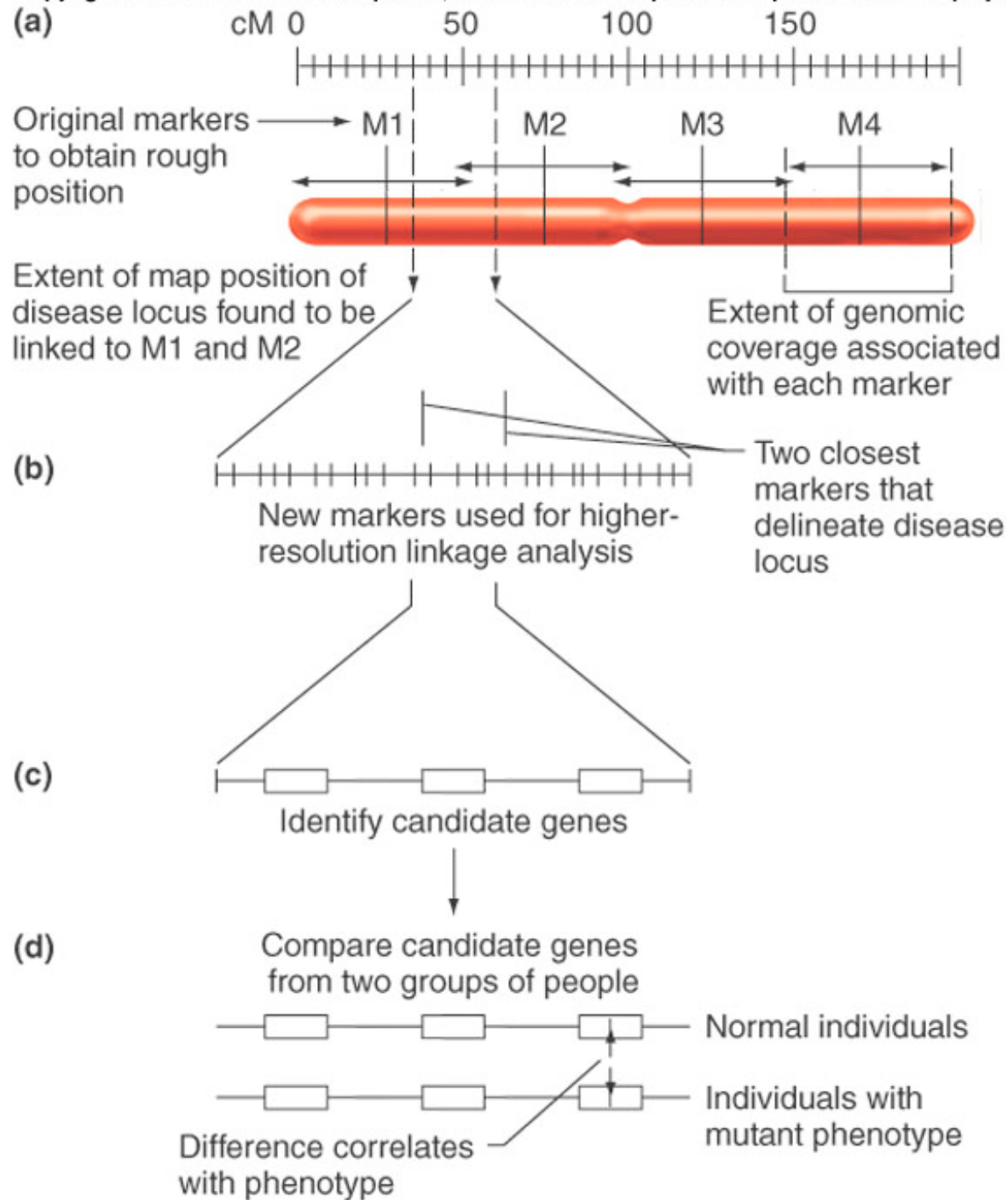


Fig. 11.17



Steps in identifying the gene in a chromosome region

Step1: Categorize genes in the region

Step2: Determine mRNA expression pattern

EST

Northern hybridization

Step3: Determine changes between wt and mut

Expression pattern

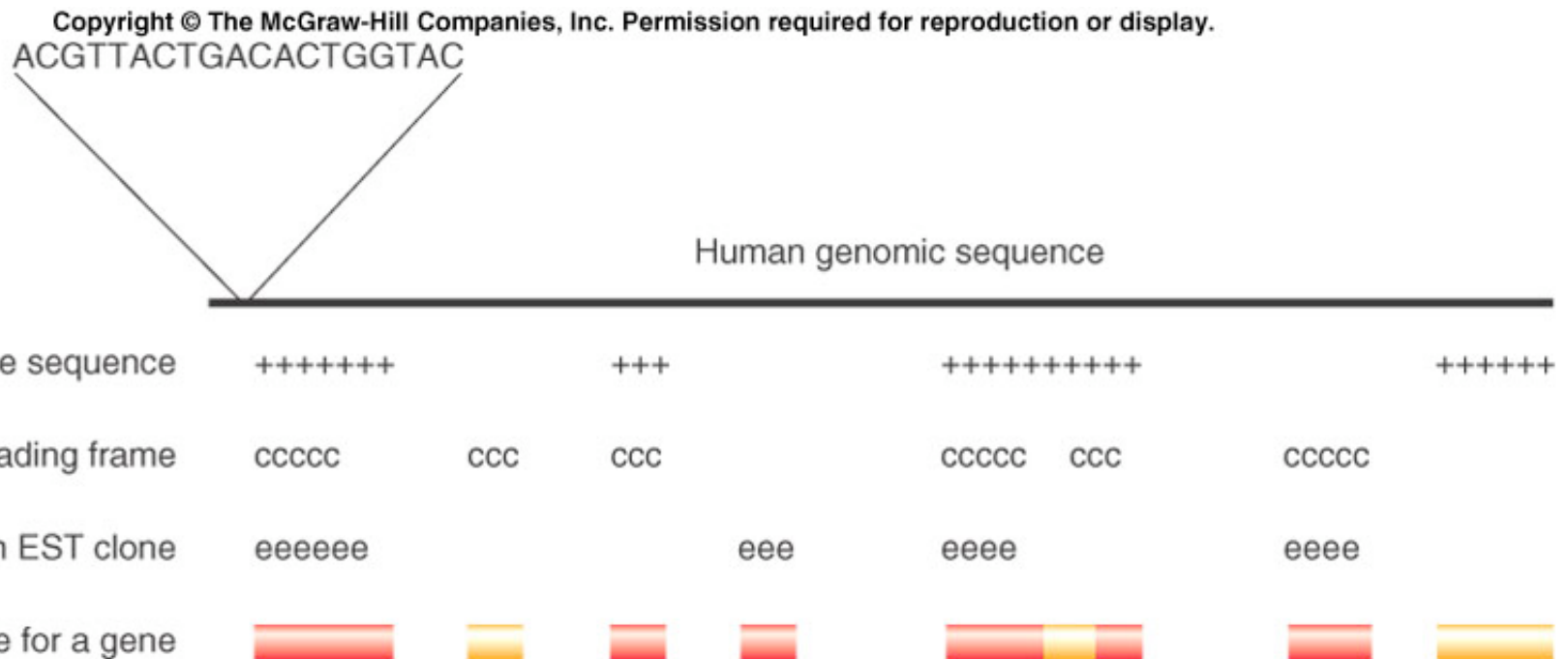
DNA sequence

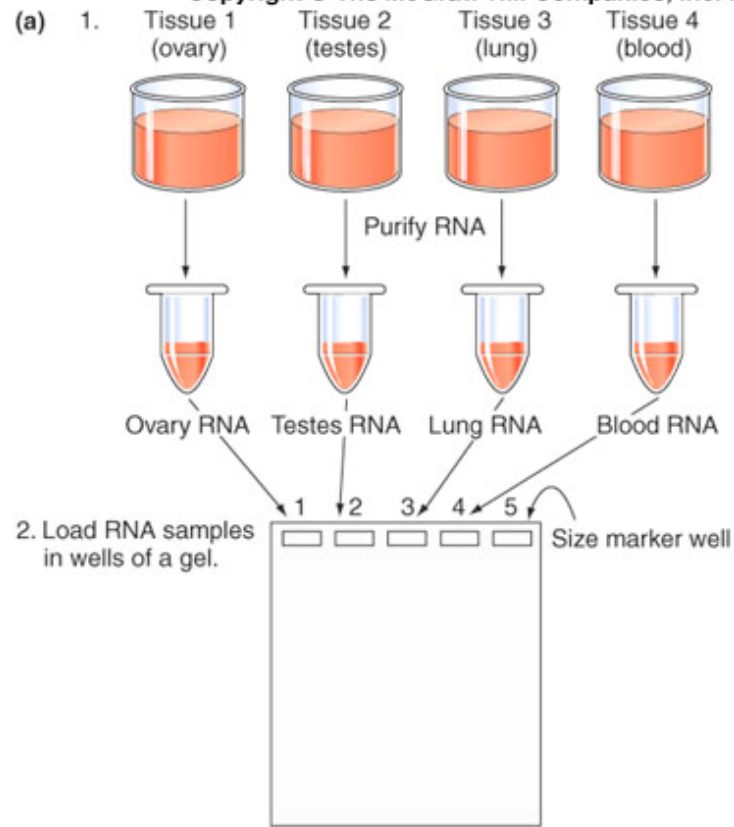
Step4: Transgenic studies (SRY example)

Case study: Cystic fibrosis

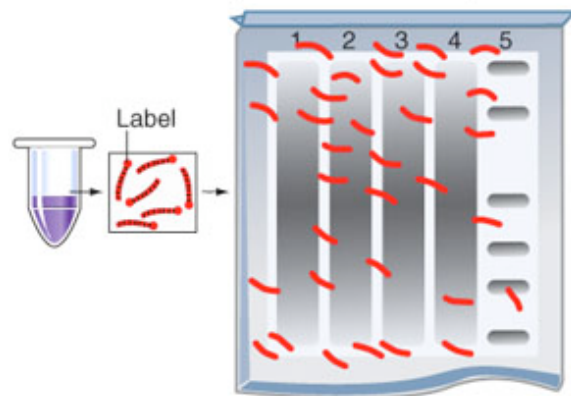
CFTR (Cystic fibrosis transmembrane conductance regulator)

Fig. 11.19

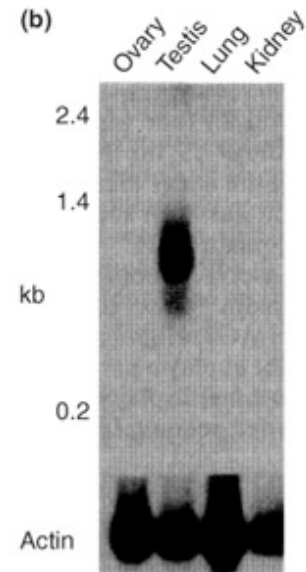
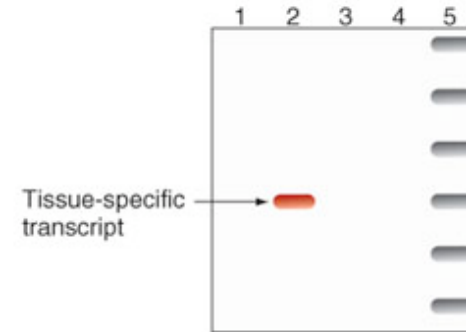




3. Separate RNA samples by gel electrophoresis.
Blot onto filter. Expose filter to labeled hybridization probe.



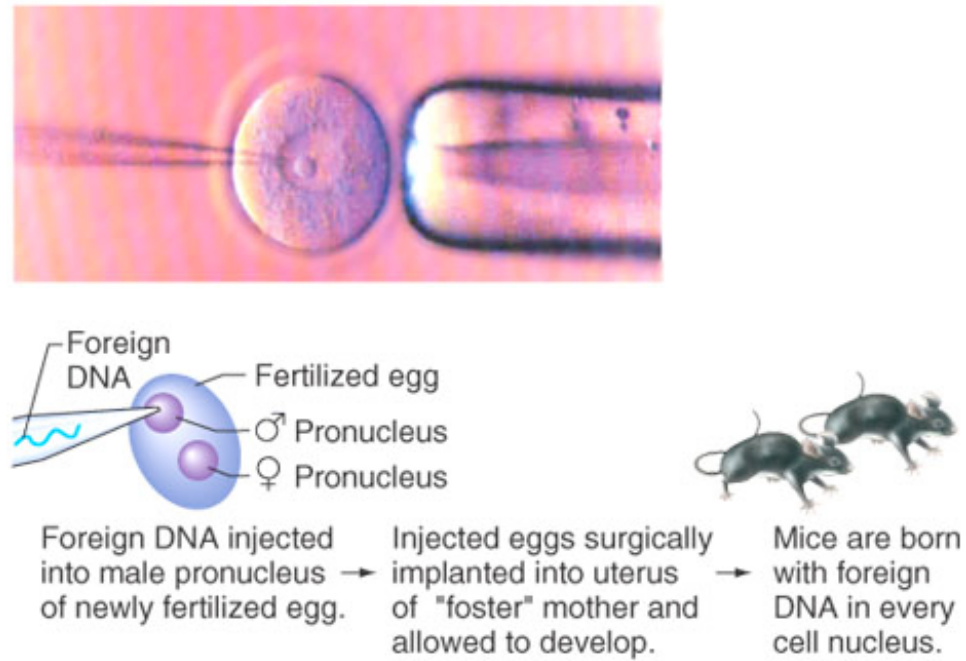
4. Wash away unhybridized probe. Make autoradiograph.



Testes-determining factor (TDF)

Fig. 11.20

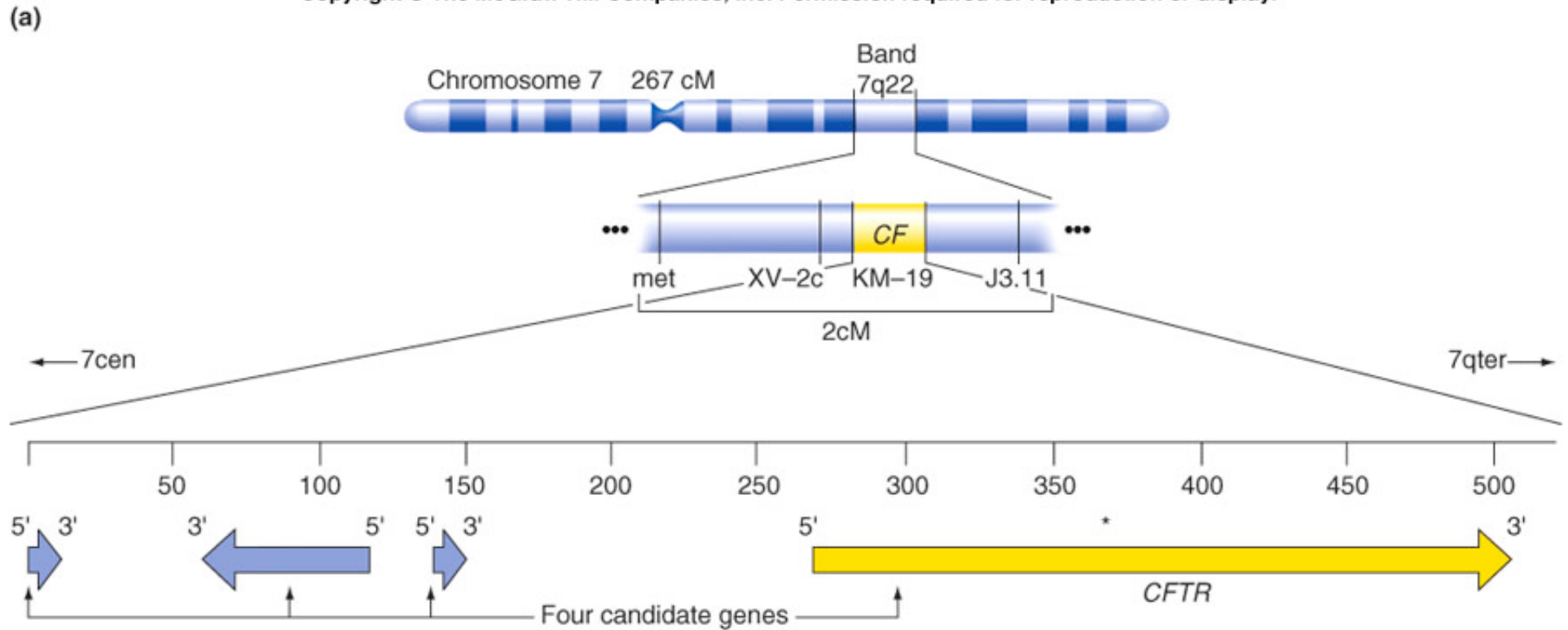
Fig. 11.21



Cloning cystic fibrosis transmembrane conductance regulator (CFTR)

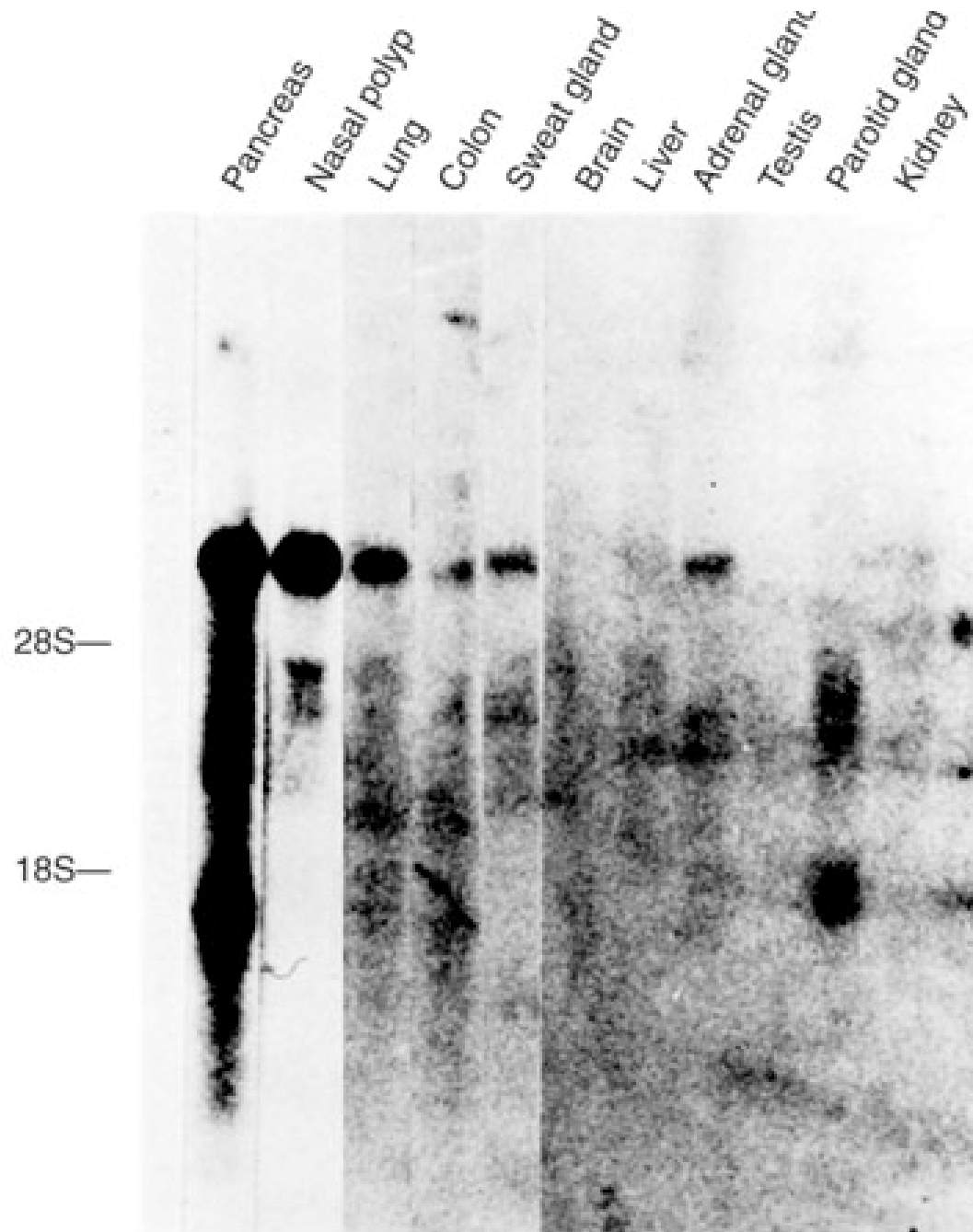
Fig. 11.22a

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Northern blots help eliminate other 3 candidate genes

Fig. 11.22b



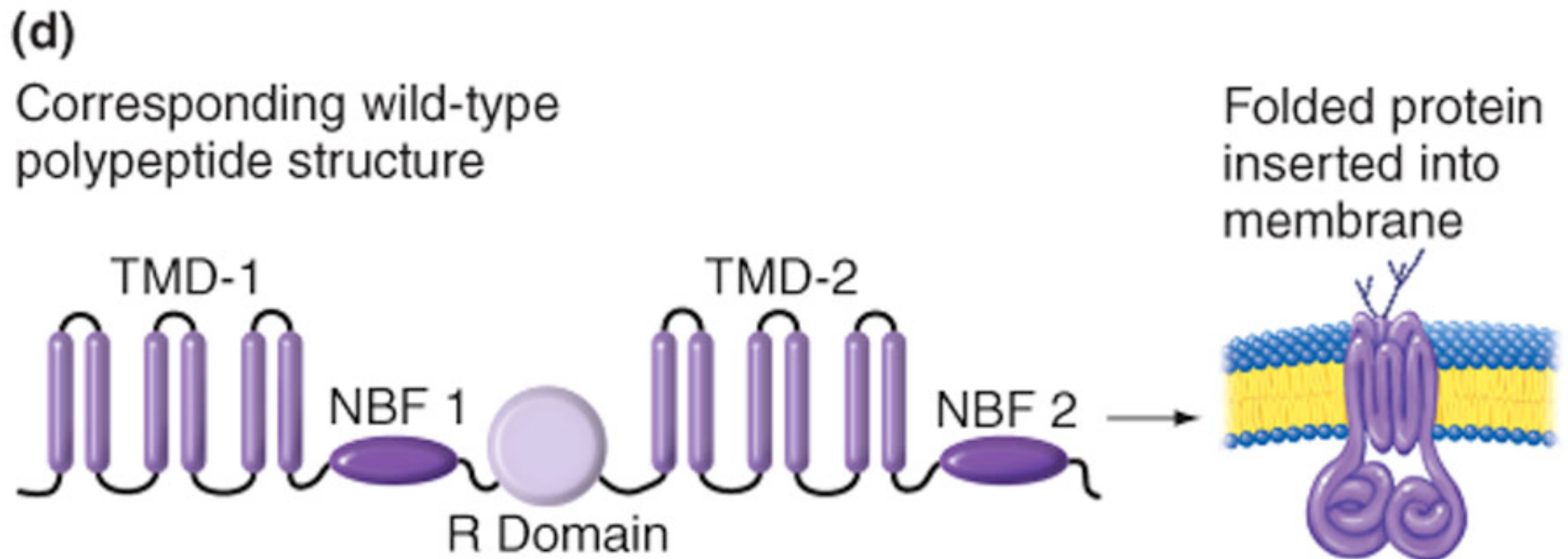
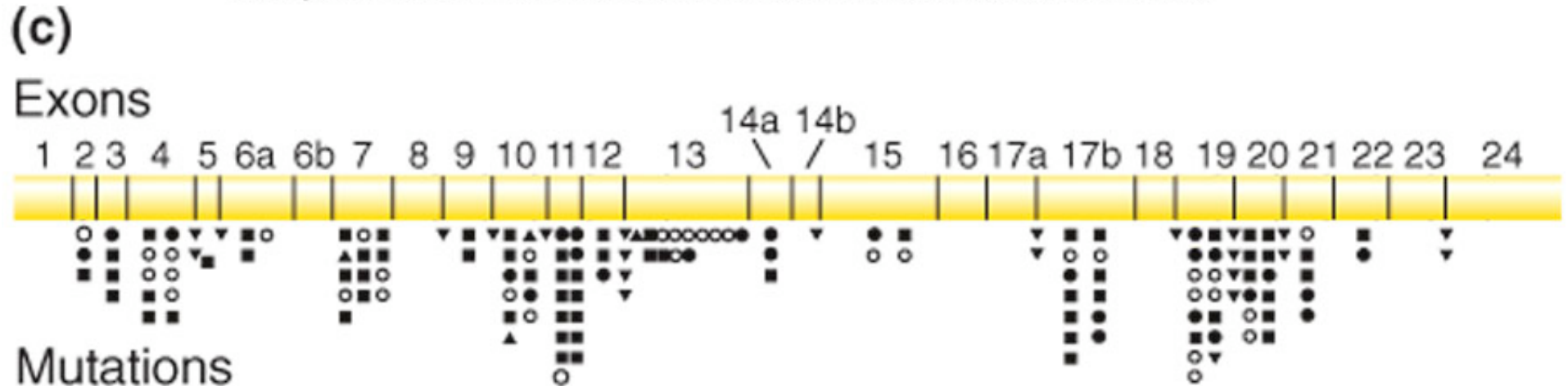


Fig. 11.22cd

CFTR

