

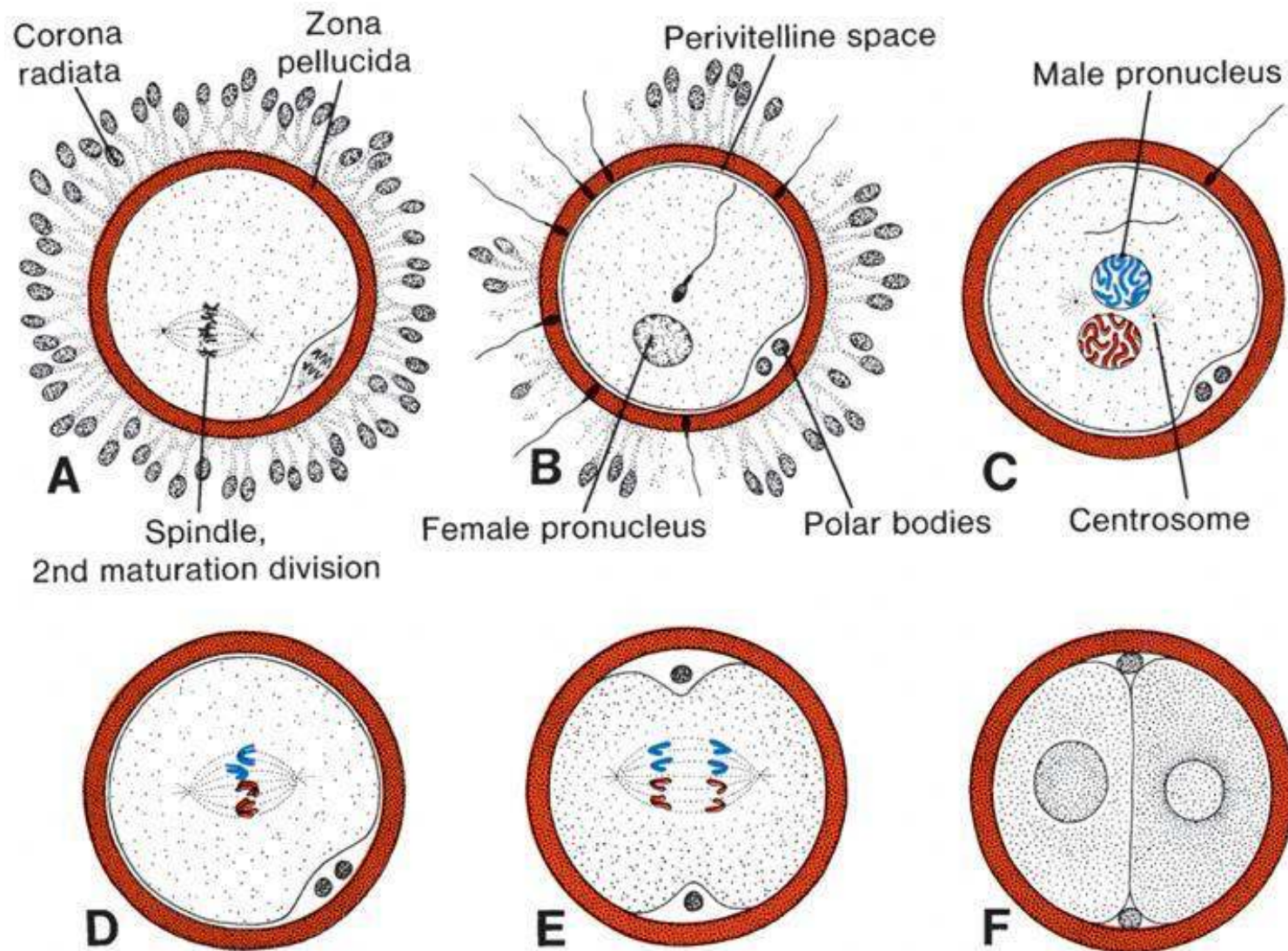


Developmental genetics

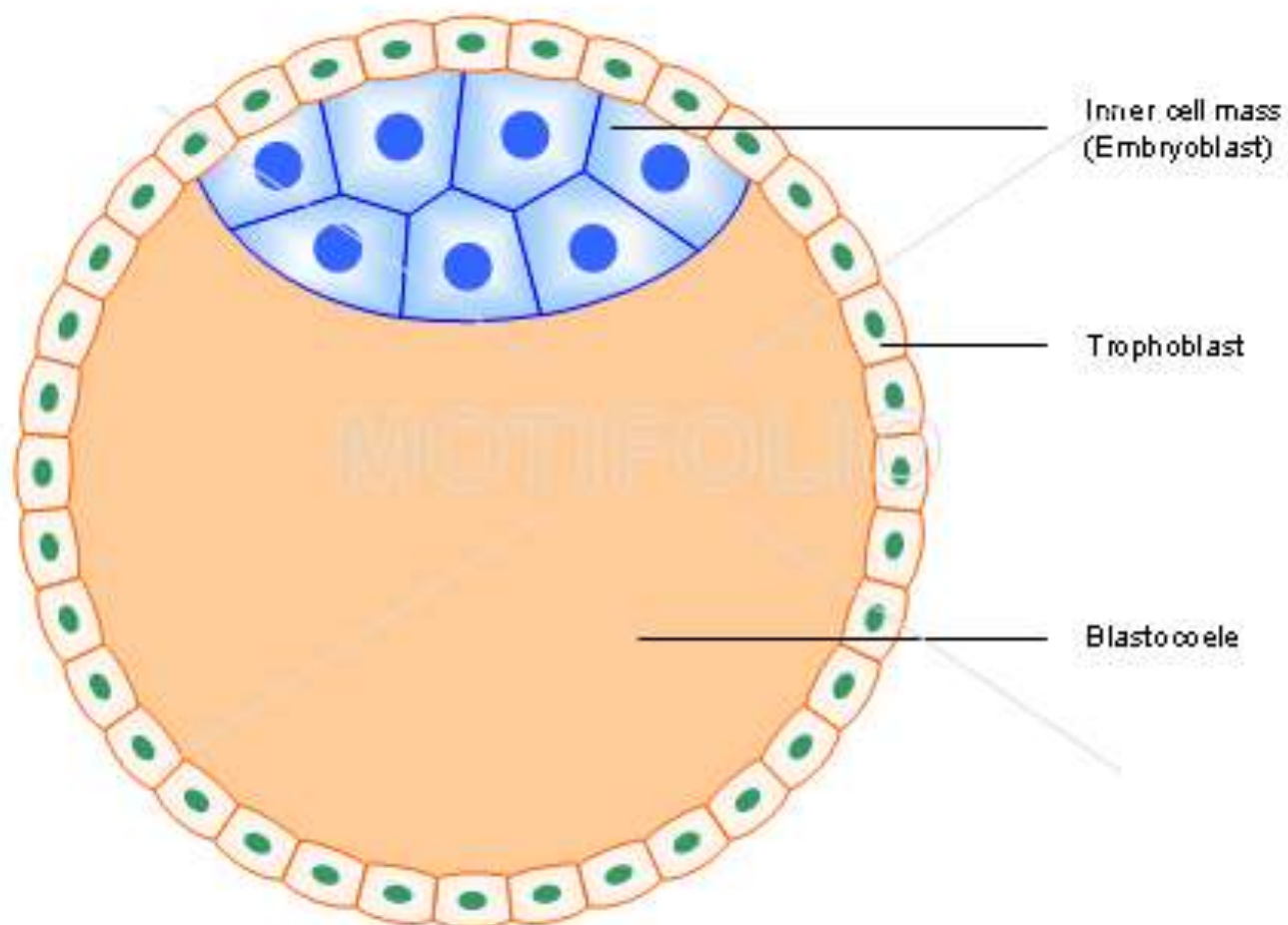
chapter 6

2018.
Molecular genetics

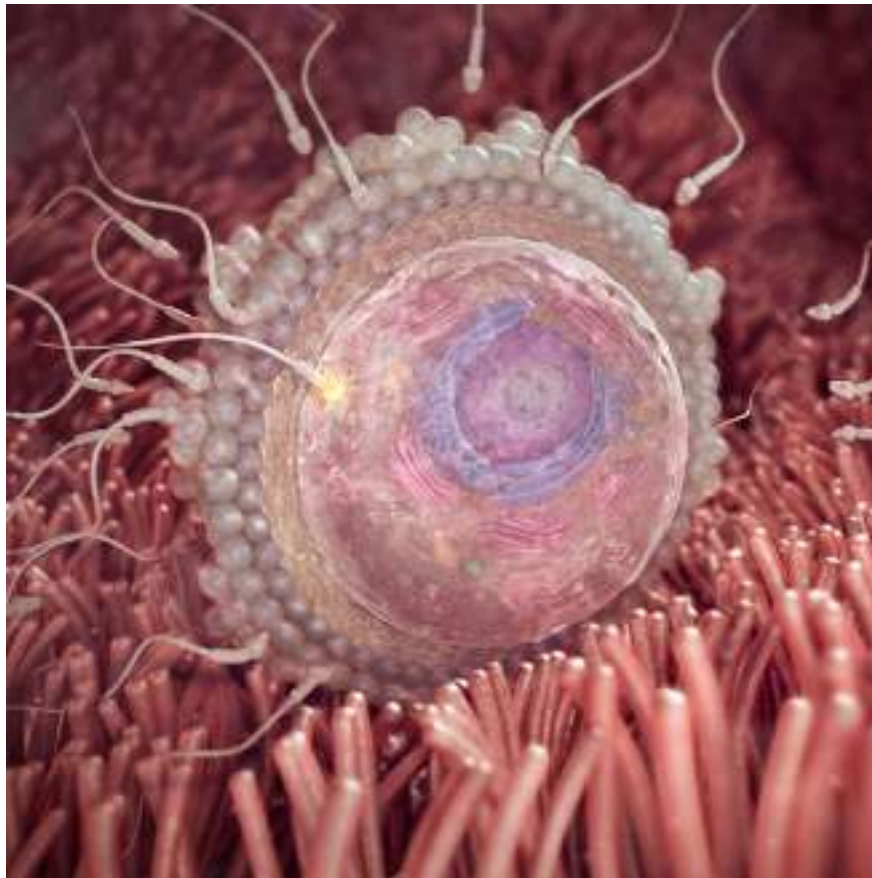
(Emery's Elements of medical genetics, 14th ed. Turnpenny P. and Ellard S.
Elsevier 2012.)



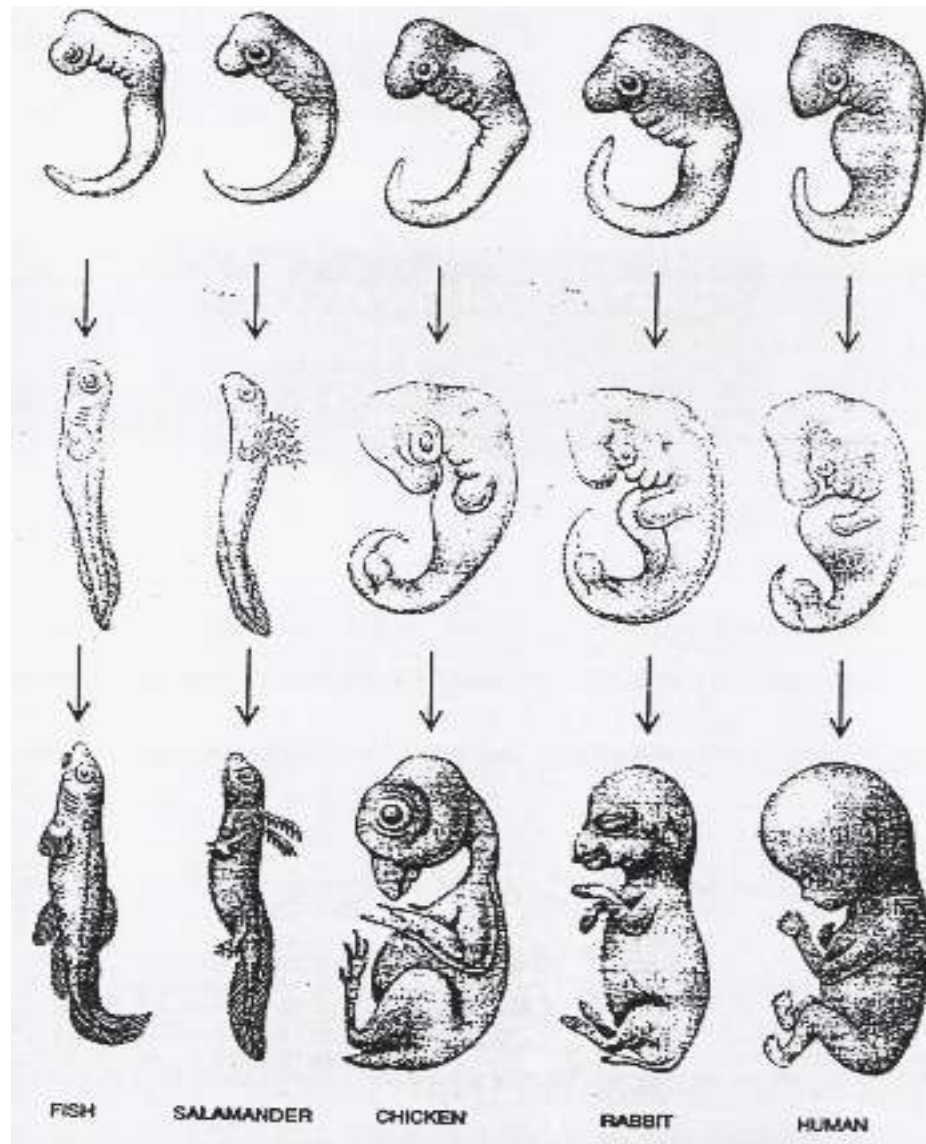
Blastocyst

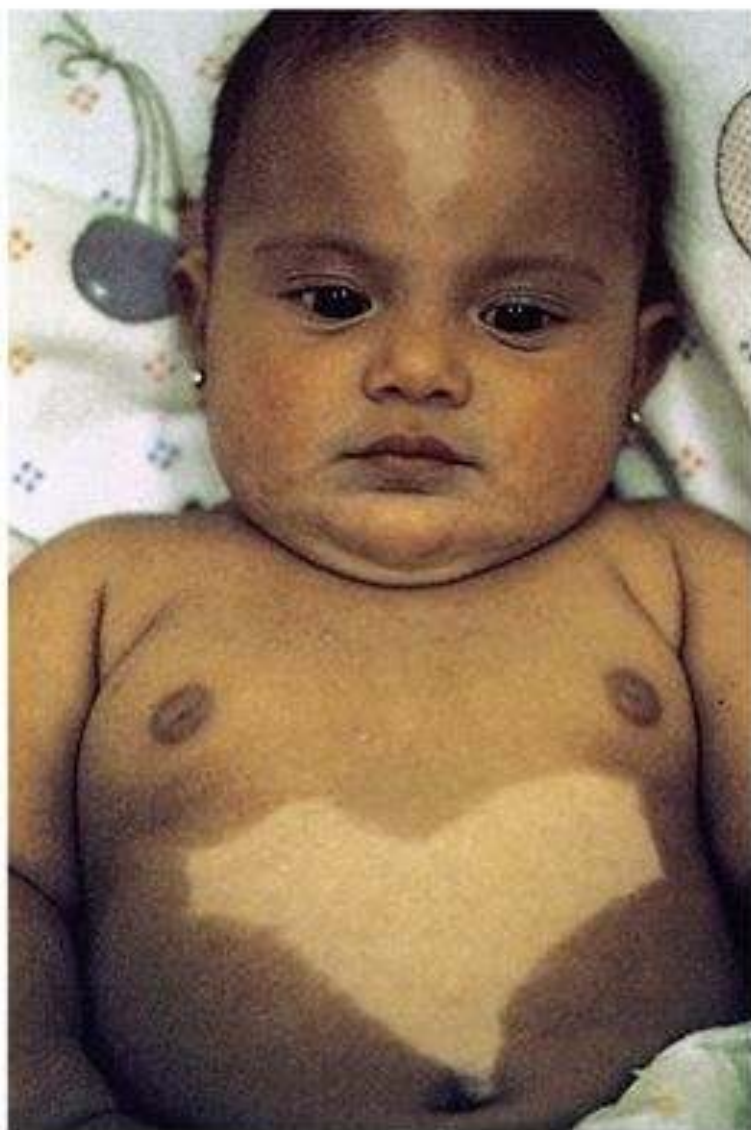


Development of human embryo



| Stage | Time from Conception | Length of Embryo/Fetus |
|---|----------------------|------------------------|
| Pre-embryonic | | |
| First cell division | 30 h | |
| Zygote reaches uterine cavity | 4 d | |
| Implantation | 5–6 d | |
| Formation of bilaminar disc | 12 d | 0.2 mm |
| Lyonization in female | 16 d | |
| Formation of trilaminar disc and primitive streak | 19 d | 1 mm |
| Embryonic stage | | |
| Organogenesis | 4–8 w | |
| Brain and spinal cord are forming, and first signs of heart and limb buds | 4 w | 4 mm |
| Brain, eyes, heart and limbs developing rapidly, and bowel and lungs beginning to develop | 6 w | 17 mm |
| Digits have appeared. Ears, kidneys, liver and muscle are developing | 8 w | 4 cm |
| Palate closes and joints form | 10 w | 6 cm |
| Sexual differentiation almost complete | 12 w | 9 cm |
| Fetal stage | | |
| Fetal movements felt | 16–18 w | 20 cm |
| Eyelids open. Fetus is now viable with specialized care | 24–26 w | 35 cm |
| Rapid weight gain due to growth and accumulation of fat as lungs mature | 28–38 w | 40–50 cm |





Trilaminar disc

Organ and Tissue Origins

Ectodermal

Central nervous system
Peripheral nervous system
Epidermis, including hair and nails
Subcutaneous glands
Dental enamel

Mesodermal

Connective tissue
Cartilage and bone
Smooth and striated muscle
Cardiovascular system
Urogenital system

Endodermal

Thymus and thyroid
Gastrointestinal system
Liver and pancreas

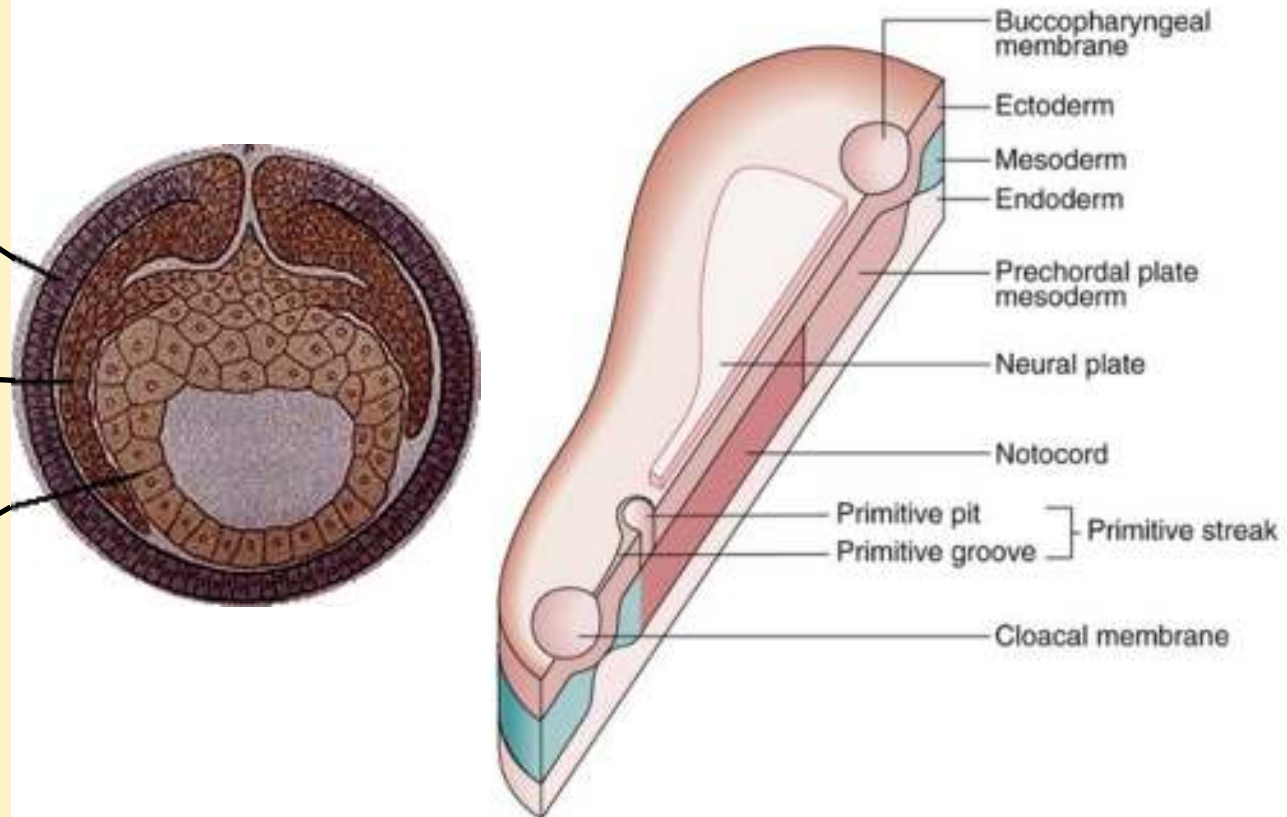
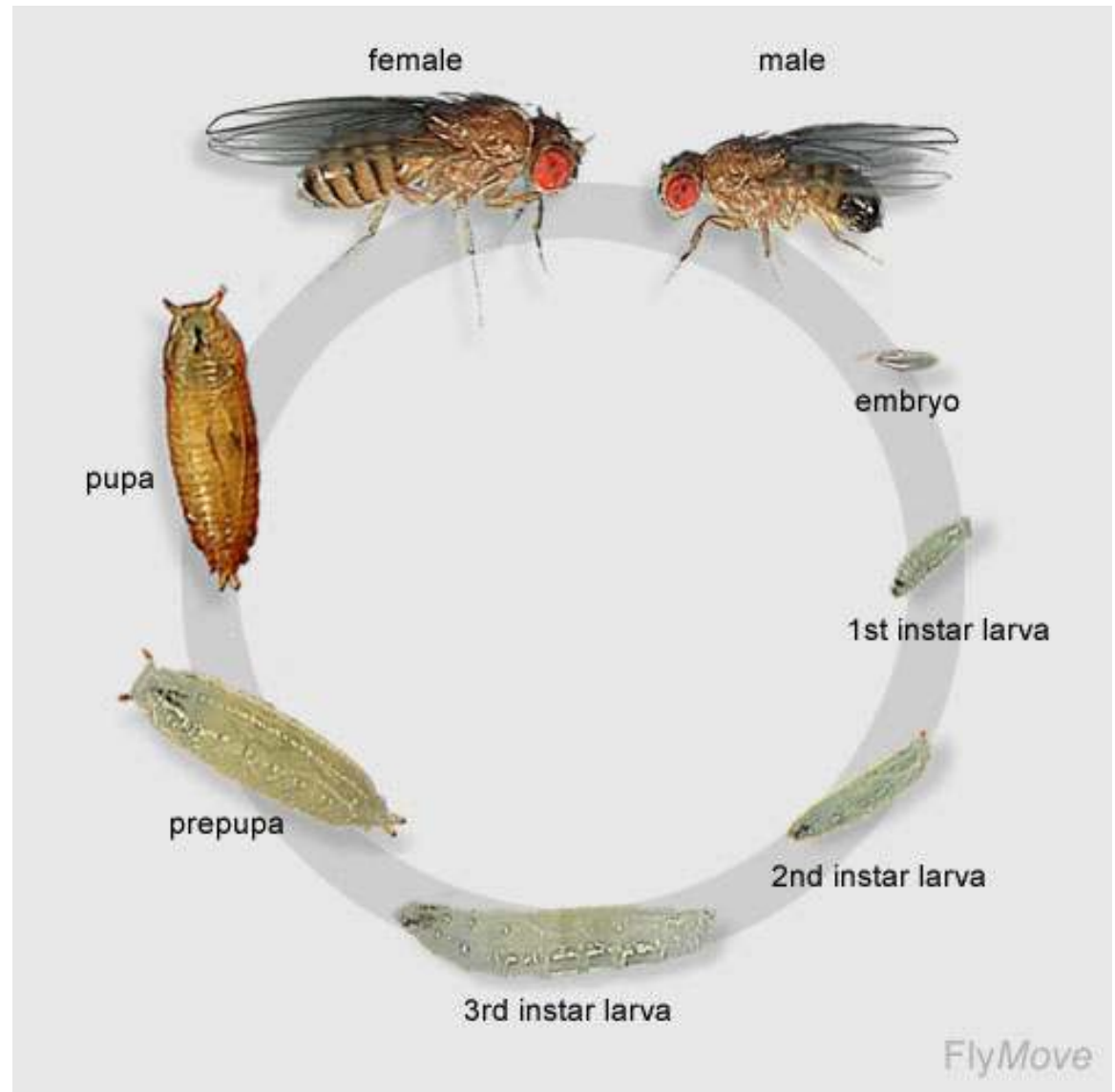


FIGURE 6.1 A schematic trilaminar disc, sectioned along the rostrocaudal axis.

Fruit fly life cycle

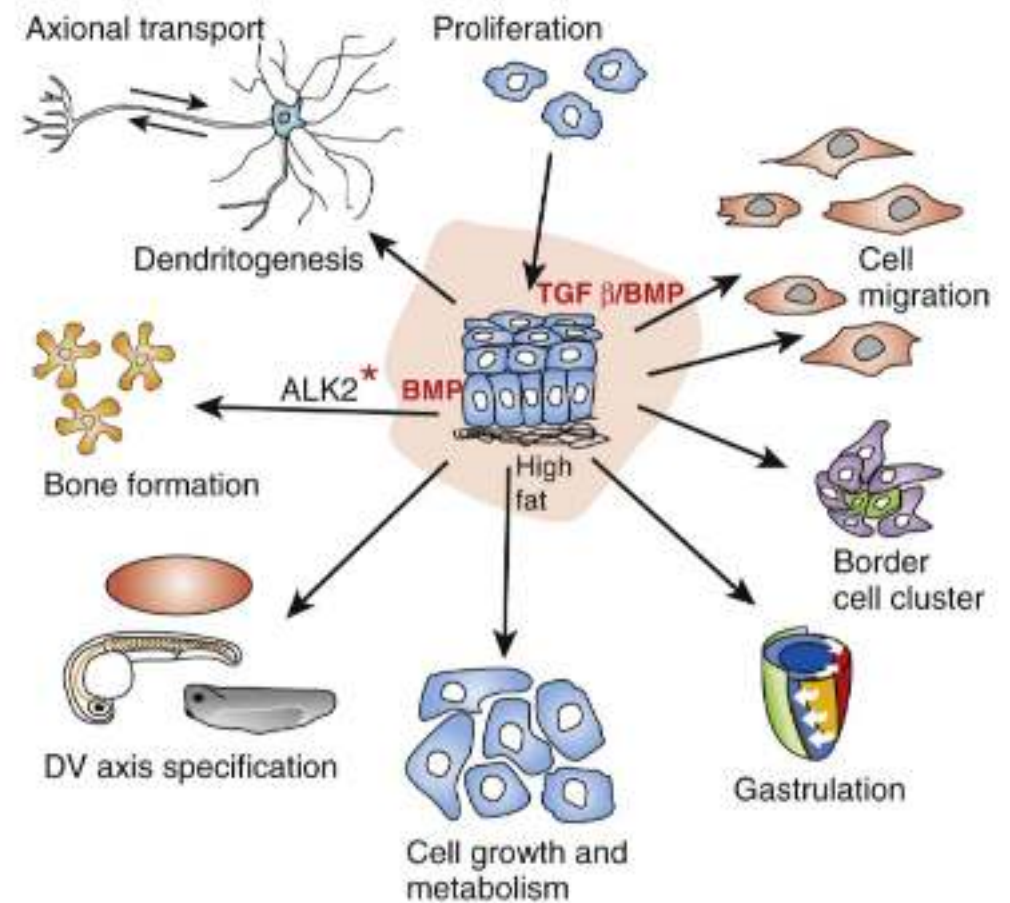
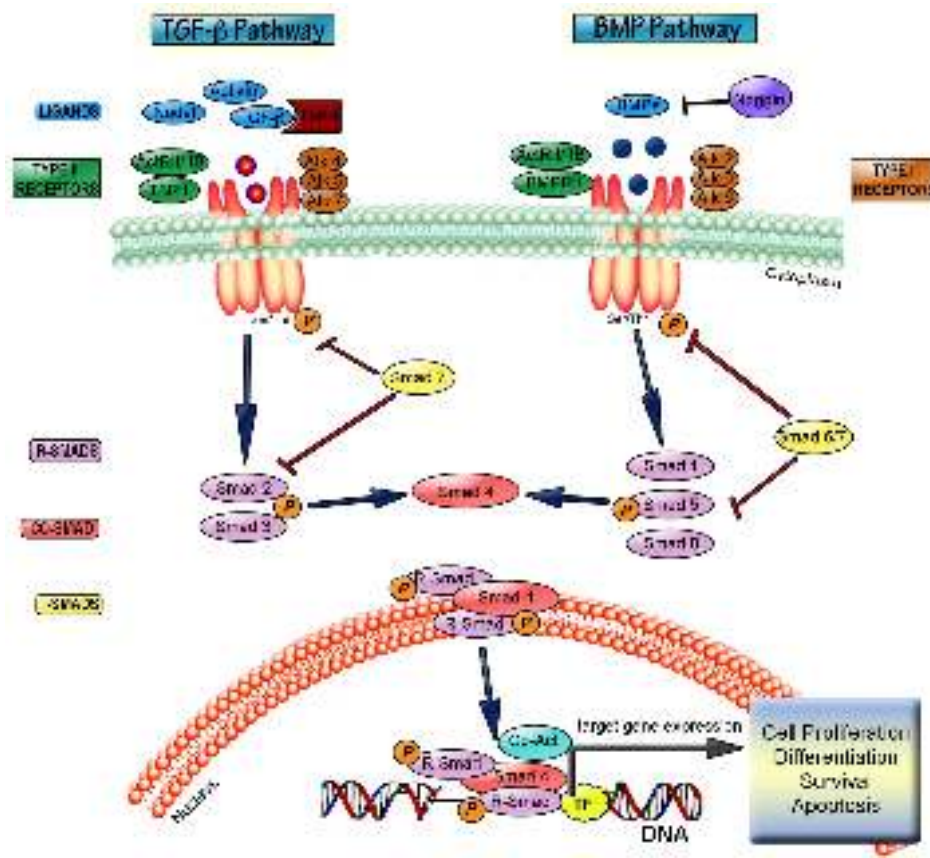


Developmental gene families in humans

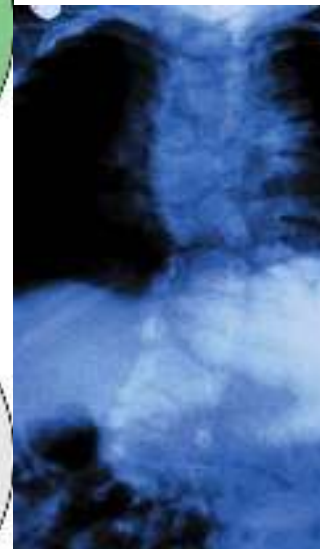
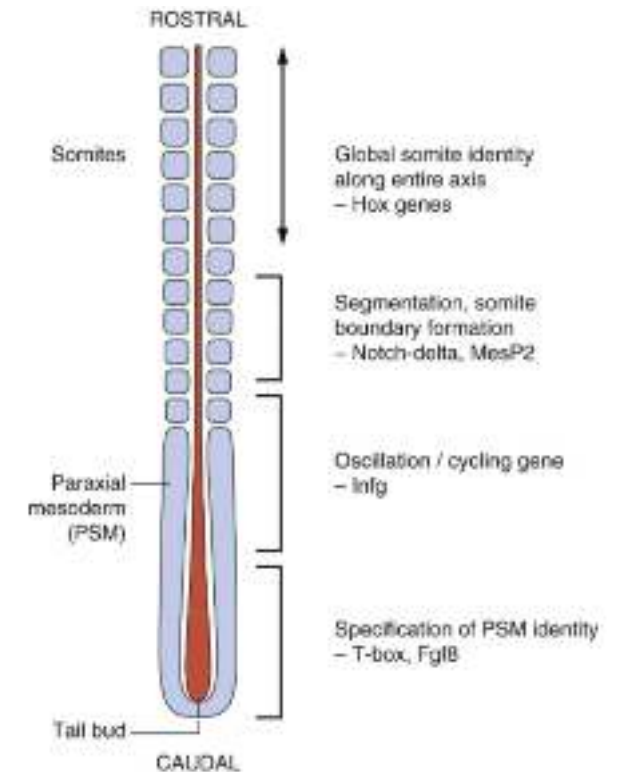
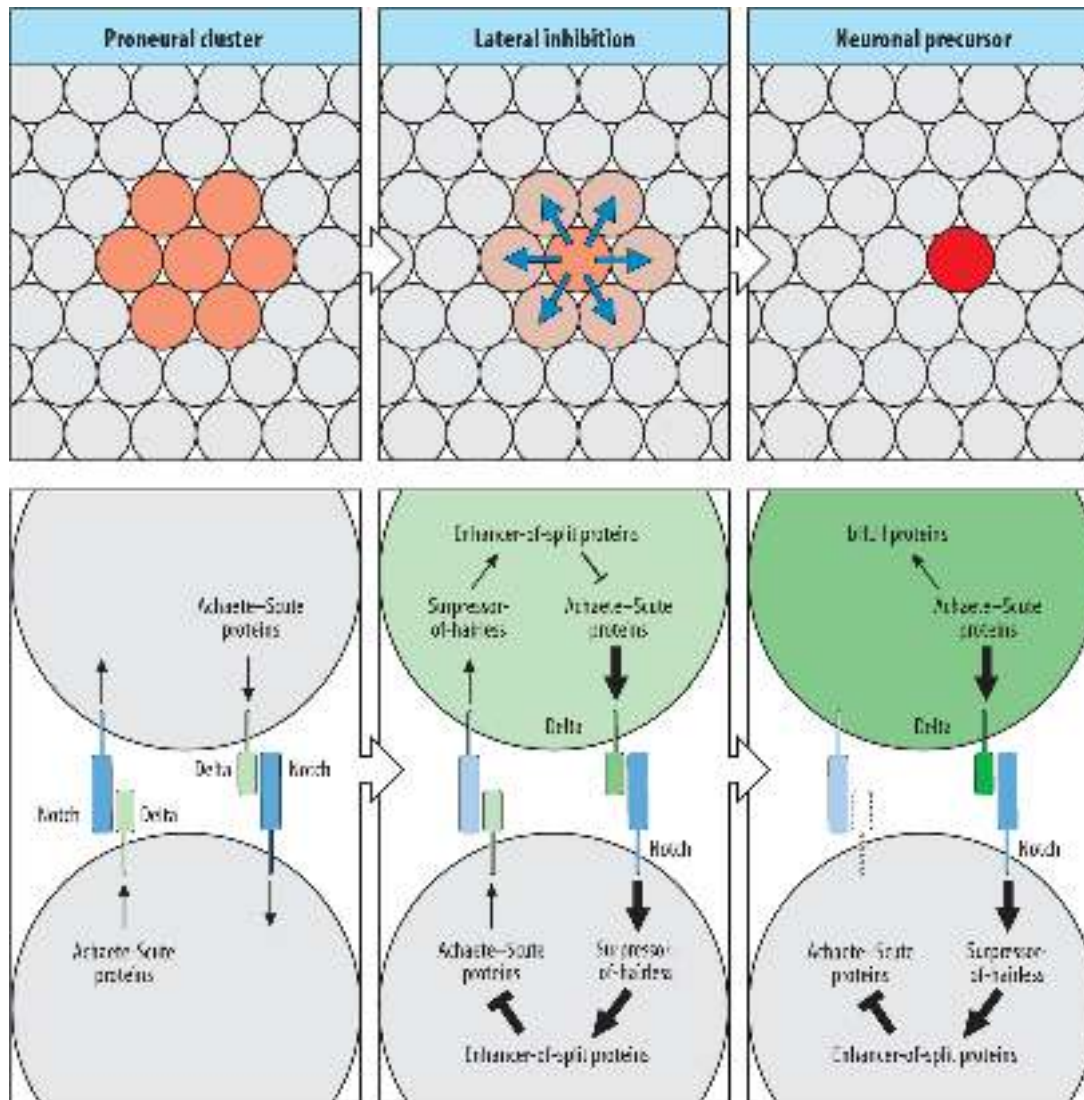
Transcription factors, transcription regulators, growth factors, cell receptors, morphogenes, cytokines...

- TGF- β superfamily of cytokines (TGF β ; BMP - SMAD)
- Wingless (Wnt)
- Hedgehog (HH), Patched, GLI - (brain)
- Nodal, FGFs, BMPs
- Notch-delta (somits)
- Homeobox
- PAX
- SOX (High mobility group) T-box genes
- Zinc finger genes,
- RET protooncogene,
- FGF receptors,.....

TGF- β genes (33 members from cytokine family) / BMP



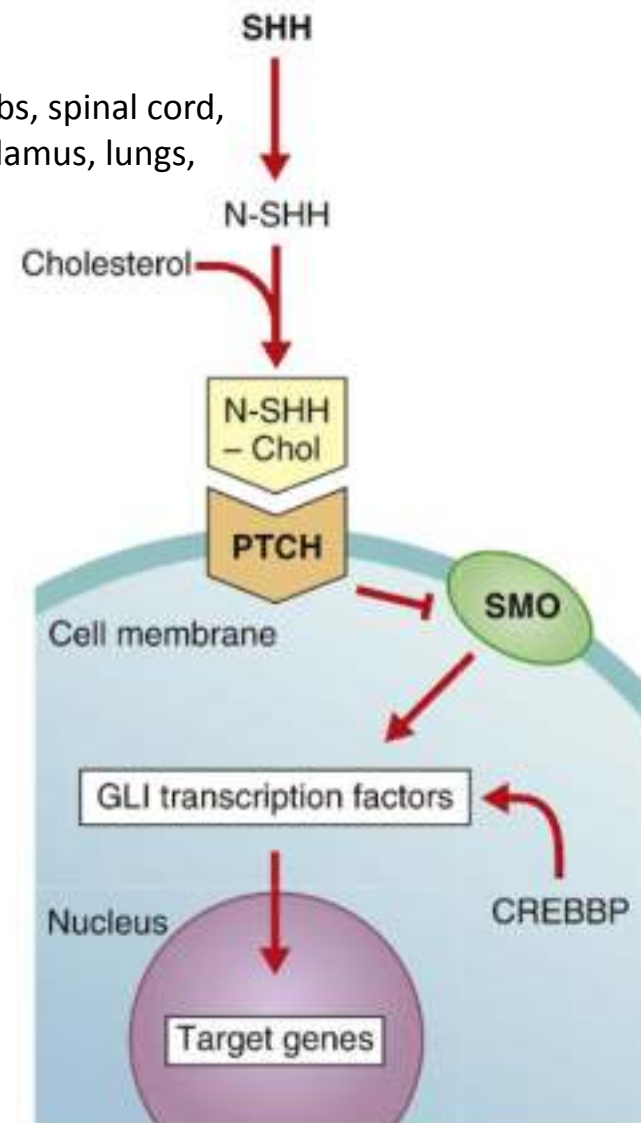
Delta-Notch signaling



Sonic hedgehog / Patched GLI

(Brain, limbs, spinal cord, axons, thalamus, lungs, teeth)

Holoprosencephaly typ3



Holoprosencephaly

Smith-Lemli-Opitz syndrome

Gorlin (nevoid basal cell carcinoma) syndrome

Some basal cell carcinomas and medulloblastomas

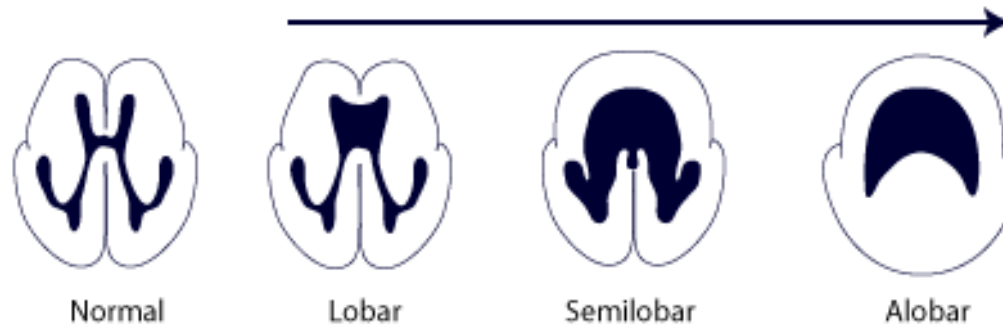
Greig cephalopolysyndactyly syndrome
Pallister-Hall syndrome

Rubenstein-Taybi syndrome

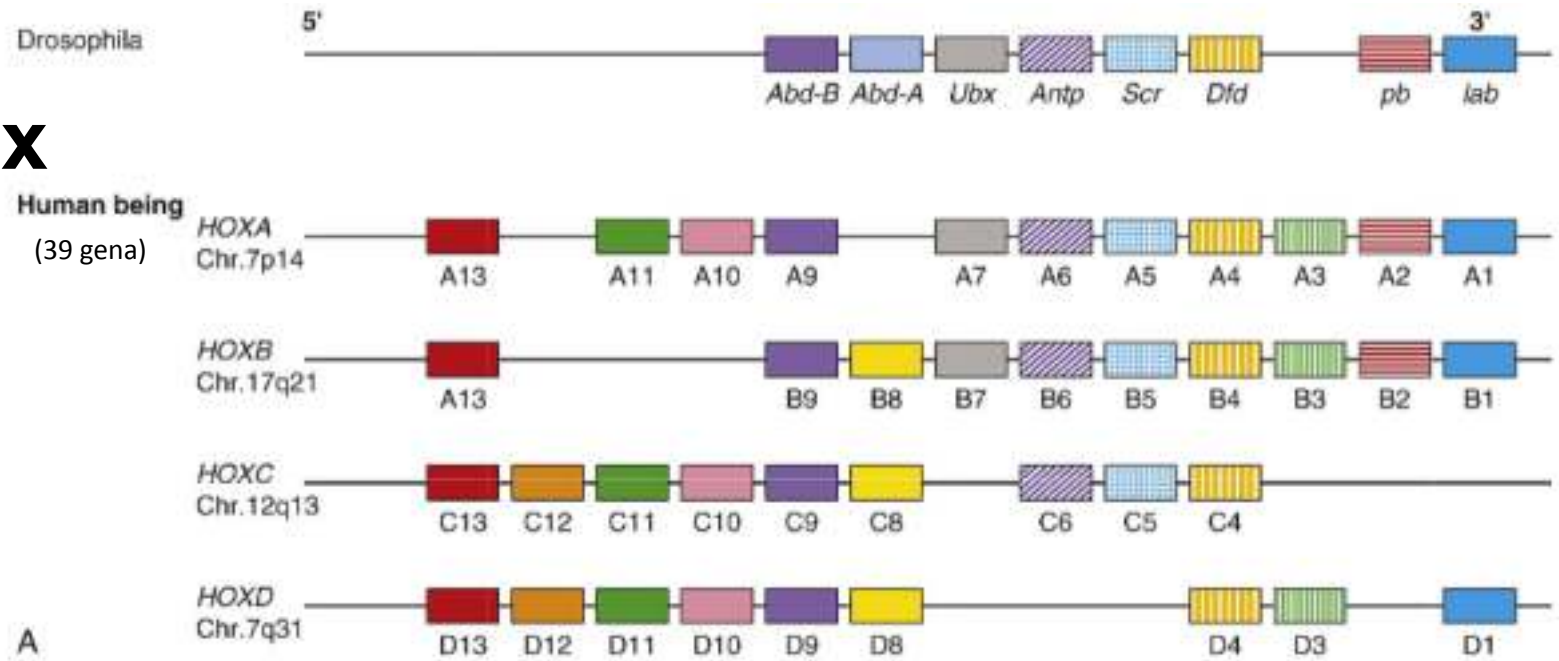
Holoprosencephaly (HH), Patched, GLI)



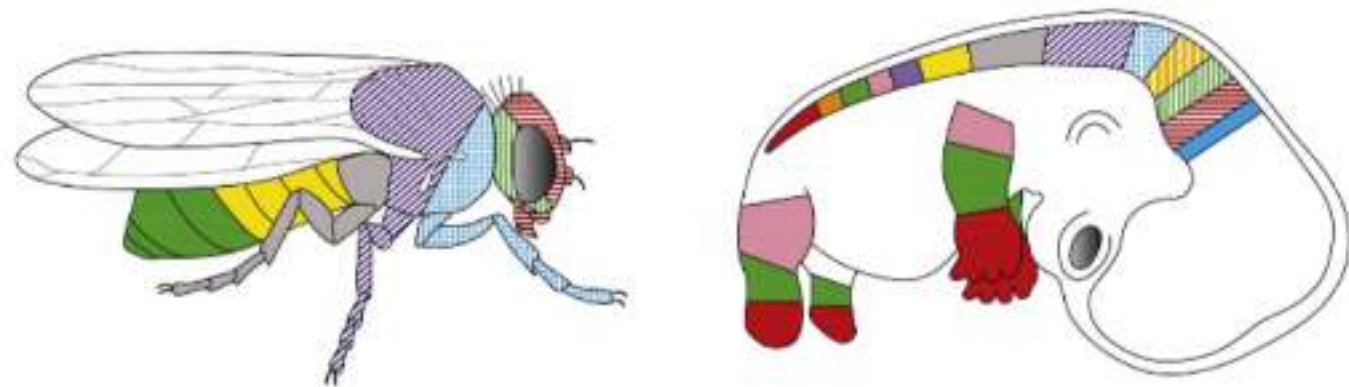
Holoprosencephaly: degree of severity



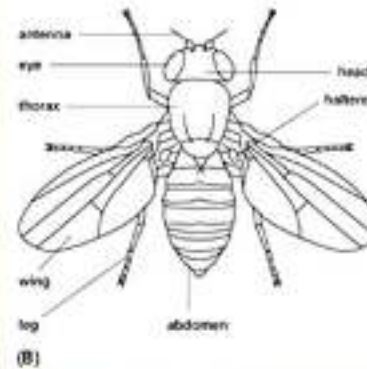
Homeobox genes



A



Homeotic mutations (egz.)



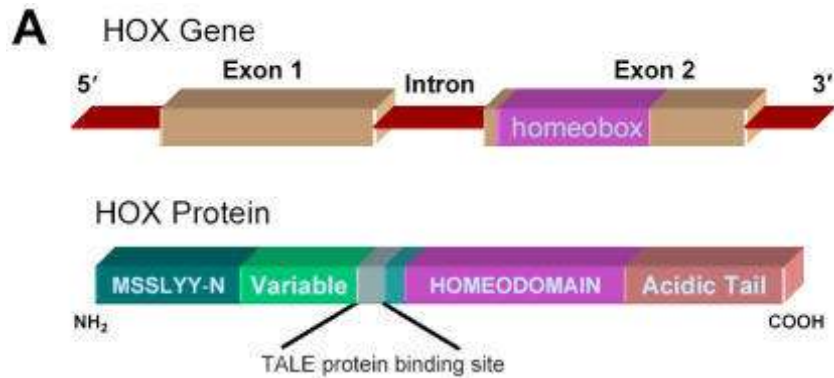
Normal adult fly



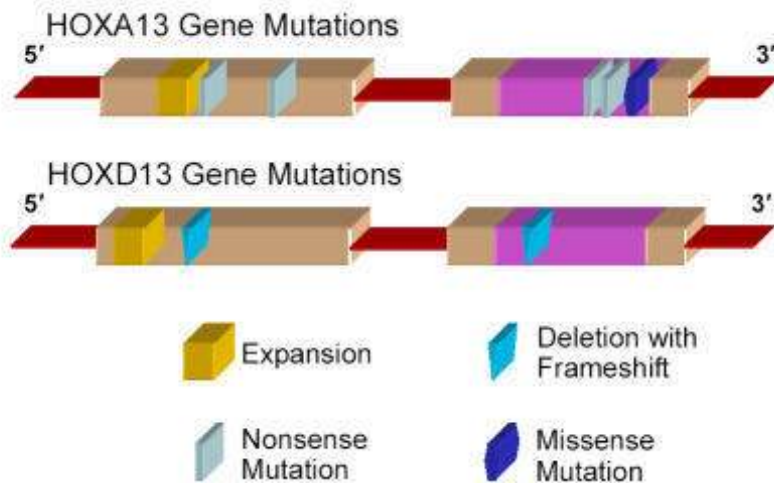
Antennapedia
mutant



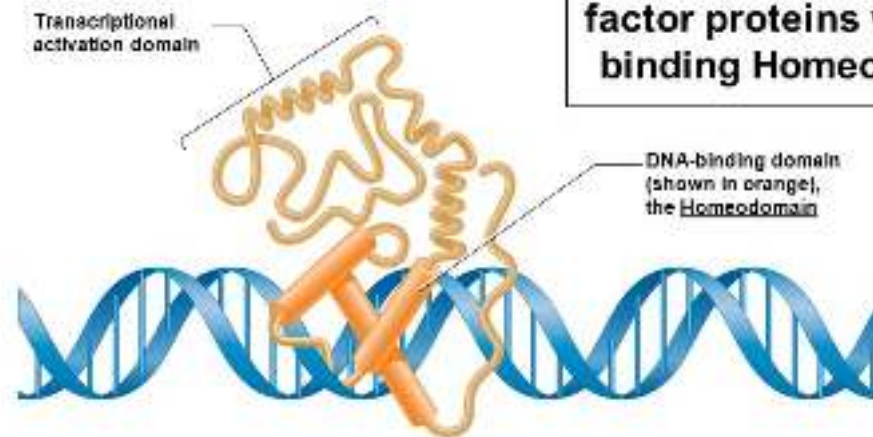
Bithorax mutant



B Disorders of limb formation



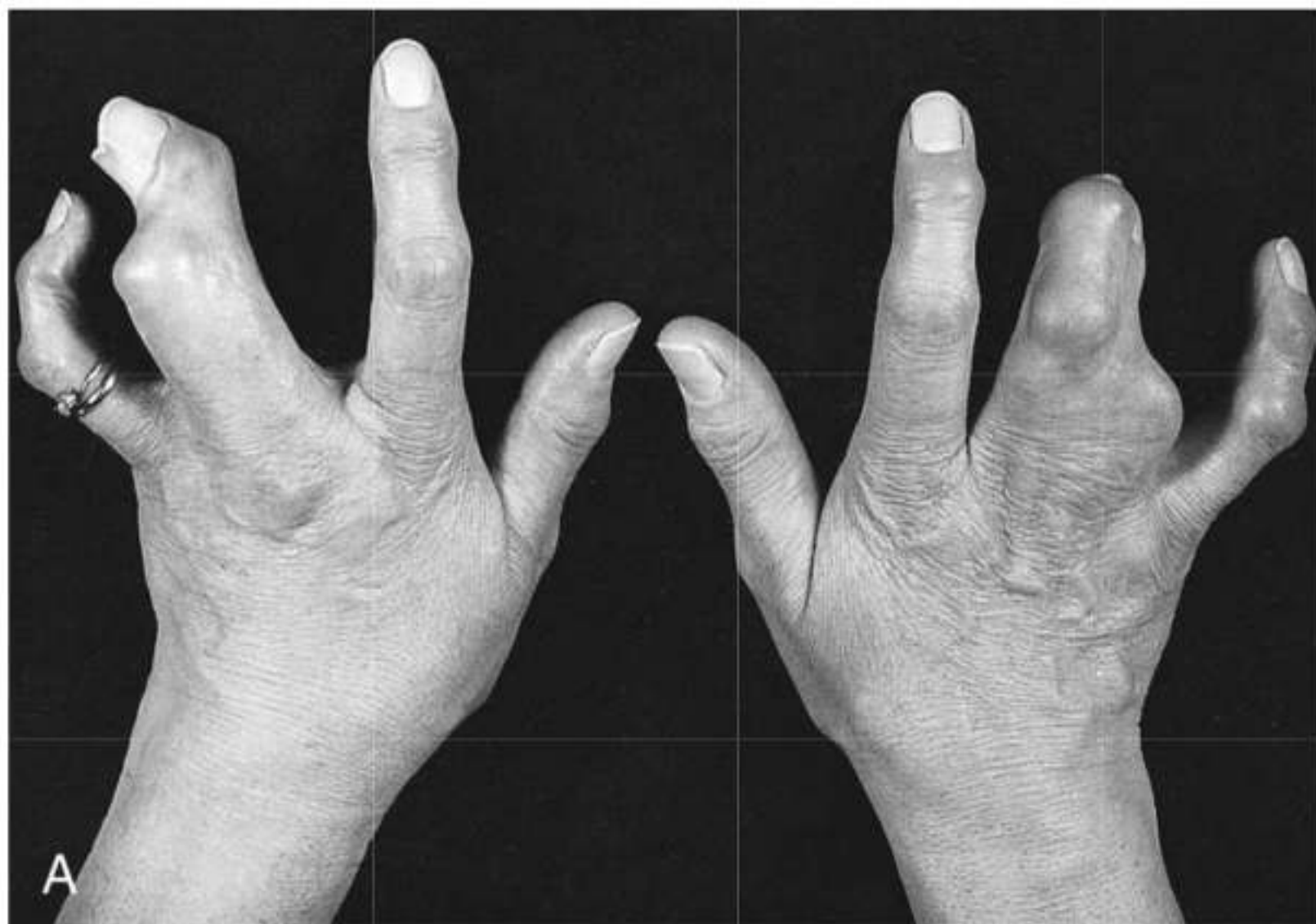
Homeotic Genes often identify transcription factor proteins with DNA-binding Homeodomains



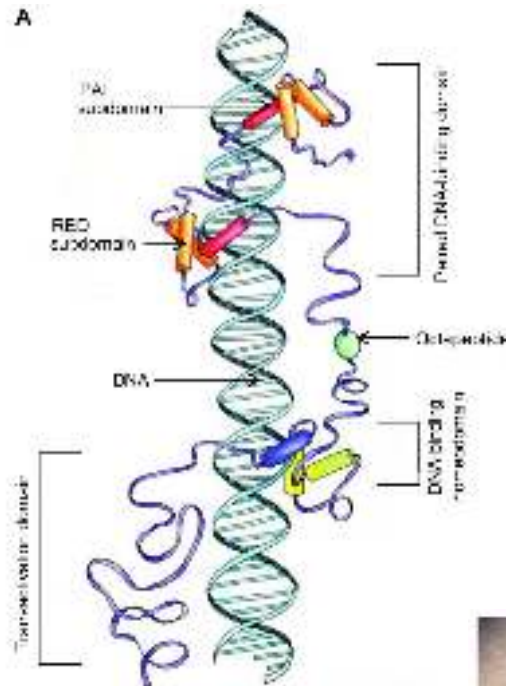
(b) Homeotic protein bound to DNA

Copyright © The McGraw-Hill Companies, Inc. Permission required for reproduction or display

Brooker, Fig 24 - 14



PAX genes



| Gene | Chromosome Location | Developmental Abnormality |
|------|---------------------|---------------------------------|
| PAX2 | 10q24 | Renal-coloboma syndrome |
| PAX3 | 2q35 | Waardenburg syndrome type 1 |
| PAX6 | 11p13 | Aniridia |
| PAX8 | 2q12 | Absent or ectopic thyroid gland |
| PAX9 | 14q12 | Oligodontia |

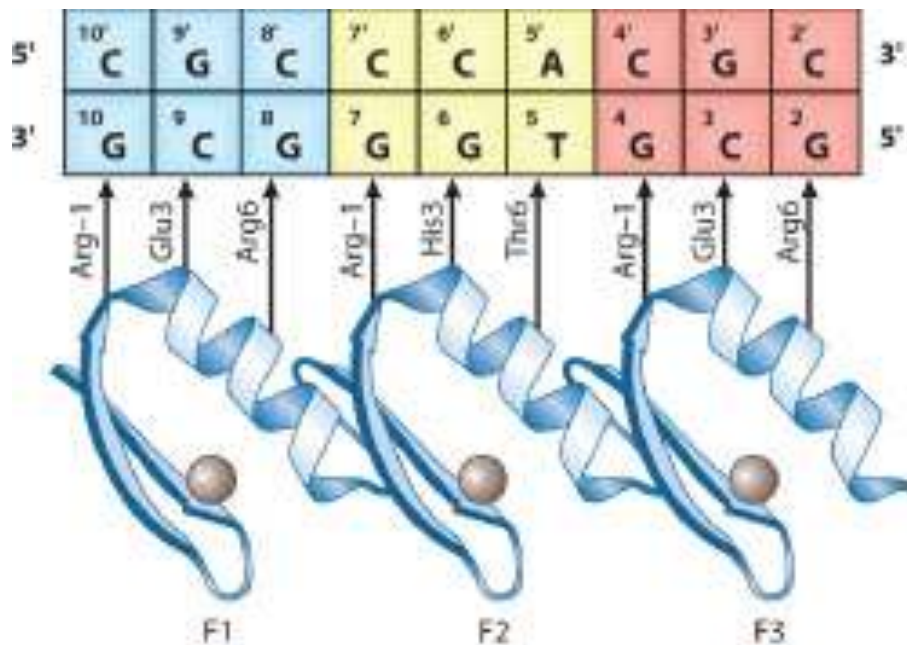
B

| | | |
|-----------|----------------------|------------------------------------|
| Group I | Pax1 Pax8 | No homeodomain Octapeptide |
| Group II | Pax2 Pax6 Pax8 | Partial homeodomain Octapeptide |
| Group III | Pax2 Pax7 | Full homeodomain Octapeptide |
| Group IV | Pax4 Pax9 | Full homeodomain No octapeptide |



Zinc finger genes

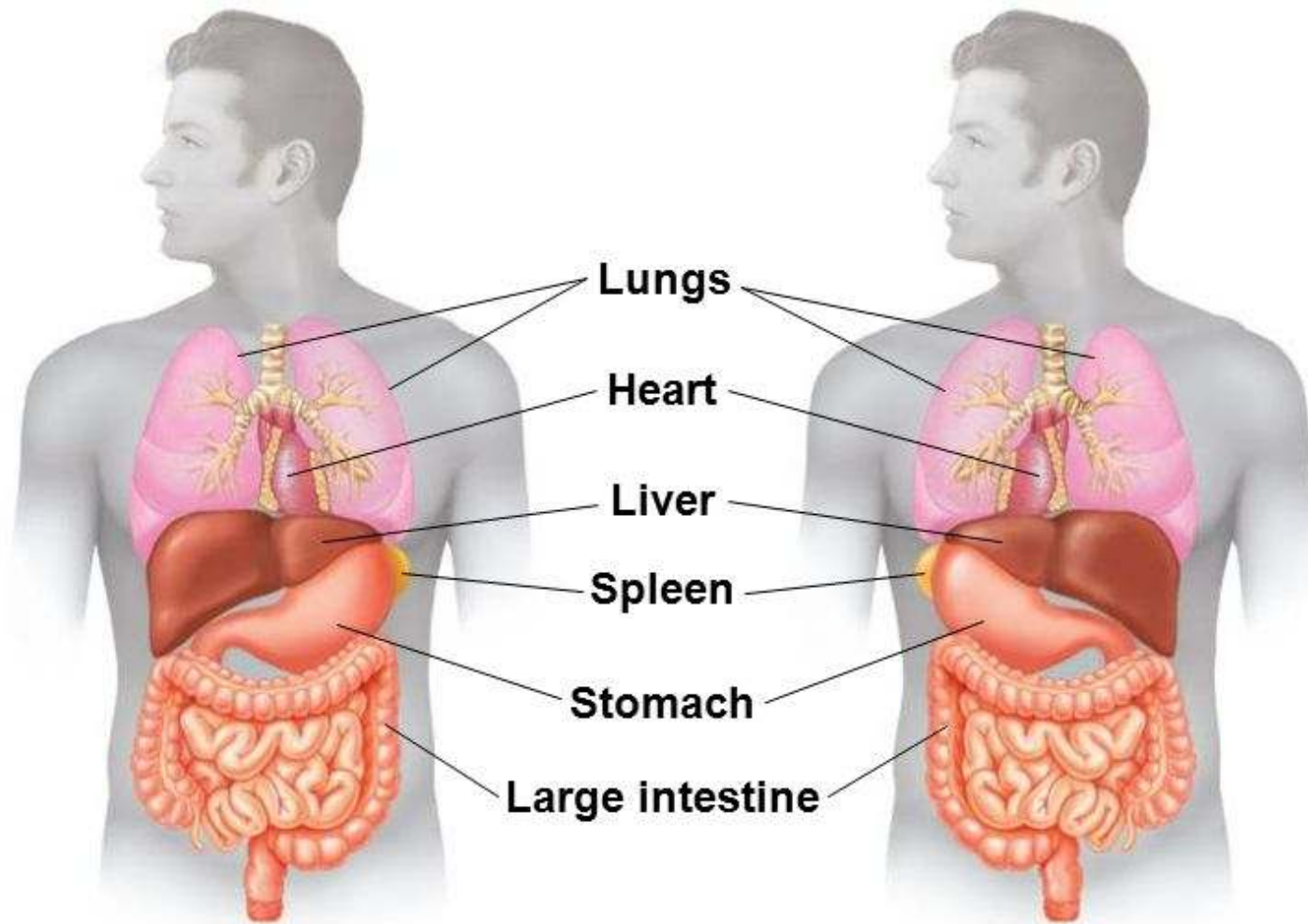
| Gene | Chromosome Location | Developmental Abnormality |
|------|---------------------|--|
| GLI3 | 7p13 | Greig syndrome and Pallister-Hall syndrome |
| WT1 | 11p13 | Denys-Drash syndrome |
| ZIC2 | 13q32 | Holoprosencephaly |
| ZIC3 | Xq26 | Laterality defects |



Abnormalities of head, hands, feet.
Laterality defects (situs inversus).

Greig syndrom (polysyndactyly)

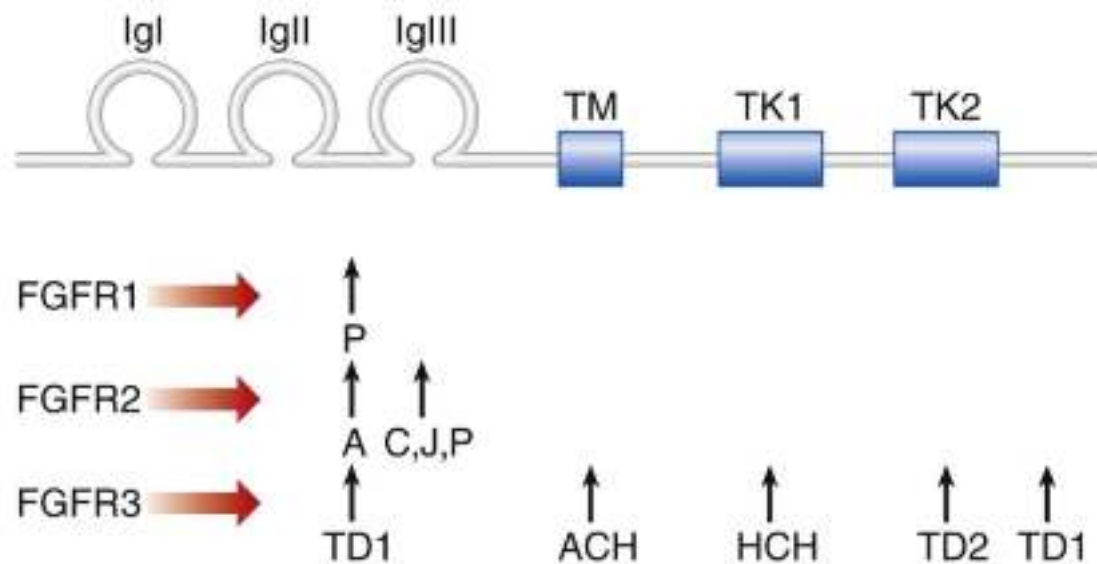




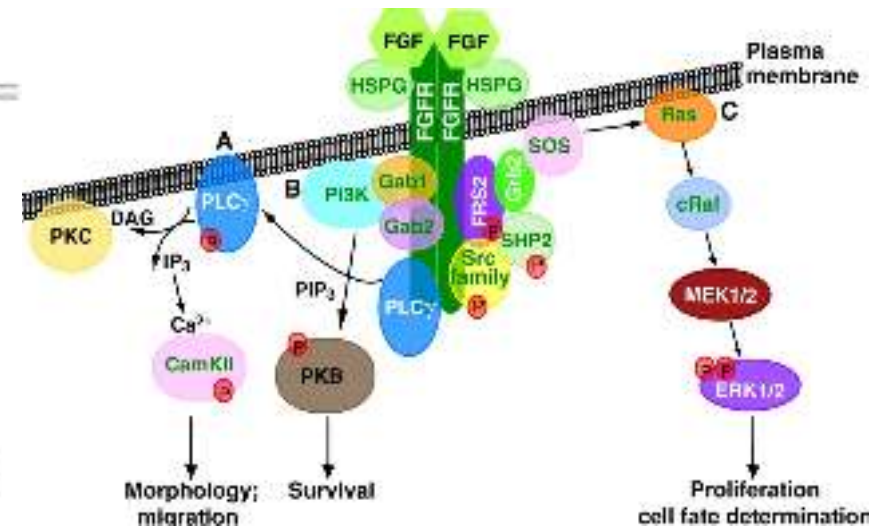
**Normal location
of internal organs**

**Location in
situs inversus**

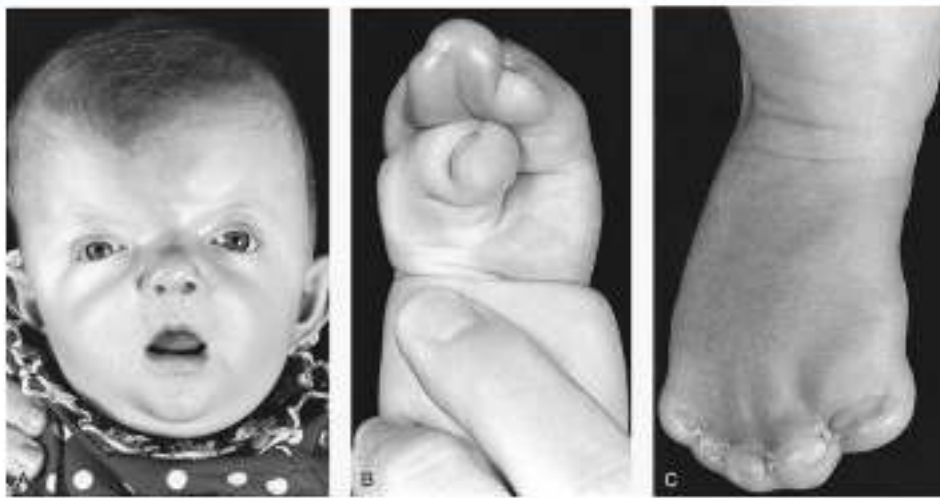
FGF receptor signaling



A – Apert
 C – Crouzon
 TK – tyrosin kinase
 Ig – immunoglobulin domene
 TM – transmembrane domene
 P – Pfeiffer
 J – Jackson Weiss
 TD – Thanatophoric dysplasia
 ACH – Achondroplasia
 HCH – Hypochondroplasia



| Gene | Chromosome | Syndrome |
|----------------------------|------------|--|
| Craniosynostosis syndromes | | |
| <i>FGFR1</i> | 8p11 | Pfeiffer |
| <i>FGFR2</i> | 10q25 | Apert Crouzon Jackson-Weiss Pfeiffer |
| <i>FGFR3</i> | 4p16 | Crouzon (with acanthosis nigricans) |
| Skeletal dysplasias | | |
| <i>FGFR3</i> | 4p16 | Achondroplasia Hypochondroplasia Thanatophoric dysplasia |



1



2



3

Different FGF signaling disorders:

- 1. Apert
- 2. Achondroplasia
- 3. Thanatophoric dysplasia
- 4. Crouzon



4

Ciliopathies

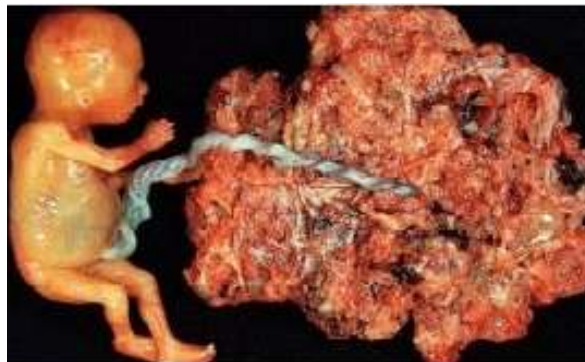
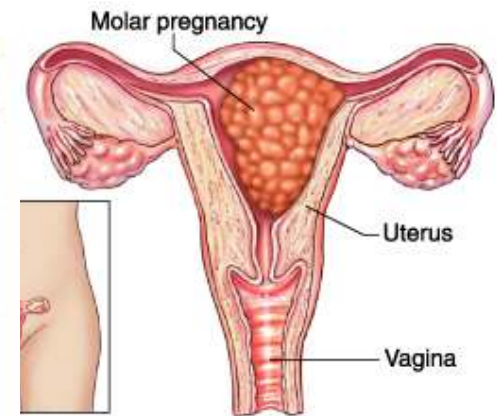
| Disease/Syndrome | Gene | Chromosome Locus | Body System(s) Affected |
|--|--------------------------|-------------------------|---|
| Alstrom syndrome | AIMS1 | 2p13 | Retina, adipose, endocrine, heart |
| Jeune asphyxiating thoracic dystrophy | JFT80 | 15q13 | Skeleton |
| Bardet-Biedl syndrome | BBS1-BBS14 | Multiple | Multisystem, including retina, kidney, skeleton |
| Cranioectodermal dysplasia (Scribner syndrome) | | | Kidney, liver |
| Ellis-van Creveld syndrome | EVCI, EVC2 | 4p16 | Skeleton, heart |
| Joubert syndrome | JBS1 (+ others) | 9q34.3 | Brain |
| Leber congenital amaurosis | GUCY2D, RPE65 (+ others) | 17p13, 11p31 (+ others) | Retina |
| McKusick Kaufman syndrome | DOSG | 20p12 | Limb, heart, urogenital tract |
| Meckel-Gruber syndrome | MKS2 (+ others) | 17q23 (+ others) | Brain, kidney, liver |
| Nephronophthisis (types 1-4) | Nephrocystin (+ others) | Multiple | Kidney |
| Cra-tacio-digital syndrome type 1 | CDR1 (+ others) | Xp.22 (+ others) | Skeleton (limb, face) |
| Polycystic kidney disease | Multiple | Multiple | Kidney |
| Primary ciliary dyskinesia (Kartagener syndrome) | Multiple | Multiple | Multi-system |
| Senior-Loken syndrome | Multiple | Multiple | Retina, kidney |
| Short rib polydactyly syndrome | DRWC2NL | 11q13 | Skeleton, kidney, urogenital tract |



Hydatiform moles

TABLE 6.9 Characteristics of Partial and Complete Hydatidiform Moles

| | Partial Mole | Complete Mole |
|--------------------------------|----------------------------|----------------------|
| No. of chromosomes | 69 | 46 |
| Parental origin of chromosomes | 23 maternal 46 paternal | All 46 paternal |
| Fetus present | Yes, but not viable | No |
| Malignant potential | Very low | High |

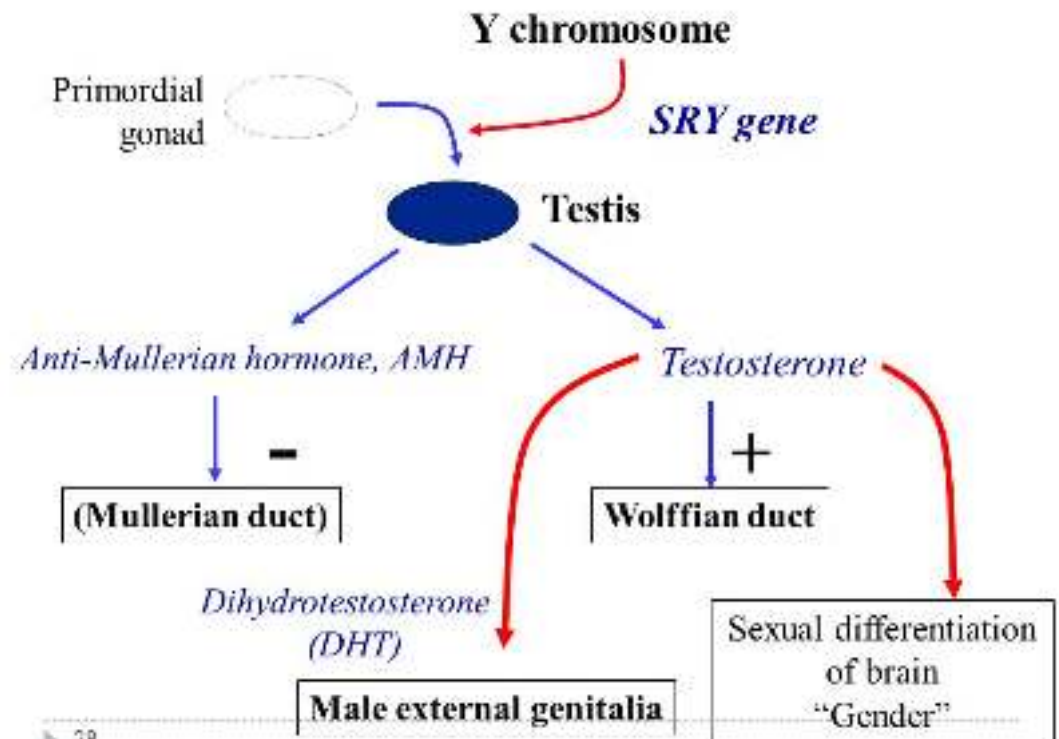
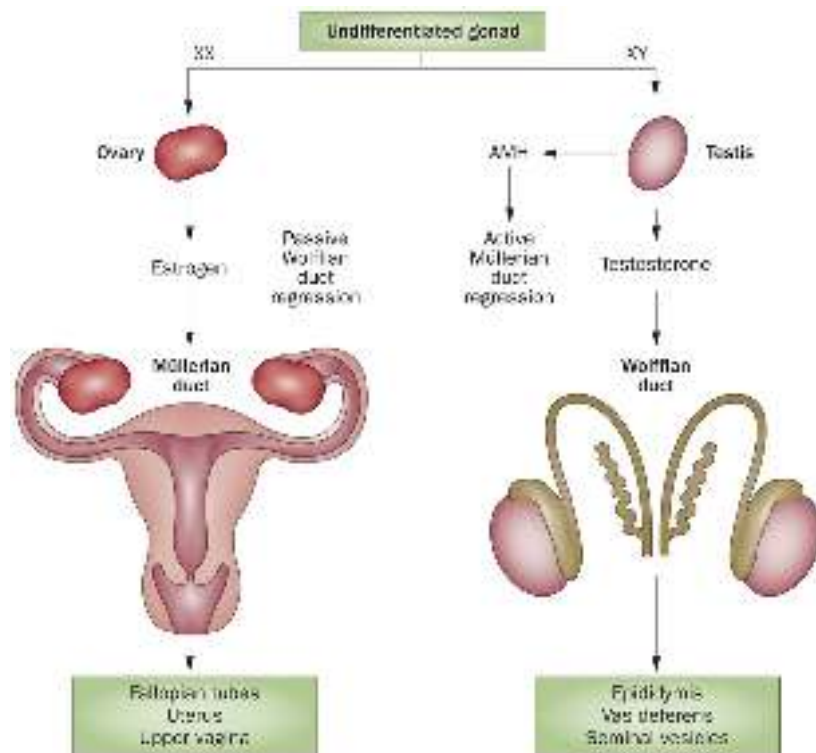


Partial Molar Pregnancy

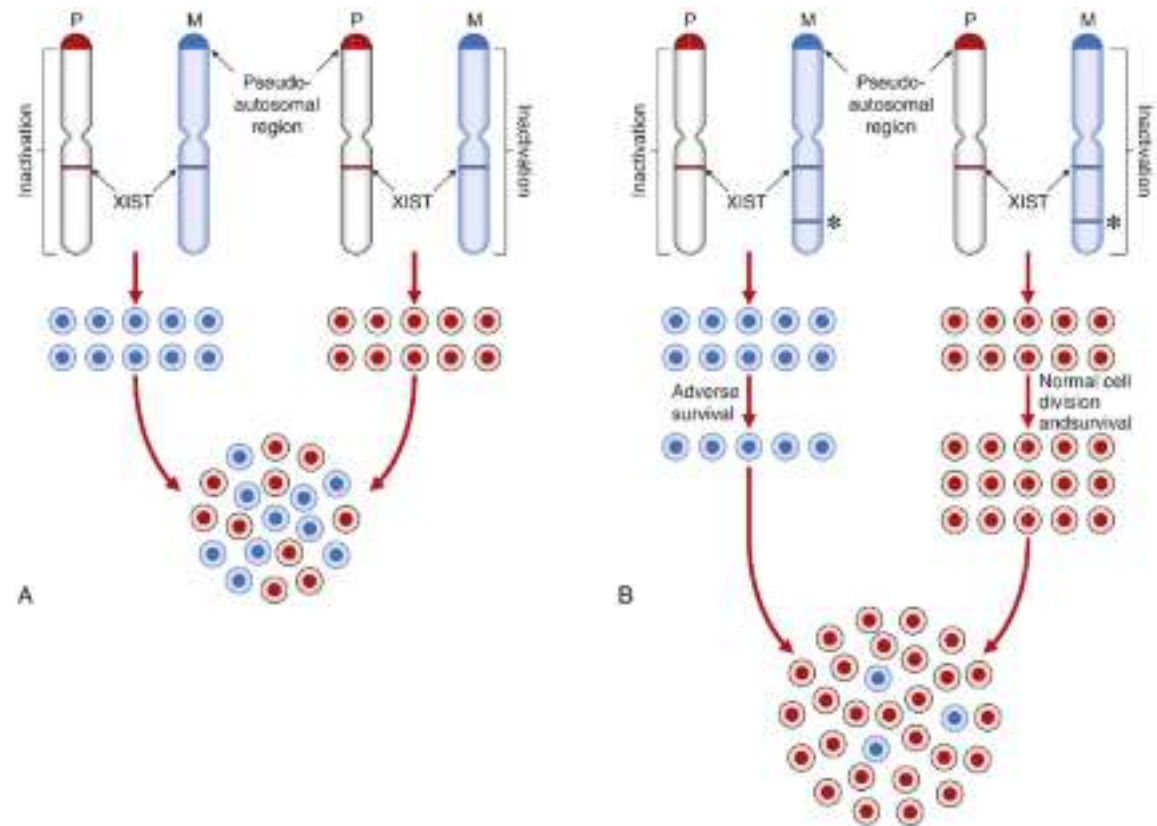
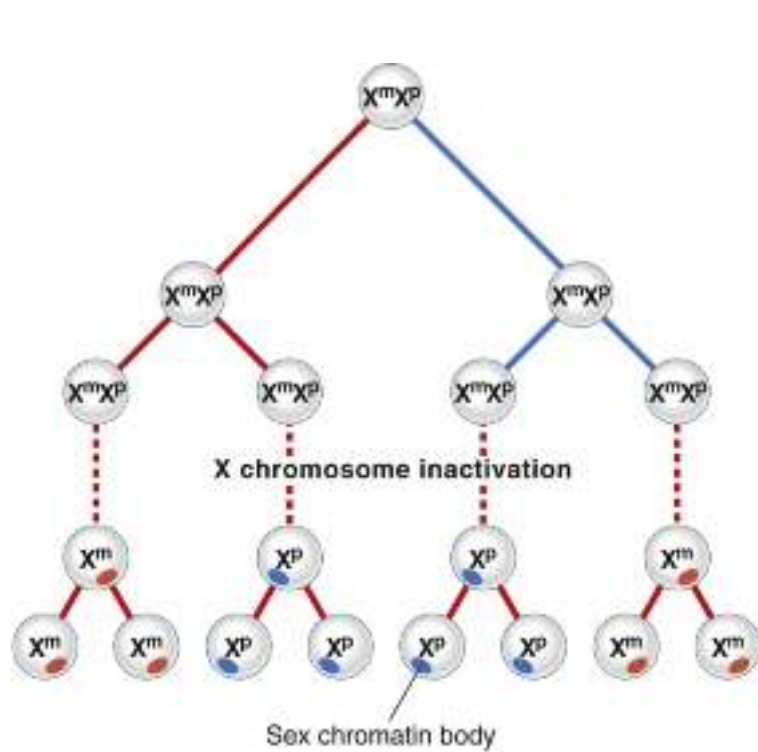


Complete Molar Pregnancy

Sex differentiation - SRY

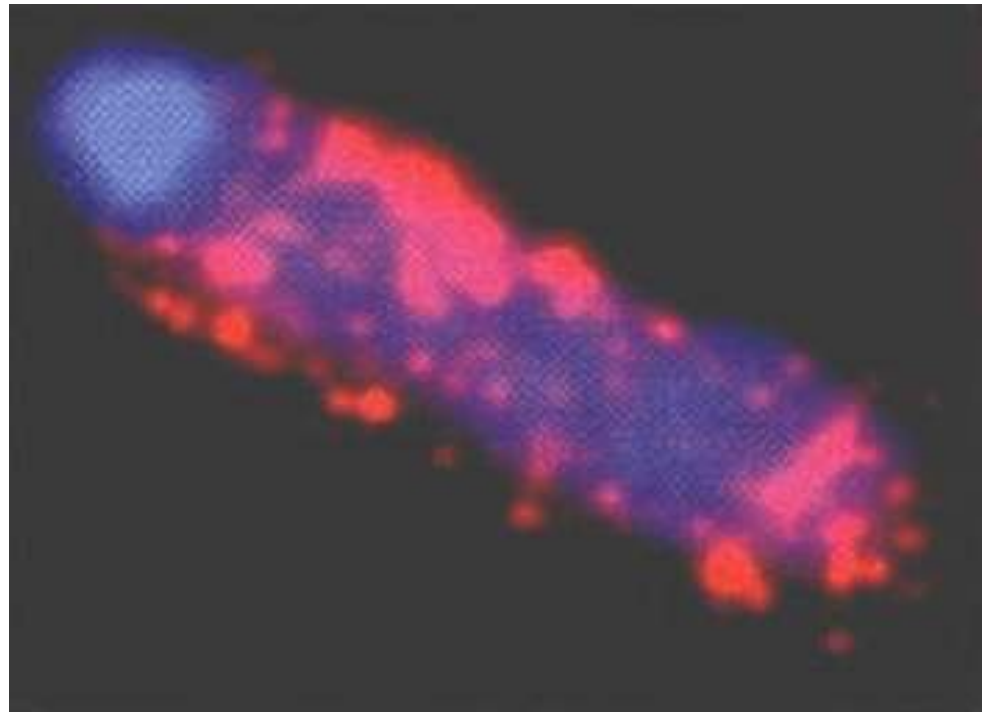


X-chromosome inactivation

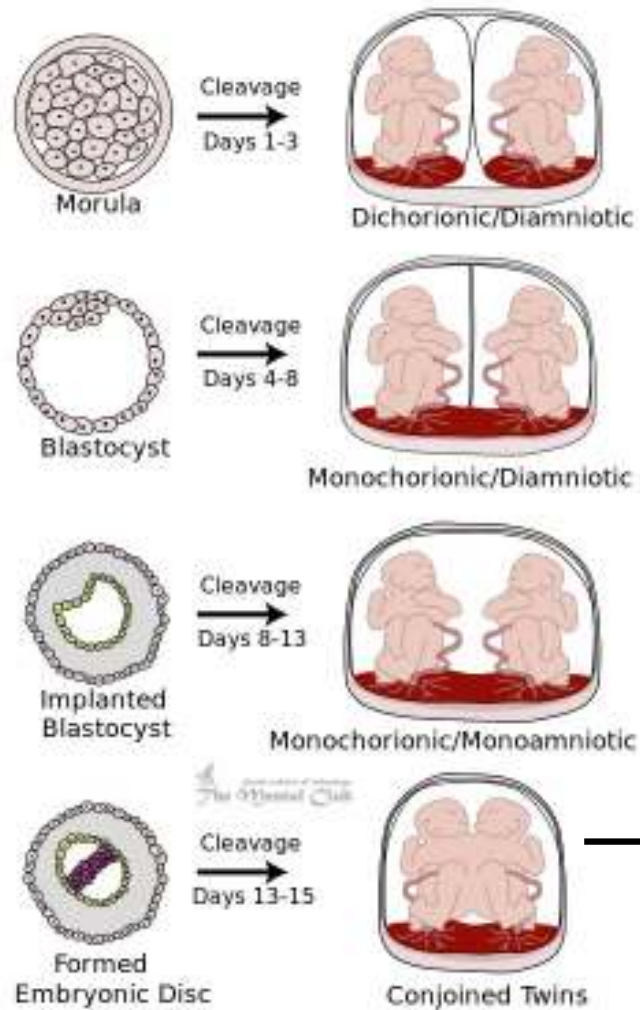


Lyonisation – X chromosome inactivation

- XIST RNA promotes methylation (epigenetic silencing) of inactivated X-chromosome
- Pseudoautosomal region remains active
- Both X chromosomes are active in sex organs

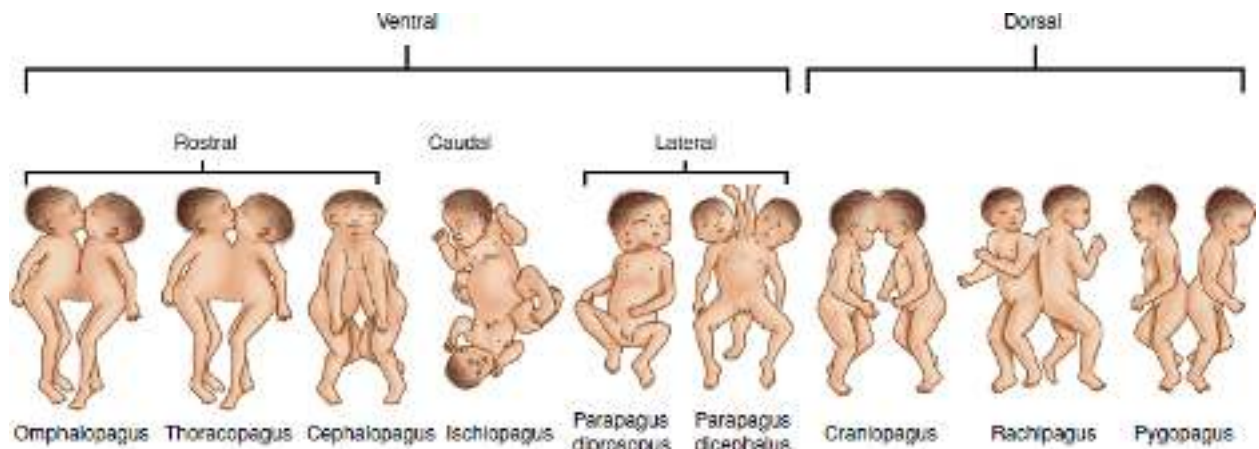
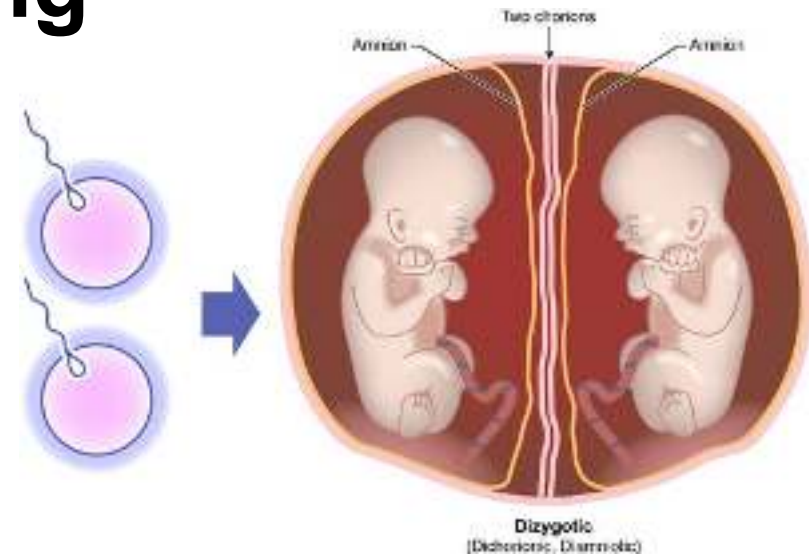


Monozygotic:



Twinning

Dizygotic:



Patterns of inheritance

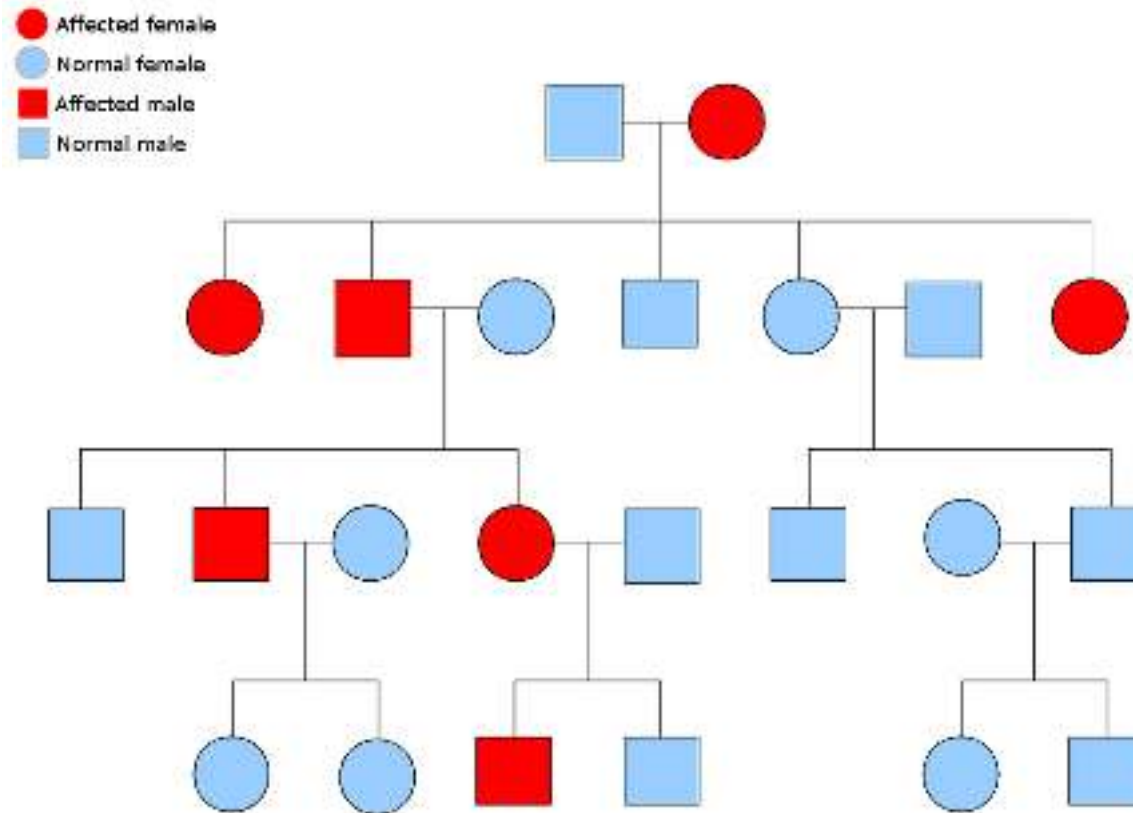
Chapter 7

Inheritance pattern

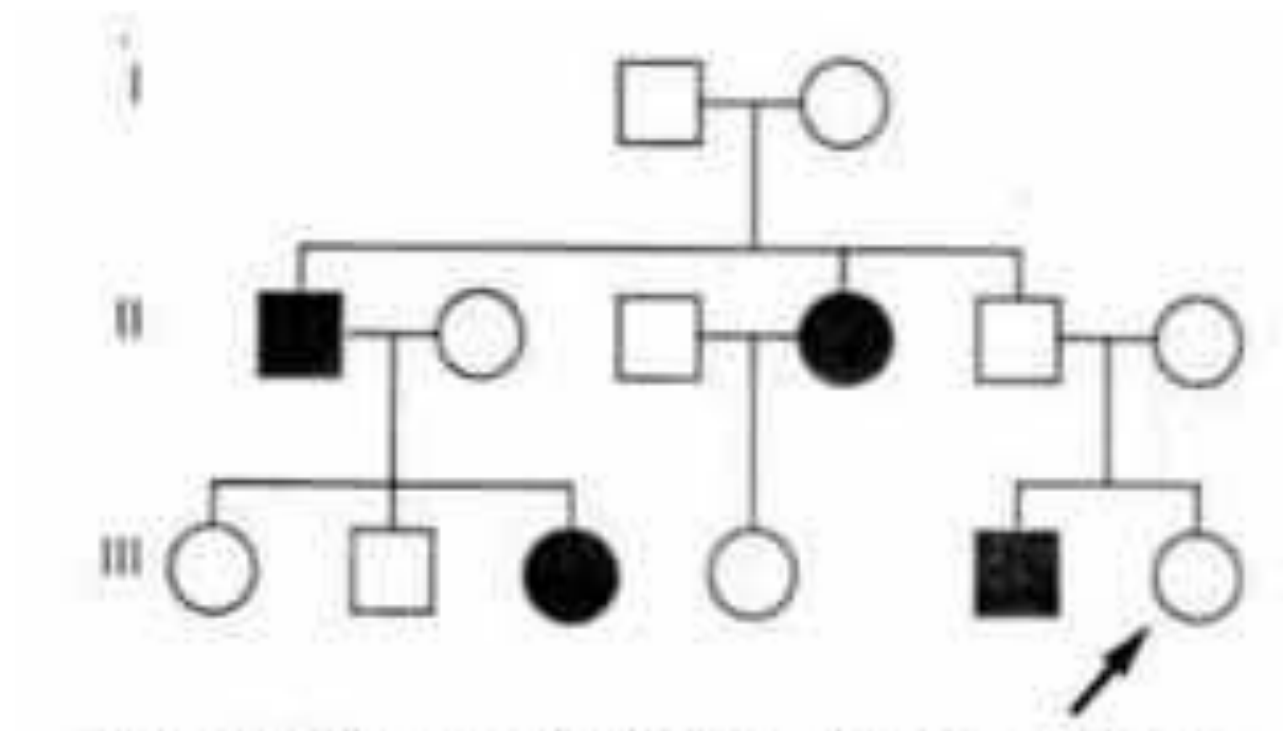
Mendelian inheritance:

- Autosomal: -dominante
 - recessive
- Sex-linked: -X linked recessive
 - X linked dominante
 - Y linked (holandric)

Autosomal dominant (50% risk)

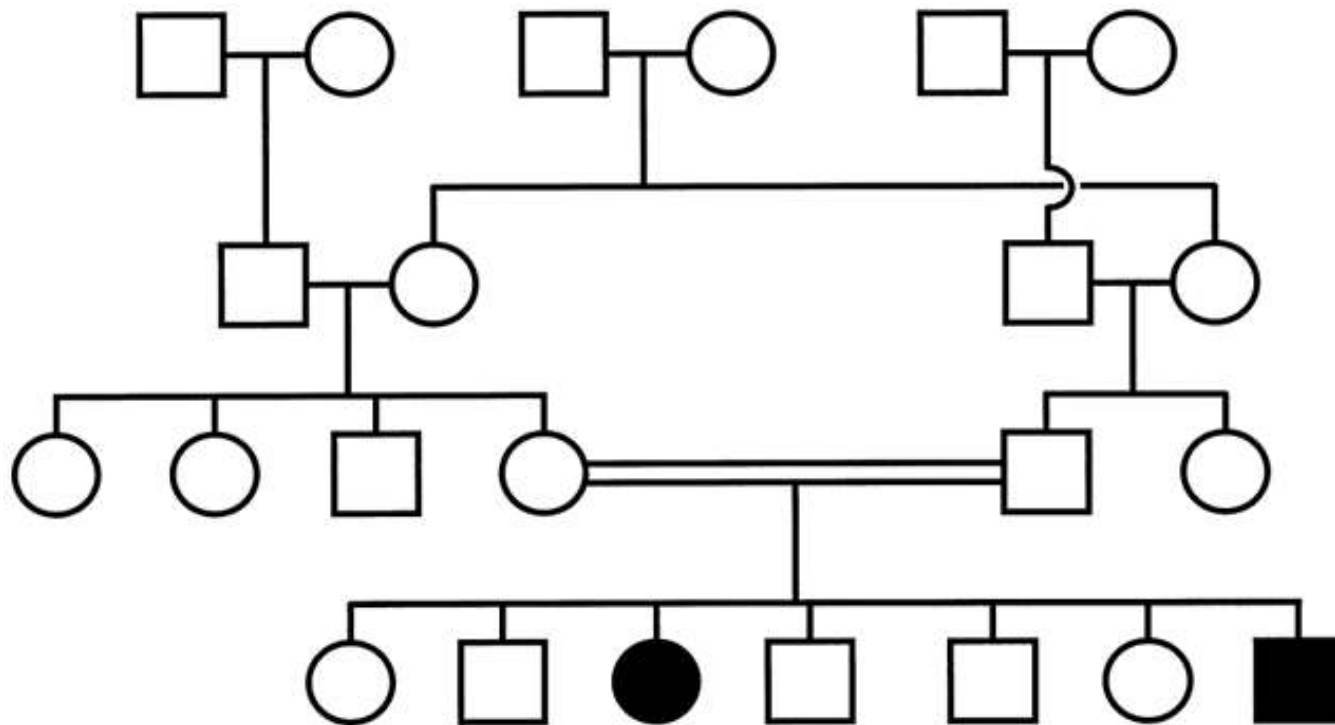


Autosomal dominante with reduced penetrance (non penetrance)

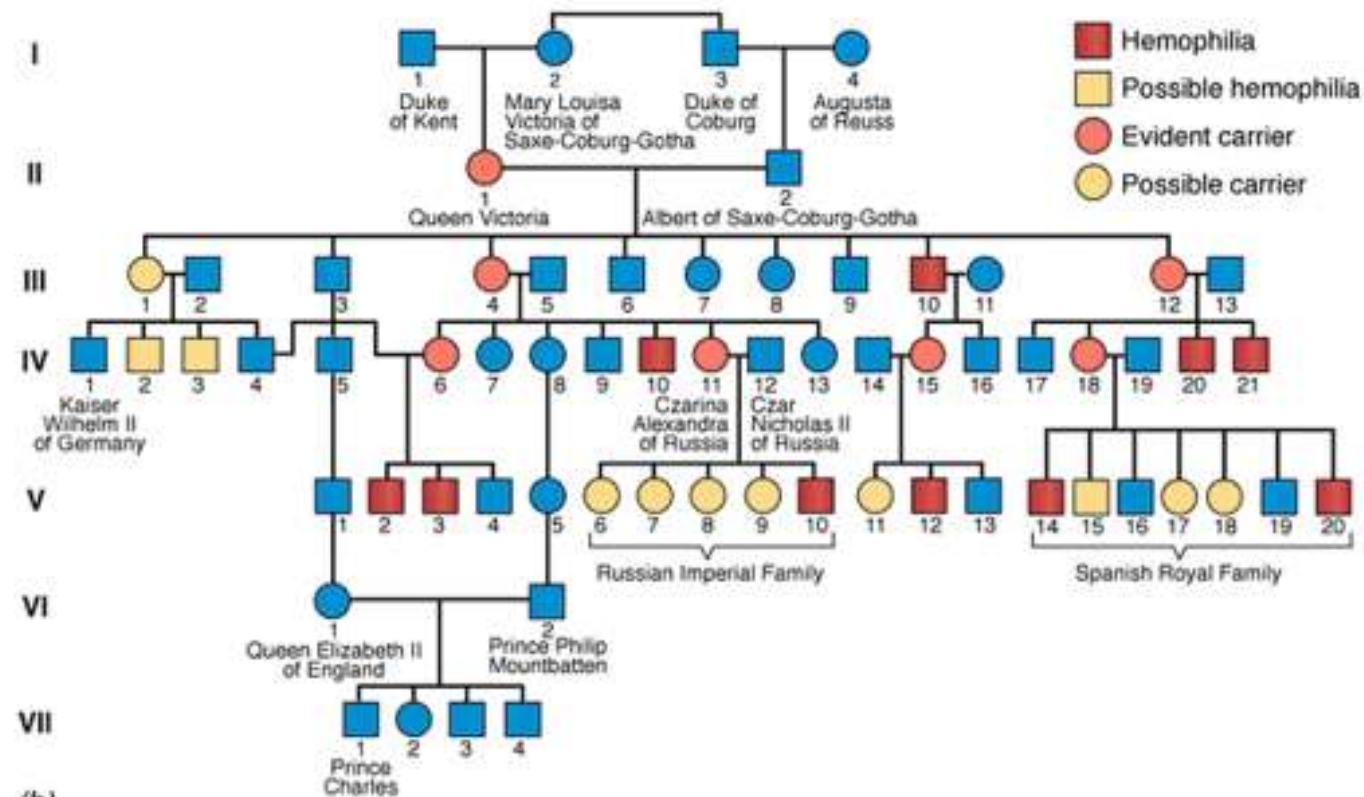


Autosomal recessive inheritance

- Consanguinity (25% risk)



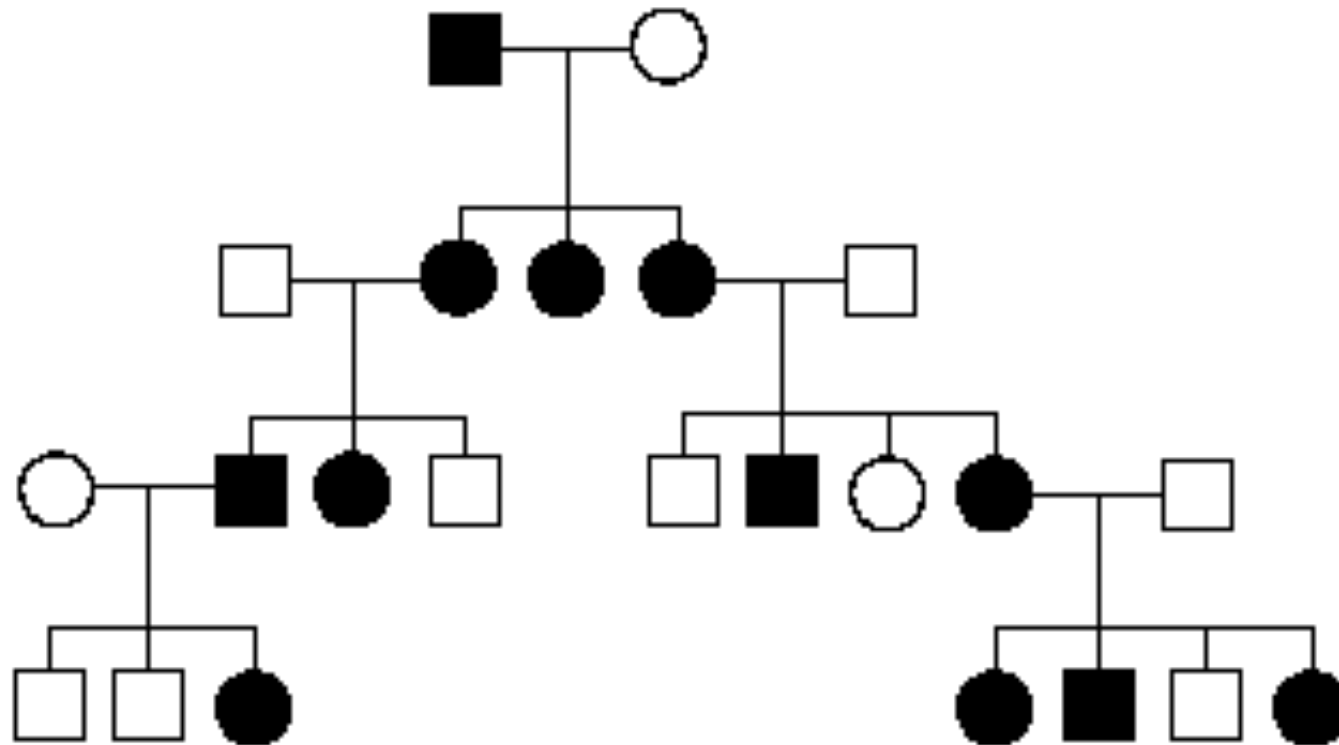
X linked recessive



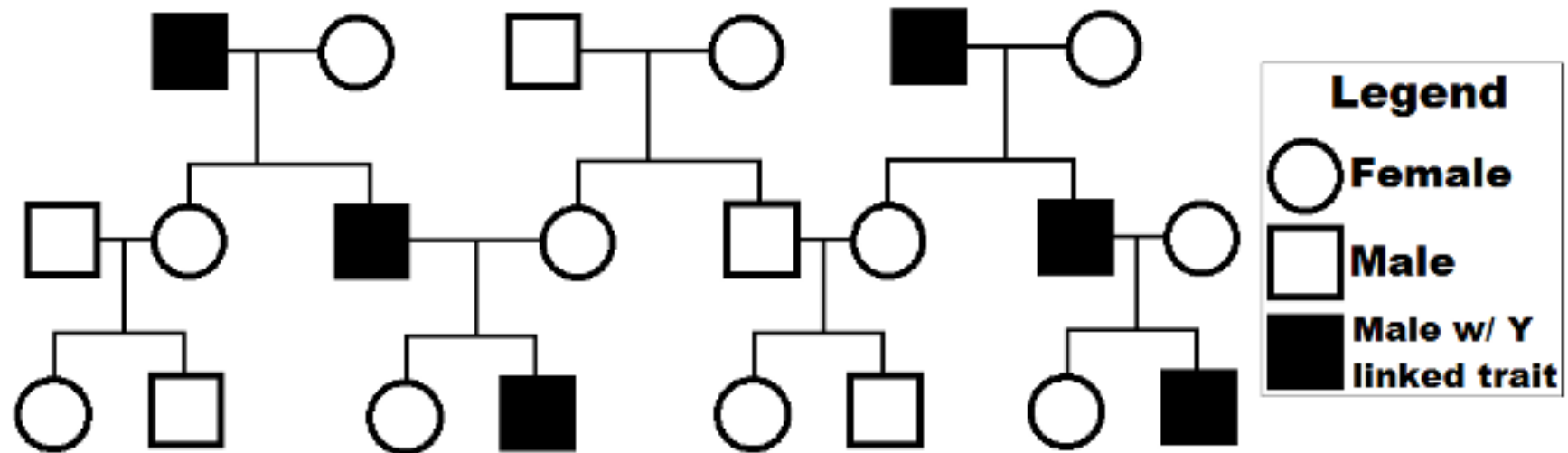
(b)

Copyright © 1997, by John Wiley & Sons, Inc. All rights reserved.

X linked dominante



Y-linked inheritance



BOX 7.1

Features that Support the Single-Gene or Mendelian Patterns of Inheritance

Autosomal Dominant

- Males and females affected in equal proportions
- Affected individuals in multiple generations
- Transmission by individuals of both sexes (i.e., male to male, female to female, male to female, and female to male)

Autosomal Recessives

- Males and females affected in equal proportions
- Affected individuals usually in only a single generation
- Parents can be related (i.e., consanguineous)

X-Linked Recessive

- Only males usually affected
- Transmitted through unaffected females
- Males cannot transmit the disorder to their sons (i.e., no male-to-male transmission)

X-Linked Dominant

- Males and females affected but often an excess of females
- Females less severely affected than males
- Affected males can transmit the disorder to their daughters but not to sons

Y-Linked Inheritance

- Affected males only
- Affected males must transmit it to their sons

Nonmendelian inheritance

- **Anticipation:** the symptoms of the genetic disorder become apparent at an earlier age with each generation
- **Mosaicism** = consists of more than one cell type (occurs during early development):
 - somatic
 - germ cells
- **Uniparental disomy:** heterodysomy or isodisomy
 - Genomic imprinting
- **Mitochondrial inheritance**
- Multifactorial diseases

Genomic imprinting

- alternative methylation pattern

Angelman (15q11-q13)

- paternal uniparental disomy
- UBE3A (ubiquitin 3 ligase), expressed from a mother is missing



Prader willy (15q11-q13)

- maternal uniparental disomy
- SNRPN (small nuclear ribonucleoprotein polypeptide N) and adjacent genes, expressed from a father, are missing



11p15 region:

- **Beckwith-Wiedemann** syndrom (paternal uniparental disomy)
 - IGF2 overexpression → overgrowth)
- **Russel-Silver** syndrom (maternal uniparental disomy) – growth retardation

Mitochondrial inheritance

- Heteroplasmy: a wide range of clinical severity
- Effects energy metabolism of a cell

