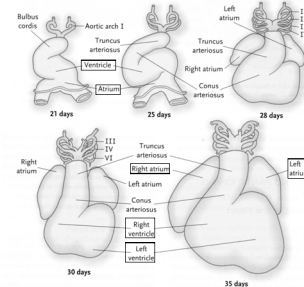


INTRODUCTION TO HUMAN HEART DEVELOPMENT

Debbie Yelon
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 Department of Cell Biology
 Skirball Institute, NYU School of Medicine

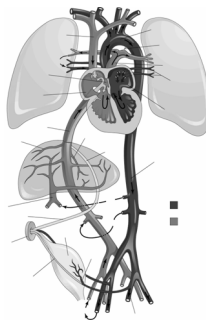
HUMAN HEART DEVELOPMENT



POSTNATAL CIRCULATION

PULMONARY CIRCULATION
 RIGHT CHAMBERS

SYSTEMIC CIRCULATION
 LEFT CHAMBERS



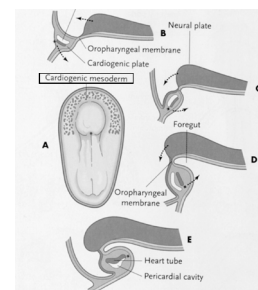
HUMAN HEART DEVELOPMENT

- HEART TUBE FORMATION
- CARDIAC LOOPING
- CHAMBER SEPTATION
- VALVE AND OUTFLOW FORMATION

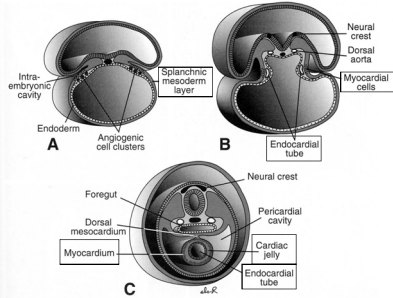
CONGENITAL HEART DISEASE

- RELATIVELY COMMON
- GENERALLY INITIATED BY EARLY DEVELOPMENTAL ERRORS
- CAN BE CAUSED BY EXPOSURE TO TERATOGENS
- CAN ORIGINATE WITH GENETIC DEFECTS

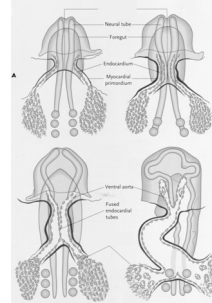
LONGITUDINAL FOLDING POSITIONS CARDIAC CELLS



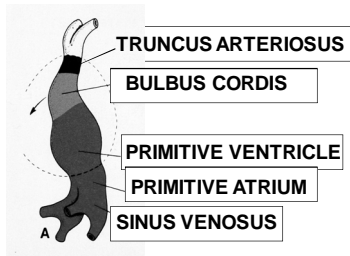
LATERAL FOLDING FACILITATES TUBE FORMATION



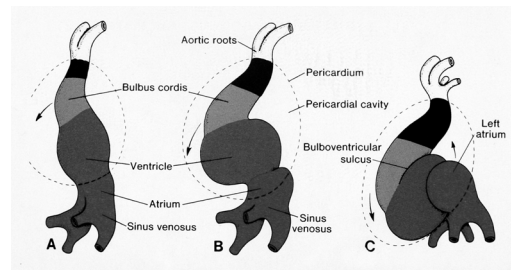
TUBE FORMATION BEGINS ROSTRALLY



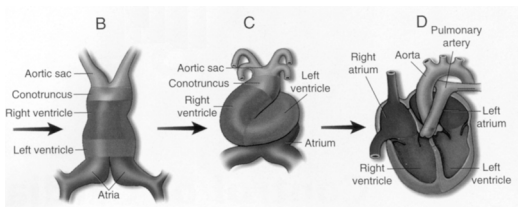
PRIMITIVE HEART TUBE



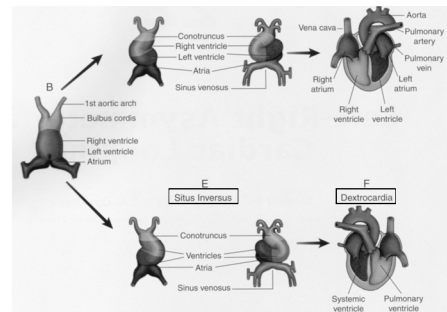
CARDIAC LOOPING



CARDIAC LOOPING



DEXTROCARDIA AND SITUS INVERSUS



GENETIC BASIS FOR HETEROTAXY

- NODAL FAMILY OF GROWTH FACTORS REQUIRED FOR ESTABLISHMENT OF LEFT-RIGHT AXIS
- *CFC1* GENE ENCODES A COMPONENT OF THE RECEPTOR FOR NODAL FACTORS
- MUTATIONS IN *CFC1* CAUSE HETEROTAXY

FROM FETAL TO POSTNATAL CIRCULATION

PARTITIONING THE HEART

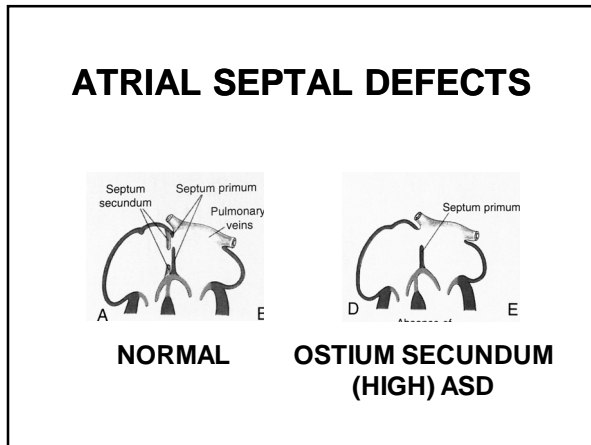
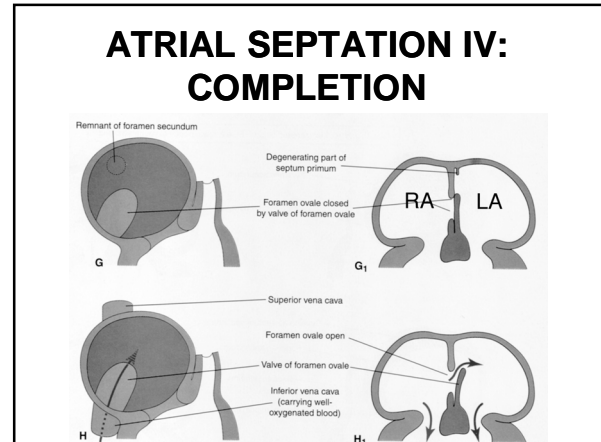
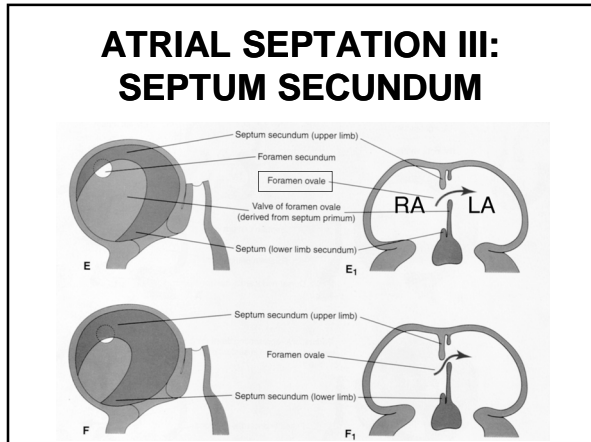
- ATRIAL SEPTATION
- VENTRICULAR SEPTATION
- ATRIOVENTRICULAR VALVE FORMATION
- DIVISION OF THE OUTFLOW TRACT

SAGITTAL SECTIONS

CORONAL SECTIONS

ATRIAL SEPTATION I: SEPTUM PRIMUM

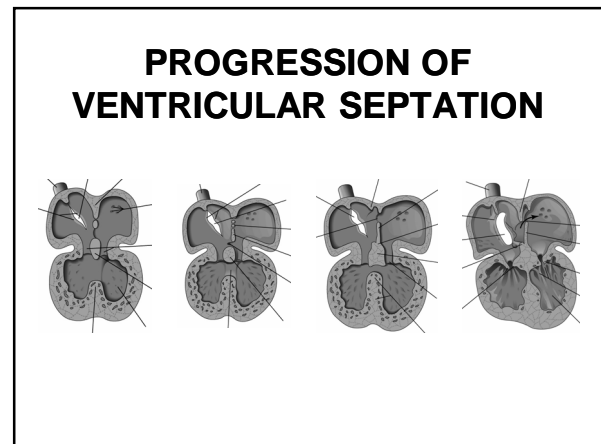
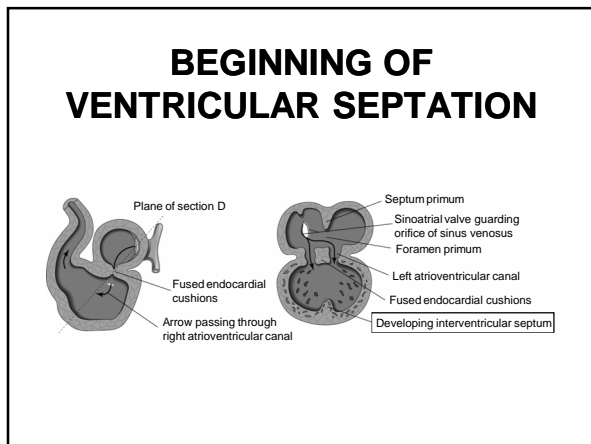
ATRIAL SEPTATION II: FORAMEN SECUNDUM



GENETIC CAUSES OF ASD

HETEROZYGOSITY OF MUTATIONS IN GENES LIKE:

- *Nkx2-5*, ENCODING A HOMEODOMAIN TRANSCRIPTION FACTOR
- *TBX5*, ENCODING A T-BOX TRANSCRIPTION FACTOR (HOLT-ORAM SYNDROME)

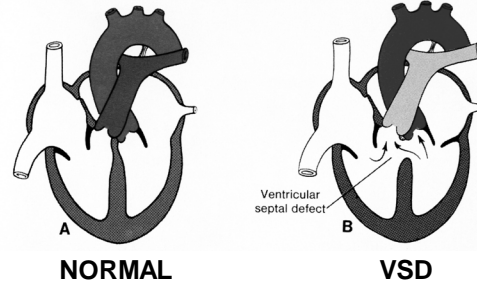


COMPLETION OF VENTRICULAR SEPTATION

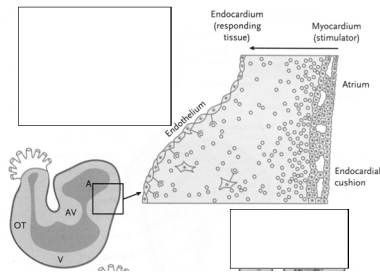
- MUSCULAR SEPTUM
- MEMBRANOUS SEPTUM
- CONOTRUNCAL SEPTUM



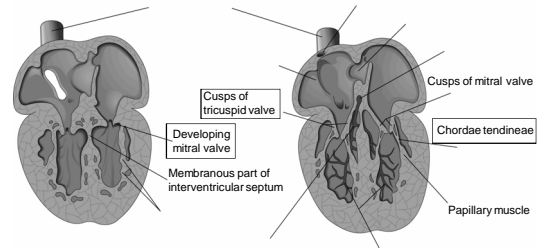
MEMBRANOUS VSD



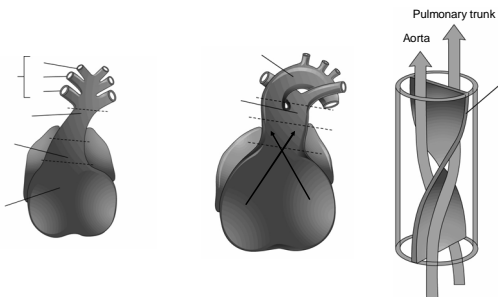
ENDOCARDIAL CUSHION FORMATION



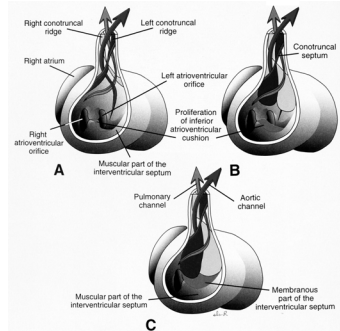
VALVE FORMATION

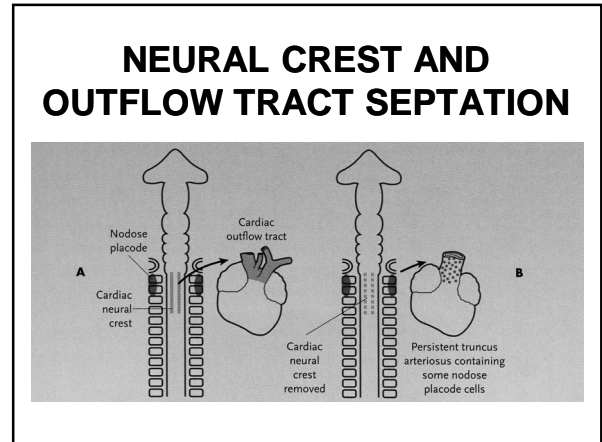
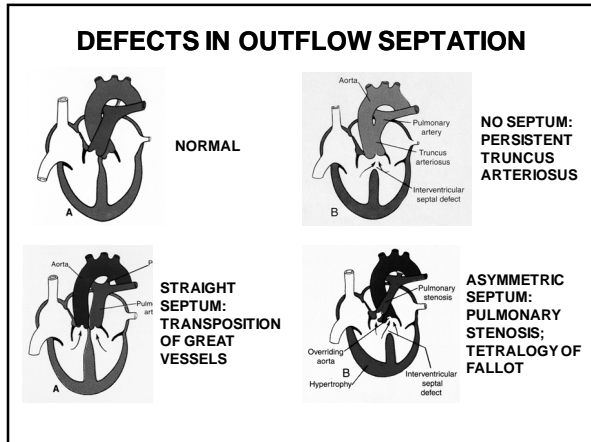


OUTFLOW SEPTATION



OUTFLOW SEPTATION



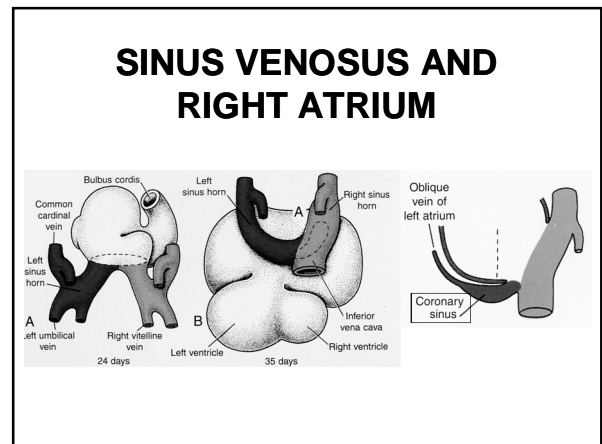


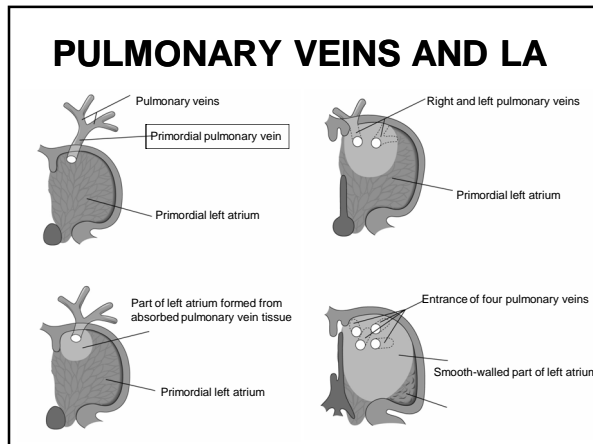
GENETIC BASIS FOR OUTFLOW DEFECTS

- *TBX1* ENCODES A TRANSCRIPTION FACTOR EXPRESSED NEAR MIGRATING NEURAL CREST CELLS
- *TBX1* MUTATION IN MICE CAUSES DEFECTS RESEMBLING DIGEORGE SYNDROME
- DELETION OF *TBX1* FOUND IN MANY DIGEORGE SYNDROME PATIENTS

HUMAN HEART DEVELOPMENT

- HEART TUBE FORMATION
- CARDIAC LOOPING
- CHAMBER SEPTATION
- VALVE AND OUTFLOW FORMATION





ATRIAL SEPTAL DEFECTS

- **OSTIUM PRIMUM (LOW) ASD**
- **OSTIUM SECUNDUM (HIGH) ASD**
- **SINUS VENOSUS ASD**