

DAND5: A newly- recognized cause of autosomal recessive laterality disorders

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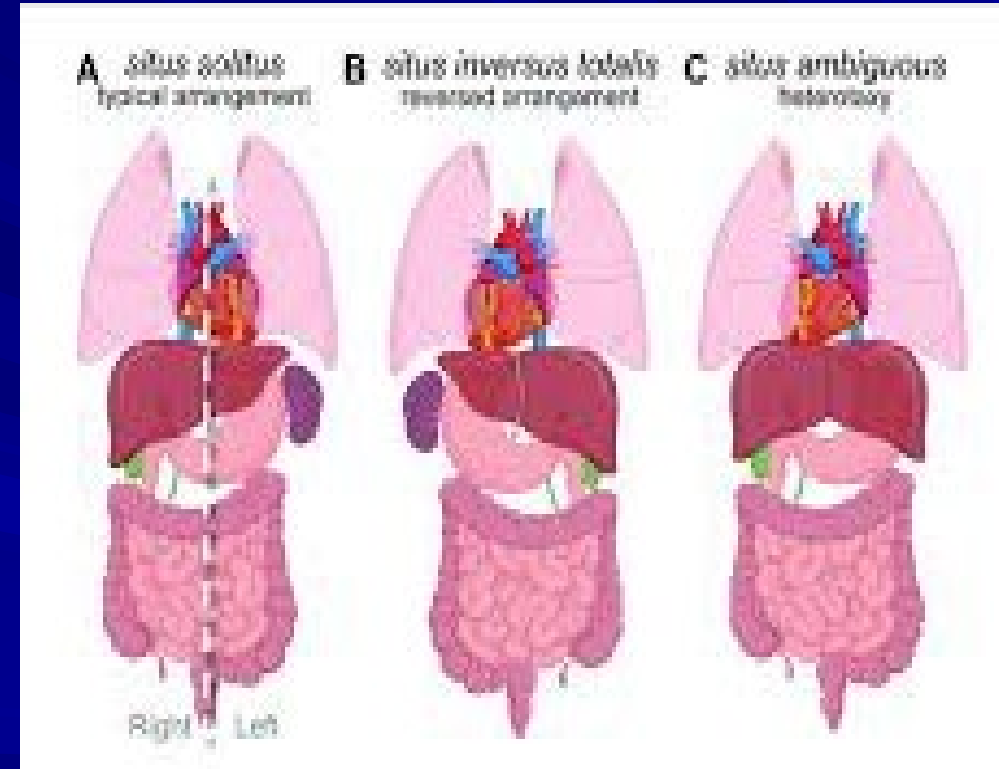
בית החולים לילדים
אדמונד ולילי ספרא

שיבא
תל השומר



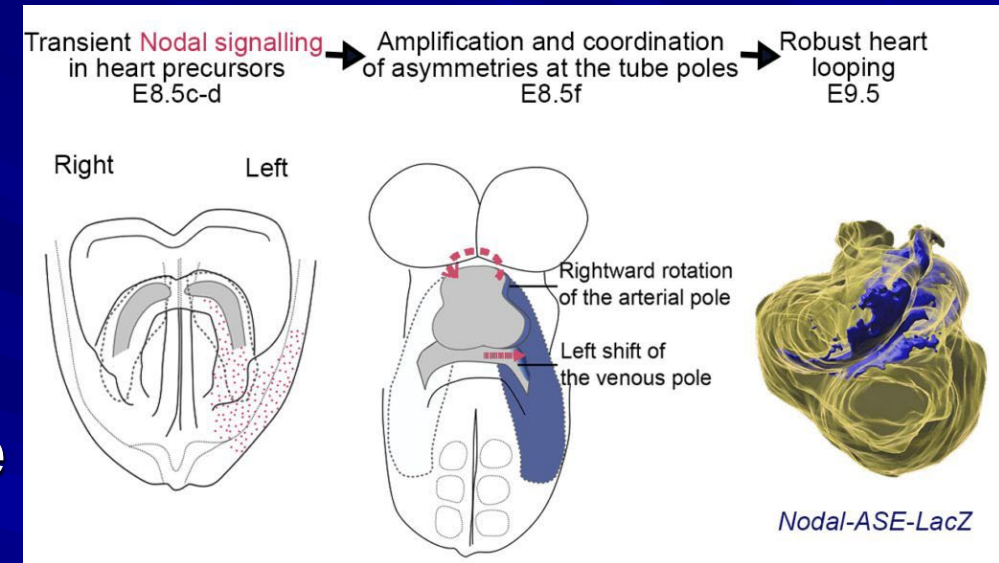
Laterality Disorders- Background:

- **Congenital heart disease-**
most common congenital malformation, ~ 1% of births
- **Laterality disorders-** ~ 1% of total CHD cases
- **Isolated/ multisystemic manifestations**
- **More than 25 genes have been implicated in all modes of inheritance.**



Laterality Disorders- Pathogenesis:

- First asymmetry in morphogenesis occurs with rightward heart looping (embryonic days 22-23)
- Involves multiple signaling molecules, of the TGF- β superfamily
 - Differential activation of Nodal on the left
 - BMP signaling on the right
 - Additional cell signaling pathways: Shh, FGF, Notch



From: *Desgrange et al., 2020 – Summary of the role of Nodal signalling in heart looping.*

Laterality Disorders- Pathogenesis:

- **Situs inversus totalis**- 3–9% accompanied by CHD

- **Situs Ambiguous- Heterotaxy**

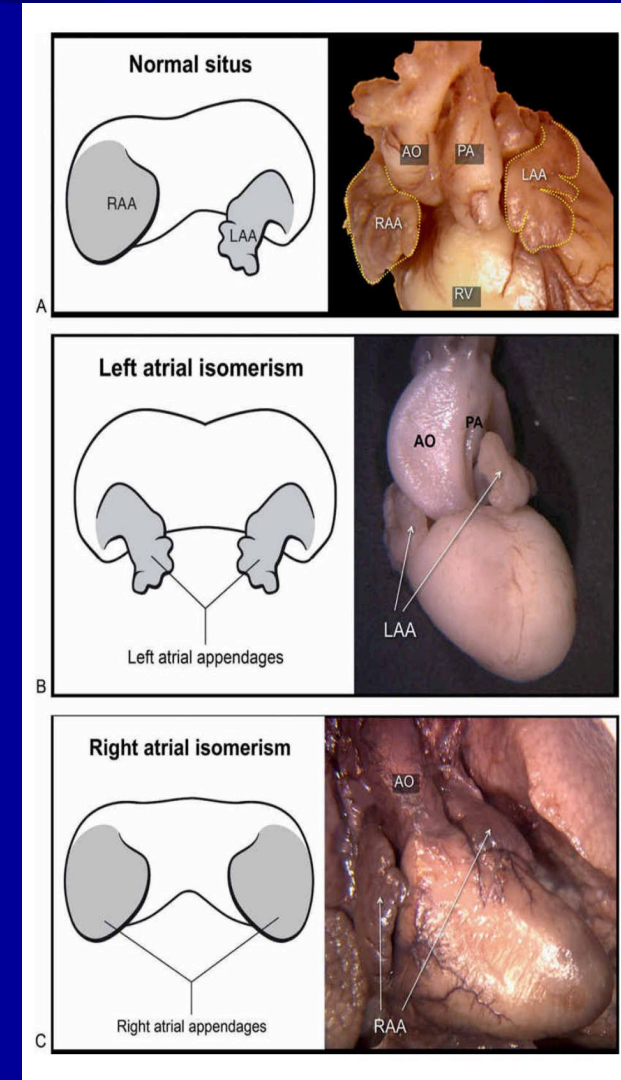
- **Right atrial isomerism**

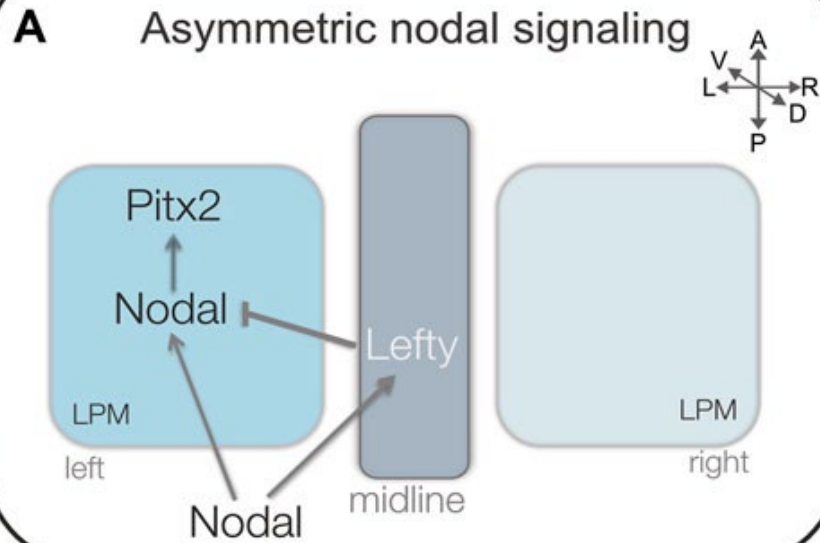
- Two morphologic Rt atria often w' single ventricle, AVSD, TGA and APVD
- Rt bronchial isomerism and asplenia

- **Left atrial isomerism**

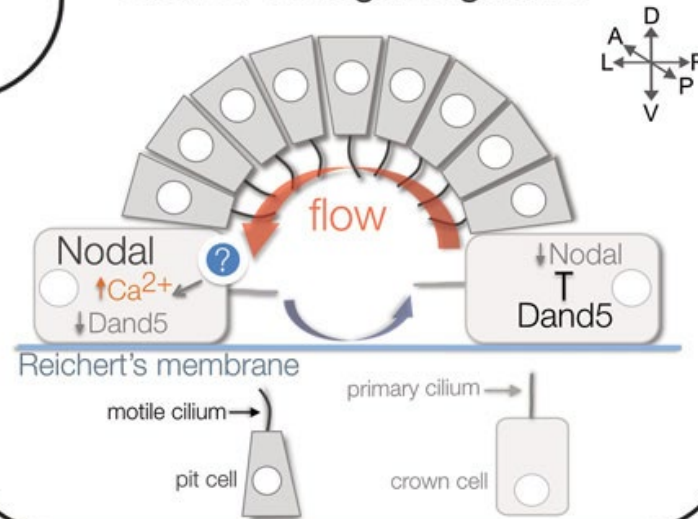
- Two morphologic Lt atria often w' ASD or AVSD.
- Absence of SA/AV node-> risk for CAVB
- Lt bronchial isomerism and polysplenia

- > 20% of cases, variable rearrangements



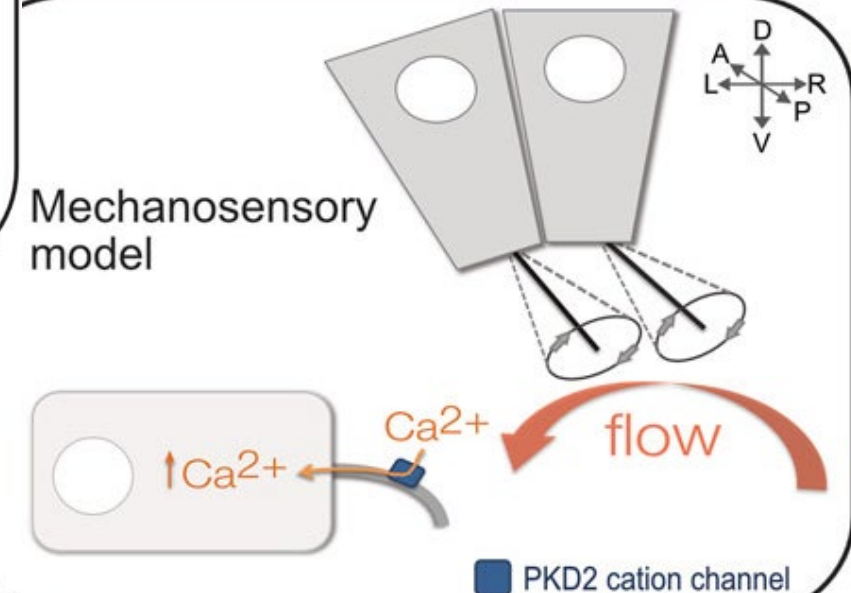


Ciliated left-right organizer



Understanding laterality disorders and the left-right organizer: Insights from zebrafish

Mechanosensory model

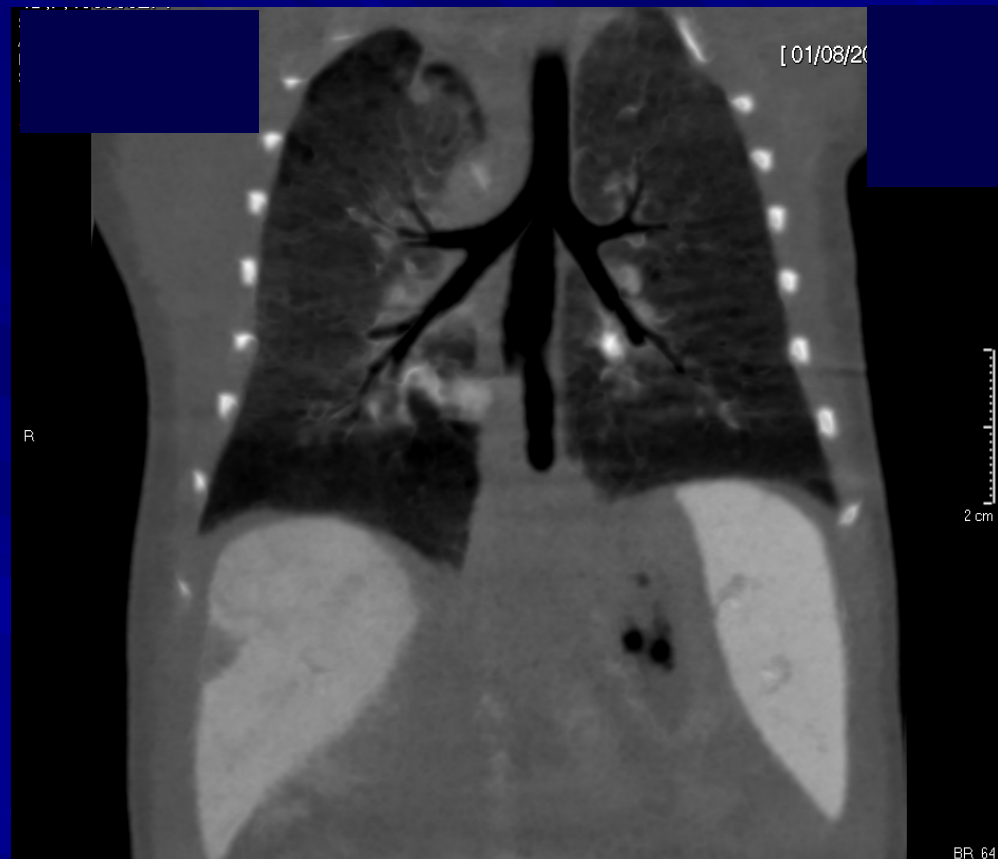
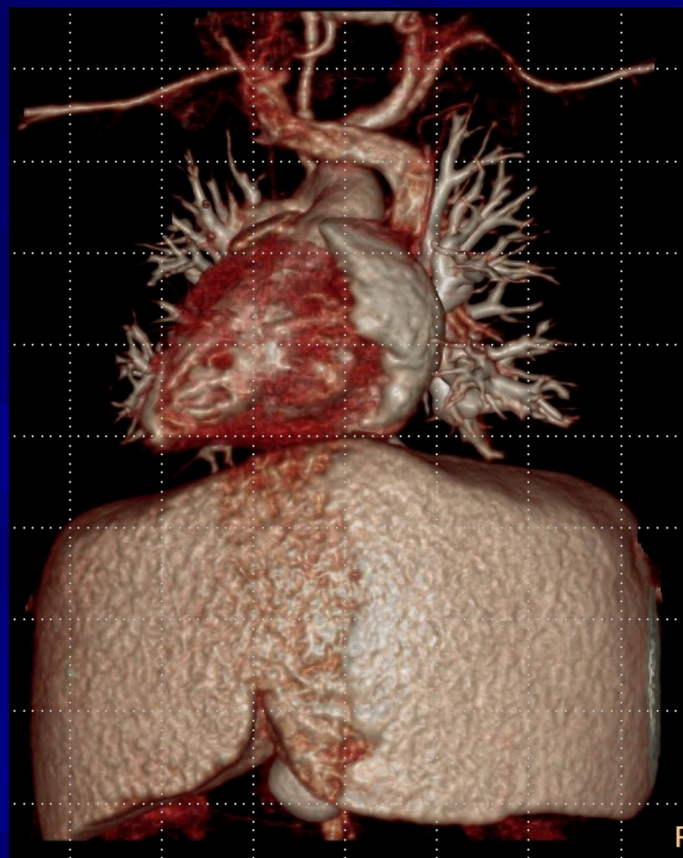
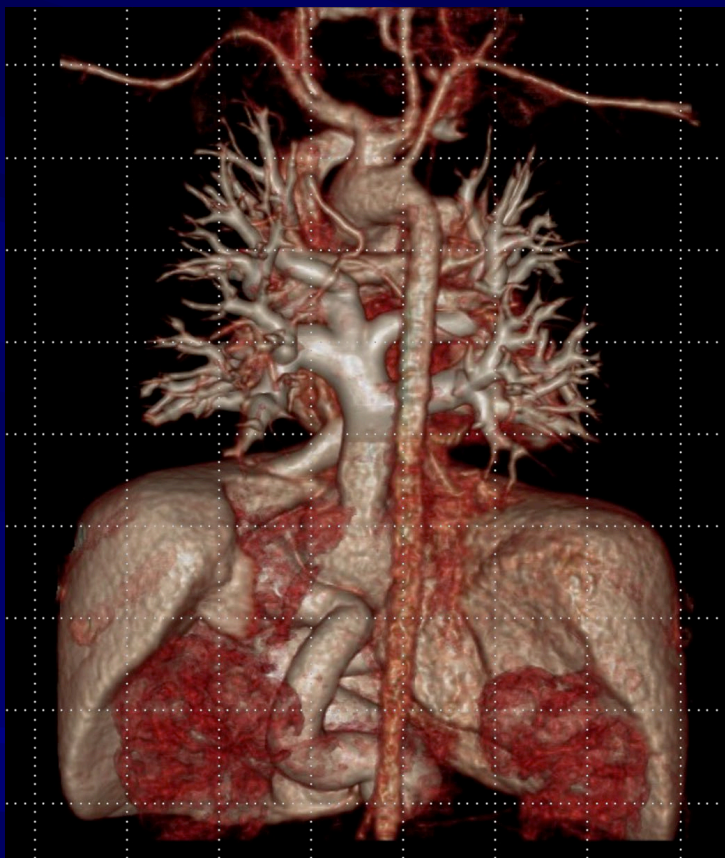


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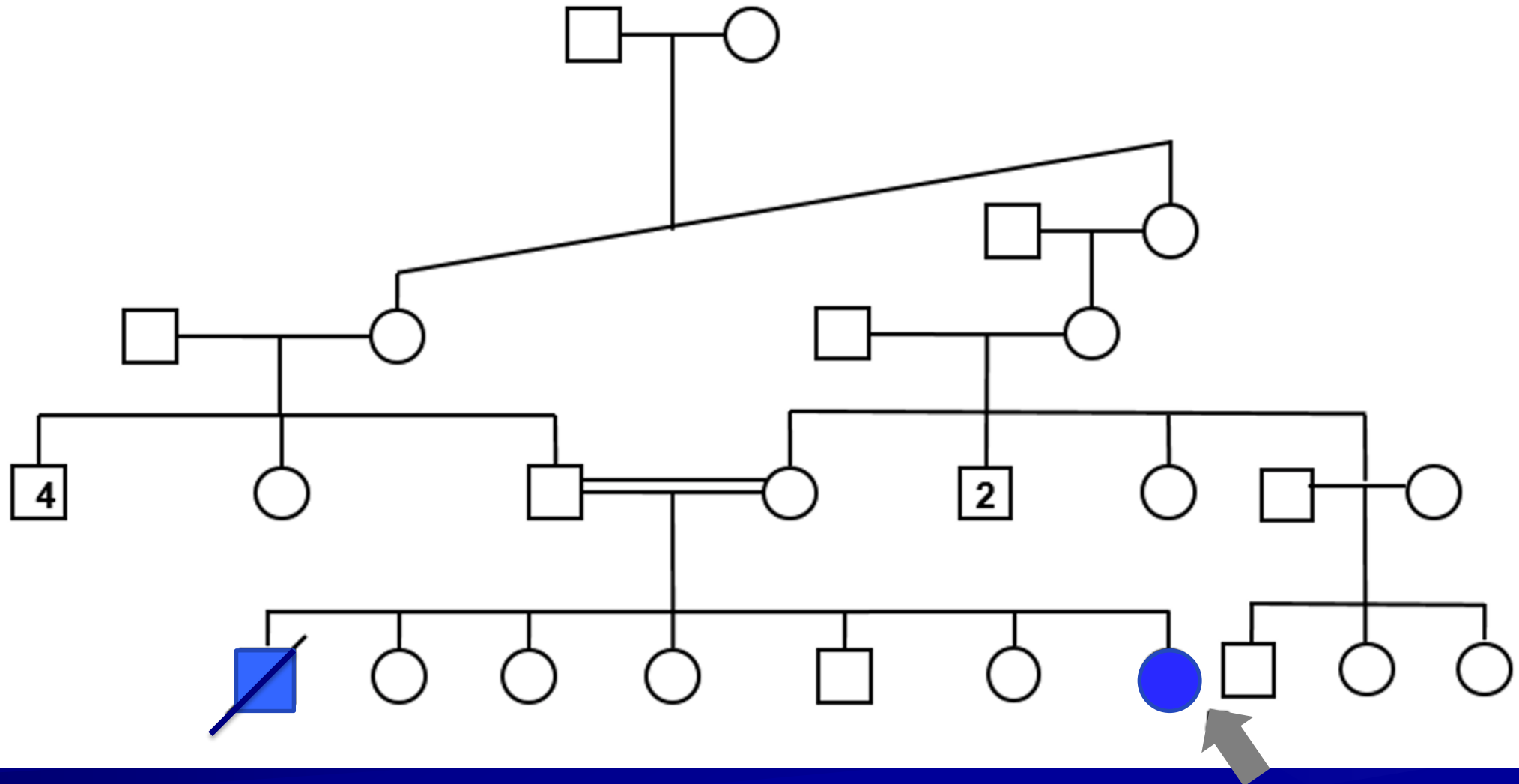
Forrest K, Barricella AC, Pohar SA, Hinman AM, Amack JD. Understanding laterality disorders and the left-right organizer: Insights from zebrafish. Front Cell Dev Biol. 2022 Dec. PMID: 36619867

Case Presentation -Patient 1:

- A 2 day old female referred to our hospital due to dextrocardia with a complex CHD
- Prenatal history uneventful
- Born term, AGA, cyanosis surrounding feeds
- Upon arrival: saturation 50% RA, tachycardia 180 and tachypnea 100, treated with Prostin
- Echo: situs inversus, unbalanced AV canal, infradiaphragmatic TAPVR, PA, asplenia
- Underwent atrial septectomy, SANO conduit, TAPVR repair

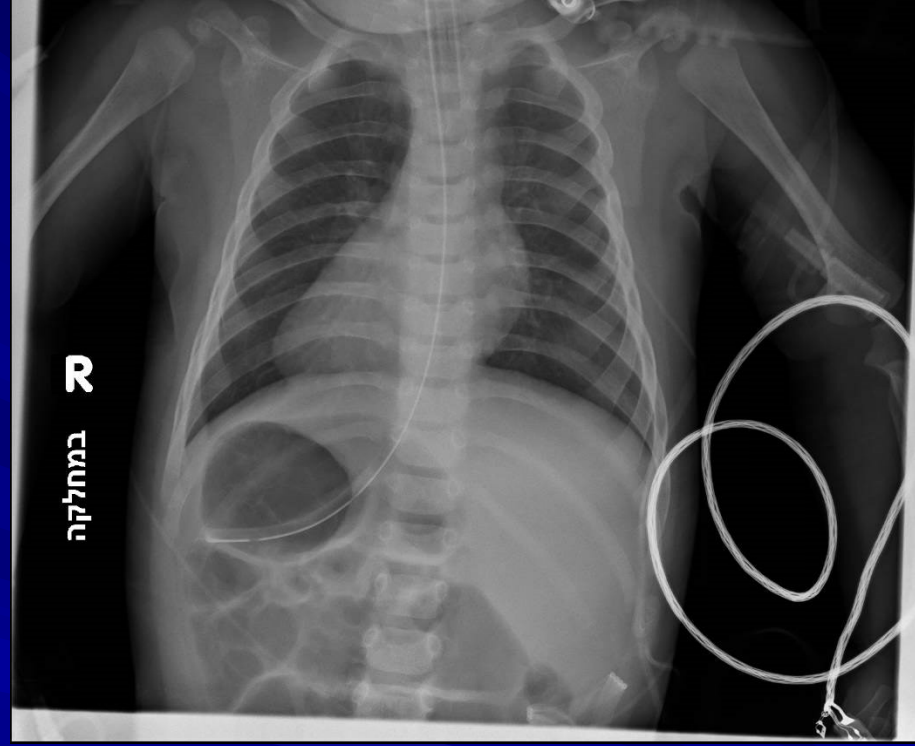
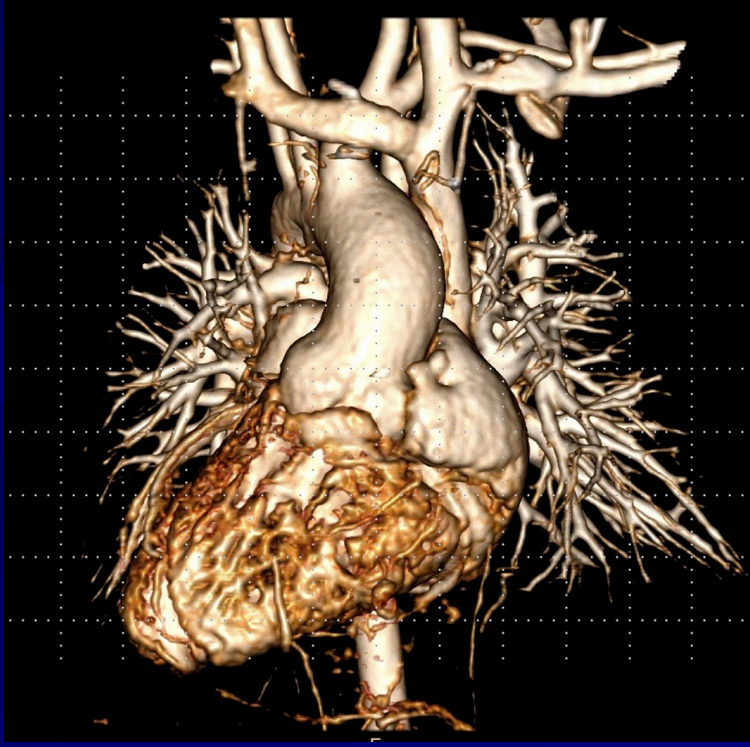


Family 1 Pedigree:



Case Presentation -Patient 2:

- A 4 months old male referred d/t dextrocardia with complex CHD
- Prenatal history uneventful
- Born term, AGA, readmitted at 5 days due to cyanosis
- Echo (Gaza)- Dextrocardia, hypoplastic LT ventricle, PA, VSD treated with Deralin
- Upon arrival: saturation 40% RA, ventilated
- Echo/ CTA- dextrocardia, atrial situs inversus, TOF, PA, non confluent pul. arteries, MAPCAS, abdominal situs inversus
- Underwent BT shunt & unifocalization of MAPCAS

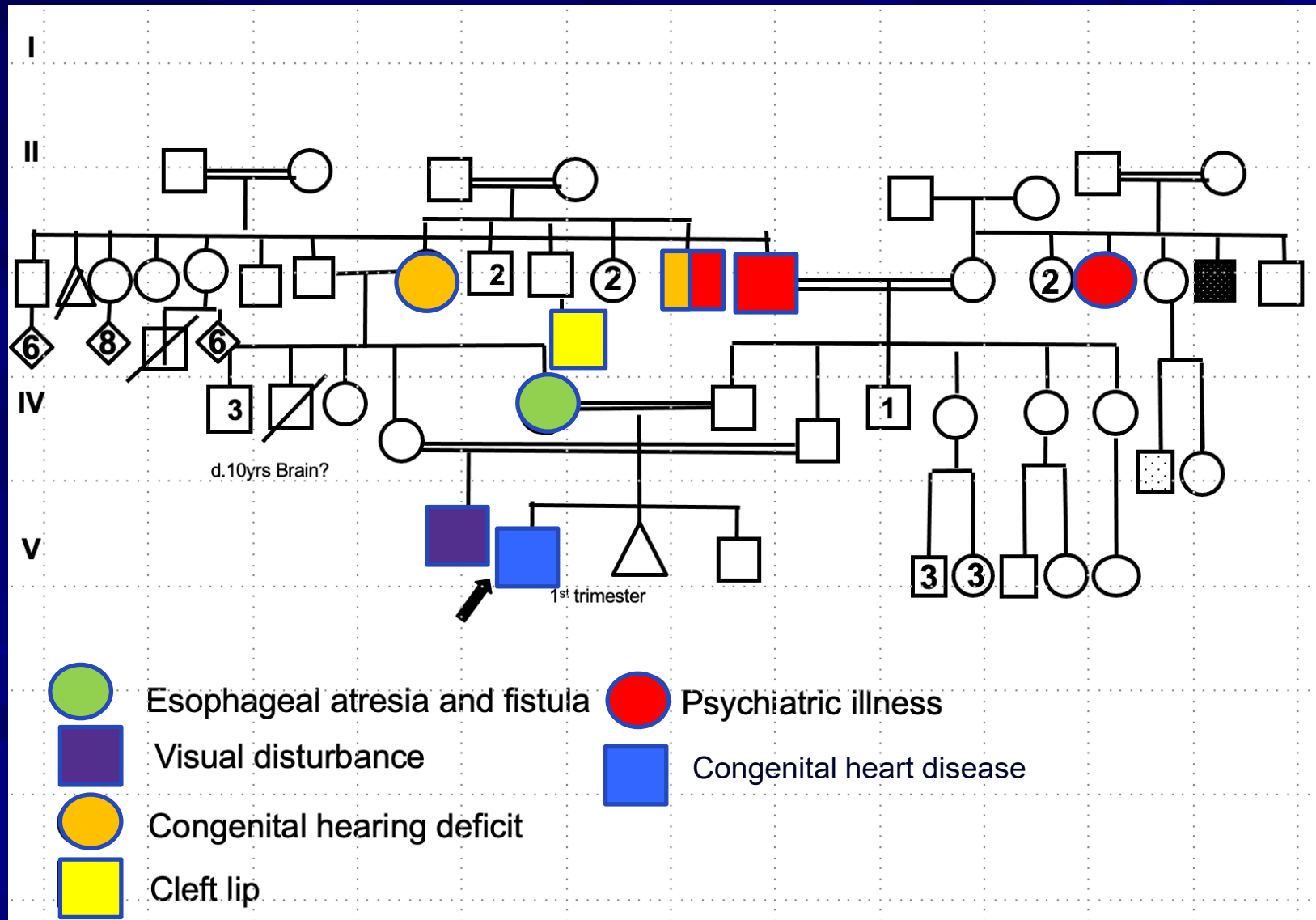


Case Presentation -Patient 2, cont'd:

- Follow up reveals global developmental delay and failure to gain weight
- Physical examination:
 - Microcephaly, -3SD
 - Weight, - 2.5 SD
 - Height- + 1.SD
- Dysmorphic facial features synophrys, long philtrum, thin upper lip, retrognathia, high arched palate



Family 1 Pedigree:



DIAGNOSIS



Exome sequencing:

- Homozygous *DAND5* variant shared by both patients:

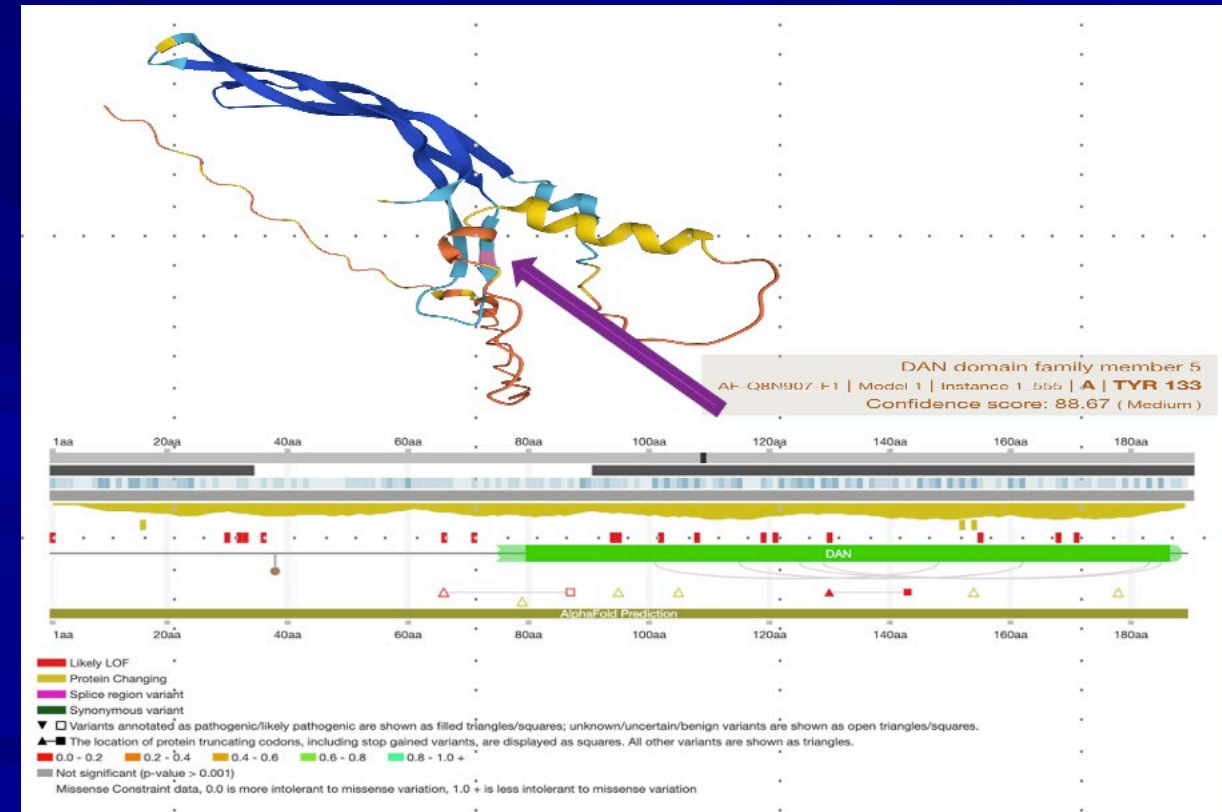
NM_152654.2: c.396_397dup,
p.Tyr133SerfsTer11 exon 2/2

- Predicted to cause early termination of translation of RNA to protein, leading to a truncated protein predicted to undergo nonsense mediated decay



DAND5- DAN DOMAIN FAMILY, MEMBER 5

- Encodes for a member of the Cerberus-related DAN protein family
- Involved in regulating organogenesis, body patterning, and tissue differentiation.
- In mouse, this protein has been shown to bind Nodal and to inhibit the Nodal signaling pathway which patterns left/right body asymmetry.



RESEARCH ARTICLE

Functional studies of *DAND5* in patients with congenital heart disease and laterality defects

Fernando Cristo^{1,2,3,4}, José M. Inácio², Rui Anjos⁷ and José A. Belo^{1,8*}

► Additional supplemental material is published online only. To view, please visit the journal online (<http://dx.doi.org/10.1186/s12881-017-0444-1>).

For numbered affiliations see end of article.

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Original Article

Whole genome sequencing in 56 patients with congenital heart disease and laterality defects

Yoav Ben-Zeev, Maayan Tal Tirosh, Odellia Alvir, Orna Shalev, Asaf Vardi

ABSTRACT
Background and congenital heart disease and laterality defects are often associated with pathogenic variants in the same genes. Objective basis of laterality defects and congenital heart disease in Arab-Muslim population.



COLD SPRING HARBOR
Molecular Case Studies

RAPID COMMUNICATION

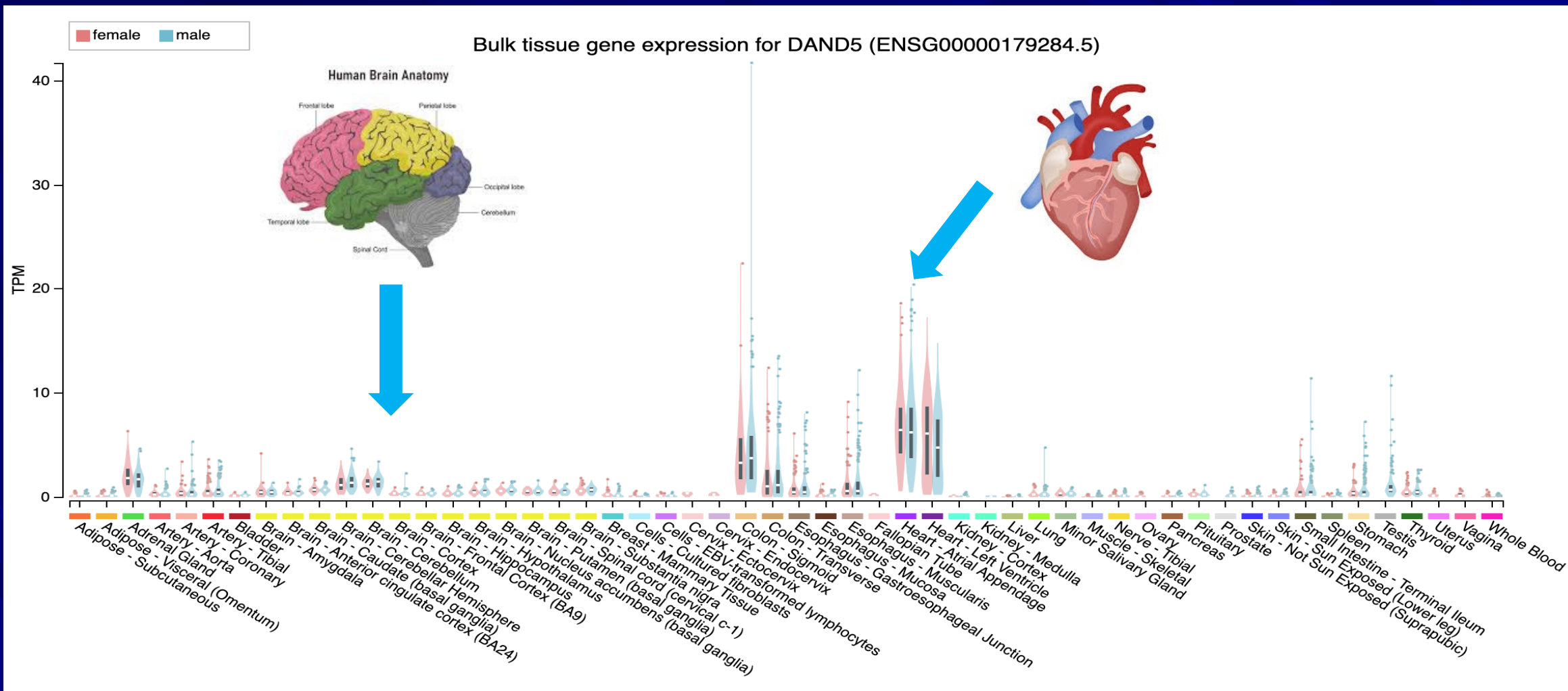
A novel biallelic loss-of-function variant in *DAND5* causes heterotaxy syndrome

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Abstract The majority of heterotaxy cases do not obtain a molecular diagnosis, although pathogenic variants in more than 50 genes are known to cause heterotaxy. A heterozygous missense variant in *DAND5*, a nodal inhibitor, which functions in early development for establishment of right-left patterning, has been implicated in heterotaxy. Recently, the first case was reported of a *DAND5* biallelic loss-of-function (LoF) variant in an individual with heterotaxy. Here, we describe a second unrelated individual with heterotaxy syndrome and a homozygous frameshift variant in *DAND5* (NM_152654.2:c.197del [p.Leu66ArgfsTer22]). Using an in vitro assay, we demonstrate that the *DAND5* c.197del variant is unable to inhibit nodal signaling when compared with the wild-type expression construct. This work strengthens the genetic and functional evidence for biallelic LoF variants in *DAND5* causing an autosomal recessive heterotaxy syndrome.

Bulk Tissue Gene Expression for DAND5



Conclusions:

- DAND5 has been suggested as a candidate gene in heterotaxia and CHD in the heterozygous state (
- Our findings confirm the association of *DAND5* variants and laterality-related CHD in biallelic state among two families.
- Our report, along with a previous report of associated GDD in an affected male may suggest a role of the gene in neuro-developmental processes.
- Further studies are required to decipher the full spectrum of *DAND5* disorders.



ד"ר יואב בולקיא



ד"ר בן פודה-שקד



ד"ר ישי סלם

- קרדיולוגיה ילדים דר' כץ ודר' תירוש והצוות
- מעבדה ביואינפורמטית - דר' בראל וצוותה
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פרופ' יאיר אניקסטר



BRING
THEM
HOME