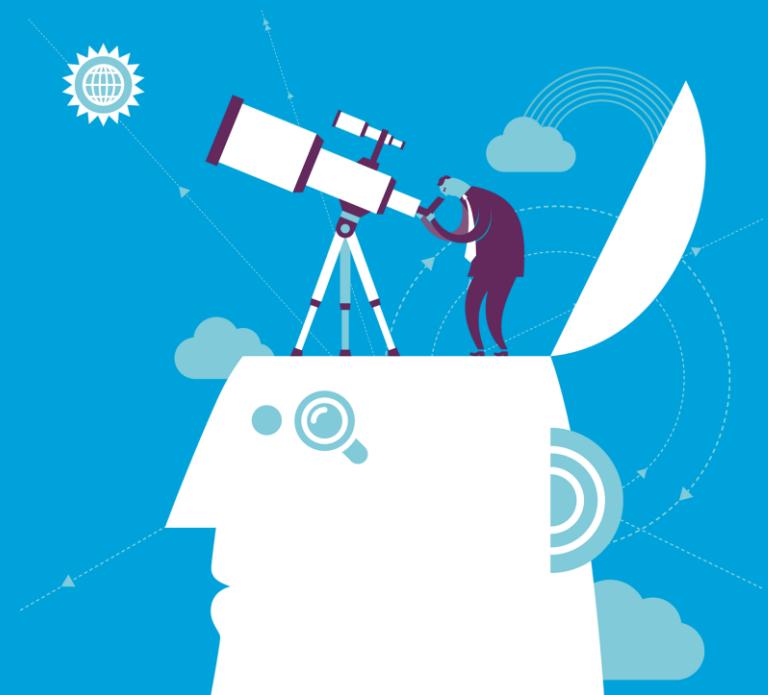


Pathway of a Rare Disease

Joubert Syndrome

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Date: 01-07-2019
Supervisor: Friederike Ehrhart



Introduction - General Information

- Rare hereditary disorder
 - Prevalence: 1 in 80,000-100,000
 - Autosomal recessive inheritance
- >30 genes
- Ciliopathy
 - Primary cilium affected
 - Wnt, SHH
- Molar Tooth Sign
- Motor/intellectual development

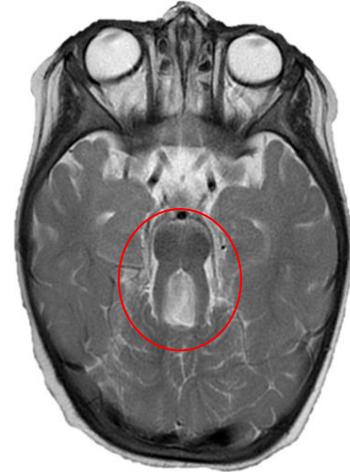


Figure 1. Molar tooth sign.

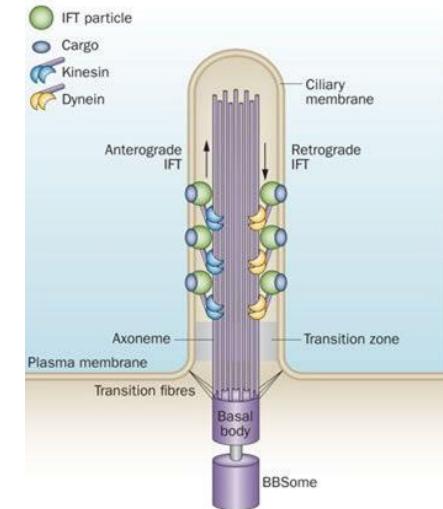
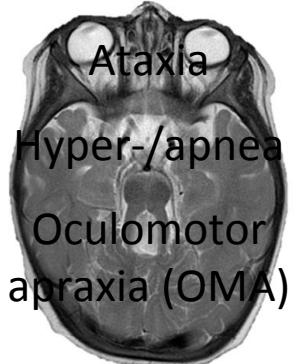


Figure 2. Primary cilium.

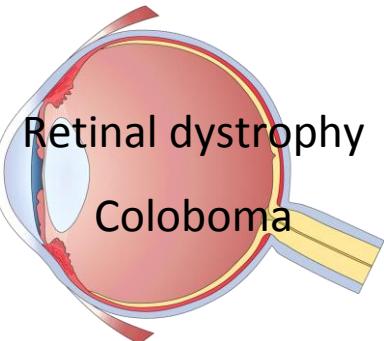
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Introduction - Phenotype

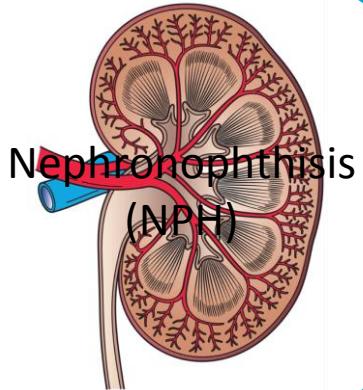
Brain



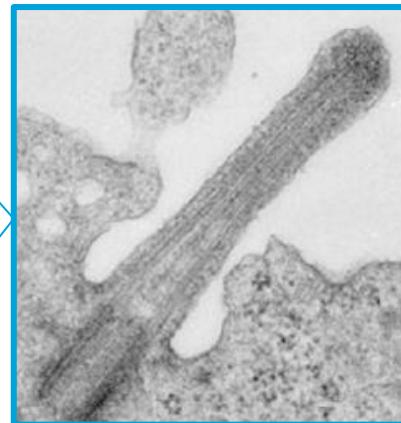
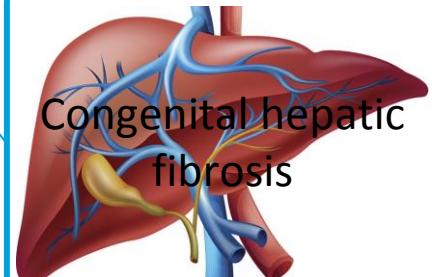
Eye



Kidney



Liver



Aim

Synthesize a visual representation of the pathways underlying Joubert Syndrome pathogenesis.

Methods - Data Collection



- Finding general information



- Overview involved genes
- Facilitation more targeted literature search
- Creation of rough outline functional units + molecular interactions



- Compile and combine known pathways
- Finding new interactions
- Finding experimental data supporting interactions



Methods - Pathway Building



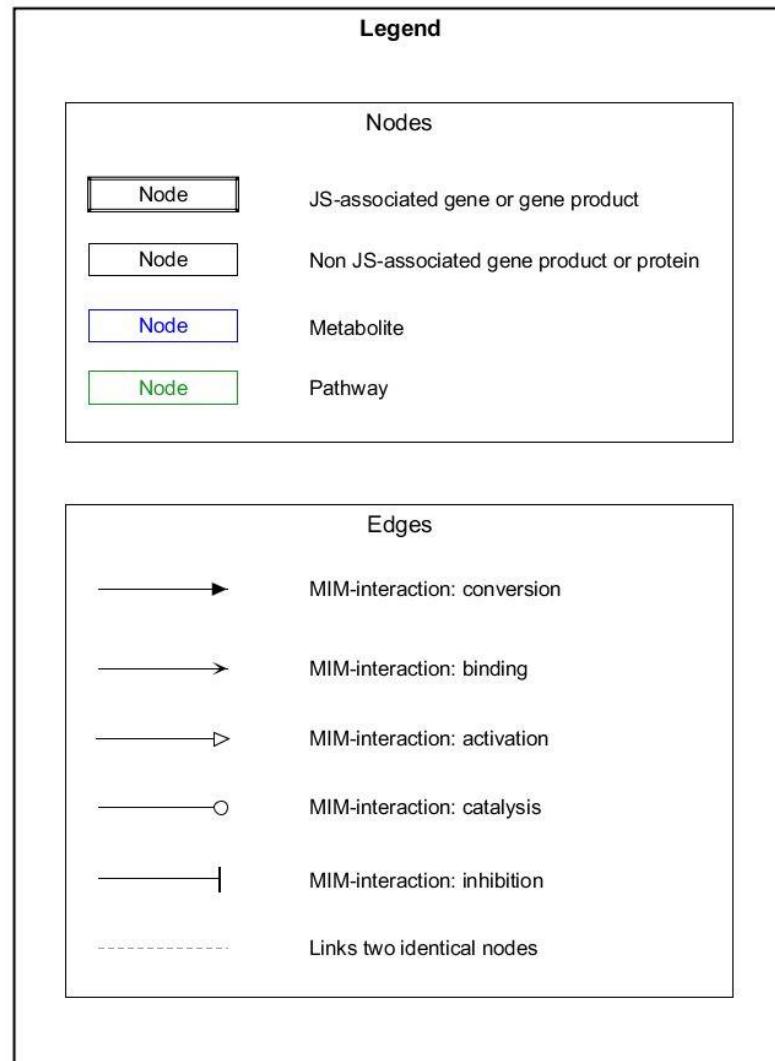
- Pathway synthesis
- Division three functional or structural areas
- MIM notation



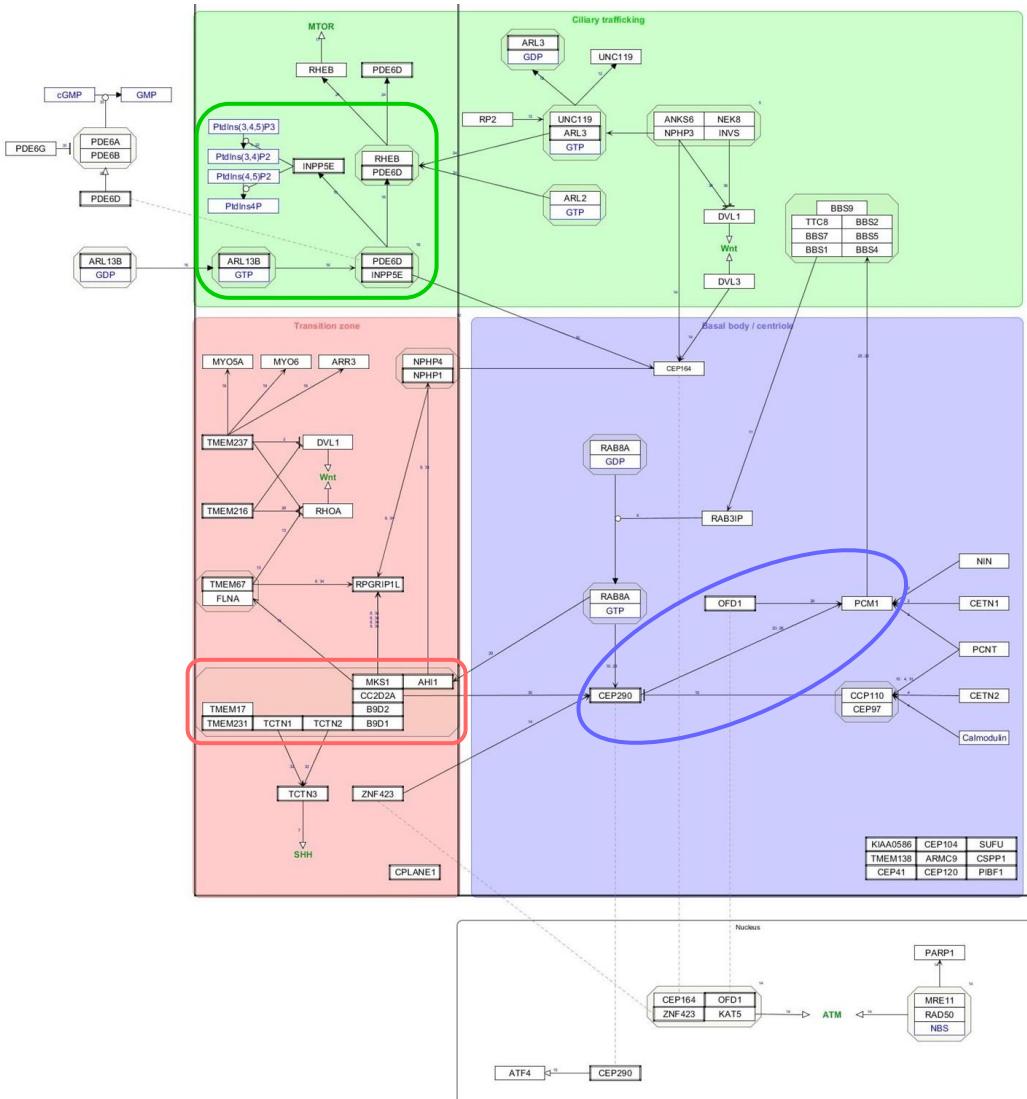
- Annotation nodes
- BridgeDb mapping database

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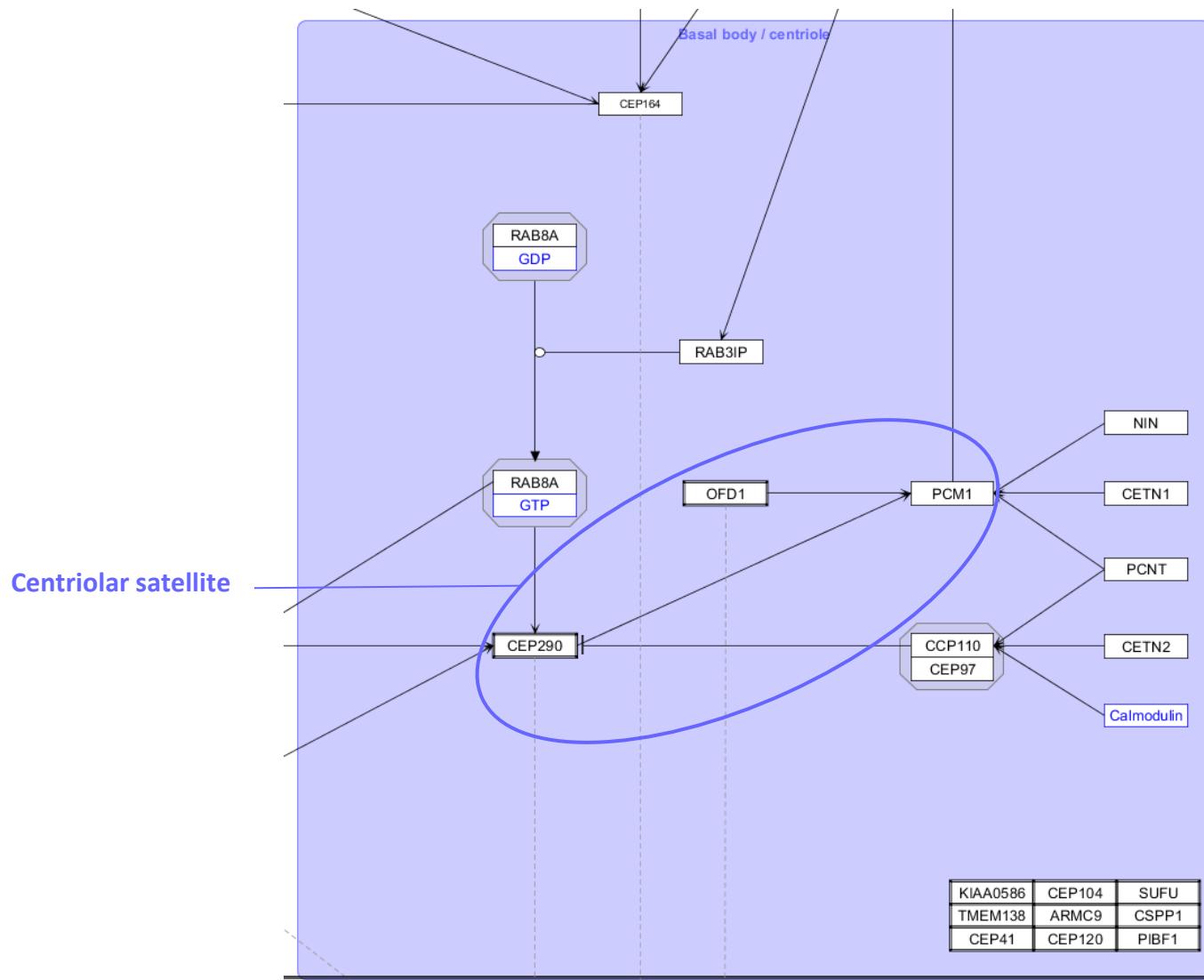
Results - Legend



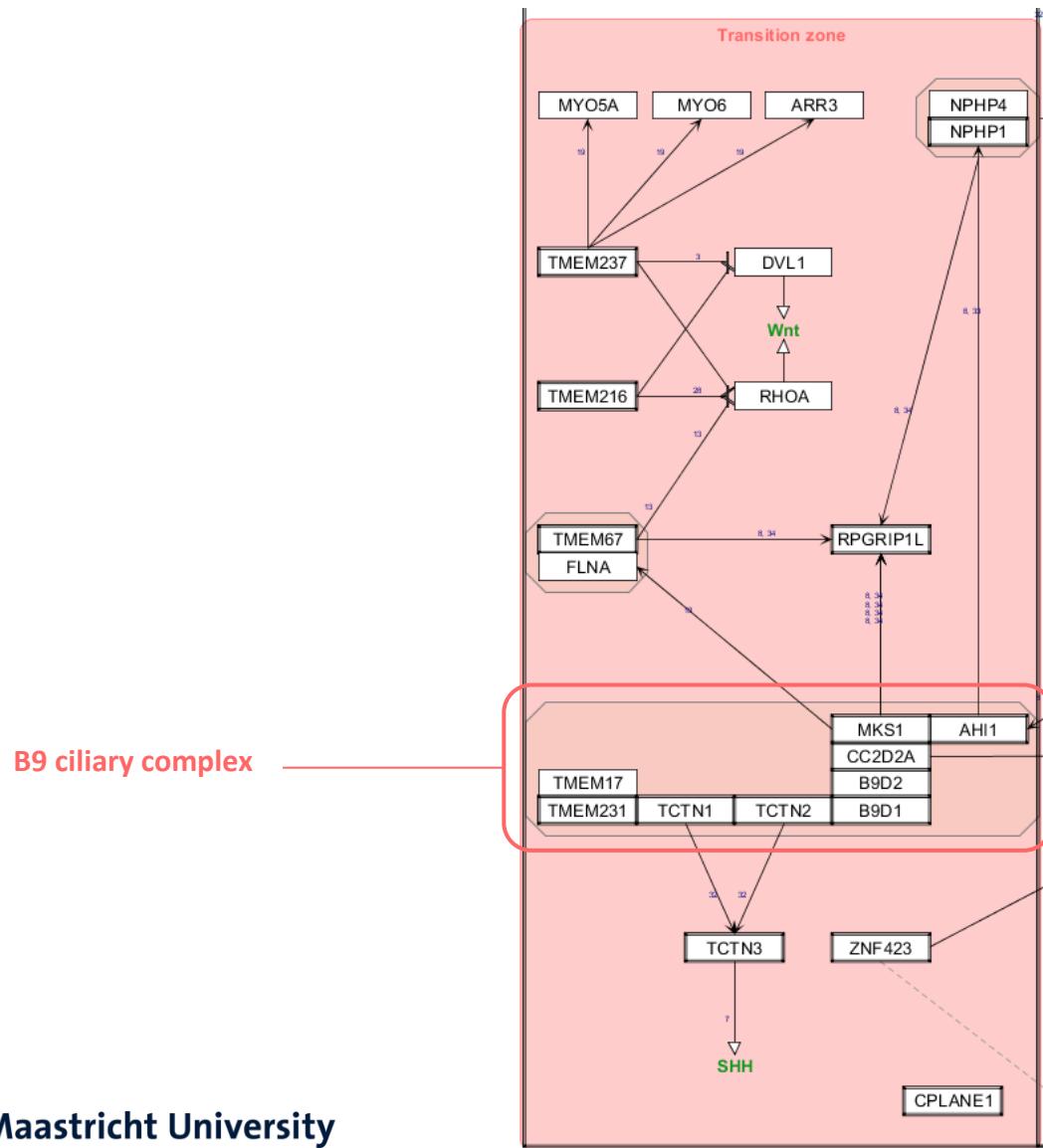
Results - Overall Pathway



Results - Basal body / Centriole

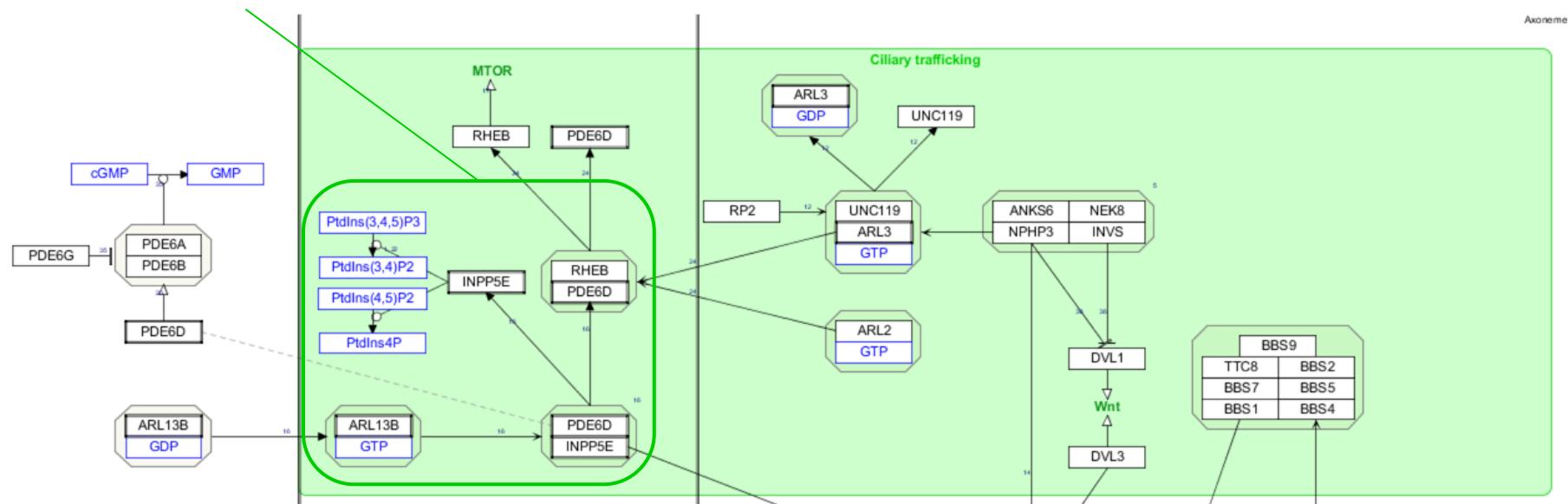


Results - Transition Zone

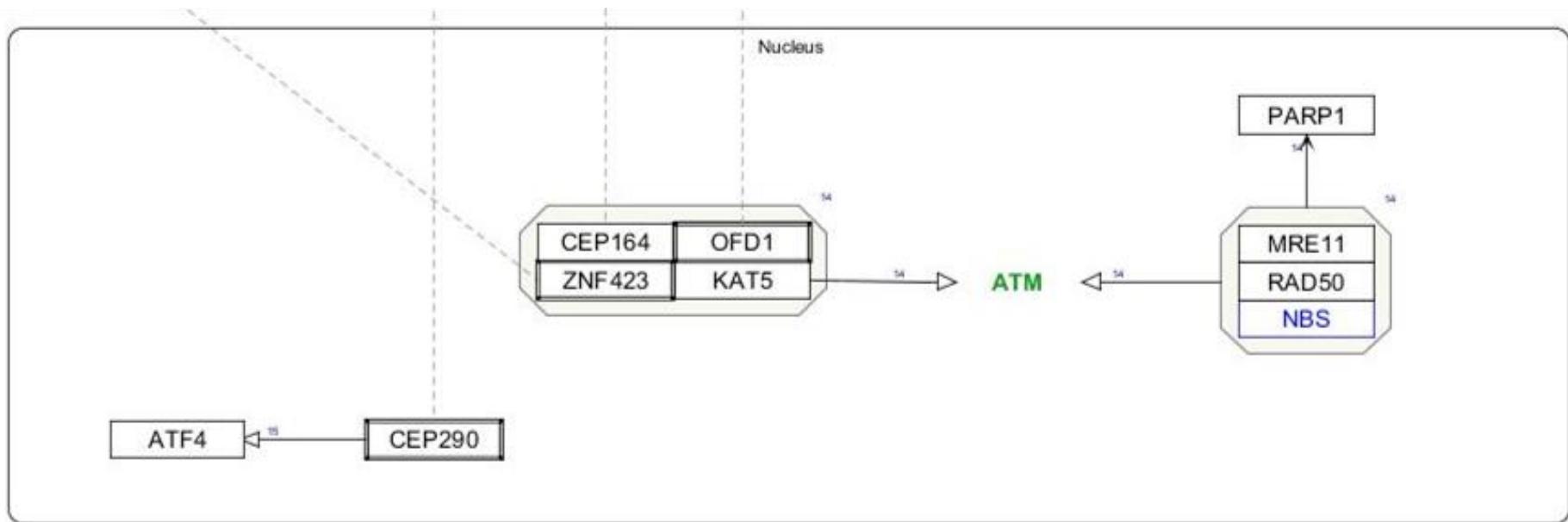


Results - Ciliary Trafficking

ARL13B-PDE6D-INPP5E
signaling network



Results - Nucleus



Conclusion

- Three structural/functional areas
- Components of special interest
 - Centriolar satellites
 - B9 ciliary complex
 - ARL13B-PDE6D-INPP5E signaling network
- Early phases of ciliogenesis
- DNA damage response

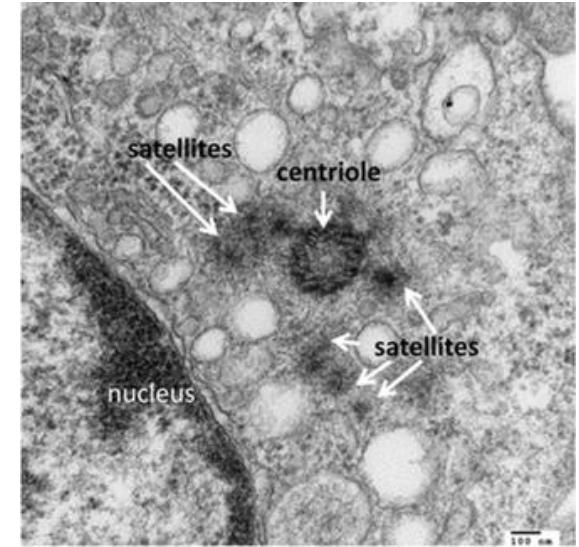


Figure 4. Microscopic image of centriolar satellites.

Discussion

- Possible new insight
 - Overview involved genes
 - Many genes are functionally connected
- Exclusion certain genes
 - Lack of knowledge/relevance
- Downstream pathways
- Links to phenotype
 - Meckel Syndrome
- DNA damage response

Gene Name	Gene Location	JS Type
ARMC9	2q37.1	JBTS30
CPLANE1	5p13.2	JBTS17
CEP120	5q23.2	JBTS31
CEP41	7q32.2	JBTS15
CSPP1	8q13.1-q13.2	JBTS21
SUFU	10q24.32	JBTS32
TMEM138	11q12.2	JBTS16
PIBF1	13q21.3-q22.1	JBTS33
KIAA0586	14q23.1	JBTS23

Thank you for your attention!

Are there any questions?