Genetics inTuberous Sclerosis Complex

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Genetics 101







DNA sequence variants



- "Normal" or "Reference" sequence
- Anything difference is considered a "variant"
- Some variants are known to be benign
- Some variants are known to be pathogenic (mutations)
- Some variants are unknown







Yang et al. Communications. January 2021









































Second random hit in the same cell





Second random hit in the same cell























Why mTOR Inhibitors can be affective against a number of symptoms in TSC

Random / Environmental Mutations are all second hits





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mTOR pathway







Variable expression

- Huge spectrum of severity
- Between families and within families

TSC1 vs TSC2



Contiguous Gene Deletion



TSC + polycystic kidney disease



Clinical Genetic Testing



Not all genetic testing is created equal

- Sanger sequencing
 - Older
 - Not reliable under ~20% mosaicism
- Next Generation Sequencing (NGS)
 - Can have a low depth read similar to Sanger
 - Can have a much higher depth read enabling it to find mosaicism under 1%
- When getting deletion data from NGS data, possibility to miss small deletions
- Different labs will analyze different areas around the genes that may or not be relevant



Current clinical genetic testing: Sequencing and deletion analysis of TSC1 and TSC2





GeneTherapy in TSC





TSC is **ALWAYS** genetic

Sometimes it is INHERITED

Other times it is sporadic



TSC Recurrence Risk



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QUESTIONS?

