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Diagnosis of Mosaic RASopathy in a Child with Rhabdomyosarcoma

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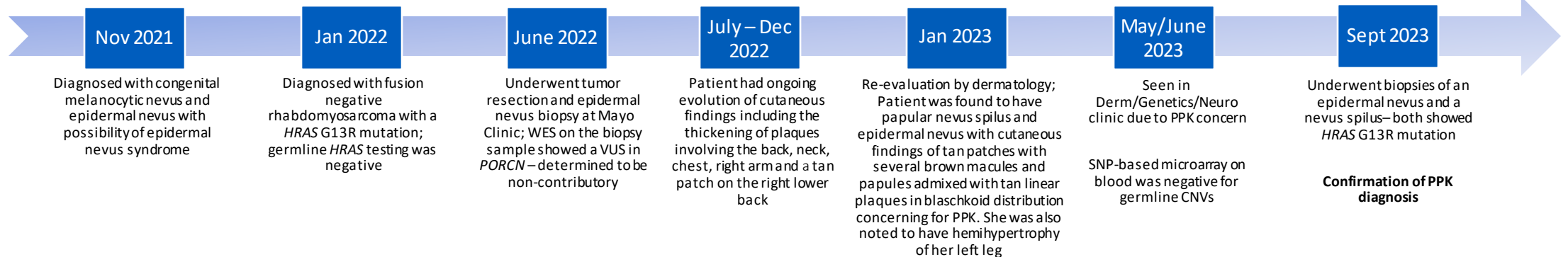
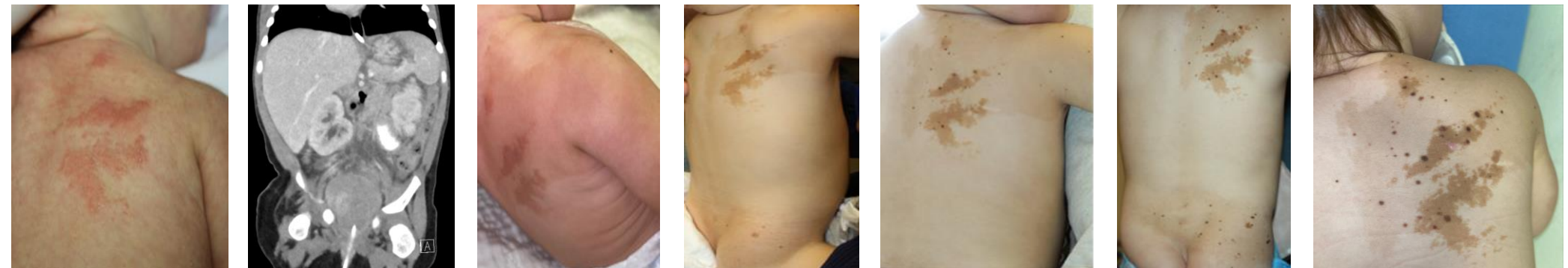
Background

- Phacomatosis pigmentokeratotic (PPK) is a rare subtype of epidermal nevus syndrome characterized by the co-existence of a sebaceous nevus and speckled lentiginous nevus described in ~30 cases in literature.
- PPK is now recognized as a mosaic RASopathy due to a post-zygotic mutation in the Ras-Raf-MEK-ERK pathway.
- RAS variants are also known to contribute to tumorigenesis in some pediatric cancers, including rhabdomyosarcoma.

Patient Presentation

- The patient is a former 32-week premature female who presented to dermatology clinic at 2 months of age for evaluation of skin-colored to pink linear plaques on her right back and shoulder and a brown macule on her right arm.
- She was diagnosed with a congenital melanocytic nevus and epidermal nevus with the possibility of having an epidermal nevi syndrome.
- At 4 months of age, she was hospitalized for obstructive renal failure secondary to a pelvic mass.
- Biopsy of the mass was diagnostic for fusion negative rhabdomyosarcoma. Genetic testing of the tumor revealed a copy-neutral loss of heterozygosity of 11p15.5 and a *HRAS* G13R pathogenic mutation.
- Germline testing for Beckwith-Wiedemann Syndrome and *HRAS* were negative, which began the journey of finding a unifying diagnosis between her skin findings and cancer diagnosis.

Clinical Course



Genetic testing performed throughout patient's clinical course	
Test	Results
Cancer Microarray on tumor sample	<ul style="list-style-type: none"> Gain of chromosome 2, 3, 5, 8, and 11 Copy-neutral loss of heterozygosity (CN-LOH) of 11p15.5
FISH analysis on tumor sample	<ul style="list-style-type: none"> <i>FOXO1</i> fusion negative
Beckwith-Wiedemann Syndrome testing on blood	<ul style="list-style-type: none"> Negative – normal methylation pattern and copy number of ICI1 and ICI2
Next Generation Sequencing (NGS) multi-gene panel on blood	<ul style="list-style-type: none"> Negative for sequence or copy number variants (CNVs) in <i>CDKN1C</i>, <i>DICER1</i>, <i>HRAS</i>, <i>NF1</i>, <i>TP53</i>
FoundationOne® Heme (comprehensive genomic profiling) on tumor sample	<ul style="list-style-type: none"> <i>HRAS</i> G13R pathogenic mutation
Comprehensive NGS Panel on blood	<ul style="list-style-type: none"> Negative for reportable variants
Whole exome sequencing (WES) on epidermal nevus biopsy sample	<ul style="list-style-type: none"> Variant of uncertain significance (VUS) in <i>PORCN</i>
Single nucleotide polymorphism (SNP)-based microarray on blood	<ul style="list-style-type: none"> Negative for germline CNVs
WashU's PIK3CA-Related Overgrowth Spectrum Disorders Panel: Comprehensive Panel via NGS on 2 skin biopsy samples	<ul style="list-style-type: none"> Sample 1 – <i>HRAS</i> G13R pathogenic mutation, VUS in <i>PIEZO1</i> Sample 2 – <i>HRAS</i> G13R pathogenic mutation, VUS in <i>HRAS</i> and <i>PIEZO1</i>

Conclusions

- Mosaic RASopathies remain a diagnostic challenge as tissue involvement can be varied and evolve over time, DNA sequencing of the blood is often negative, and phenotypes can depart from germline RASopathies.
- With only case reports and small case series describing an association between cutaneous mosaic RASopathies and rhabdomyosarcoma, there is a need for further awareness of this association as it may impact treatment decisions and necessitate multidisciplinary care.
- Unfortunately, the risk of malignancies for patients with mosaic RASopathies is unknown and therefore cancer screening guidelines are not well established for this patient population.