

Pediatric Haematology and Oncology, Frankfurt, Germany, ⁷⁶Division of Hematology and Oncology, The Hospital for Sick Children, University of Toronto, Toronto, ON, Canada

BACKGROUND: Constitutional mismatch repair deficiency syndrome (CMMRD) is a severe cancer predisposition syndrome resulting in early onset central nervous system (CNS) and other cancers. International guidelines for surveillance exist but no study has systematically evaluated the efficacy of this protocol. **METHODS:** We surveyed all confirmed CMMRD patients in the International Replication Repair Deficiency Consortium. A surveillance protocol consisting of frequent biochemical, endoscopic and imaging (CNS and total body MRI) studies were employed. Survival analyses and efficacy of each method were assessed. **RESULTS:** Surveillance data were collected from 105 CMMRD individuals from 41 countries. Of the 193 malignant tumors, CNS malignancies were the most common (44%). The surveillance protocol uncovered 49 asymptomatic tumors including 16 glioblastomas and medulloblastomas. Five-year overall survival was 89% for tumors discovered by surveillance, and 61% for symptomatic tumors ($p < 0.004$). Similarly, 5-year survival was 82%±11% and 24%±6% for surveillance and non-surveillance of brain tumors ($p = 0.005$). Yearly total body and q6 month brain MRI detected asymptomatic cancers in all but 3 symptomatic CNS gliomas. These were tumors uncovered when time between scans was >6 months as per protocol. Finally, of the low grade tumors identified asymptotically, 5 were low grade gliomas. All of the low grade gliomas, which were not resected transformed to high grade tumors at a median of 1.6 ± 0.9 years. **CONCLUSION:** These data support a survival benefit in CMMRD patients undergoing a surveillance protocol. Adherence to protocol and resection of lower grade lesions may improve survival for patients with CNS tumors.

RARE-18. GENETIC EVALUATION IN PATIENTS WITH CHOROID PLEXUS TUMORS

Milena Oliveira, Nasjla Silva, Andrea Cappellano, Daniela Almeida, Sergio Cavalheiro, Patricia Dastoli, Frederico Silva, and Fernanda Lima; IOP/GRAACC/UNIFESP, São Paulo, São Paulo, Brazil

INTRODUCTION: Choroid plexus tumors (CPT) are rare intraventricular neoplasms of epithelial origin. They usually occur in the 2nd year of life, corresponding to 0.4–0.6% of intracranial tumors in this age group. They are sub classified, according to WHO 2016, in choroid plexus carcinoma (CPC), atypical choroid plexus papilloma (ACPP) and choroid plexus papilloma (CPP). Li-Fraumeni syndrome (LFS) is present in 50% of patients with CPC. In Brazil, the TP53 p.R337H mutation affects 0.3% of the population in the South/Southeast. **OBJECTIVE:** Evaluate the incidence of genetic mutations in patients with choroid plexus tumors and therefore the importance of genetic evaluation. **PATIENTS AND METHODS:** Between 1992–2019, 38 patients were diagnosed with CPT in our institution, 23 with CPC. From 2012, 21 patients were referred for genetic evaluation, 16 of which had CPC (2 had previously CPP). Positive family history for neoplasms was present in 87.5%; 37.5% compatible with LFS, 50% of them with mutations. All the patients with positive, but unspecific, family history of neoplasms, had pathogenic mutation. The molecular investigation of the TP53 gene in patients with CPC was performed and positive in 56.2%: R337H (5 patients), R110C, R158H, H179R, R196* (1 patient each). Of those with R337H, p53 protein immunohistochemistry resulted in 90–100% positivity. One of the patients with CPP that evolved to CPC had the H179R mutation. Clinical course was similar among them, and with those without mutations. **CONCLUSION:** These results confirm the need for genetic evaluation in patients with choroid plexus tumors for adequate therapeutic management and long-term follow-up.

RARE-19. PEDIATRIC HIGH GRADE GLIOMA WITH DNA REPAIR PATHWAY ABERRATIONS, CLINICAL CHARACTERISTICS AND OUTCOME

Muhammad Baig, David McCall, Tyler Moss, David Sandberg, Gregory Fuller, Susan McGovern, Arnold Paulino, Amer Najjar, Joya Chandra, Soumen Khatua, and Wafik Zaky; MD Anderson Cancer Center, Houston, TX, USA

DNA mismatch repair machinery is an integral part of the human genome and its defect has been involved in tumorigenesis and treatment resistance. Heterozygous monoallelic germline loss of function in MLH-1, MSH-2, MSH-6 or PMS-2 is involved in Lynch syndrome, whereas biallelic mutations cause constitutional mismatch repair deficiency (CMMRD) which is associated with hematologic malignancies and glioblastoma. We report here the clinical characterization and molecular analyses of 7 patients who presented with gliomas and MMR machinery aberrations. Two patients had a clinical diagnosis of NF-1 with dermatologic stigmata, of whom one patient has CMMRD and the other has Lynch syndrome. Two patients had a known family history of Lynch syndrome upon their diagnosis of glioma. Three patients with high-grade glioma and negative family history, 2 had

germline mutations in MMR genes, and one had numerous mutations including MMR genes with microsatellite instability. Patients were initially treated with chemotherapy and radiation for high-grade gliomas (HGG); 5/7 had progression. Median time to progression was 12 months (range: 5–52), and median time from progression to death was 7 months (range: 2–25). One patient had low-grade glioma initially but progressed to HGG and is currently on therapy. Another patient treated with temozolomide and radiation is currently receiving maintenance therapy without any disease recurrence. Although the literature data on brain tumors with MMR deficiency is limited, these consistently show that MMRD-associated gliomas are treatment-resistant and have a dismal outcome. Collaborative efforts are needed to better understand this subgroup of pediatric HGG and to define optimal treatment strategy.

RARE-20. MALIGNANT PERIPHERAL NERVE SHEATH TUMOR OF A CRANIAL NERVE IN AN INFANT WITH NEUROCUTANEOUS MELANOSIS

Lacey Carter, Naina Gross, Rene McNall-Knapp, and Jo Elle Peterson; University of Oklahoma Health Sciences Center, Oklahoma City, OK, USA

At one month of age, a female presented with a giant congenital nevus along lower back and thighs and hydrocephalus. A ventriculoperitoneal shunt was placed. An MRI was done at six months, initially reported as normal. At eleven months of age, five months after original MRI, patient presented with dysconjugate gaze and lethargy. MRI showed new $3.8 \times 3.7 \times 3.4$ cm right cerebellopontine angle mass extending into Meckel's cave and foramen ovale along with leptomeningeal disease extending from the mass along the entire length of the spinal cord. Retrospective review of prior MRI revealed subtle leptomeningeal enhancement concerning for neurocutaneous melanosis (NCM). Given the leptomeningeal disease, family elected for open biopsy and debulking of lesion instead of aggressive resection. Histologically, the mass showed hypercellular spindle cell neoplasm with mitotic activity and necrosis mixed with remnants of normal cranial nerve. GFAP was negative, excluding a glioma. HMB-45, MITF, panmelanoma, and Melan-A were negative, excluding melanoma. A negative myogenin stain ruled out ectomesenchymoma. S-100 protein and SOX-10 positivity with variable loss of staining for trimethylation of histone H3 K27 were indicative of malignant peripheral nerve sheath tumor (MPNST). Given the course of the mass, trigeminal nerve MPNST was presumed. Given the poor prognosis of intracranial MPNST and NCM, family elected to forgo treatment and was discharged with hospice. She died 25 days after surgery. Cranial nerve MPNST is rare. MPNST in patients with NCM has not previously been reported to our knowledge.

RARE-21. CANCER SPECTRUM IN GERMLINE SUFU MUTATION CARRIERS: A COLLABORATIVE PROJECT OF THE SIOPE HOST GENOME WORKING GROUP

Léa Guerrini-Rousseau¹, Sebastian Waszak², Franck Bourdeaut³, Olivier Delattre³, Nicola Dikow⁴, Christelle Dufour¹, Amar Gajjar⁵, Jacques Grill¹, Steffen Hirsch⁴, Saskia Hopman⁶, David Jones⁷, Majoline Jongmans⁶, Andrey Korshunov⁴, Christian Kratz⁸, Lucie Lafay-Cousin⁹, Julien Masliah³, Till Milde¹⁰, Paul Northcott⁵, Kristian Pajtlér⁷, Stefan Pfister⁷, Stéphanie Puget¹¹, Marie Agnès Rame Collonge¹², Giles Robinson¹³, Eric Sariban¹⁴, Nicolas Sevenet¹⁵, Miriam Smith¹⁶, Dominik Sturm¹⁰, Hélène Zattara¹⁷, Pascale Varlet¹⁸, Gareth Evans¹⁹, and Laurence Brugères¹; ¹Gustave Roussy, Villejuif, France, ²EMBL, Heidelberg, Germany, ³Institut Curie, Paris, France, ⁴Medicine University, Heidelberg, Germany, ⁵St. Jude, Memphis, TN, USA, ⁶UMC, Utrecht, Netherlands, ⁷KITZ, Heidelberg, Germany, ⁸MH, Hannover, Germany, ⁹Alberta Children Hospital, Calgary, AB, Canada, ¹⁰DKFZ, Heidelberg, Germany, ¹¹APHN Neckar, Paris, France, ¹²CHU, Besançon, France, ¹³St. Jude, Memphis, TN, USA, ¹⁴Hôpital Universitaire des Enfants Reine Fabiola, Bruxelles, Belgium, ¹⁵Institut Bergonié, Bordeaux, France, ¹⁶University, Manchester, United Kingdom, ¹⁷APHM, Marseille, France, ¹⁸St. Anne Hospital, Paris, France, ¹⁹St. Mary's Hospital, Manchester, United Kingdom

BACKGROUND: Little is known about cancer risk associated with pathogenic germline *SUFU* variants. **METHODS:** Data of all previously published and 25 still unpublished patients with a pathogenic germline *SUFU* mutation were compiled. **RESULTS:** 124 patients in 67 families were identified, most of them ascertained after the occurrence of a medulloblastoma (MB) or as part of Gorlin syndrome cohorts. Overall, 30 patients were healthy carriers and 94 patients developed a total of 129 tumors (up to 4 tumors/patient): 68 MBs, always as first tumor (median age at diagnosis: 1.5yr [0.1–5]), 22 patients with at least 1 basal cell carcinoma (BCC) (median 10/patient) (median age at first BCC: 43yr, [17–52]), 15 meningiomas (median age 43yr, [13–72]), 7 ovarian stromal/fibrous tumors (median age 12yr [5–34]), and 17 other tumors including 5 sarcomas (median age: 50yr [7–79]). Median age at last follow-up was 30yr. Nineteen patients died, including 11 from MB. Second malignancies were diagnosed