GENOMIC PROFILING OF PEDIATRIC TUMORS: A SINGLE INSTITUTION EXPERIENCE

Aubrey Swilling¹, Robin Pham¹, Tyler Hamby², Anish Ray³

¹Texas College of Osteopathic Medicine, UNT Health Science Center, Fort Worth, TX ²Department of Research Operations, Cook Children's Health Care System, Fort Worth, TX ³Department of Pediatric Hematology/Oncology, Cook Children's Health Care System, Fort Worth, TX

PURPOSE

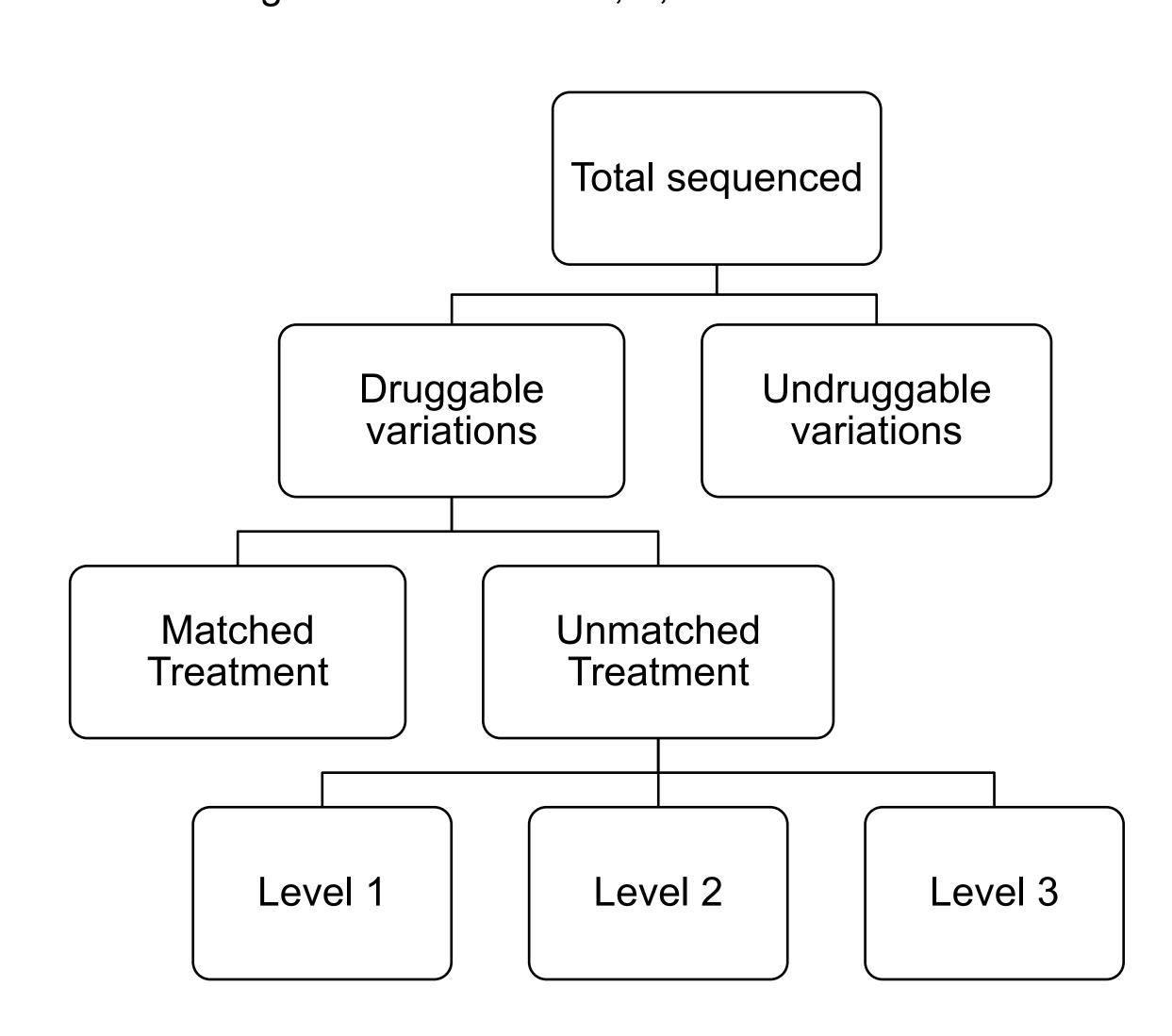
• The goal of this project is to study the prevalence of oncogenic variants and potential utility of targeted therapy among pediatric cancers.

BACKGROUND

- Outcomes remain poor for relapsed malignancies.
- Whole genome sequencing in an emerging tool to identify driver genetic alterations.¹⁻⁵

METHODS

- Retrospective chart review was conducted on 82 patients (2013-2019) at Cook Children's Medical Center.
- Tumor samples were sent to Foundation Medicine, Inc., which identified:
 - Genetic variants and available targeted therapies
 - Programmed death-ligand 1 (PD-L1) expression
 - Tumor mutation burden (TMB)
 - Microsatellite instability (MSI)
- The druggable recommendations were then stratified by clinical significance to level 1, 2, and 3.



RESULTS

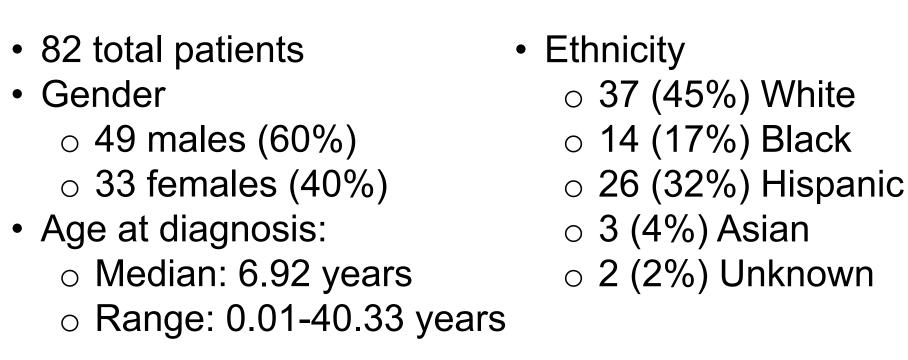
Level

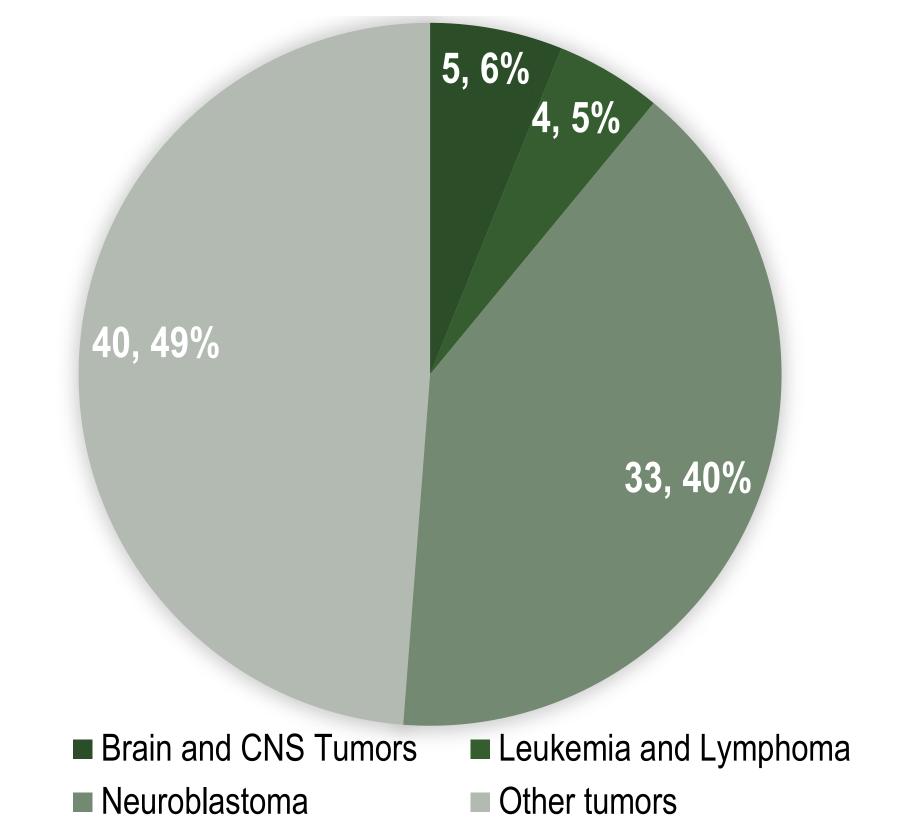
Brain/CNS Tumors

41

Variable

Diagnosis





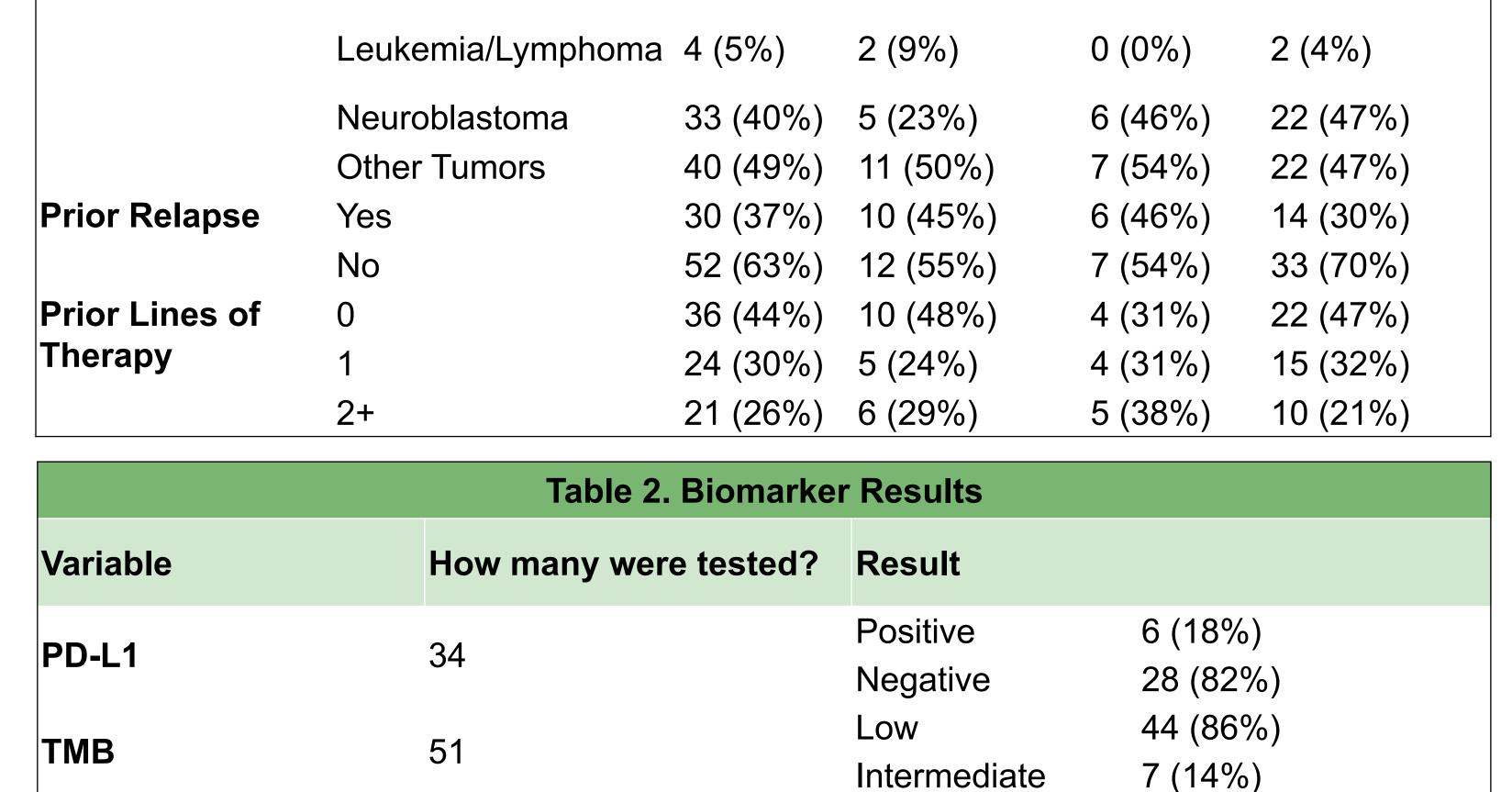


Table 1. Descriptive statistics for diagnostic and treatment variables stratified by

alteration group

(N=82)

Druggable (N=35)

Unmatched

4 (18%)

Matched

41 (100%)

0 (0%)

(N=13)

Undruggable

(N=47)

1 (2%)



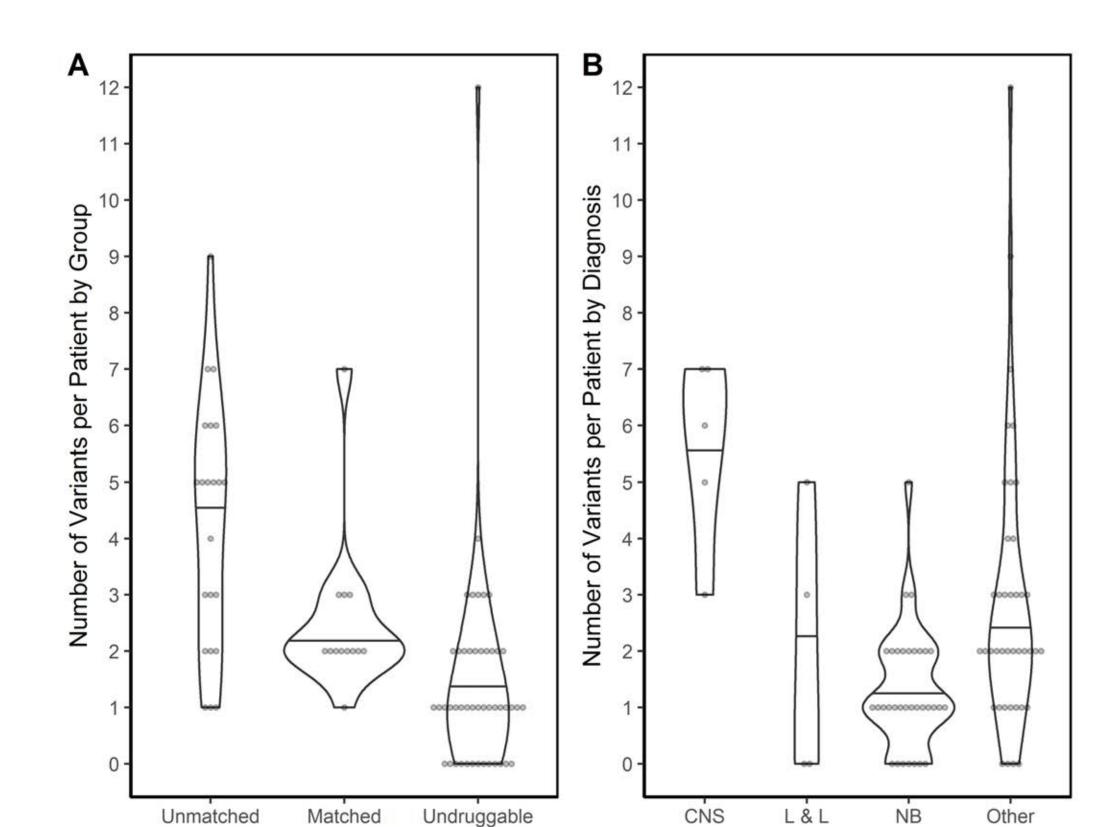
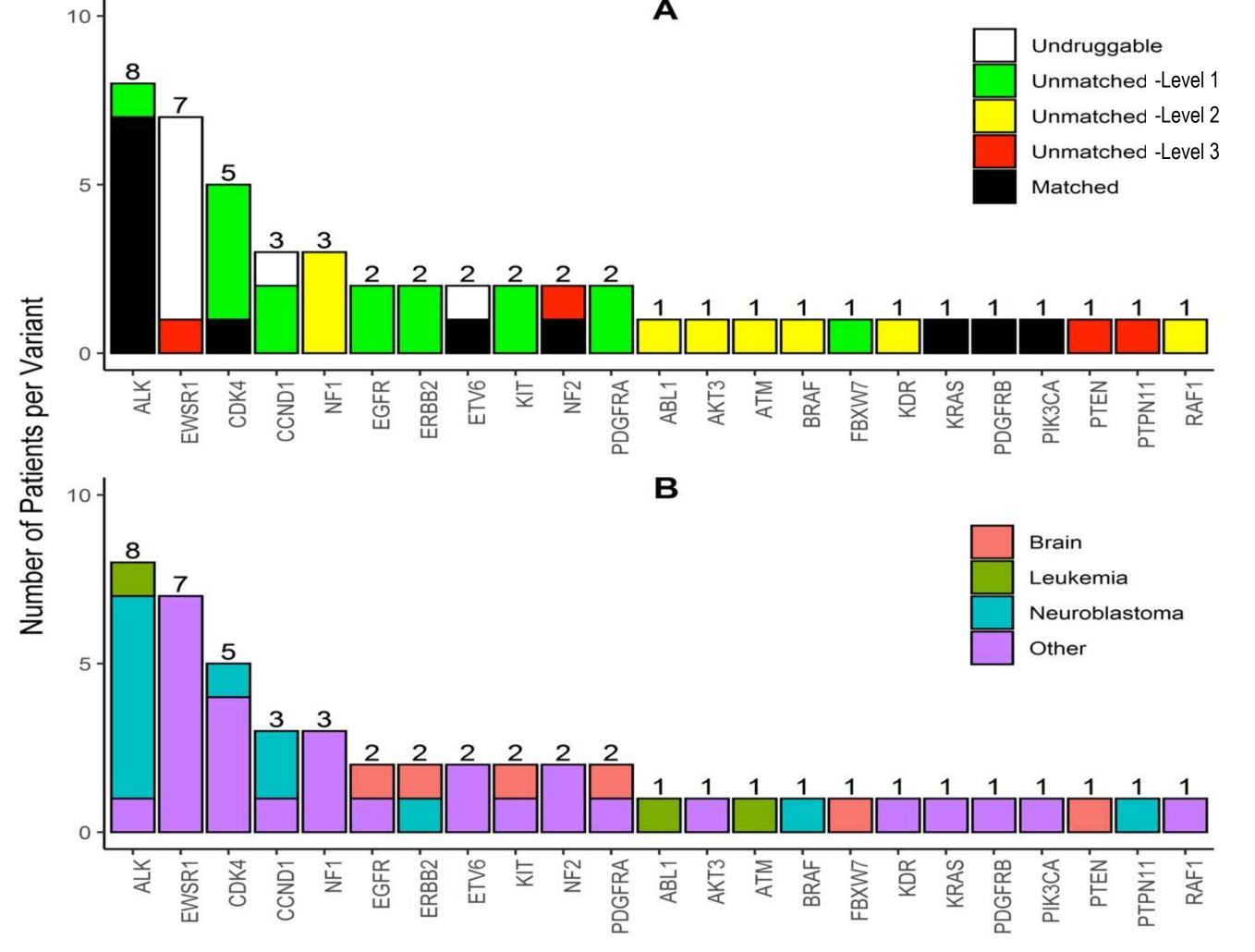


Fig. 2: Violin and dot plots of number of mutations per patient stratified by (A) mutation group and (B) diagnosis. Horizontal lines indicate median values.

NB: Neuroblastoma
L & L: Leukemia and lymphoma

CNS: CNS and brain tumors **Other**: Other solid tumors



Stable

Unstable

Fig. 3: Bar plot of number of patients per mutation stratified by (A) Alteration group and (B) Diagnosis for all druggable mutations.

RESULTS

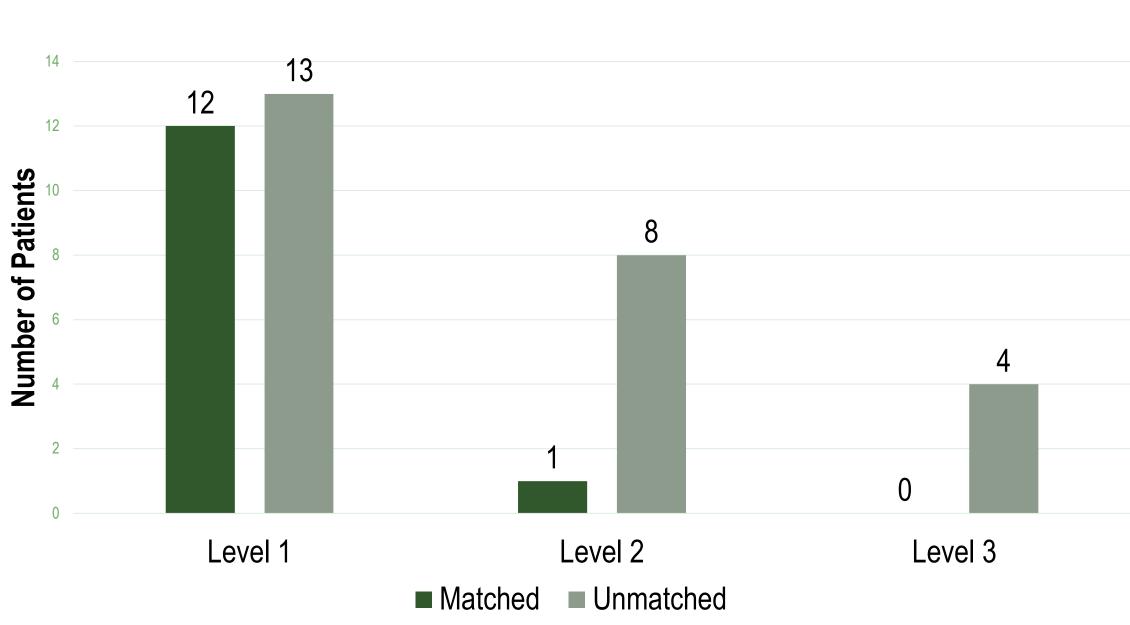


Fig. 4: Bar plot of matched and unmatched variants stratified by level 1, 2, and 3.

DISCUSSION

- Our study showed 35 of 82 (43%) patients had ≥ 1 potentially targetable alterations, which is comparable to results from other pediatric studies⁶⁻¹¹.
- 13 patients received targeted therapy; 12/13 were Level 1 recommendations, and all were off-label drug use.
- Three patients experienced adverse effects from targeted therapy.
- Of those positive for PD-L1, only one patient received pembrolizumab.

CONCLUSION

 This study highlights how tumor profiling can be utilized to identify potential molecular targets and biomarkers in pediatric cancers.

REFERENCES

- Dorris, K., et al., A comparison of safety and efficacy of cytotoxic versus molecularly targeted drugs in pediatric phase I solid tumor oncology trials. Pediatr Blood Cancer, 2017. 64(3).
 Dorris, K. M. O., et al. "A Comparison of Safety and Efficacy of Cytotoxic versus Molecularly Targeted Drugs in Pediatric Phase I Solid Tumor Oncology Trials." Journal of Clinical Oncology, vol. 29, no. 15_suppl, 2011, pp. 9543–9543., doi:10.1200/jco.2011.29.15_suppl.9543.
 Hirshfield, Kim M. "Clinical Actionability of Comprehensive Genomic Profiling for Management of Rare 9.
- Hirshfield, Kim M. "Clinical Actionability of Comprehensive Genomic Profiling for Management of Rare or Refractory Cancers." *The Oncologist*, vol. 21, 2016, pp. 1315–1325.
 Grobner, S.N., et al., *The landscape of genomic alterations across childhood cancers.* Nature, 2018. 555(7696): p. 321-327.
- 555(7696): p. 321-327.
 Ma, X., et al., Pan-cancer genome and transcriptome analyses of 1,699 paediatric leukaemias and solid tumours. Nature, 2018. 555(7696): p. 371-376.
 Harris, M.H., et al., Multicenter Feasibility Study of Tumor Molecular Profiling to Inform Therapeutic Decisions in Advanced Pediatric Solid Tumors: The Individualized Cancer Therapy (iCat) Study. JAMA
- Parsons, D.W., et al., *Diagnostic Yield of Clinical Tumor and Germline Whole-Exome Sequencing for Children With Solid Tumors.* JAMA Oncol, 2016.

 Worst, B.C., et al., *Next-generation personalised medicine for high-risk paediatric cancer patients The INFORM pilot study.* Eur J Cancer, 2016. **65**: p. 91-101.

 Pincez, T., et al., *Feasibility and clinical integration of molecular profiling for target identification in pediatric solid tumors.* Pediatr Blood Cancer, 2017. **64**(6).

 Mody, R.J., et al., *Precision medicine in pediatric oncology: Lessons learned and next steps.* Pediatr Blood Cancer, 2017. **64**(3).

 Oberg, J.A., et al., *Implementation of next generation sequencing into pediatric hematology-oncology*

practice: moving beyond actionable alterations. Genome Med, 2016. 8(1): p. 133.

