In this issue of Modern Pathology, Duan et al. present their work on primary adult sellar SMARCB1-deficient tumors. This entity represents one further group in the ever expanding universe of SMARCB1 deficient tumors:

While the most prominent entity that displays a homozygous loss of the SMARCB1 are undoubtedly rhabdoid tumors (RT), the last years have seen molecular characterizations of further (extra and intracranial) SMARCB1 or SMARCA4 deficient entities including SMARCB1 deficient chordomas, cribriform neuroepithelial tumors and renal medullary carcinomas.

But even when only considering RT, it is increasingly clear that differences between eMRT (extracranial malignant RT) and ATRT (Atypical teratoid RT) that may exist at the morphological and immunohistochemical level blur at the molecular level: Both the loss of SMARCB1 but also a shared methylation-phenotype emerge as a unifying feature of ATRT and eMRT.

An important differential diagnosis to RT include epithelioid sarcomas. These rare sarcomas—most of which also display a homozygous loss of SMARCB1/INI1—predominantly arise in the acral parts of the body, but can also occur in intraabdominally or very rarely even at the skull basis. In contrast to RT, other cytogenetic abnormalities (such as the loss of 8p in about 50% of all cases, Fig. 1) have been described.

The distinction between RT and epithelioid sarcomas is clinically relevant as the latter—unlike RT—are usually not amenable to cytostatic therapy and an early resection should be attempted.

Duan et al. investigate eight adult, sellar SMARCB1 deficient tumors and elucidate their molecular makeup, aiming at differences and commonalities to RT and epithelioid sarcomas as well as SMARCB1 deficient chordomas.

In a methylation analysis (based on EPIC arrays), these tumors cluster with the ATRT-MYC subgroup. While this similarity has been described before, the authors now confirm previous analysis in a relatively large cohort given the rarity of this tumor type.

They also find CD34—a marker otherwise mostly absent in adult RT—to be present in most of the tumor samples. Unfortunately, this marker is also commonly present in epithelioid sarcomas and thus unsuitable for clear, immunohistochemical distinction of both entities. With regards to a further immunohistochemical characterization, it is notable that the authors cannot confirm SALL4 and ERG to be good discriminatory markers that distinguish between Epithelioid Sarcoma and RT as it has been otherwise proposed.

Overall, the publication by Duan et al. represents an important piece to define the landscape of SMARCB1 deficient tumors. An unresolved issue that remains to be elucidated is the female predominance among patients with sellar adult, RT.

Along a similar line, the localization of these tumors is enigmatic: The predominance of sellar tumors among adult ATRT is intriguing given that this localization is not prominent in pediatric rhabdoid tumor series. One may speculate that these tumors may evolve from other cells of origin than pediatric RT. Some evidence (such as the lack of CD34+ in other adult RT) indeed points to sellar, adult RT as a separate nosological entity. However, a more comprehensive review that would include whole genome and transcriptomic data still remains elusive. Also, the currently published number of sellar adult rhabdoid tumor cases seems to small to reliably infer that there are not genetic lesions beyond SMARCB1.

Other clinically interesting features such as a potential endocrine dysfunction owing to the localization close to the pituitary gland remain to be investigated systematically. Also, a potential adaption of treatment strategies to the adult patient cohort remains to be implemented.

Overall, while diagnostic features of sellar, RT emerge and their methylomic similarity to ATRT-MYC becomes more and more apparent, the clinical characterization of this “riddle inside an enigma” would be an important step forward to ultimately improve the bleak prognosis for adult rhabdoid tumor patients.

Pascal D. Johann

✉ Swabian Children’s Cancer Center, 86156 Augsburg, Germany.
✉ email: pascal.johann@uk-augsburg.de

Received: 4 June 2022 Revised: 3 July 2022 Accepted: 15 July 2022 Published online: 21 September 2022

Fig. 1 Synopsis on epithelioid sarcomas and rhabdoid tumors. An overview on important clinical and molecular characteristics in epithelioid sarcomas versus adult, sellar rhabdoid tumors.
REFERENCES
1. Fahiminiya, S., Witkowski, L., Nadaf, J., Carrot-Zhang, J., Goudie, C., Hasselblatt, M.
et al. Molecular analyses reveal close similarities between small cell carcinoma of the
ovary, hypercalcemic type and atypical teratoid/rhabdoid tumor. Oncotarget 7, 1732–1740 (2016).
2. Hasselblatt, M., Thomas, C., Hovestadt, V., Schrimpf, D., Johann, P., Bens, S. et al.
Poorly differentiated chordoma with SMARCB1/INI1 loss: a distinct molecular
entity with dismal prognosis. Acta Neuropathol. 132, 149-151 (2016).
3. Johann, P. D., Hovestadt, V., Thomas, C., Jeibmann, A., Hess, K., Bens, S. et al.
Cribiform neuroepithelial tumor: molecular characterization of a SMARCB1-
deficient non-rhabdoid tumor with favorable long-term outcome. Brain Pathol. 27, 411-418 (2017).
4. Msaouel, P., Walker, C. L., Genovese, G. & Tannir, N. M. Molecular hallmarks of
renal medullary carcinoma: more to c-MYC than meets the eye. Mol. Cell Oncol. 7, 1777060 (2020).
5. Jamshidi, F., Bashashati, A., Shumansky, K., Dickson, B., Gokgoz, N., Wunder, J.S.
et al. The genomic landscape of epithelioid sarcoma cell lines and tumours. J. Pathol. 238, 63-73 (2016).
6. Chun, H. E., Johann, P.D., Milne, K., Zapatka, M., Buellesbach, A., Ishaque, N., et al.
Identification and Analyses of Extra-Cranial and Cranial Rhabdoid Tumor Mole-
cular Subgroups Reveal Tumors with Cytotoxic T Cell Infiltration. Cell Rep. https://
doi.org/10.1016/j.celrep.2019.10.013 (2019).
7. Lian, D. W., Lee, H.Y., Tan, K.K., Sito, Y.Y., Yap, W.M. & Chua, K.L. Proximal, type
epithelioid sarcoma arising in the base of the skull: a diagnostic challenge. J. Clin.
Pathol. 63, 472–474 (2010).
8. Koelsche, C., Schrimpf, D., Stichel, D., Sill, M., Sahm, F., Reuss, D. et al. Sarcoma
classification by DNA methylation profiling. Nat. Commun. 12, 498 (2021).
9. Johann, P. D., Bens, S., Oyen, F., Wagener, R., Giannini, C., Perry, A. et al. Sellar
Region Atypical Teratoid/Rhabdoid Tumors (ATRT) in Adults Display DNA
Methylation Profiles of the ATRT-MYC Subgroup. Am. J. Surg. Pathol. 42, 506-
511 (2018).
10. Kohashi, K., Yamada, Y., Hotokebuchi, Y., Yamamoto, Y., Taguchi, T., Iwasato, Y.
et al. ERG and SALL4 expressions in SMARCB1/INI1-deficient tumors: a useful tool
for distinguishing epithelioid sarcoma from malignant rhabdoid tumor. Hum.
Pathol. 46, 225-230 (2015).
11. Chan, V., Maro, A., Findlay, J. M., Schmitt, L. M. & Das, S. A Systematic Review of
Atypical Teratoid Rhabdoid Tumor in Adults. Front. Oncol. 8, 567 (2018).

AUTHOR CONTRIBUTIONS
PDJ conceived, drafted and edited the article.

FUNDING
Open Access funding enabled and organized by Projekt DEAL.

COMPETING INTERESTS
The author declares no competing interests.

ADDITIONAL INFORMATION
Correspondence and requests for materials should be addressed to Pascal D. Johann.

Reprints and permission information is available at http://www.nature.com/reprints

Publisher’s note Springer Nature remains neutral with regard to jurisdictional claims in
published maps and institutional affiliations.

Open Access This article is licensed under a Creative Commons
Attribution 4.0 International License, which permits use, sharing,
adaptation, distribution and reproduction in any medium or format, as long as you give
appropriate credit to the original author(s) and the source, provide a link to the Creative
Commons licence, and indicate if changes were made. The images or other third party
material in this article are included in the article’s Creative Commons licence, unless
indicated otherwise in a credit line to the material. If material is not included in the
article’s Creative Commons licence and your intended use is not permitted by statutory
regulation or exceeds the permitted use, you will need to obtain permission directly
from the copyright holder. To view a copy of this licence, visit http://
creativecommons.org/licenses/by/4.0/.

© The Author(s) 2022