



Genomic characterization of metastatic pituitary neuroendocrine carcinoma (PitNET) in two cases

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Introduction

- Primary pituitary neuroendocrine (PitNET) carcinomas are exceptionally rare, with only a few case reports in the literature.¹⁻⁶
- We present two patients with high-grade pituitary lesions who developed metastases isolated to the central nervous system.
- In these cases, recurrence despite treatment was common, underscoring their difficulty to treat.

Methods and Materials

- Whole exome sequencing (WES) was performed on all available lesions and matched blood samples.
- Target capture was performed with IDT xGen Exome Research Panel Version 2 + GOAL region spike-ins.
- Sequencing was performed with Illumina NovaSeq 600.
- Downstream analysis to identify somatic single nucleotide variations (SNVs), short insertion-deletions (INDELs), and copy number variations (CNVs) was carried out following the Genome Analysis Toolkit (GATK) Best Practices Guidelines.
- A literature search was conducted to identify similar cases for comparison.

Patient 1

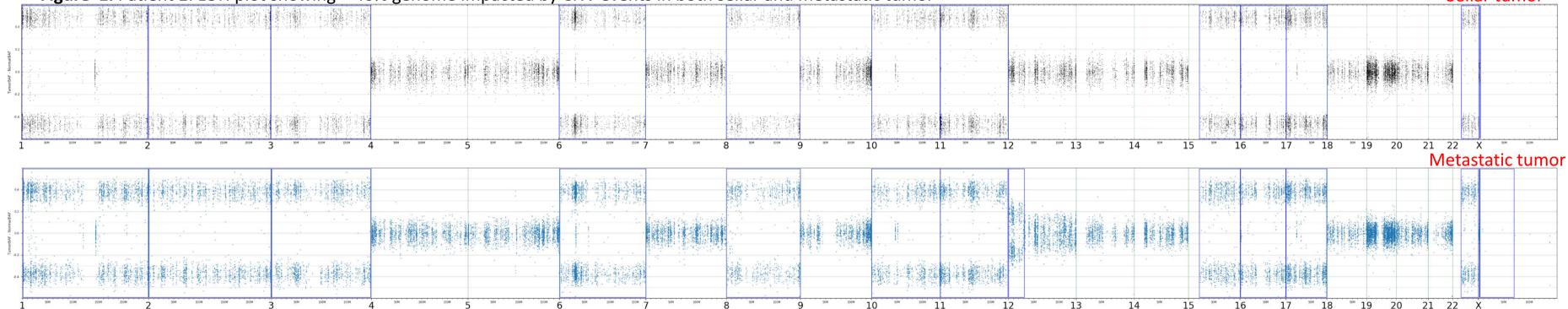
Clinical presentation:

- 47-year-old male with pituitary tumor resected twice before developing recurrent pituitary lesion.
- Treated with re-resection via EEA and adjuvant radiotherapy (50.4 Gy).
- Histological assessment revealed pituitary adenoma with Ki-67 index 4-5%.
- New extrasellar brain lesions identified on follow-up MRI after 2.5 years.
- Surgical pathology of one of the new lesions revealed neuroendocrine carcinoma.

Genomic profile (somatic WES):

- Both the sellar and metastatic tumors demonstrated **>40% of the genome altered by CNVs.**
- Both the primary and recurrent tumor had **DAXX: frameshift mutation (K437Tfs).**
 - A nuclear protein involved in transcriptional repression, apoptosis, chromatin organization, telomere maintenance, and genomic instability.

Figure 1. Patient 1: LOH plot showing > 40% genome impacted by CNV events in both sellar and metastatic tumor



Patient 2

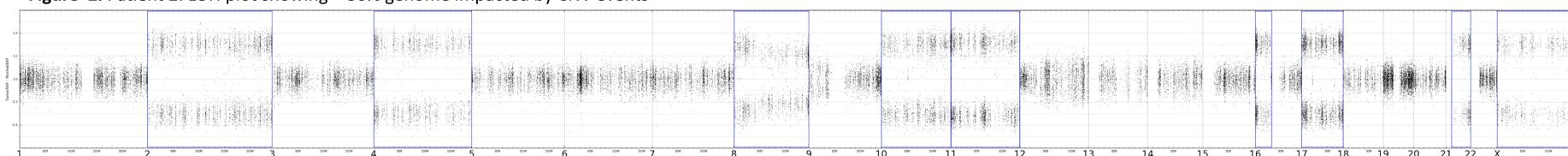
Clinical presentation:

- 26-year-old female with prolactinoma on cabergoline presented with pituitary lesion causing hydrocephalus.
- Required resection via EEA. Histology showed neuroendocrine tumor with Ki-67 10-15%. Patient was lost to follow-up.
- Lesion recurred in 6 months which required re-resection and adjuvant radiotherapy (50.4 Gy).
- She developed multiple CNS metastases including brain and intradural extramedullary spinal lesions.

Genomic profile (somatic WES):

- **36% of the genome altered by CNVs.**
- Had a deleterious **TP53: missense mutation (R209Q).**
 - Consistent with the well-established role of TP53 loss in promoting chromosomal instability.

Figure 2. Patient 2: LOH plot showing > 36% genome impacted by CNV events



Discussion & Conclusions

- Metastatic PitNET is rare, and deleterious somatic alterations in key genomic stability regulators may underlie the aggressive biology of these otherwise typically benign tumors.
- Future work should emphasize early diagnosis and aggressive management (including the role of CSI) in the management of patients with disease limited to the CNS.

Contact

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