NTRK and S100 staining to report NTRK fusion sarcoma of the cervix

Sarah Chiang, MD

Memorial Sloan Kettering Cancer Center





Disclosure of relevant financial relationships

No relevant financial relationships





NTRK fusion-positive uterine sarcoma is a distinct clinicopathologic entity

Median age 32 y

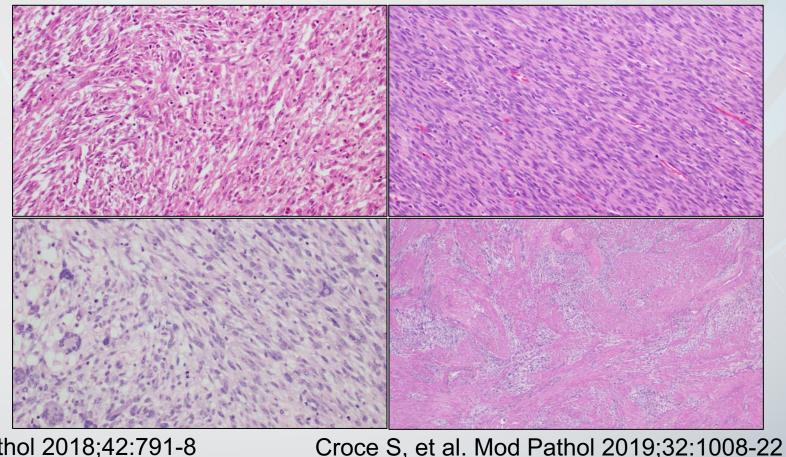
FIGO stage

IA: 7 (50%)

IB: 6 (43%)

IIB: 1 (7%)

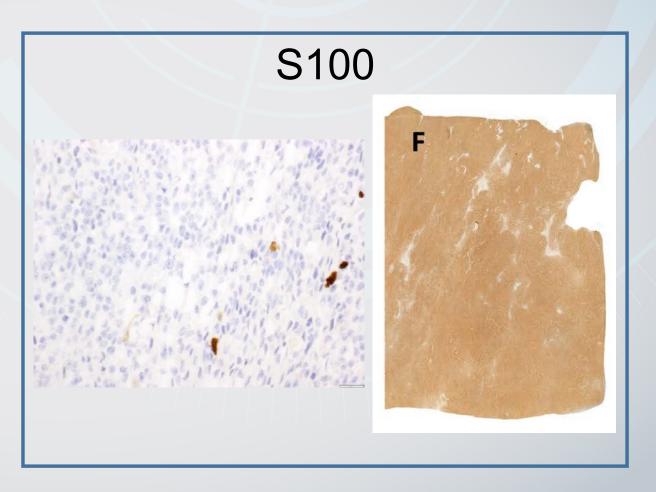
Cervix origin 93%



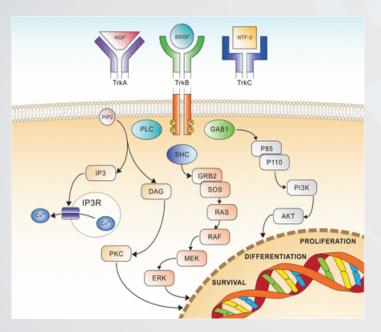
Chiang S, et al. Am J Surg Pathol 2018;42:791-8 Croc Hodgson A, et al. Int J Gynecol Pathol 2020;Epub ahead of print Rabban JT, et al. Histopathol 2020;Epub ahead of print

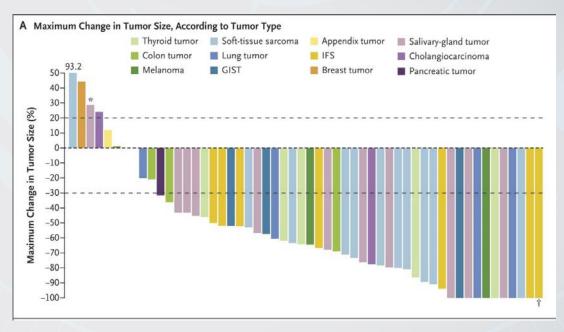
NTRK fusion-positive uterine sarcomas lack evidence of cellular differentiation

Antibody	Staining pattern
Desmin	_
SMA	+ (focal)
ER	_
PR	_
CD34	+/-
S100	+ (varied extent)
SOX10	_
H3K27me3	+



Uterine sarcoma subtype is defined by NTRK fusion and targetable by Trk inhibition





One patient with SPECC1L-NTRK3 fusion positive uterine sarcoma s/p polypectomy, doxorubicin/ifosfamide (5 cycles) and pelvic radiation, developed pleural metastasis and had complete radiographic response to larotrectinib

Chiang S, et al. Am J Surg Pathol 2018;42:791-8 Drilon A, et al. N Engl J Med. 2018;378:731-9 Croce S, et al. Mod Pathol 2019;32:1008-22

Rabban JT, et al. Histopathol 2020; Epub ahead of print Hodgson A, et al. Int J Gynecol Pathol 2020; Epub ahead of print

Pan-Trk immunohistochemistry is a potential screen for *NTRK* fusion

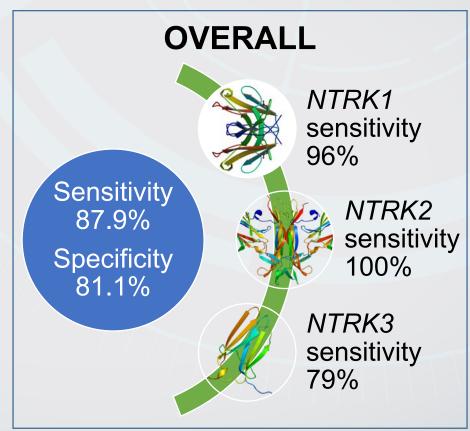
Pan-Trk monoclonal antibody clone EPR 17341 (Abcam, Roche/Ventana)

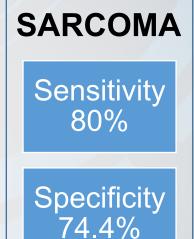
Reactive to homologous region of TrkA, B and C near C terminus

Expressed in testis, neural tissue, smooth muscle

>1% of tumor cell staining

= positive result



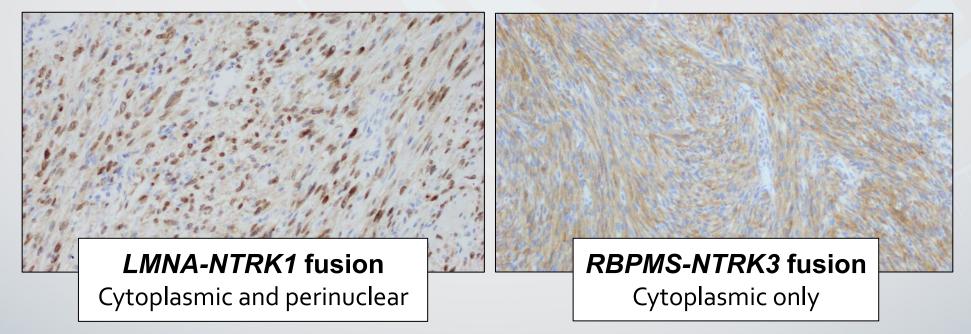


Trk staining patterns vary due to different fusion partners

Uterine sarcoma fusions:

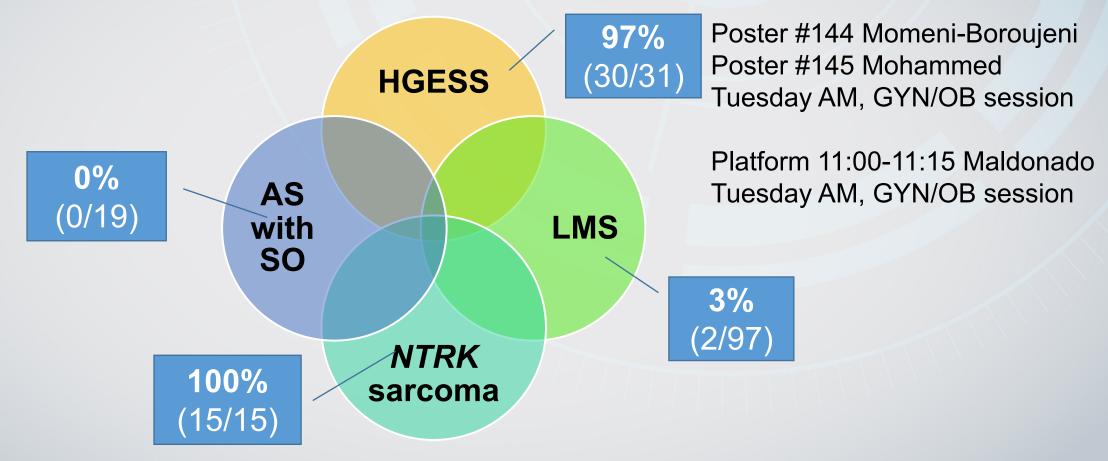
NTRK1: TPM3, LMNA, TPR, SPECC1L

NTRK3: RBPMS, EML4

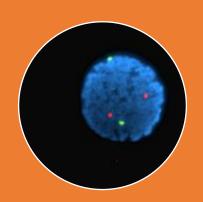


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Trk expression may be seen in morphologic mimickers



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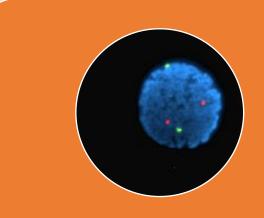
FISH



DNA-based NGS



RNA-based NGS



FISH

PRO

Commercially available probes Requires less material Short TAT

CON

False-negatives with NTRK1

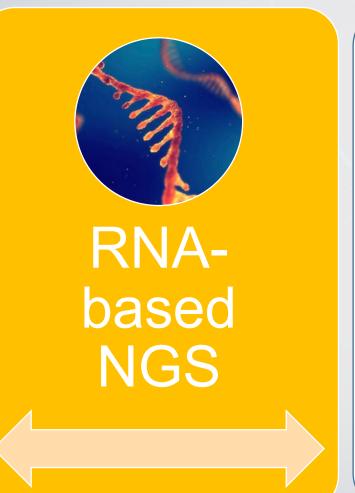


PRO

Assessment of other genetic alterations Monitor for resistance mutations

CON

Longer TAT
Requires more material
Requires intronic coverage
Cannot determine functional expression
Less sensitive than RNAseq



PRO

Intronic coverage not needed
Detects functionally transcribed fusions
Permits fusion discovery
More sensitive than DNAseq

CON

Limited by RNA quality Longer TAT Requires more material

Algorithm for NTRK fusion detection in uterine sarcoma depends on available test modalities

RNA sequencing is preferred

 Comprehensive fusion testing increasingly used as first-line test for sarcoma

Pan-Trk immunohistochemistry should be used with caution

- Lower specificity in sarcoma
- Trk expression observed in HGESS and uterine LMS

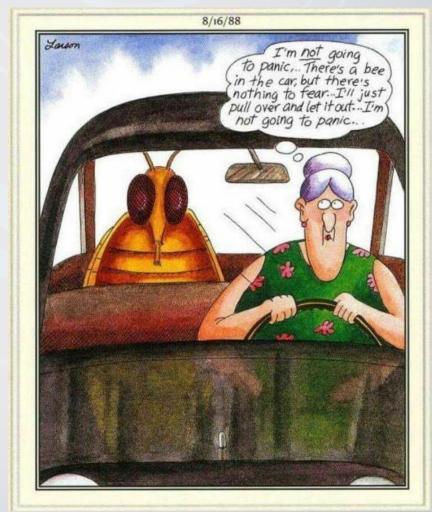
Stakes are high in diagnosing *NTRK* fusion-positive uterine sarcomas

Rare disease

Morphologic heterogeneity

No quick and easy screening test

Treatment implications



Join the uterine sarcoma profiling initiative!



Goal: Integrated genetic, epigenetic and pathological diagnosis for uterine sarcoma

- Network of gynecologic pathology departments sharing data and samples
- Collaborators can send clinical or research cases for profiling
 - MSK-IMPACT: matched tumor-normal DNA NGS of 468 genes
 - MSK Solid Fusion Assay: RNA NGS of 68 genes
 - Methylation MSK/NYU/Stanford: whole genome DNA methylation for molecular diagnosis and subclassification
- Share unclassifiable cases to establish new tumor entities

E-mail: chiangs@mskcc.org