Serial plasma and CSF cell-free tumor DNA (cf-tDNA) tracking in diffuse midline glioma patients undergoing treatment with ONC201



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Disclosures

• Nothing to disclose

DMG is pathologically defined but driven by mutations in H3

Glioma – grade 2-4 (i.e., infiltrates healthy brain)

- Any part of the midline of the brain
 (brainstem, thalamus, and spinal cord)
- Most common in younger patients

Behavior, prognosis, and treatments are driven by the H3 K27M mutation

- $\odot~75\%$ of H3 K27M occur in the H3F3A gene
- OS of H3 K27-mutant DMG: ~12 m



^a H3 K27M immunohistochemistry: positive in tumors harboring the H3F3A mutation (75% of cases)

ACVR1, activin A receptor type 1; ATRX, ATRX chromatin remodeler; DAXX, death domain associated protein; DMG, diffuse midline glioma; FOXG1, forkhead box G1; H3, Histone 3; H3F3A, Histone 3 F3A; H3 K27M, Histone 3 lysine to methionine substitution at residue 27; m, months; OLIG1, oligodendrocyte transcription factor 1; OLIG2, oligodendrocyte transcription factor 2; OS, overall survival; TP53, tumor protein p53.

Monitoring treatment response in DMG is complicated by pseudoprogression

- Tumors often increase in size early in the course of treatment
 - May be tumor progression
 - May be "pseudo-progression" radiationrelated swelling/necrosis
- Some non-CNS tumors displaying multi-pronged clinical responses are tracked by molecular analyses of plasma
- For brain tumors (glioma), MRI is the only approved diagnostic
 - $\circ~$ Serial biopsy in HGG carries significant risk



Tiwari et al, Cleveland Clinic Consult QD, 2019

CNS, central nervous system; DMG, diffuse midline glioma; HGG, high grade glioma; MRI, magnetic resonance imaging.

Droplet digital PCR to diagnosis and quantify H3K27M cf-tDNA





Stallard et al (Koschmann), *Acta Neuropath Comm*, 2018 Bruzek et al (Koschmann), *Clinical Cancer Research*, 2020



ONC201: DRD2 antagonist and ClpP agonist



DRD2/3 antagonist ClpP agonist

Oral capsules, administered weekly

Pharmacodynamic-based dose without DLT

Crosses blood-brain barrier



AKT, protein kinase B; ATF4, activating transcription factor 4; CHOP, C/EBP-homologous protein; ClpP, caseinolytic protease P; DLT, dose-limiting toxicity; DRD2/3, dopamine receptor D2/D3; ERK, extracellular-regulated kinase.

Allen et al, *Science Translational Medicine*, 2013; Ishizawa et al, *Science Signaling*, 2016; Kline et al, *Science Signaling*, 2016; Ishizawa et al, *Cancer Cell*, 2019; Prabhu et al, *Neoplasia*, 2020

H3 K27M-mutant gliomas exhibit sensitivity to ONC201

Adult phase 2 study in recurrent GBM -ONC201 (NCT02525692)

- Well tolerated, achieved intratumoral 0 concentrations that exceeded preclinical efficacy thresholds
- Primary endpoint of PFS6 at 5% by RANO Ο was not achieved
- However, 1 patient enrolled with the H3 0 K27M mutation had a complete regression of enhancing multifocal lesions that remained durable for >1.5 years

8 weeks 16 weeks Arrillaga-Romany et al, Neuro-Oncology, 2020





GBM, glioblastoma multiforme; H3.3, H3 histone variant; H3 K27M, Histone 3 lysine to methionine substitution at residue 27; H3 K27m3, Histone 3 residue lysine 27 tri-methylation; PFS6, progression-free survival at 6 months; RANO, Response Assessment in Neuro-Oncology.

Kawakibi et al, Society of Neuro-Oncology, 2019, Unpublished



Other arms (UM patients only)







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Serial plasma and CSF H3.3 K27M cf-tDNA monitoring on ONC201 treatment



H3.3K27M cf-tDNA "spikes" can precede tumor progression



UMICH-PED20: 10 yo DMG-H3.3K27M



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H3.3K27M cf-tDNA can help differentiate pseudo-progression and pseudo-response (bevacizumab)



Conclusions

- Serial CSF collection is feasible and useful in management of glioma
- ddPCR is highly sensitive and specific for H3K27M tDNA in CSF and plasma
- Change in tumor area / VAF over time don't correlate but "patterns" of change demonstrate potential clinical utility
 - Sustained reduction in tDNA correlates with / predicts long term response
 - Increase in tDNA correlates with progression may predict progression in 2-4 months
 - H3K27M tDNA patterns less affected by pseudo-progression and bevacizumab than MRI

cf-tDNA, cell-free tumor DNA; CSF, cerebrospinal fluid; ddPCR, droplet digital polymerase chain reaction; H3 K27M, Histone 3 lysine to methionine substitution at residue 27;

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