

AG-120 (ivosidenib), a first-in-class mutant IDH1 inhibitor, promotes morphologic changes and upregulates liver-specific genes in IDH1 mutant cholangiocarcinoma

Yuko Ishii¹, Carlie Sigel², Maeve A Lowery^{2,3}, Lipika Goyal⁴, Camelia Gliser¹, Liewen Jiang¹, Susan Pandya¹, Bin Wu¹, Sung Choe¹, Vikram Deshpande⁴

¹*Agios Pharmaceuticals, Inc., Cambridge, MA, USA;*

²*Memorial Sloan Kettering Cancer Center, New York, NY, USA [at time of work];*

³*Trinity College, Dublin, Ireland [current];*

⁴*Massachusetts General Hospital, Boston, MA, USA*

Disclosure information

Disclosure information

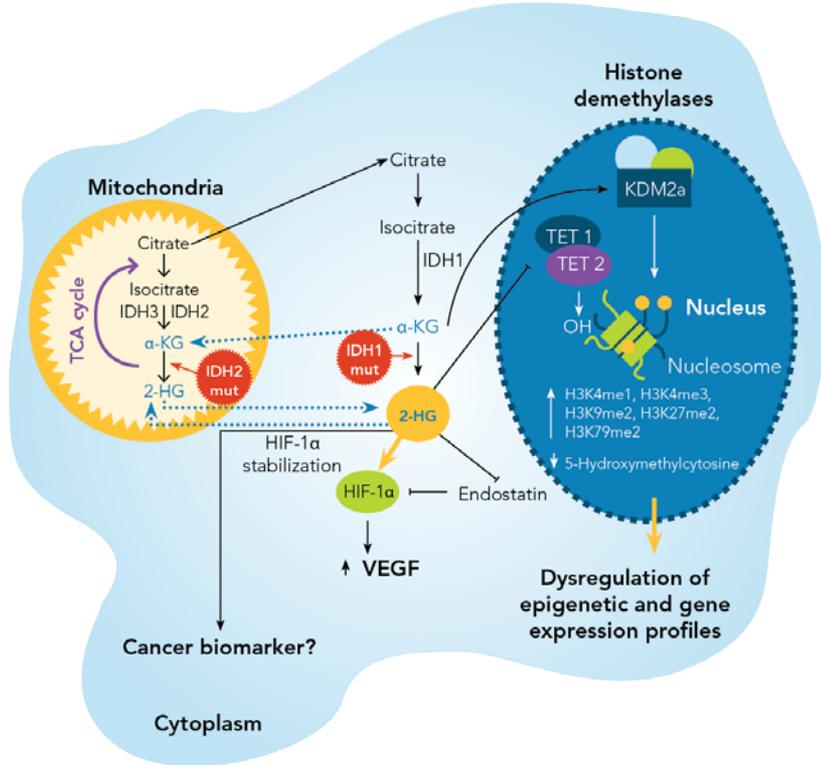
- This study was funded by Agios Pharmaceuticals.
- YI, CG, LJ, SP, BW, SC: Agios Pharmaceuticals – employment and stockholder. CS: Agios Pharmaceuticals – travel expenses. MAL: Agios Pharmaceuticals – advisor/board member; Celgene – advisor/board member. LG: Ribon Therapeutics – honorarium recipient; DebioPharm – consultant/independent contractor. VD: Agios Pharmaceuticals – consultant/independent contractor; Advance Cell Diagnostics – grants/research support recipient; Affymetrix – grants/research support recipient.
- Editorial assistance was provided by Susanne Vidot, PhD, CMPP, Excel Scientific Solutions, Horsham, UK, and supported by Agios.

I will discuss the following off label use and/or investigational use in my presentation:

ClinicalTrials.gov NCT02073994: Study of Orally Administered AG-120 in Subjects With Advanced Solid Tumors, Including Glioma, With an IDH1 Mutation

IDH1 mutations in cholangiocarcinoma

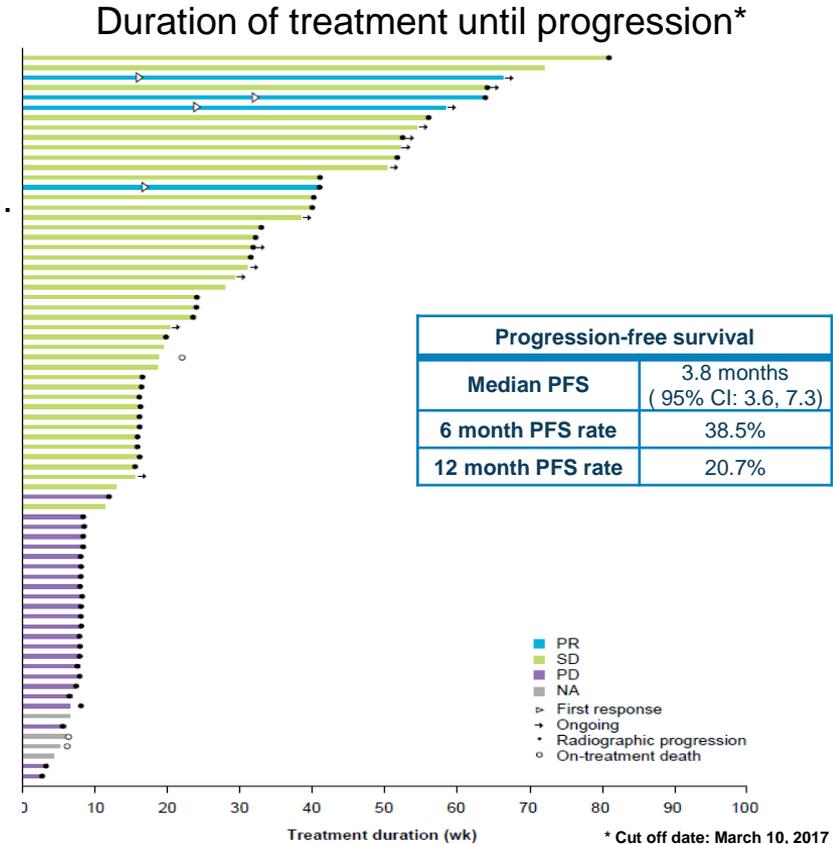
- Mutations in the isocitrate dehydrogenase 1 (IDH1) gene are detected in 13–15% of cholangiocarcinoma (CC).¹⁻³
 - ~25% of intrahepatic CC
- The mutant IDH1 (mIDH1) enzyme produces the oncometabolite D-2-hydroxyglutarate (2-HG),^{4,5} which leads to epigenetic dysregulation and a block in cellular differentiation.⁶⁻⁹
- AG-120 (ivosidenib) is a first-in-class, oral, potent, reversible, selective inhibitor of the mIDH1 enzyme.¹⁰⁻¹²



1. Goyal L et al. *Oncologist* 2015;20:1019-27. 2. Borger DR et al. *Oncologist* 2012;17:72-9. 3. Kipp BR et al. *Hum Pathol* 2012;43:1552-8. 4. Ward PS et al. *Cancer Cell* 2010;17:225-34. 5. Dang L et al. *Nature* 2009;462:739-44. 6. Saha SK et al. *Cell Cycle* 2014;13:3176-82. 7. Saha SK et al. *Nature* 2014;513:110-4. 8. Lu C et al. *Nature* 2012;483:474-8. 9. Xu W et al. *Cancer Cell* 2011;19:17-30. 10. de Botton S et al. *Haematologica* 2015;100(s1):214;P563. 11. Fan B et al. *Haematologica* 2015;100(s1):218;P572. 12. Fan B et al. *Blood* 2015;126(23):A1310.

AG-120 in mIDH1 cholangiocarcinoma

- AG120-C-002 (ClinicalTrials.gov NCT02073994), a first-in-human phase 1 study, assessed AG-120 in patients with mIDH1 advanced solid tumors.
 - 73 patients with mIDH1 CC (median 2 prior lines of therapy).
- AG-120 was well tolerated and associated with a favorable safety profile.
 - no dose-limiting toxicities or treatment-related deaths^{13,14}
- AG-120 demonstrated encouraging clinical activity in this heavily pre-treated mIDH1 CC population.^{13,14}
- The exploratory objectives included the assessment of morphological and molecular changes in serial tumor biopsy samples.

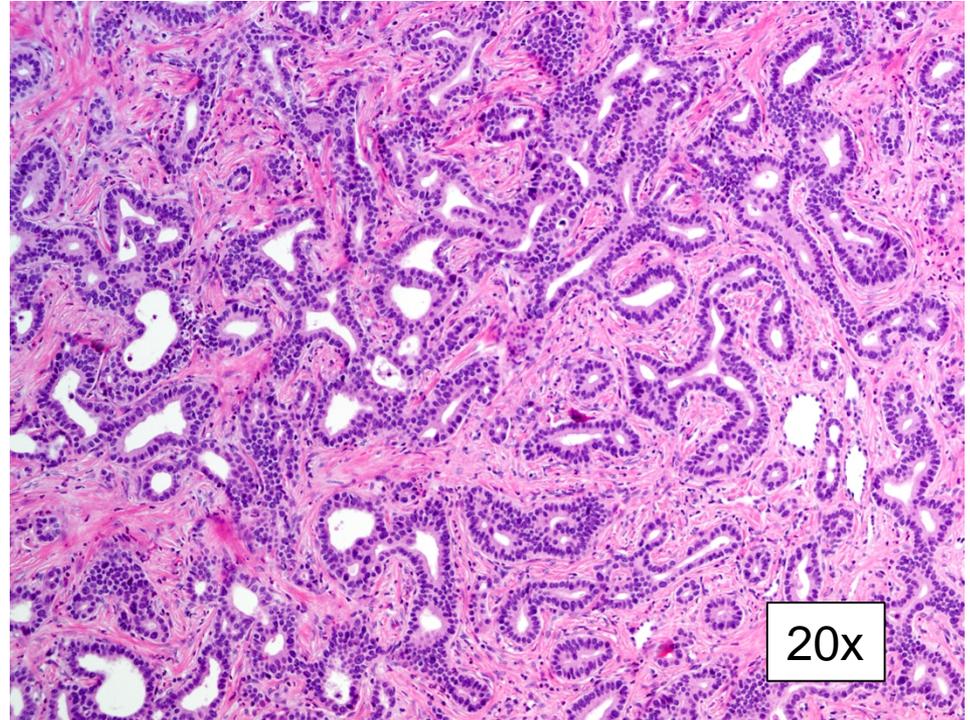


¹³Lowery MA et al. *J Clin Oncol* 2017;35(Suppl):Abstr 4015. ¹⁴Lowery MA et al. *ASCO Annual Meeting* 2017: Poster 4015.

Histological characteristics of mIDH1 CC

- A cholangiolar pattern was defined as being composed of glands with an antler-horn configuration and angulated shapes, and lined with low cuboidal epithelium.¹⁵⁻¹⁷
- Cholangiolar histology is associated with better clinical outcomes and survival rates in patients with ICC.^{15,19}
- Untreated mIDH1 ICCs often show heterogeneous histoarchitecture.
 - 61% of tumors lack a dominant pattern¹⁸
 - Cholangiolar histology is commonly present in mIDH1 CC, but often to a limited extent (median 10% cholangiolar histology).^{15,18}
- Tumor phenotype and morphologic differentiation in CC patients treated with AG-120 have not previously been examined.

Cholangiolar pattern¹⁹



Sample and data summary

Sample Collection

(n = number of patients with samples at baseline and ≥ 1 on-treatment time point)

Procedure

(n = number of patients with data available at baseline and ≥ 1 on-treatment time point)

Morphology

Hematoxylin and eosin (H&E) stained slides from FFPE tissue (n = 27)

Blinded evaluation of architectural, cytologic, and stromal patterns by two gastrointestinal pathologists (n = 17^a)

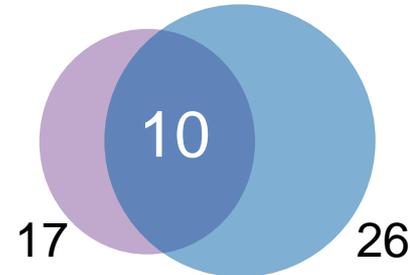
Gene Expression

Fresh frozen biopsies (n = 38)

Tumor content and assay quality control

Personalis[®] ACE Transcriptome[™] RNAseq platform (n = 26^b)

10 patients have both morphology and gene expression data available.



^aIncludes 16 patients dosed at 500 mg QD and one patient dosed at 100 mg BID

^bIncludes 22 patients dosed at 500 mg QD, two patients dosed at 1200 mg QD, and two patients dosed at 300 mg QD

Baseline to post-dose morphologic changes in AG-120-treated mIDH1 CCs

- The percentage of tumor with a cholangiolar pattern was recorded. A baseline to postdose increase was defined as a $\geq 20\%$ increase in cholangiolar histology.
- The volume of cytoplasm in tumor cells was semi-quantitatively assessed.
- These morphologic changes were not associated with AG-120 dose level. All patients had plasma 2-HG reduction upon AG-120 treatment, regardless of post-dose morphology.²⁰

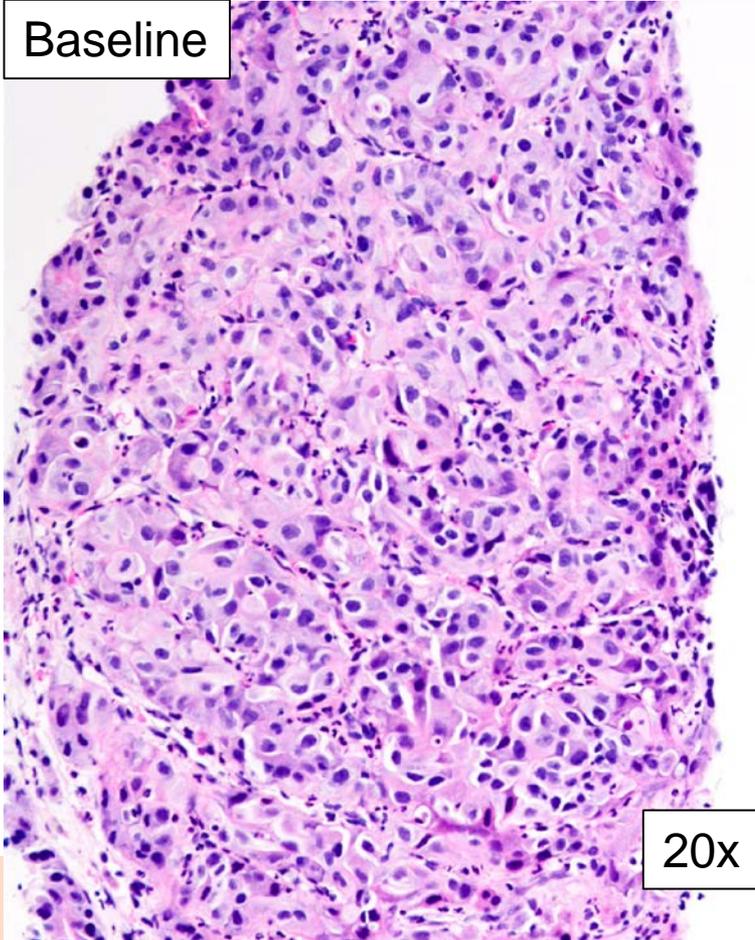
	Morphology data available	Increase in cholangiolar histology	Cytoplasmic reduction	Cholangiolar and cytoplasmic changes
Number of patients	17	5	9	4
By treatment response^a				
PR	3	1	3	1
SD	12	4	6	3
PD	2	–	–	–

15. Liu JY et al. *Mod Pathol* 2014;27:1163-73. 20. Fan B et al. *J Clin Oncol* 2017;35(Suppl):Abstr 4082.

^aTreatment response measured according to Response Evaluation Criteria in Solid Tumors (RECIST) v1.1 (PR = partial response; SD = stable disease; PD = progressive disease). The clinical data were based on a cutoff date of May 12, 2017.

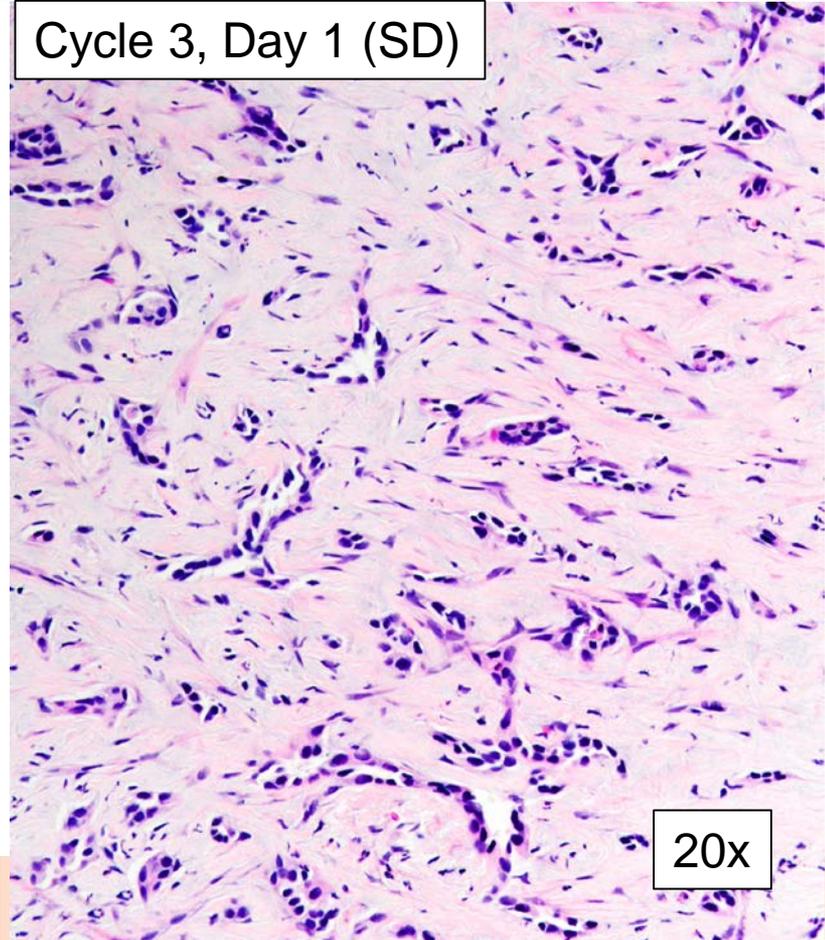
Example 1: Increased cholangiolar histology and decreased cytoplasm upon AG-120 treatment

Baseline



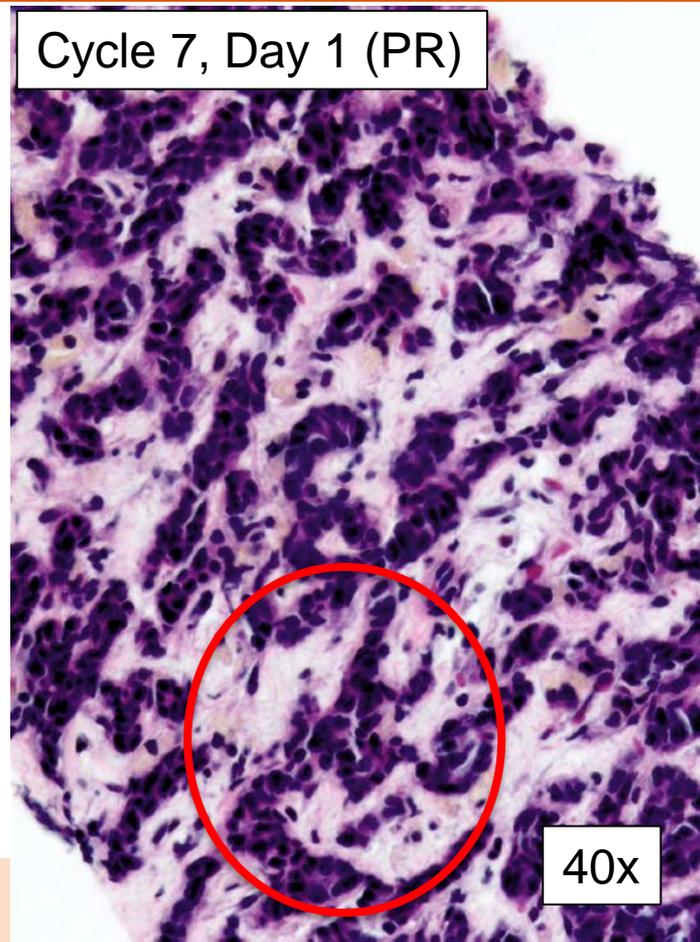
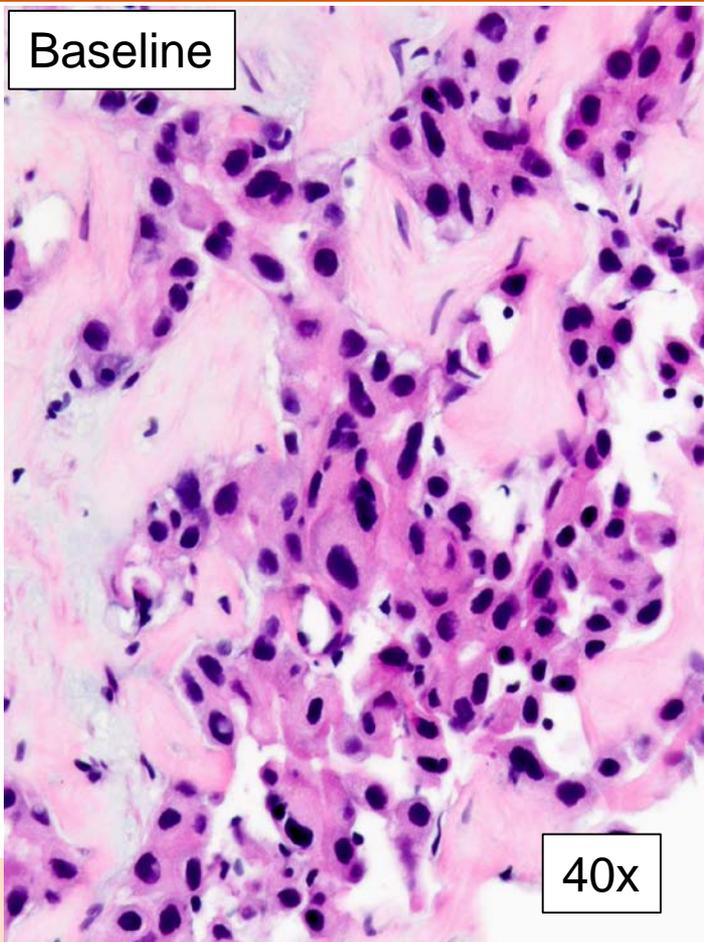
20x

Cycle 3, Day 1 (SD)

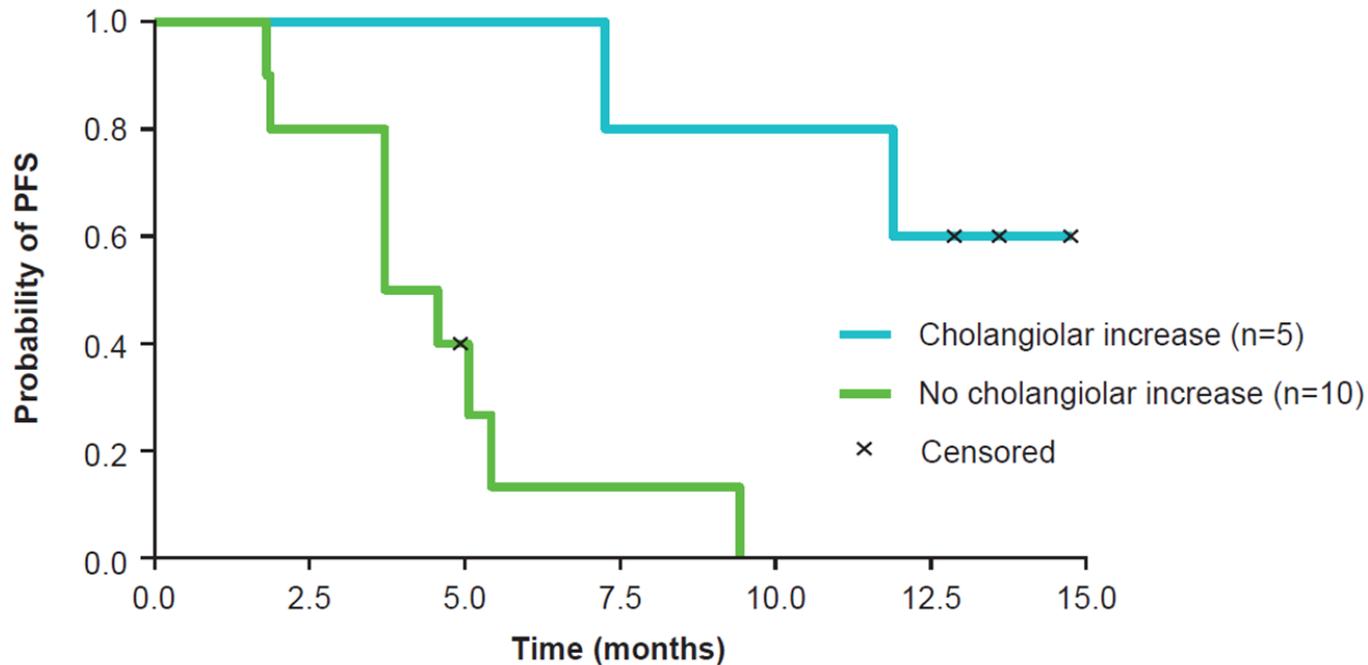


20x

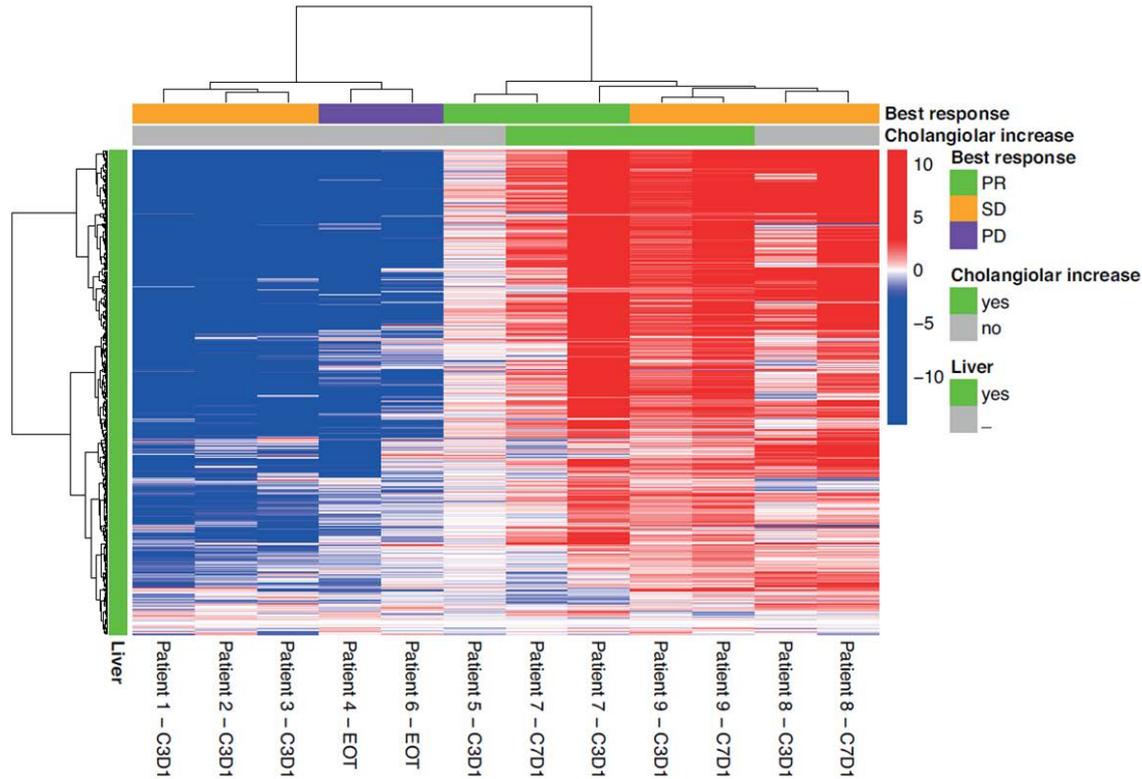
Example 2: Increased cholangiolar histology and decreased cytoplasm upon AG-120 treatment



Increased cholangiolar pattern seems to be associated with increased PFS

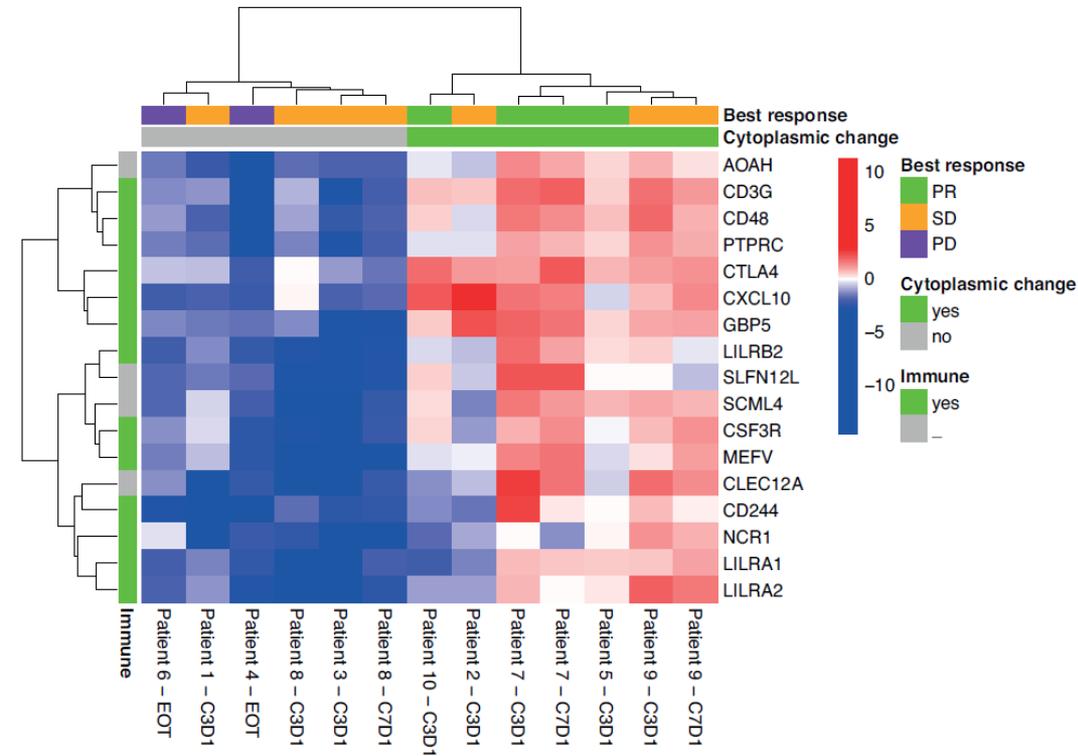


mIDH1 CCs with cholangiolar increase show upregulation of a broad set of adult liver-specific genes



- Preclinical studies have shown IDH1 mutations to block hepatocyte differentiation and promote biliary cancers.^{6,7}
- Gene expression data were available for two patients with observed cholangiolar pattern increase ($\geq 20\%$).
- Both showed increased expression of liver specific genes (N = 485), derived from two sources:
 - Farshidfar et al. (2017)²¹
 - Hsiao et al. (2001)²²

Patients with cytoplasmic decrease show increased expression of immune-response related genes

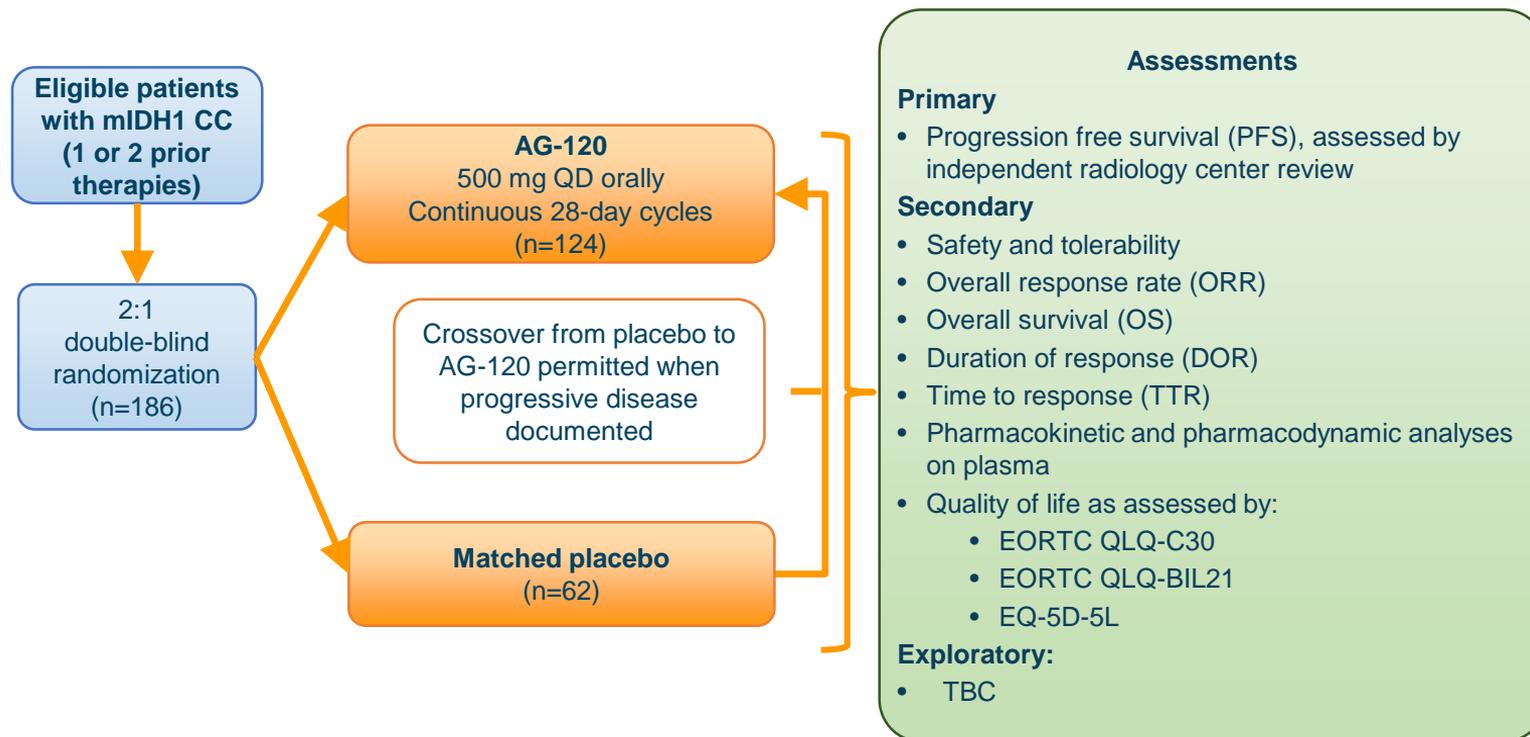


- Gene expression data were available for five patients with observed cytoplasm decrease.
- These five patients showed upregulation of multiple immune response-related genes, including CXCL10, CD3G, and CTLA4.
- In preclinical studies, IDH1m glioma showed lower expression of the chemokine CXCL10, and combined IDH1m inhibitor / vaccine treatment resulted in increased CXCL10 expression and CD8 T cell infiltration.²³

Conclusions

- This is the first demonstration that AG-120 treatment may induce morphologic and molecular changes in a subset of mIDH1 CCs.
- Increased cholangiolar histology seems to be associated with increased PFS.
- Tumors with increased cholangiolar histology showed upregulation of genes associated with mature liver cells.
- The increased expression of immune response related genes in some tumors suggests a potential rationale for AG-120 in combination with immunotherapies.
- Given the limited sample size of this dataset, additional studies are warranted to explore the biological and clinical significance of these observations.
- AG-120 is under further evaluation in an ongoing, global, phase 3, randomized, placebo-controlled study in previously treated mIDH1 CC (ClarIDHy; ClinicalTrials.gov NCT02989857).

Phase 3 ClarIDHy Trial Design



Acknowledgments

- We would like to thank the patients taking part in this study.
- Dr. Nabeel Bardeesy at MGH/Broad Institute provided consultations on gene sets.