

AG-270 Data at 2019 AACR-NCI-EORTC International Conference 2019

October 27, 2019



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Today's Agenda

- Opening Remarks Jackie Fouse, Ph.D., Chief Executive Officer
- Preclinical AG-270 Data Kevin Marks, Ph.D., Vice President, Head of Biology
- AG-270 Phase 1 Results Chris Bowden, M.D., Chief Medical Officer
- Q&A Keith T. Flaherty, M.D., Director of Clinical Research MGH Cancer Center



2019 Key Milestones & Data Presentations Position Agios for Long-term Value Creation



Key 2019 Milestones

- ✓ FDA approval and commercialization of monotherapy TIBSOVO® in untreated AML
- ✓ Initiate AG-636 Phase 1 dose-escalation trial in lymphoma in 1H 2019
- ✓ Complete AG-270 Phase 1 dose-escalation and select go forward dose
- ✓ Initiate expansion arms in the AG-270 Phase 1 study in Q3 2019
- Achieve proof-of-concept for mitapivat in thalassemia in 2H 2019
- Submit sNDA for TIBSOVO® in second line or later cholangiocarcinoma by YE
- Initiate Phase 3 INDIGO study of vorasidenib in low grade glioma by YE
- Complete enrollment in PK deficiency pivotal trials ACTIVATE-T and ACTIVATE by YE



Key Upcoming Data Presentations

- Updated data from the perioperative study of ivosidenib and vorasidenib accepted for presentation at the SNO Annual Meeting
- Data from IDH and PKR programs have been accepted for presentation at ASH, including:
 - New data from the extension phase of the Phase 2 DRIVE PK study of mitapivat in adults with PK deficiency
 - Important translational data from the Phase 1 study of TIBSOVO® and azacitidine in frontline AML





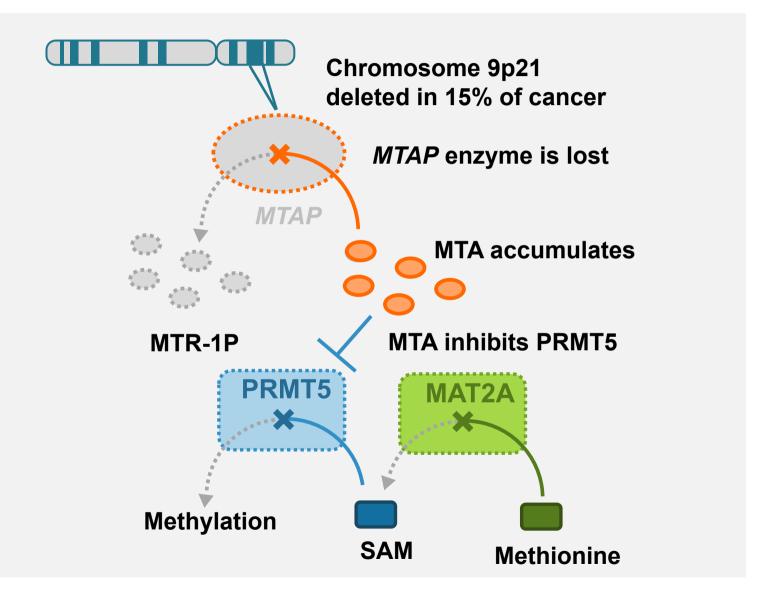
Targeting MAT2A in *CDKN2A/MTAP*-deleted Cancers

Kevin Marks, Vice President and Head of Biology



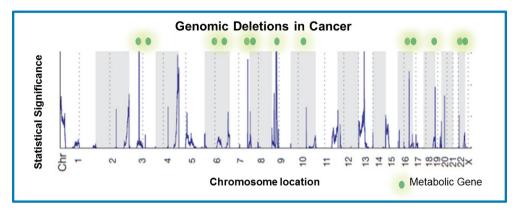
A Key Insight: Deletion of *MTAP* Makes Cancers Vulnerable to Targeting of MAT2A

- 1. MTAP deletion
- 2 Substrate MTA accumulates
- Partial inhibition of PRMT5
- Sensitivity to a 'second hit': targeting MAT2A starves PRMT5 of its substrate

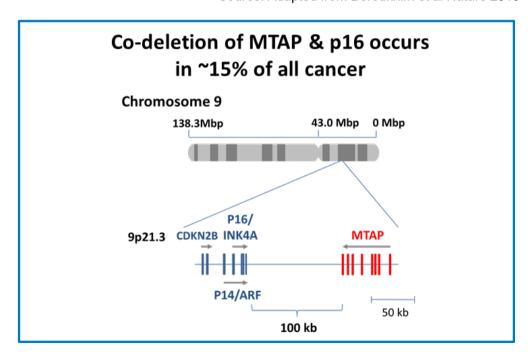


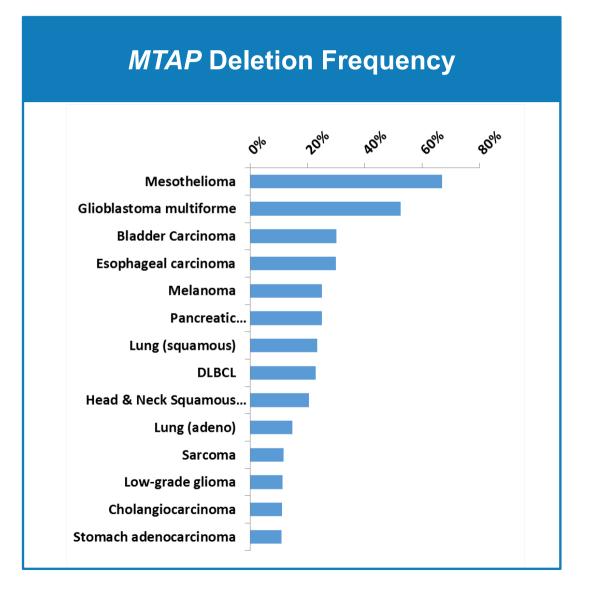


MTAP Deletions Occur in ~15% of All Cancers



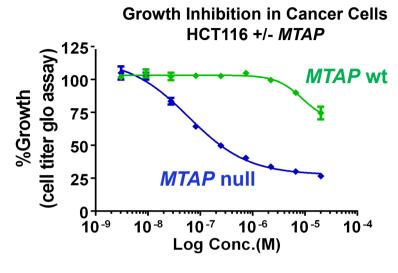
Source: Adapted from Beroukhim et al Nature 2010

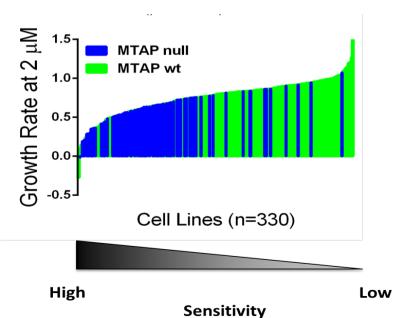


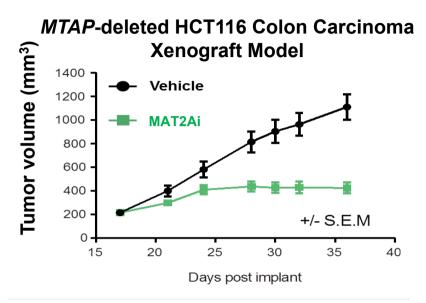


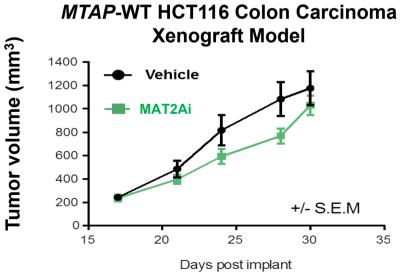


Agios MAT2A Inhibitors Selectively Impact Proliferation of *MTAP*-null Cancers



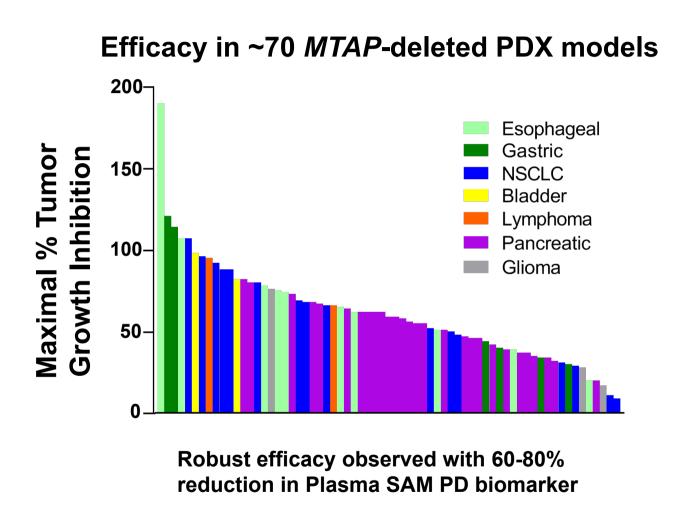


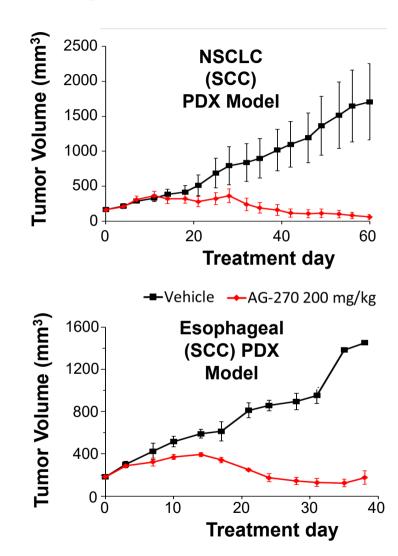






MAT2A Inhibitor AG-270 Possesses Broad Activity in 'Mouse Clinical Trial' Using Patient Derived Xenograft Models



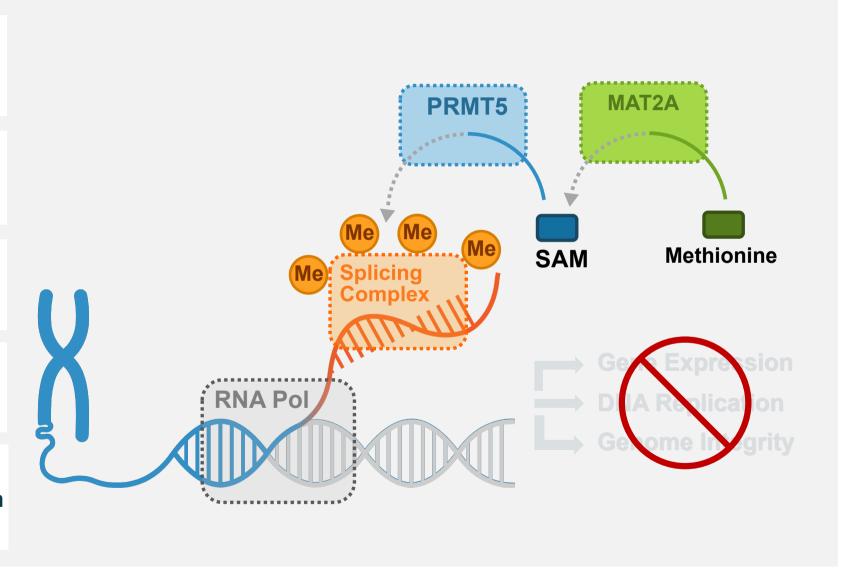


N=3 per model; established tumors treated at 200 mpk AG-270 QD



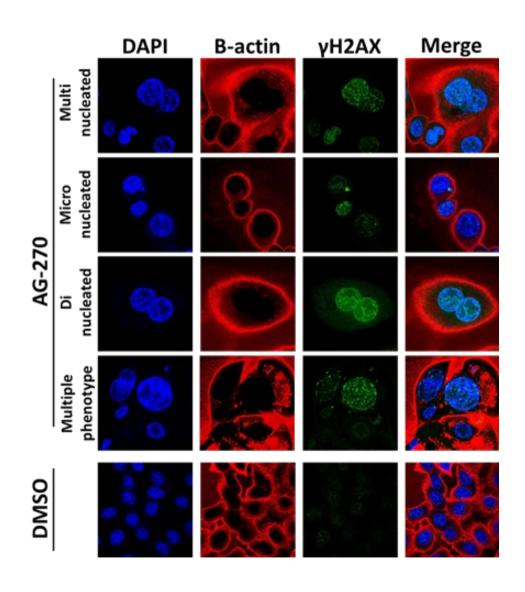
Mechanistic Understanding of the Pathway Downstream of MAT2A

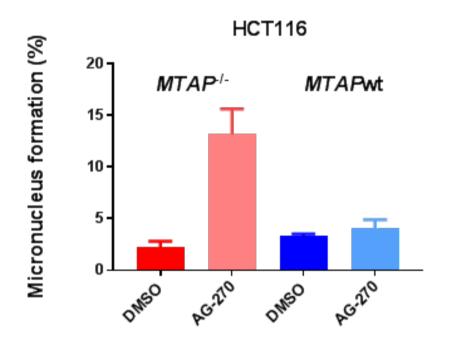
- RNA splicing concurrent with transcription
- 2 Splicing complex requires PRMT5
- MAT2A inhibition blocks splicing
- Defects in gene expression, DNA replication, genome integrity
- 5. DNA repair and cell cycle defects, leading to actionable combination partners including taxanes





AG-270 Treatment Induces Substantial Mitotic Defects in HCT116 *MTAP*-/- cells

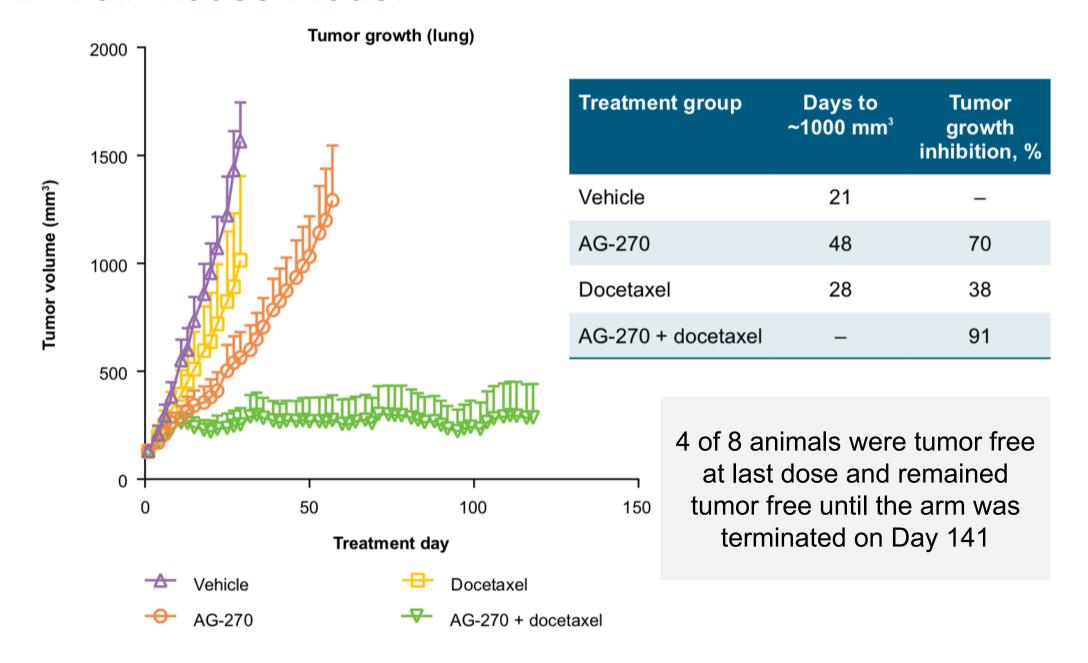




- Single Agent AG-270 treatment leads to DNA damage (γH2AX) and micronuclei formation
- Effects are selectively observed in *MTAP*-/-cells and not in *MTAP*-wt cells

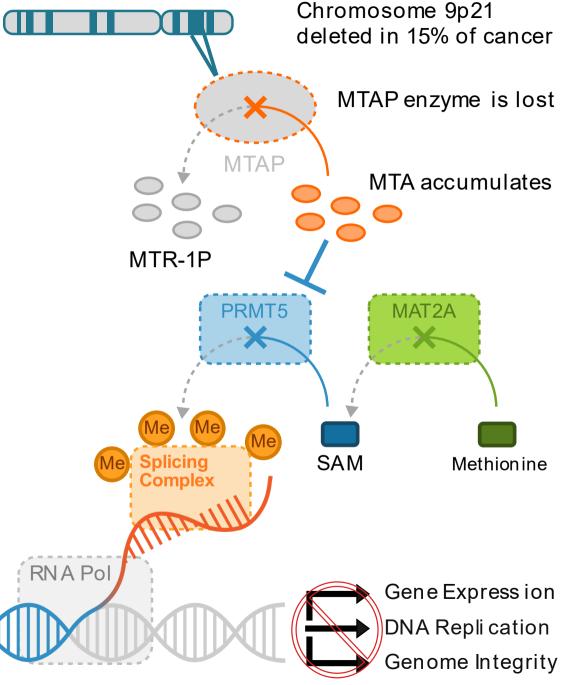


AG-270 Enhanced Docetaxel Treatment in an NSCLC (SCC) *MTAP*-null Mouse Model





MAT2A Inhibition Leads to DNA Damage and Cell Cycle Defects, Leading to Strong Synergy with Anti-mitotic Taxanes





A phase 1 trial of AG-270 in patients with advanced solid tumors or lymphoma with homozygous *MTAP* deletion

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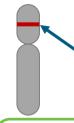
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Study design

Phase 1, open label, multicenter study (ClinicalTrials.gov NCT03435250) Adult patients Arm 1. n=39: Continuous oral AG-270 QD or BID in 28-**Primary objective:** day cycles, until disease progression or unacceptable MTD of AG-270 with: toxicity Secondary objectives: Advanced solid Dose escalation guided by a Bayesian logistic regression Safety and tolerability tumors or model lymphoma PK and profiling of potential PK/PD without effective metabolites sampling: Single standard C1D8 C1D15 • PD (changes in circulating dose D-3 treatment SAM and methionine options **Assessments** concentrations) Homozygous PK/PD: Antitumor activity CDKN2A or Plasma SAM and methionine concentrations **Exploratory objective:** MTAP deletion Tumor biopsies for SDMA assessment by IHC • PD in tumor tissue (changes (Figure 3) before the start of treatment and at C2D1 in SDMA methyl marks). Efficacy: · Disease status every 2 cycles

BID = twice daily; C = cycle; D = day; IHC = immunohistochemistry; MTD = maximum tolerated dose; PK/PD = pharmacokinetics/pharmacodynamics; QD = once daily

MTAP and CDKN2A deletion for patient selection



MTAP and CDKN2A within 100 kbp of each other on chromosome 9p21

- Chr9p21 deleted in ~15% of cancers1
- MTAP loss commonly coincides with CDKN2A loss²

Analysis of Cancer Genome Atlas data

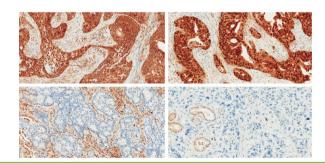
| Cancer type | CDKN2A deletion, % | MTAP deletion, % | Tumors with CDKN2A/MTAP co-deletion, % |
|------------------------|-----------------------|---------------------|--|
| Pancreatic | 28 | 25 | 88 |
| DLBCL | 31 | 23 | 73 |
| Esophageal and gastric | 25 | 19 | 76 |
| Lung (all) | 23 | 19 | 82 |

Required for patient enrollment: evidence of homozygous *CDKN2A* deletion by local testing (e.g. next-generation sequencing) or of homozygous MTAP deletion by a central IHC assay

IHC assay optimized for MTAP protein expression in FFPE tumor tissue, with <20% MTAP-positive cells as the cutoff value for MTAP deletion

NSCLC (SCC)
MTAP-positive tumor cells = 100%

NSCLC (adenocarcinoma)
MTAP-positive tumor cells = 0%



Pancreatic (adenocarcinoma)
MTAP-positive tumor cells = 100%

Pancreatic (adenocarcinoma) MTAP-positive tumor cells = 0%

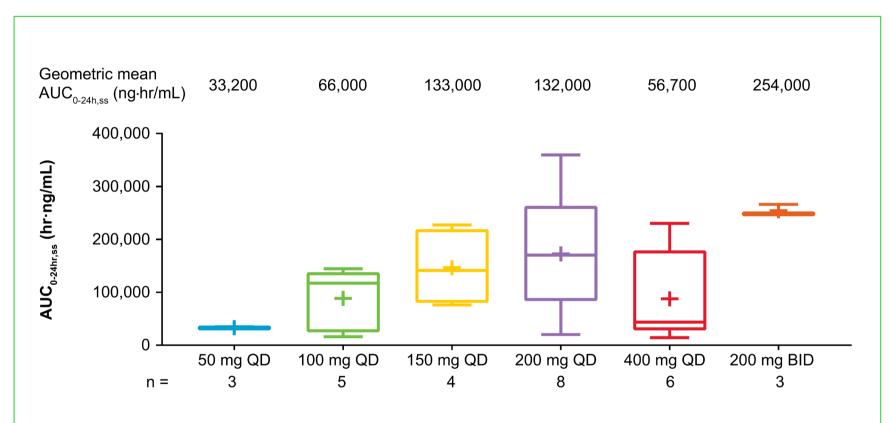
DLBCL = diffuse large B-cell lymphoma; FFPE = formalin-fxed paraffn-embedded; NSCLC = non-small-cell lung cancer; SCC = squamous cell carcinoma

Patient Characteristics

| Dose | 50 mg QD | 100 mg QD | 150 mg QD | 200 mg QD | 400 mg QD | 200 mg BID | | |
|---|--|-----------|-----------|-----------|---|------------|--|--|
| Patients, n | 3 | 7 | 6 | 11 | 6 | 6 | | |
| Baseline char | Baseline characteristic | | | N=39 | | | | |
| Age, median (ran <60, n (%) ≥60, n (%) | ge), years | | | (| 65 (32–87) 17 (44) 22 (56) | | | |
| Male sex, n (%) | | | | | 21 (54) | | | |
| Enrollment on the CDKN2A deletion, MTAP deletion, | on, n (%) | | | | 34 (87) 5 (13) | | | |
| evaluable for MT/ | n <i>CDKN2A</i> deletion A <i>P</i> deletion by IHC th <i>CDKN2A</i> deletio | , n (%) | on by | | 22 (56) 15 (68) | | | |
| Primary tumor type Bile duct cancer Pancreatic cance Mesothelioma NSCLC Other cancer type | eer | | | | 7 (18) 7 (18) 4 (10) 4 (10) 17 (44) | | | |
| Number of lines of One Two Three or more | of prior therapy, n (| %) | | | 12 (31) 9 (23) 18 (46) | | | |
| aCDKN2A status based on local testing, MTAP status by IHC performed centrally | | | | | | | | |

Pharmacokinetics

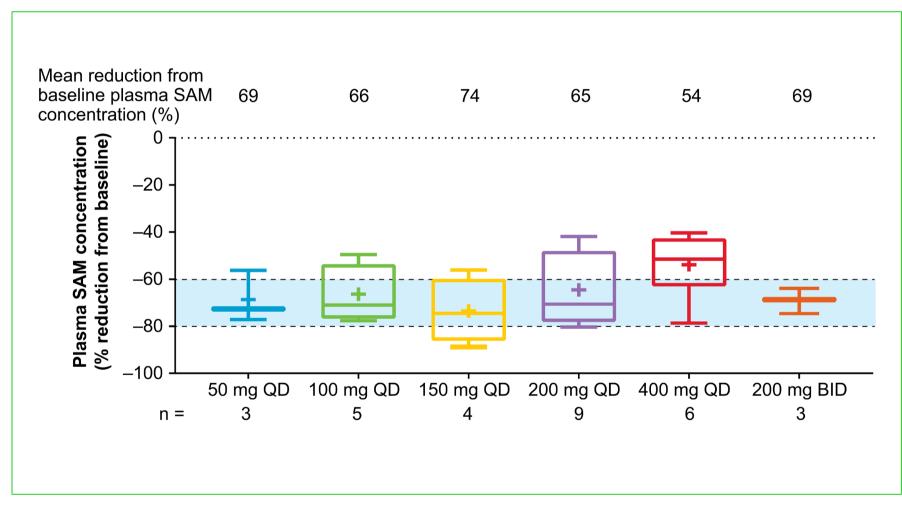
 $AUC_{0.24hr}$ = AUC from 0 to 24 hr; ss = steady state



Box denotes 25th to 75th percentiles, horizontal bar the median, and + the mean, with whiskers extending to the minimum and maximum values

- Mean exposure increased in an approximately dose-proportional manner between 50 mg QD and 200 mg QD
- Mean exposure was lower at 400 mg QD than 200 mg QD, possibly secondary to a reduction in oral bioavailability
- Due to this observation, a dose of 200 mg BID was evaluated, which increased steady-state area under the plasma concentration-time curve (AUC) by 1.9-fold relative to a dose of 200 mg QD.

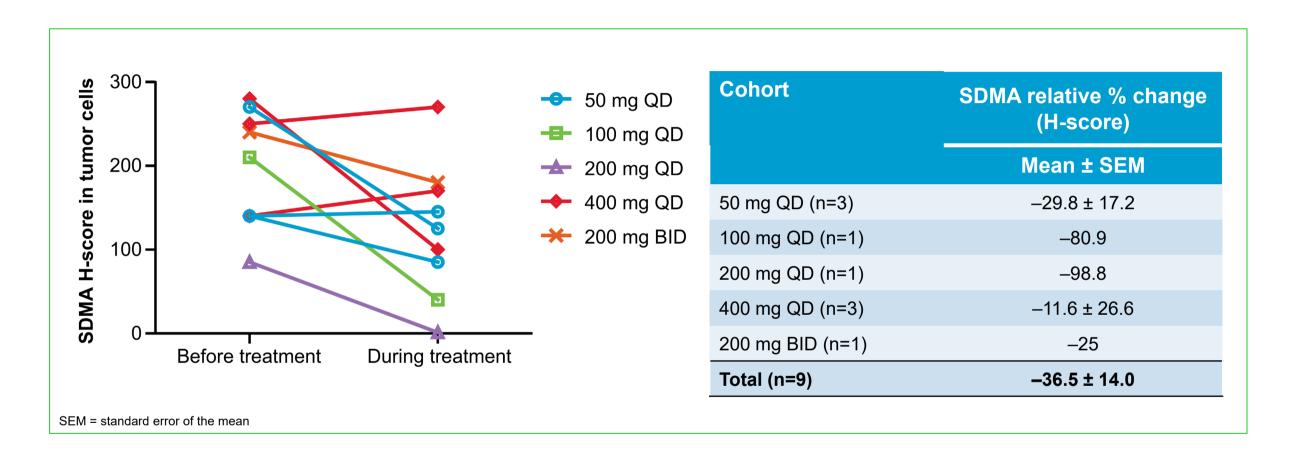
Reductions in plasma SAM concentration at steady state (C1D15)



- Plasma SAM concentration at C1D15 decreased by 65–74% across doses of 50–200 mg QD and 200 mg BID
 - The lower reduction in plasma SAM concentration (~54%) observed at 400 mg QD is consistent with the lower AG-270 exposure observed at this dose
- Average reductions in plasma SAM concentration are within the range associated with maximum tumor growth inhibition in preclinical models (60–80%)

Box denotes 25th to 75th percentiles, horizontal bar the median, and + the mean, with whiskers extending to the minimum and maximum values

SDMA expression by IHC in paired pre- and post-dose tumor biopsies



- Analysis of nine paired tumor biopsies by IHC showed decreases in levels of SDMA residues, consistent with MAT2A inhibition
- The average (min, max) H-score reduction compared with baseline was 36.5% (–98.8%, +21.4%).

Summary of AEs and dose-limiting toxicities by dose cohort

| | 50 mg QD n=3 | 100 mg QD n=7 | 150 mg QD n=6 | 200 mg QD n=11 | 400 mg QD n=6 | 200 mg BID n=6 | Total N=39 |
|--|---------------------------------|---------------------------------|--|--|---------------------------------|--|---|
| Patients with any AG-270–related AE, n (%) | 3 (100) | 4 (57) | 4 (67) | 6 (55) | 4 (67) | 5 (83) | 26 (67) |
| Most common (>10%) AG-270–related AE, n (%) Increased blood bilirubin Fatigue Decreased platelet count Rash | 0 2 (67) 0 2 (67) | 1 (14) 3 (43) 1 (14) 0 | 2 (33) 1 (17) 1 (17) 2 (33) | 4 (36) 1 (9) 1 (9) 1 (9) | 0 1 (17) 0 0 | 3 (50) 1 (17) 3 (50) 1 (17) | 10 (26) 9 (23) 6 (15) 6 (15) |
| Patients with grade 3 or higher AG-270–related AE, n (%) Increased blood bilirubin Decreased neutrophil count Decreased platelet count Decreased white blood cell count Lymphopenia Anemia Rash Liver injury | 0 0 0 0 0 0 0 | 0 0 0 0 0 0 0 | 1 (17) 1 (17) 1 (17) 1 (17) 1 (17) 1 (17) 1 (17) 1 (17) | 2 (18) 1 (9) 1 (9) 0 0 0 0 | 0 0 0 0 0 0 0 | 4 (67) 2 (33) 0 2 (33) 0 0 0 1 (17) 2 (33) | 7 (18) 4 (10) 2 (5) 3 (8) 1 (3) 1 (3) 1 (3) 2 (5) 2 (5) |
| Dose-limiting toxicities, n (%) Increased blood bilirubin Decreased neutrophil count Decreased platelet count Rash Acute liver injury | 0 0 0 0 | 0 0 0 1 (14) 0 | 1 (17) 0 0 1 (17) 0 | 0 1 (9) 0 0 | 0 0 0 0 | 0 0 1 (17) 1 (17) 2 (33) | 1 (3) 1 (3) 1 (3) 3 (8) 2 (5) |

Summary of AEs and dose-limiting toxicities

- Generalized erythematous rash in three patients, treated at 100 mg QD, 150 mg QD and 200 mg BID:
- Onset during second week of treatment, resolved <1 week after AG-270 interruption
- Successful rechallenge at a lower dose in two patients.

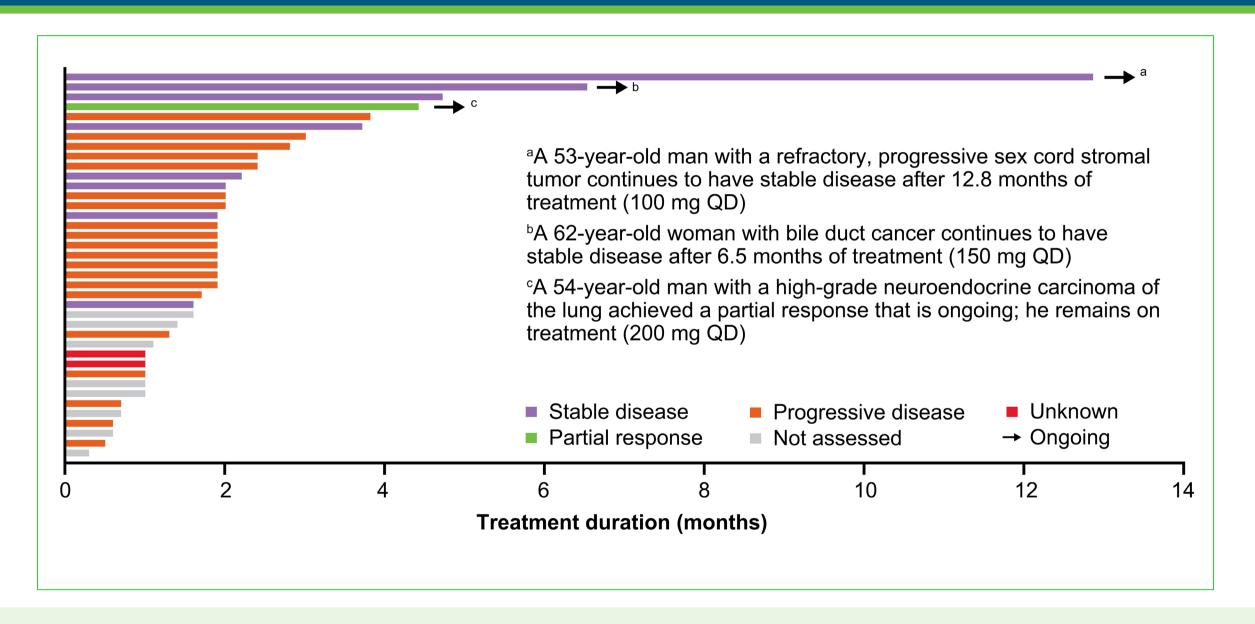
QD cohorts

- Increases in unconjugated bilirubin, starting at 100 mg QD:
 - Consistent with UGT1A1 inhibition, exposure-dependent, reversible.
- Mild myelosuppression, starting at 200 mg QD:
 - Most consistently manifested as reversible thrombocytopenia (with or without leukopenia/anemia).

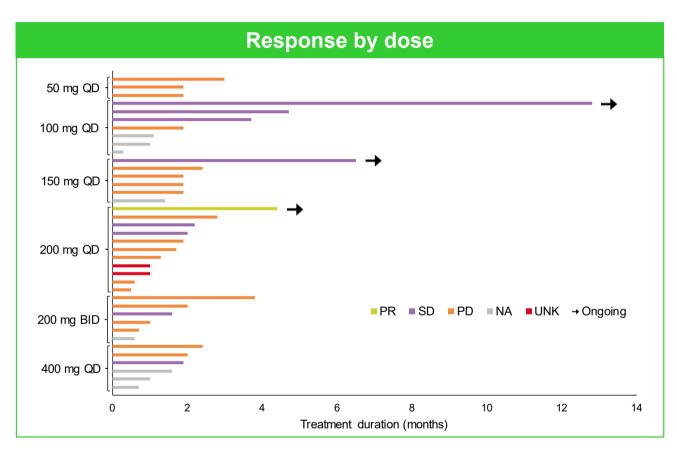
200 mg BID cohort

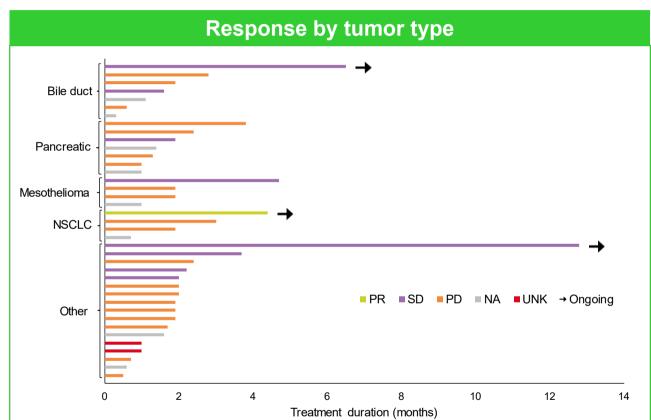
- Reversible acute liver injury in two of six patients:
 - Asymptomatic grade 3 and 4 increases in alanine aminotransferase, aspartate aminotransferase, and total bilirubin
 - Outpatient treatment with oral steroids, leading to complete resolution
 - Not clearly related to higher AG-270 systemic exposure.
- Grade 3 and 4 thrombocytopenia in two of six patients.
- MTD was determined to be 200 mg QD.

Duration of treatment and best overall response in patients receiving AG-270



Duration of treatment and best overall response in patients receiving AG-270

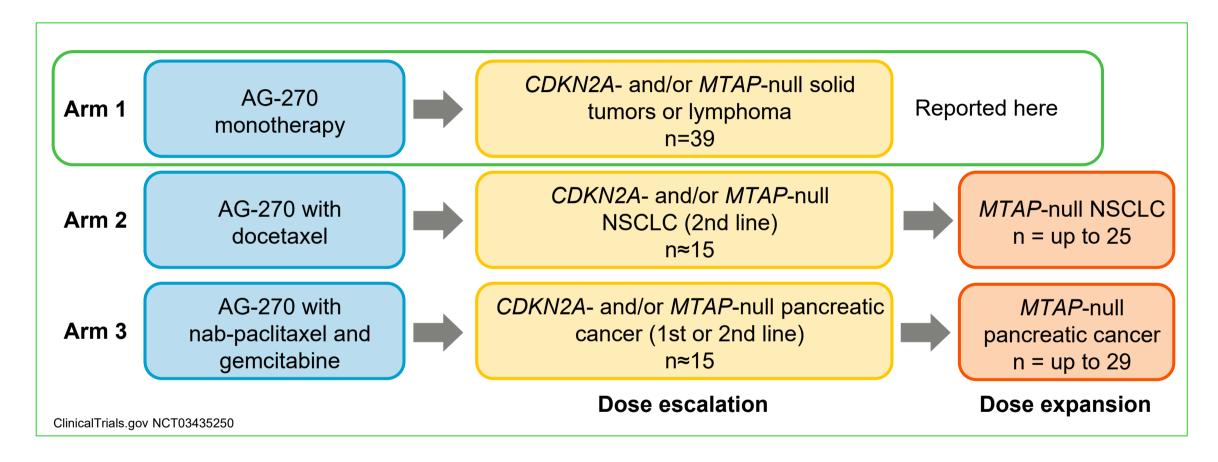




Conclusions

- AG-270 is the first MAT2A inhibitor to be evaluated in humans.
- The MTD was determined to be 200 mg QD.
 - DLTs included transient diffuse rashes, neutropenia and thrombocytopenia, and reversible acute liver injury.
- AG-270 generates reductions in plasma SAM concentration and in levels of tumor SDMA at welltolerated doses.
- Average reductions in plasma SAM concentration were similar between 50 and 200 mg QD, and within the range associated with maximum tumor growth inhibition in preclinical models (60–80%).
- Objective tumor response was uncommon in this group of patients with treatment-refractory malignancies.
 - However, a confirmed partial response was observed in a patient with a high-grade neuroendocrine carcinoma of the lung and two patients experienced prolonged stable disease of more than 6 months.

Two arms of the Phase 1 trial combining AG-270 with taxanes currently enrolling patients



Dosing with AG-270 in the combination arms will start at 100 mg QD and can be increased to 200 mg QD

Q&A

