Abstract #151606

COP 151606: ML-0207/ASP8731: A Novel BACH1 Inhibitor That Induces Fetal Hemoglobin in Treatment of Sickle Cell Disease

Review Category 113: 113. Hemoglobinopathies, Excluding Thalassemia—Basic and Translational Science (20297)

Abstract Text

Sickle cell disease (SCD) is a genetic disorder caused by a point mutation in the β globin subunit resulting in hemoglobin S (HbS). Following deoxygenation of red blood cells, HbS forms polymers that can promote hemolysis and the release of free heme that cause pro-oxidative and pro-inflammatory stress, vaso-occlusive pain crises, and ischemia-reperfusion pathophysiology. Heme also functions as an intracellular activator of antioxidant and globin gene expression. Heme binds to the transcriptional repressor BTB and CNC homolog 1 (BACH1), which relieves BACH1's repression of gene transcription. The release of BACH1 repression increases the binding of nuclear factor erythroid 2-related factor 2 (NRF2) to antioxidant response elements (ARE) and the cellspecific transcription of antioxidant genes such as heme oxygenase-1 (HMOX1), glutathione reductase (GR), solute carrier family 7 member 11 (SLC7A11), and NAD(P)H dehydrogenase [quinone] 1 (NQO1). We have previously shown that pharmacologic activation of the NRF2 pathway in SCD mice provides protection against heme-induced vascular occlusion, is anti-inflammatory, and decreases hepatic necrosis. NRF2 activation also promotes erythroid expression of the A-gamma (HBG1) and Ggamma (*HBG2*) globins, which are subunits of hemoglobin F (HbF) that replace β^{s} globins and thus increase HbF and decrease HbS in red blood cells. Thus, BACH1 inhibitors have the potential to increase expression of antioxidant and HbF genes and prevent or reduce SCD-related pathophysiology, resulting in reduction of hemolysis, inflammation, and vaso-occlusive pain crises. Mitobridge is currently developing ML-0207/ASP8731, a highly potent, selective small molecule inhibitor of BACH1 capable of activating the Nrf2 pathway in human and murine models and investigated the ability of ML-0207 to modulate antioxidant and anti-inflammatory genes and induce HbF in human translational cellular models and a preclinical murine model of SCD. ML-0207 induced mRNA expression of Nrf2 target genes HGB1, HBG2, HMOX1, SLC7A11, GCLM, and NQO1 in human bone marrow-derived CD34+ cells differentiated to erythrocytes. We observed 2-fold increases in both the percentage and number of CD71+/HbF+ erythrocytes by FACS using 1 µM ML-0207 and 10 µM HU compared to DMSO control (Figure 1A). The combination of ML-0207 and HU induced significantly

more HbF+ erythrocytes compared to each drug alone (Figure 1B). In a single healthy CD34+ donor non-responsive to 10 μ M HU, we observed ML-0207 was able to significantly induce CD71+/HbF+ cells at 1 & 3 μ M (Figure 1C). In Townes SCD mice, there were significant increases in heme oxygenase 1 and decreases in VCAM-1, ICAM-1, and decreases in phospho-p65 NF- κ B protein. Furthermore, we observed a significant reduction in hemin-induced vaso-occlusion and an increase in the percentage of F-cells. The increases in F-cells were accompanied by increases in blood A-gamma globin and erythrocytes and decreases in leukocytes. Taken together, these data support BACH1 inhibitors as potential novel and effective treatments for SCD patients.

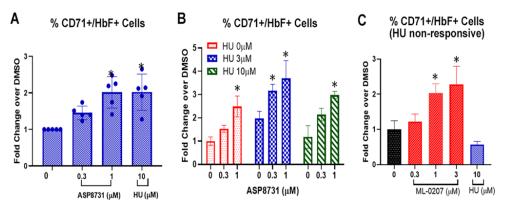


Figure 1. BACH1 Inhibitor ML-0207/ASP8731 Induces HbF Cells. Human bone marrow-derived CD34+ cells were incubated with the indicated concentrations of ASP8731, HU, vehicle (DMSO), or ASP8731 + HU and differentiated into erythroid cells. Differentiated cells were stained with fluorescent antibodies to CD71 and HbF. The percent CD71+/HbF+ cells relative to CD71+/HbF+ cells was calculated for each treatment and expressed relative to DMSO-treated cells as fold change. (A) Data is mean ± standard deviation (SD) of 5 donors. (B) Data is mean ± SD of 2 donors, each treatment done in triplicate per donor. (C) Data is mean ± SD of triplicate measurement from one donor. For all data (A-C), *p<0.001, one way ANOVA followed by Tukey's multiple comparison.

Presenter has relevant financial relationship(s) to disclose: Yes.