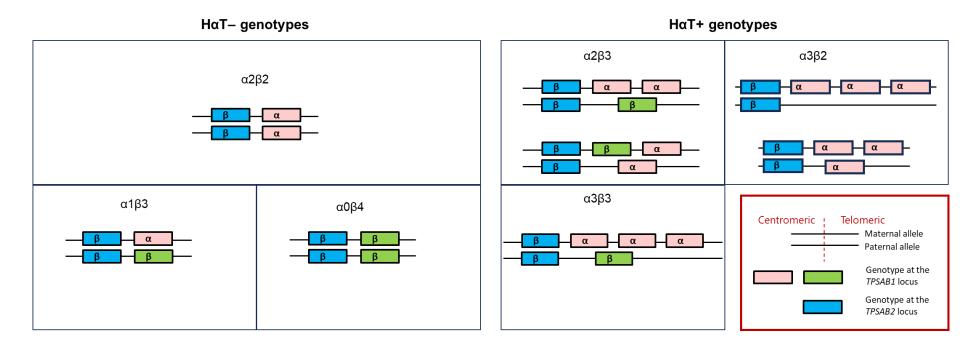
Analysis of Hereditary Alpha Tryptasemia and Association With Baseline Characteristics in Patients With Indolent Systemic Mastocytosis Enrolled on the PIONEER Study

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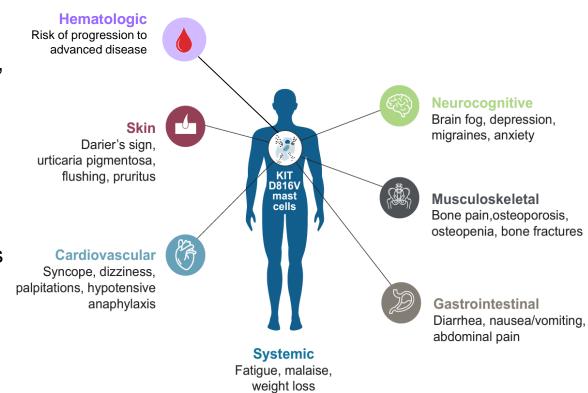
Hereditary α-tryptasemia (HαT)

- HαT is a genetic trait defined by an increased TPSAB1 gene copy number coding for alpha tryptase. This leads to elevated serum levels of tryptase, a protein produced by mast cells (MCs)^{1–4}
 - TPSAB1 encodes for the α allele (which may be present in multiple copies) or the β allele
 - TPSAB2 encodes for the β allele only²
- HαT can be associated with symptoms of MC diseases such as anaphylaxis, gastrointestinal symptoms, and skin symptoms^{1–4}



Patients with SM have higher rates of $H\alpha T$ than the general population

- HαT is present in 4–6% of the general population^{1–3}
- Systemic mastocytosis (SM) is a clonal mast cell disease, driven by the KIT D816V mutation
 - Patients with SM may experience life-long debilitating symptoms and disease progression^{4–6}
 - Elevated tryptase is one of the minor diagnostic criteria although not all patients with SM experience high serum tryptase levels^{4–6}
- Presence of HαT may overlap with SM and having both is associated with increased risk of severe anaphylaxis^{1–3}



Recent studies have found that HαT is present in 9–18% of patients with SM¹-³

PIONEER allowed the evaluation of the relationship between $H\alpha T$ and SM

- PIONEER (NCT03731260): Randomized, double-blind, placebo-controlled study in patients with indolent systemic mastocytosis (ISM)
 - Avapritinib is a potent and highly selective oral therapy targeting KIT D816V, the underlying driver of SM
 - The safety, efficacy, and quality of life in patients with ISM receiving avapritinib plus best supportive care (BSC) compared with patients receiving placebo plus BSC were evaluated
- A total of 250 patients were screened for HαT (positive, HαT+; negative, HαT–) by digital polymerase chain reaction detection of copy number variations of the *TPSAB1* gene on the tryptase locus^a
- Baseline objective measures of disease burden were recorded;
 KIT D816V variant allele fraction (VAF), serum tryptase levels,
 bone marrow (BM) MCs, and skin MCs

The cohort of patients enrolled on the PIONEER study represent one of the largest, most well characterized populations of patients with ISM to date

Here, we use this rare opportunity to compare baseline objective disease and symptom burden in H α T+ *versus* H α T- patients with ISM

Patient baseline characteristics were similar between groups

In patients screened, 11% (28/250) were $H\alpha T$ +

- Of the HαT+ patients, 14 had the α2β3 genotype, 12 had the α3β2 genotype, and two had the α3β3 genotype
- The median age of patients who were HαT+ was slightly younger than that of patients who were HαT– (P=0.01) and prior therapies were similar across groups

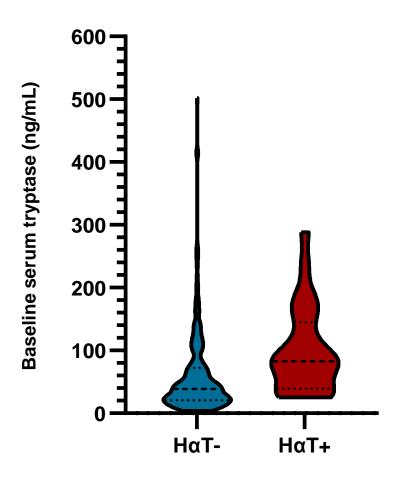
	Patients		
	HαT+ (N=28)	HαT- (N=222)	
Age (years), median (range)	46 (21–64)	51 (18–79)	
Female, n (%)	22 (79)	161 (73)	
Corticosteroid use for SM, n (%)	3 (11)	17 (8)	
Prior TKI therapy, n (%)	1 (4)	16 (7)	
Prior cytoreductive therapy, n (%)	4 (14)	27 (12)	

Baseline tryptase and KIT D816V VAF were significantly different between H α T+ and H α T- patients

- Baseline serum tryptase was significantly higher in HαT+ patients
- Baseline KIT D816V VAF in the peripheral blood was significantly lower in HαT+ patients
- BM MC burden was similar in patients with and without HαT
- No patients with HαT had palpable spleens or palpable livers
- Additionally, history of anaphylaxis was similar in patients with and without HαT

	Patients		Dyelie	
	HαT+ (N=28)	HαT- (N=222)	P-value	
Median serum tryptase, ng/mL (range)	82.6 (25.2–288.0)	38.6 (3.6–501.6)	0.0001	
Serum tryptase >20 ng/mL, n (%)	28 (100)	171 (77)	NA	
Median KIT D816V VAF, % (range)	0.06 (0.0–16.2)	0.40 (0.0–41.3)	0.0004	
Median BM MC burden, % (range)	6.0 (2.0–30.0)	7.0 (1.0–70.0)	ns	
MC aggregates present, n (%)	20 (71)	177 (80)	NA	
Median lesional skin MC density, counts/mm² (range)	336 (83–4300)	489 (53–2870)	ns	
Median non-lesional skin MC density, counts/mm² (range)	114 (61–1337)	130 (10–659)	ns	
Palpable spleens, n (%)	0 (0)	4 (2)	ns	
Palpable livers, n (%)	0 (0)	12 (6)	ns	
Anaphylaxis, n (%)	5 (18)	37 (17)	ns	

Baseline serum tryptase levels were higher in the HαT+ group



	Patients		P-value
	HαT+ (N=28)	HαT- (N=222)	r-value
Serum tryptase levels, median ng/mL (range)	82.6 (25.2–288.0)	38.6 (3.6–501.6)	0.0001

- No HαT+ patients (n=28) with moderate-to-severe ISM had a baseline serum tryptase of <20 ng/mL
- Of the HαT– patients with moderate-to-severe ISM, 23% (n=222) had a baseline serum tryptase of <20 ng/mL

Regardless of $H\alpha T$ status, patients had similar response to avapritinib therapy^a

	HαT+ patients	HαT– patients
Mean percent change in serum tryptase at 24 weeks	−51% (n=18)	-45% (n=116)
Mean percent change in KIT D816V VAF at 24 weeks	−54% (n=11)	-56% (n=98)
Mean percent change in ISM-SAF TSS at 24 weeks	−32% (n=17)	-33% (n=114)

Conclusions

- The prevalence of HαT in the PIONEER population was 11%, which is consistent with previously reported frequencies of HαT in patients with ISM
- Compared to patients without HαT, patients with HαT had significantly higher baseline tryptase values and significantly lower KIT D816V VAF in the peripheral blood
 - There were no differences in the rate of anaphylaxis or ISM-SAF TSS at baseline
- Regardless of HαT status, patients had similar symptom and disease biomarker reductions after treatment with avapritinib

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