The CXCL12/CXCR4 Pathway as a Potential Target of Tipifarnib: Preliminary Results from an Open-Label, Phase II Study in Relapsed or Refractory Peripheral T-cell Lymphoma

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BACKGROUND

Tipifarnib:

- Potent and highly selective inhibitor of Farnesyl Transferase (FT)
- FT catalyzes post-translational attachment of farnesyl groups required for localization of signaling molecules to the inner cell membrane.

CXCL12

- Chemokine that is essential for T cell homing to lymphoid organs and the bone marrow, and for the maintenance of immune cell progenitors.
- CXCL12 has been postulated to signal in part through HRAS, a signaling protein that is uniquely dependent on farnesylation for activity.

METHODS

This is a Phase 2 study designed to investigate the antitumor activity of tipifarnib in patient (pts) with relapsed or refractory PTCL.

Primary Objective: ORR by IWC and/or mSWAT Secondary Objectives: PFS, DOR, safety/tolerability **Exploratory Objectives:** Biomarkers

Expansion Cohorts: AITL, CXCL12+ PTCL

- Based on observed antitumor activity and biomarker evaluation in stages 1 and 2, the study is being amended to include expansion cohorts in AITL (enrolling) and CXCL12+ PTCL (N = 12 each).
- Tipifarnib 300 mg twice daily (bid) on days 1 21 of 28-day treatment cycles.

Clinical trial information: NCT02464228

CONCLUSIONS

- Encouraging activity of tipifarnib was observed in PTCL pts, particularly in those with tumors of AITL histology and high CXCL12 expression.
- Study is being amended to continue enrollment of pts with CXCL12+ **PTCL.** Enrollment in the AITL cohort is ongoing.
- Tipifarnib was generally welltolerated. Most common treatmentrelated AEs (grade \geq 3) were hematology related, including neutropenia, thrombocytopenia, leukopenia, anemia and febrile neutropenia.

CLINICAL BENEFIT FROM TIPIFARNIB IS ASSOCIATED WITH CXCL12 GENOTYPE



CLINICAL BENEFIT FROM TIPIFARNIB IS ASSOCIATED WITH CXCL12 EXPRESSION

CXCL12 expression was a specific biomarker of clinical benefit in patients receiving tipifarnib







RESULTS

RNA Seq Experiment#1

Pt ID	2002	2004	3002	4001	2001	4004
HIS	NOS	NOS	NOS	NOS	NOS	AITL
RESP	PD	PD	PD	SD	SD	PR
CXCL12	469	759	613	2659	1160	3265
CXCL13	4602	303	351	1729	1039	25355
					•	
VCAM1	3643	2328	1776	4111	1118	7359
PDGFRA	898	1538	1453	1038	440	1080
ANGPT1	0	21	188	84	0	28
TEK	495	634	791	734	138	691
GAPDH	17369	10574	9485	12101	6328	16189

RNA Seq Experiment#2

2005	2006	4005	4006	4007	6003	2007
NOS	NOS	NOS	NOS	NOS	NOS	AITL
PD	PD	PD	PD	PD	SD	PR
						1
1570	325	834	1211	1081	3728	14076
		•				•
16876	47	715	10	2088	424	1176
2874	1255	5746	546	5344	4785	29134
1415	74	2037	2872	684	856	6227
34	43	24	63	22	93	769
560	192	207	990	436	389	4542
	,					
45614	22595	15801	112381	17003	27244	10403
	2005 NOS PD 1570 16876 2874 1415 34 560 45614	2005 2006 NOS NOS PD PD 1570 325 16876 47 2874 1255 1415 74 34 43 560 192 45614 22595	200520064005NOSNOSNOSPDPDPD157032583416876477152874125557461415742037344324560192207456142259515801	2005200640054006NOSNOSNOSNOSPDPDPDPD157032583412111687647715102874125557465461415742037287234432463560192207990456142259515801112381	20052006400540064007NOSNOSNOSNOSNOSPDPDPDPDPD1570325834121110811687647715102088287412555746546534414157420372872684344324632256019220799043645614225951580111238117003	200520064005400640076003NOSNOSNOSNOSNOSPDPDPDPDPD1570325834121110811570325834121110813728168764771510208842428741255574654653444785141574203728726848563443246322935601922079904363894561422595158011123811700327244

CXCL12 3UTR

reference rs2839695

Days

600

N=15⁴ P=0.038

500

007

Total Treated AITL ALCL, ALK PTCL, NO

Prior Lines of

Total Discon Reasons for Progressiv Withdrawa

SAFETY & TOLERABILITY

- (16%)
- 2 of the study.

High CXCL12 expression may result from CXCL13-induced CXCL12 upregulation and angiogenesis



Results based on preliminary data as of 22 Nov 2017.

- 1. Eight subjects carried rs2839695; one subject presented a novel 3UTR variant, and 8 subjects carried reference 3UTR CXCL12 sequences
- 2.Last prior therapy: Chemotherapy (7),
- brentixumab (4), HDACi (2) lenalidomide (2), ruxolitinib (1)
- 3.Median overall PFS for tipifarnib (5th line) in subjects with NGS data was 53 days. Median
- overall PFS for these subjects on their last prior was 82 days
- 4.Low CXCL12 expression was observed in
- tumors samples carrying the CXCL12
- rs2839695 3UTR variant
- 5.SPD data not available for 2 pts: 004-006, 004-

PATIENT DISPOSITION

d	N (%)	19 (100)					
	n (%)	3 (16)					
<pre>K negative</pre>	n (%)	1 (5)					
S	n (%)	15 (79)					
of Therapy	Median (Range)	4 (1 – 7)					
tinuations	n (%)	17 (89)					
Discontinuation:							
ve Disease	n (%)	16 (94)					
al of Consent	n (%)	1 (6)					

Toxicities were consistent with the known safety profile of tipifarnib.

Grade \geq 3 drug-related TEAEs occurring in \geq 10% of pts were hematology related: neutropenia (74%), thrombocytopenia (58%), leukopenia (47%), anemia and febrile neutropenia (32% each) and lymphopenia

Myelosuppression was manageable with treatment interruption, dose reductions, growth factor or transfusion support.

The dose regimen was amended to 300 mg bid on days 1-21 of 28 day treatment cycles due to the observed tolerability profile of the alternate week regimen tested in stages 1 and