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BACKGROUND

- C-C motif chemokine receptor 8 (CCR8) is a G-protein coupled receptor that highly expressed on tumor-infiltrating T regulatory cells (Tregs)¹
- Intratumoral CCR8+ Tregs are highly immunosuppressive, enriched in the tumor microenvironment (TME) and frequently associated with a lack of response to immunotherapy, therefore selective depletion of intratumoral Tregs is an attractive immunotherapy strategy²
- Strong scientific rationale for targeting CCR8 in head and neck squamous cell carcinoma (HNSCC) and other solid tumors:
- CCR8+ Tregs are most abundant in HNSCC, cervical squamous cell carcinoma (SCC), and gastric cancer, and broadly expressed in several solid tumors, including colorectal cancer (CRC) and lung cancer
- Radiation and PD-1 blockade on Tregs are known to increase Tregs, supporting the use of anti-CCR8 with anti-PD-1 agents
- CHS-114 is a selective, cytolytic anti-CCR8 monoclonal antibody (mAb) with no off-target binding³
- CHS-114 has the potential to overcome Treg-mediated immune suppression within the TME by recruiting T cells, turning cold tumors to hot, and enhancing antitumor immunity when combined with other immuno-oncology (IO) agents such as the anti-PD-1 antibody, toripalimab⁴



isotope control) was calculated. EC_{50} values of CHS-114 were calculated based on 4-1BB induction on NK cells. C-D) PBMCs were treated with increasing concentrations of CHS-114 at 37°C for 16 hours. The frequency of B7-H3 and CD83 expressing monocytes was determined by flow cytometry. EC₅₀ values of CHS-114 were calculated based on B7-H3 frequency (C) and CD83 frequency on monocytes (D). Based on the above, in vitro pharmacology data target trough levels for the expected pharmacological activity in participants treated with CHS-114 in clinical trials were defined.





Density (y-axis) vs frequency (x-axis) of CCR8+ Treas (of total FOXP3-Treas) in tumor microarrays from 12 types of solid tumors evaluated by a mIF assay. Tumor types in which clinical benefit has been observed with anti-CCR8 antibodies (CHS-114-101 study or competitor trials) are highlighted (red circles). For evaluating CCR8+ Tregs, CCR8 and FOXP3 staining was performed.





- A Phase 1, first-in-human (FIH), open-label single-agent and combination dose trial was initiated to evaluate CHS-114 in patients with advanced solid tumors and HNSCC. During monotherapy dose escalation (Stage 1a):⁵
- CHS-114 demonstrated an acceptable safety profile in heavily pre-treated patients with advanced solid tumors, with no DLTs reported to date and generally low-grade treatment-emergent adverse events (TEAEs)
- There were no complete or partial responses, but 47.4% of patients had stable disease (SD); one patient with SD >12 months remains on study
- Pharmacokinetic (PK) exposure increased with dose, was approximately dose proportional, and the elimination appeared linear with a half-life of about 10 days (range 9-17 days)
- Depletion of peripheral CCR8+ Tregs was observed (>85% in the periphery) and depletion was maintained over the dosing interval, establishing proof of mechanism; CHS-114 did not deplete non-CCR8+ Tregs and effector CD4 T cells (proof of specificity) • Two doses were selected for dose optimization based on safety, peripheral CCR8+ Treg depletion, PK and biomarker data

METHODS

- A Phase 1, FIH, open-label single-agent and combination dose trial to evaluate CHS-114 in patients with advanced solid tumors and HNSCC (NCT05635643)
- Here we report CHS-114 monotherapy and combination results in the HNSCC patients enrolled in the trial as of the January 24, 2025, data cutoff date



Phase 1 study of anti-CCR8 antibody CHS-114 with and without anti-PD-1 antibody toripalimab in patients with advanced solid tumors

cytotoxicity **(GZMB, C)** in PBMC samples were measured by Cytek FACS assay at indicated time points. An increase in IFNy and CD8+ T cell activation (Ki67, GZMB) compared t pre-treatment levels was observed through one dosing cycle for participants enrolled in CHS-114-101 monotherapy dose escalation and HNSCC dose expansion (only select participants from HNSCC dose expansion). Furthermore, IFNy induction and CD8+ T cell activation was sustained at DL5 and higher doses supportive of a pharmacologically and biologically active dose. Participants with longitudinal samples are plotted (C1D1 and C2D1); N's based on number of participants at baseline. Thick solid and dashed lines represen median values for groups defined in the legend; thin dashed lines represent individual

HNSCC Demographics & Baseline Characteristics				
	CHS-114 Monotherapy (n=14†)	CHS-114 + Toripalimab (n=7)		
Age, median years (range)	67 (42, 88)	67 (49, 79)		
Gender, n (%) Female Male	3 (21.4) 11 (78.6)	2 (28.6) 5 (71.4)		
Race, n (%) American Indian or Alaska Native Asian Black or African American Native Hawaiian or Other Pacific Islander White	0 0 2 (14.3) 0 12 (85.7)	0 0 0 7 (100.0)		
Unknown Other Not Reported	0 0 0	0 0 0		
ECOG, n (%) 0 1	4 (28.6) 10 (71.4)	5 (71.4) 2 (28.6)		
Median time since initial diagnosis, months (range)	25.5 (11, 184)	21 (15, 84)		
Lines of prior systemic therapy, n (%) 0 1-2 3-4 ≥5	1 (7.1) 7 (50.0) 5 (35.7) 1 (7.1)	0 5 (71.4) 2 (28.6) 0		
PD-L1 expression*, n (%) Positive Negative Not Done	13 (92.9) 1 (7.1) 0	7 (100.0) 0 0		

Patients with HNSCC from stage 1a (n=2) and stage 1b (n=12 *PD-L1 expression measured by combined positive score (CPS) or tumor proportion score (TPS



Confirmed Partial Response in Target Lesions



HNSCC Time on Treatment



Safety Summary in HNSCC			
AE Summary	CHS-114 Monotherapy (n=14)	CHS-114 + Toripalimab (n=7)	
Treatment-emergent adverse event (TEAE), n (%)	12 (85.7)	7 (100.0)	
Treatment-related ⁺ AE, n (%)	7 (50.0)	7 (100.0)	
Grade ≥3 TEAE, n (%)	4 (28.6)	5 (71.4)	
Grade ≥3 treatment-related† AE, n (%)	1 (7.1)	4 (57.1)	
Serious treatment-emergent adverse event (TESAE), n (%)	3 (21.4)	3 (42.9)	
Treatment-related ⁺ SAE, n (%)	0	0	
TEAE leading to study drug(s) discontinuation*, n (%)	0	4 (57.1)	
Treatment-related ⁺ AE leading to CHS-114 discontinuation [*] , n (%)	Ο	2 (28.6)	
TEAE leading to death, n (%)	0	0	
Treatment-related ⁺ AE leading to death, n (%)	0	0	

or monotherapy with HNSCC. treatment-related AEs are CHS-114-related AE; For stage 2 subjects, treatment-related AE are CHS-114 and/or toripalimab related AEs. *For stage 2 subjects, treatment discontinuation are CHS-114 and/or toripalimab discontinuation



Follow-up 3



LA: ~0.5 cm

RECIST Assessment:

- Target lesions: 40.0% reduction in the target lesions at the time of confirmed PR at follow-up 3
- Non-target lesions: 2 out of 4 non-target lesions disappeared after 2 treatment cycles
- No new lesions

Responder Demographics:

- Oropharyngeal squamous cell carcinoma (OPSCC)
- P16 positive, PD-L1 CPS 4, and TMB 8 mut/Mb
- CDKN2A, EGFR. Mutations: NOTCH, PIK3CA
- Patient was diagnosed at Stage 4 and refractory to 3 prior lines of systemic therapy including anti-PD-1 (~12 mo from 1L αPD-1 plus chemo)

RESULTS











- CHS-114 with and without toripalimab had a manageable safety profile in HNSCC patients
- remodeling and establishing proof of mechanism
- drugs such as T cell engagers and bispecifics

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CONCLUSIONS

• Two doses were selected for dose optimization based on safety, peripheral CCR8+ Treg depletion, PK and biomarker data; these two doses are supported by the observed immune activation in blood and CCR8+ Treg depletion in tumor

• CHS-114 with toripalimab had promising antitumor activity in HNSCC that warrants continued exploration

• In on-treatment tumor biopsies, CHS-114 depleted CCR8+ Tregs and increased CD8+ T cells in the TME indicating favorable TME

- CHS-114 administration leads to a substantial increase in CD8+ T cells in the TME providing a strong rationale for combining with other

• All together, these data support further evaluation of CHS-114 in combination with other drugs including toripalimab; a second-line (2L) HNSCC CHS-114 with toripalimab dose optimization study is ongoing