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Positive Impact of CT1812 (Zervimesine) Treatment on Plasma Biomarkers in Lower p-tau217 Subgroup Aligns with Clinical Benefits in Mild-to-Moderate AD Patients

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## Disclosures

### **Presenter Disclosures:**

• Mary Hamby, PhD is an employee of Cognition Therapeutics

### **Product Disclosure:**

- CT1812 (zervimesine\*) is an investigational therapeutic that has not been approved for any use by the US Food and Drug Administration or any other health authority
- Plans for subsequent clinical trials have not yet been reviewed by FDA or EMA

\*CT1812 assigned USAN name: zervimesine



# Zervimesine (CT1812) Mechanism of Action

Investigational, oral, small molecule oligomer antagonist

- Preclinical and clinical evidence that zervimesine acts to displace Aβ oligomers from synapses, facilitating clearance of Aβ oligomers in the cerebrospinal fluid (CSF)
- Proposed *synaptoprotective* mechanism of action to slow further neuronal injury / loss
- MoA distinct from anti-amyloid immunotherapies





### Phase 2 Safety and Efficacy Study in Adults with Mild-tomoderate Alzheimer's Disease



SHINE COG0201 study (NCT03507790) partially funded by \$31M NIA grant R01AG058660



Full safety and tolerability data are available in presentations from CTAD / AAIC on our website



udy protocol.

# Summary of SHINE Safety and Tolerability findings

Favorable safety profile vs placebo, AEs well balanced between arms, no ARIA

- Zervimesine demonstrated a favorable safety and tolerability profile
- Most TEAEs were mild or moderate in severity
- Similar percentages of adverse events in treated (76.5%) and placebo (78%) groups
- No discontinuations due to AEs in the 100mg dose group
- Most discontinuations were in 300mg group and all the reportable liver enzyme elevations were in 300mg group

#### **Adverse Events**

Zervimesine	76.5%					
Placebo	78.0%					
Serious AEs						
Zervimesine	4.9%					
Placebo	10.0%					
Deaths <sup>†</sup>						
Zervimesine	0					
Placebo	1 (cancer)					



### Plasma p-tau217 Reflects Brain Amyloid and Tau Burden

- Plasma p-tau217 is a blood-based biomarker representing AD pathology (amyloid plaques and tau)<sup>1</sup>
  - The degree of AD brain pathology can be indirectly assessed by measuring levels of plasma p-tau217<sup>1</sup>
- Given zervimesine MoA, we hypothesized that a larger treatment effect may be observed in participants with less AD pathology / lower tau
- SHINE included a prespecified subgroup analysis defined by median plasma p-tau217 (AlzPath, 1 pg/ml) at baseline



- Prior data indicate that individuals with lower AD pathology have greater response to amyloid-based therapies, eg:
  - Donanemab TRAILBLAZER 2
    - iADRS: 36% slowing in low tau tercile
    - iADRS: 21% slowing in high tau tercile



### Baseline Characteristics of Below/Above median p-tau217

### Reflects expected baseline characteristics based on mITT population

	mITT population (n=150)	Below median* p-tau217 Cohort (n=69)	Above or equal to median* p-tau217 Cohort (n=69)
Percent (%) female	60	59.4	58
Percent (%) white	96	94.2	97.1
Percent (%) non-Hispanic or Latino	92	89.9	97.1
ApoE4 Status: n (%) <ul> <li>Percent ApoE4 carriers</li> <li>Percent ApoE4 non-carriers</li> </ul>	91 (61) 59 (39)	42 (60.9) 27 (39.1)	43 (62.3) 26 (37.7)
Percent (%) concomitant AChEi or NMDA use	62.7	55.1	68.1
Mean age (range)	72.7 (51-85)	72.6 (51-84)	72.8 (53-85)
MMSE at baseline mean (range)	21.37 (13-29)	21.94 (14-29)	20.83 (13-28)
Plasma p-tau217 mean (range) in pg/mL	1.07 (0.2-3.5)	0.66 (0.2 - 1.0)	1.53 (1.0-3.5)
CSF neurofilament light chain mean (range) in pg/mL	1217.67 (220.0-2850.0)	994.70 (220.0 - 1840.0)	1389.88 (513.0 - 2850.0)



### SHINE Cognitive Endpoints: ADAS-Cog 11

Preservation of cognition in participants in below-median plasma p-tau217<sup>†</sup> subgroup





#### Above median p-tau217 (N=69)



\* ADAS-Cog 11 mITT in the pooled dose group vs placebo was the first of the ordered secondary efficacy endpoints

<sup>†</sup> Median plasma p-tau217 level is 1.0pg/mL a<u>t baseline</u>



## **SHINE Cognitive Endpoints:** *MMSE*

Preservation of MMSE in participants in below-median plasma p-tau217\* subgroup



#### mITT population (n=150)

Below median p-tau217 (n=69)





### **SHINE Functional Endpoints:** ADCS-ADL and -CGIC

Function and global impression preserved in below-median plasma p-tau217\* subgroup





ADCS-CGIC Below median plasma p-tau217 (n=69)





### Larger Biomarker Effect Observed in Below Median ptau217 Group vs mITT

Plasma bion	narker (ng/L)	mITT (N=150)	< median p-tau217 (N=69)	> median p-tau217 (N=69)	
	LS mean (SE)	-0.188 (0.2393)	-0.64 (0.326)	0.24 (0.322)	
Αβ42	95% CI	(-0.66, 0.29)	(-1.289, 0.018)	(-0.400, 0.885)	
	p-value	0.4343	0.0565	0.4539	
	LS mean (SE)	-8.272 (5.0557)	-12.27 (7.296)	-1.07 (6.793)	More robust effect
Αβ40	95% Cl	(-18.28, 1.74)	(-26.869, 2.334)	(-14.653, 12.523)	sizes observed
	p-value	0.1044	0.0980	0.8759	
	LS mean (SE)	-3.2 (10.53)	-28.35 (11.687)	23.07 (16.789)	
GFAP	95% CI	(-24, 18)	(-51.769, -4.933)	(-10.544, 56.679)	measured in below
	p-value	0.7593	0.0186	0.1748	median p-tau217
	LS mean (SE)	-1.44 (1.130)	-2.67 (1.503)	-0.22 (1.707)	subgroup, save for
NFL	95% CI	(-3.7, 0.8)	(-5.681, 0.339)	(-3.637, 3.200)	brain-derived tau
	p-value	0.2057	0.0809	0.8987	
	LS mean (SE)	-0.053 (0.0569)	-0.10 (0.079)	0.02 (0.089)	
p-Tau217	95% CI	(-0.17, 0.06)	(-0.264, 0.056)	(-0.154, 0.203)	
	p-value	0.3581	0.1976	0.7884	
Brain- derived tau	LS mean (SE)	-0.168 (0.0987)	-0.11 (0.089)	-0.19 (0.149)	Trand p. (0.10
	95% CI	(-0.36, 0.03)	(-0.293, 0.070)	(-0.491, 0.104)	Significant $p < 0.10$
	p-value	0.0907	0.2208	0.1987	



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### **Biomarker impact on neuroinflammation: Plasma GFAP**

Significant impact (p<0.05) observed in population with lower plasma p-tau217 at baseline





### **Biomarker Impact on Neurodegeneration: Plasma NfL**

Noticeable trend in dampening neurodegeneration





# Trend Towards Change in p-Tau217 May Suggest Slowing of AD Progression





## **Up to 123% Percent Slowing on Assessments**

Biomarker findings align with strong, consistent efficacy signals across measures





Pooled

### **SHINE Study: Summary and Conclusions**

Baseline plasma p-tau217 biomarker identified strong zervimesine-treatment responder group

- Zervimesine was found to be generally safe and well-tolerated
- Robust cognitive and functional impact observed in the pre-specified below-median plasma p-tau217 subgroup
- Prominent effects on recognized biomarkers of neuroinflammation (GFAP), neurodegeneration (NfL) and amyloid biology (Aβ42) consistent with favorable clinical outcomes
- This study enabled dose selection and a biomarker-defined responder population (low plasma pTau217) to be identified for Ph3

SHINE trial supports advancing zervimesine to Phase 3 in Alzheimer's disease population defined by plasma p-tau217





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