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BIOMARKER IDENTIFICATION IN RELAPSED/REFRACTORY NON-GERMINAL CENTER B-CELL-LIKE DIFFUSE LARGE B-CELL LYMPHOMA TREATED WITH ZANUBRUTINIB

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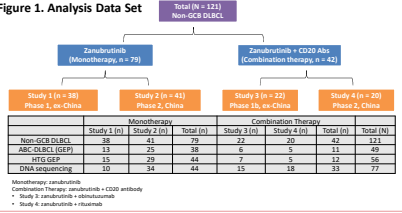


INTRODUCTION

- The non-germinal center B-cell-like (non-GCB) subtype of diffuse large B-cell lymphoma (DLBCL) is associated with poor clinical outcomes.¹
- Zanubrutinib, a highly selective covalent Bruton's tyrosine kinase (BTK) inhibitor, was specifically engineered to decrease toxicities and improve tumor tissue distribution.²
- Inhibitors of BTK have established therapeutic activity in mantle cell lymphoma, chronic lymphocytic leukemia, and Waldenström macroglobulinemia and have shown modest activity in DLBCL.³
- Biomarker identification has gradually become the focus of DLBCL research.
- Here we report zanubrutinib efficacy and biomarker identification in relapsed/refractory (R/R) non-GCB DLBCL from four clinical studies.

METHODS

- 121 R/R DLBCL patients from 4 zanubrutinib studies were included in analysis
- 2 Monotherapy (zanubrutinib alone) studies and 2 Combination Therapy (Zanubrutinib plus anti-CD20 antibody) studies were included
- Similar eligibility criteria and response assessment criteria across different studies
- Patients distribution and analysis data set shown in Figure 1
- Patient assessments
- Response: objective response rate (ORR) according to Lugano classification 2014
- GEP subtyping: performed by HTG EdgeSeq DLBCL Cell of Origin Assay using tumor FFPE samples collected before study drug treatment
- RNA-expression: gene expression readout by GEP subtyping assay were further analyzed by R package limma for correlation with response to zanubrutinib treatment
- DNA-sequencing: tumor FFPE samples were tested by next generation sequencing (NGS) and Chi-square test was used to evaluate the association between mutations and ORR



RESULTS

- The unadjusted ORR in non-GCB DLBCL was similar across the four studies with an average of 30%. Median PFS of the four zanubrutinib studies ranged from 2.8m to 4.9m, and median OS ranged from 8.4m to 11.8m. (Table 1)

Table 1. Efficacy of Zanubrutinib in Non-GCB DLBCL

	Monotherapy			Combination Therapy			Total (N=121)
	Study 1 (n=38)	Study 2 (n=41)	Total (n=79)	Study 3 (n=22)	Study 4 (n=20)	Total (n=42)	
ORR, n (%)	12 (31.6%)	12 (29.3%)	24 (30.4%)	5 (22.7%)	7 (35.0%)	12 (28.6%)	36 (29.8%)
95% CI *	(17.50, 41.86)	(16.13, 45.54)	(20.53, 41.75)	(7.82, 45.37)	(15.39, 59.22)	(15.72, 44.58)	(21.79, 38.74)
mPFS (Months) ^b	3.5	2.8	2.8	4.9	3.5	3.5	3.1
95% CI	(1.97, 5.02)	(2.56, 5.45)	(2.60, 4.88)	(1.71, 10.58)	(2.69, 5.55)	(2.69, 5.6)	(2.73, 4.93)
mOS (Months) ^b	11.8	8.4	8.8	11.8	10.3	11.6	9.9
95% CI	(5.09, 22.11)	(4.80, NE)	(5.52, 14.92)	(3.52, 16.53)	(5.39, NE)	(6.77, 15.21)	(6.80, 13.17)

Monotherapy: zanubrutinib
 Combination Therapy: zanubrutinib + CD20 antibody
 * Study 3: zanubrutinib + obinutuzumab
 * Study 4: zanubrutinib + rituximab
 ORR: Objective Response Rate
 mPFS: median Progression-Free Survival
 mOS: median Overall Survival
 * CI was calculated using the Clopper-Pearson method.
^b Estimated by Kaplan-Meier method with 95% CIs estimated using the method of Brookmeyer and Crowley.
 Data cut: Study 1 - September 9, 2019; Study 2 - August 31, 2019; Study 3 - August 31, 2019; Study 4 - May 11, 2019

- For 49 patients with GEP-confirmed activated B-cell (ABC) DLBCL classification, the ORR tended to be higher than non-GCB DLBCL although the number was small. The ORR was comparable for monotherapy (42%) and combination therapy (46%) for those with ABC-DLBCL (Table 2).

Table 2. Unadjusted ORR of Zanubrutinib in ABC-DLBCL

	Monotherapy			Combination Therapy			Total (N=49)
	Study 1 (n=13)	Study 2 (n=25)	Total (n=38)	Study 3 (n=6)	Study 4 (n=5)	Total (n=11)	
ORR, n (%)	7 (53.8%)	9 (36.0%)	16 (42.1%)	3 (50.0%)	2 (40.0%)	5 (45.5%)	21 (42.9%)
95% CI	(25.13, 80.78)	(17.97, 57.48)	(26.31, 59.91)	(11.81, 88.19)	(5.27, 85.44)	(16.75, 77.66)	(28.82, 57.79)

Monotherapy: zanubrutinib
 Combination Therapy: zanubrutinib + CD20 antibody
 * Study 3: zanubrutinib + obinutuzumab
 * Study 4: zanubrutinib + rituximab
 ORR: Objective Response Rate
 CI was calculated using the Clopper-Pearson method.
 Data cut: Study 1 - September 9, 2019; Study 2 - August 31, 2019; Study 3 - August 31, 2019; Study 4 - May 11, 2019

- For the 56 non-GCB patients with HTG gene expression profiles, PAX5 expression was higher in monotherapy responders than non-responders (Figure 2A), and PIM1, BCL2, and FOXP1 expression was higher in combination therapy responders than non-responders (Figure 2B).

Figure 2A. Genes Enrichment Analysis by Response to Zanubrutinib Monotherapy

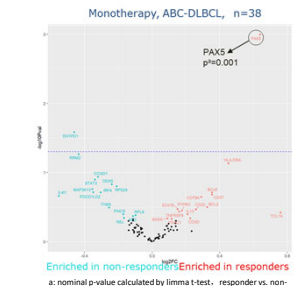
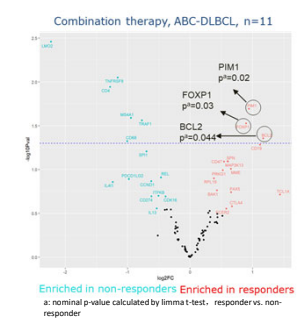
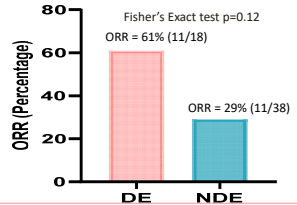


Figure 2B. Genes Enrichment Analysis by Response to Zanubrutinib Combination Therapy



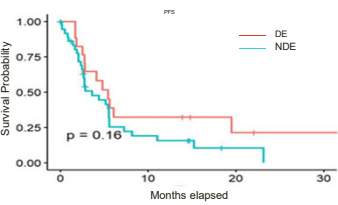
- Patients with MYC and BCL2 double-expressor DLBCL tended to have higher ORRs (11/18, 61% vs 11/38, 29%; P = 0.12) (Figure 3A) and longer progression-free survival (5.4 months vs 3.6 months; P = 0.16) (Figure 3B) and overall survival (10 months vs 7 months, P = 0.32) (Figure 3C).

Figure 3A. Correlation of BCL2/MYC Expression With Best of Response to Zanubrutinib



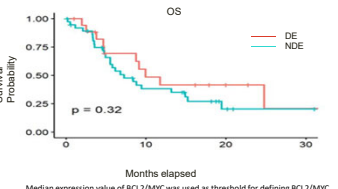
Median expression value of BCL2/MYC was used as threshold for defining BCL2/MYC high or low
 DE: BCL2 and MYC double high expression by GEP
 NDE: other than BCL2 and MYC double high expression by GEP

Figure 3B. Correlation of BCL2/MYC Expression With Progression-Free Survival



Median expression value of BCL2/MYC was used as threshold for defining BCL2/MYC high or low
 DE: BCL2 and MYC double high expression by GEP
 NDE: other than BCL2 and MYC double high expression by GEP

Figure 3C. Correlation of BCL2/MYC Expression With Overall Survival



Median expression value of BCL2/MYC was used as threshold for defining BCL2/MYC high or low
 DE: BCL2 and MYC double high expression by GEP
 NDE: other than BCL2 and MYC double high expression by GEP

- Patients with MYC and BCL2 double-expressor For the 77 patients with NGS panel data, non-GCB DLBCL with CD79B mutations (n = 25) showed significantly higher ORR than patients without CD79B mutations (n = 52) in the pooled analysis (60% vs 26.9%; P = 0.005). All three patients with NOTCH1 mutations responded to zanubrutinib monotherapy. The adjusted results showed a similar signal (Table 3).

Table 3. Correlation of CD79B/NOTCH1 Mutations With Response to Zanubrutinib

	Monotherapy (n=79)		Combination Therapy (n=42)		Total (n=121)
	n	%	n	%	
CD79B					
CD79B ^{mut}	9/27	33.3	6/8	75.0	15/29 51.7
CD79B ^{wt}	9/27	33.3	5/25	20.0	14/29 48.3
Difference in ORR (95% CI) *	18.6 (10.20, 46.87)		50.0 (15.75, 78.76)		33.1 (8.46, 53.57)
Fisher's ^b	0.0001		0.0001		0.0001
Adjusted P-value ^c	0.0001		0.0001		0.0001
NOTCH1					
NOTCH1 ^{mut}	3/3	100.0	2/5	40.0	5/8 62.5
NOTCH1 ^{wt}	15/41	36.6	9/28	32.1	24/69 34.8
Difference in ORR (95% CI) *	65.4 (4.48, 76.54)		7.9 (27.98, 49.62)		27.7 (4.78, 54.80)
Fisher's ^b	0.0001		0.7104		0.2206
Adjusted P-value ^c	0.0001		0.0001		0.0001

Monotherapy: zanubrutinib
 Combination Therapy: zanubrutinib + CD20 antibody (obinutuzumab or rituximab)
 * Difference 95% CI were based on Miettinen and Nurminen method.
^b Chi-square test is used to compute p-value.
^c Multivariate regression model is used to compute p-value with adjustment for baseline covariates and

CONCLUSIONS

- Zanubrutinib alone or in combination with an anti-CD20 antibody (obinutuzumab or rituximab) showed activity in the overall non-GCB DLBCL population.
- The retrospective biomarker analysis identified subsets of patients (such as PAX5 high or with CD79B mutations) with higher response rates to zanubrutinib treatment.

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