

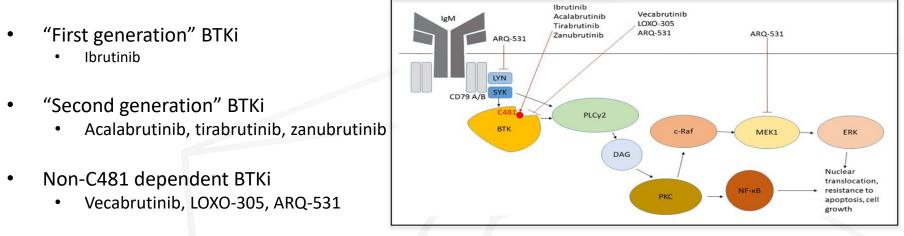
BTK Leu528Trp - a Potential Secondary Resistance Mechanism Specific for Patients with Chronic Lymphocytic Leukemia Treated with the Next Generation BTK Inhibitor Zanubrutinib

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\*contributed equally to the research

#### BTK inhibitors in CLL

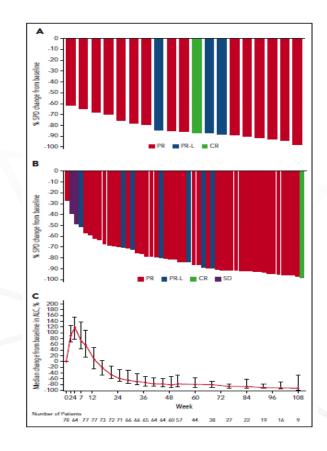
- Highly efficacious class of agent in frontline and relapsed/refractory CLL (Byrd *et al*, NEJM 2014; Shanafelt *et al*, NEJM 2019; Woyach *et al*, NEJM 2018)
- BTKi covalently bind to Cys481 residue in BTK resulting in blocking of enzymatic activity



(Bond & Woyach, 2019)

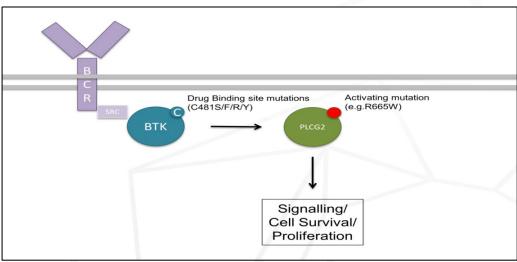
### Zanubrutinib in CLL

- Second generation BTK inhibitor
- Efficacious in treatment-naïve and relapsed/refractory CLL/SLL (Tam *et al*, Blood 2019)
- Multiple clinical trials in a variety of B-cell lymphoma subtypes ongoing
- Greater selectivity for BTK (over EGFR, ITK, and TEC) than ibrutinib (Tam et al, Blood 2019)



### Resistance to BTK inhibitors in CLL

- BTK inhibitor (ibrutinib) resistance mechanisms
  - (i) Drug-binding site Cys481 mutations (Cys481Ser, Cys481Phe/Arg/Tyr) (Woyach *et al*, NEJM 2014)
  - (ii) Downstream activating PLCG2 mutations (Liu et al, Blood 2015)





 To investigate possible resistance mechanisms to the second generation BTK inhibitor zanubrutinib (ZANU) in patients with CLL

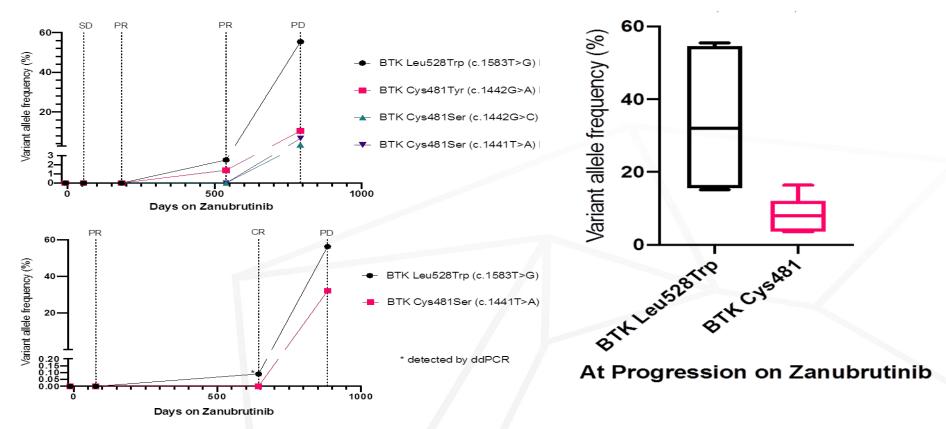
### Cohort

- 38 patients with relapsed/refractory CLL treated with ZANU on clinical trials (NCT02343120, NCT02569476, NCT03336333, NCT02795182) at three centres in Melbourne, Australia
- Four of 38 patients had CLL progression on ZANU (time to progression 5, 26, 29 and 48 months)
- Amplicon next generation sequencing (NGS)
  - Targeted amplicon sequencing (sensitivity approx 3-5% VAF)
    - ARAF, BCL2, BIRC3, BRAF, BTK (exon 11, 15, 16), CARD11, CD79B, CXCR4, DNMT3A, EZH2, FOXO1, FYN, ID3, IDH1, IDH2, JAK3, KRAS, MAP2K1, MYD88, NOTCH1, NRAS, PHF6, PLCG1, PLCG2(exon 16, 19-20, 24, 27-28), RHOA, RUNX1, SF3B1, STAT3, STAT5B, STAT6, TCF3, TP53, XPO1

#### BTK Leu528Trp detected in 4/4 patients with progressive CLL on zanubrutinib

Patient ID	Pre-ZANU	Post-ZANU	
CLLZ1	TP53 c.659A>G; p.(Tyr220Cys)	TP53 c.659A>G; p.(Tyr220Cys)	
		BTK c.1441T>A; p.Cys481Ser BTK c.1583T>G; p.Leu528Trp	
CLLZ2	BRAF c.1799T>A; p.(Val600Glu) NOTCH1 c.7541_7542del; p.(Pro2514Argfs*4)	BRAF c.1799T>A; p.(Val600Glu) NOTCH1 c.7541_7542del; p.(Pro2514Argfs*4)	
		BTK c.1441T>A; p.(Cys481Ser) BTK c.1442G>C; p.(Cys481Ser) BTK c.1583T>G; p.(Leu528Trp)	
CLLZ3	No mutations detected	TP53 c.1125_1140del; p.(Ser376Lysfs*41)	
		BTK c.1441T>A; p.(Cys481Ser)	
		BTK c.1442G>C; p.(Cys481Ser)	
		BTK c.1442G>A; p.(Cys481Tyr)	
		BTK c.1583T>G; p.(Leu528Trp)	
CLLZ4	BRAF c.1406G>C; p.(Gly469Ala)	BRAF c.1406G>C; p.(Gly469Ala)	
	XPO1 c.1711G>A; p.(Glu571Lys)	XPO1 c.1711G>A; p.(Glu571Lys)	
		BTK c.1442G>C; p.(Cys481Ser) BTK c.1583T>G; p.(Leu528Trp)	

### Leu528Trp is detectable in zanubrutinib treated patients before clinical CLL progression



Leu528Trp identified in 3 out of 34 patients on zanubrutinib in steady state

 ddPCR performed on 34 patients without disease progression but persistent measurable disease on zanubrutinib

• BTK Leu528Trp detected in 3 out of 34 patients (VAF <1%)

### BTK Leu528Trp and Cys481 mutations are present in different cells in zanubrutinib progressors



EXON 16

EXON 15

### BTK Leu528Trp and Cys481 mutations are present in different cells in zanubrutinib progressors



**EXON 16** 

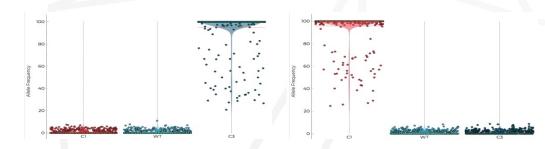
EXON 15

### BTK Leu528Trp and Cys481 mutations are present in different cells in zanubrutinib progressors



#### EXON 16





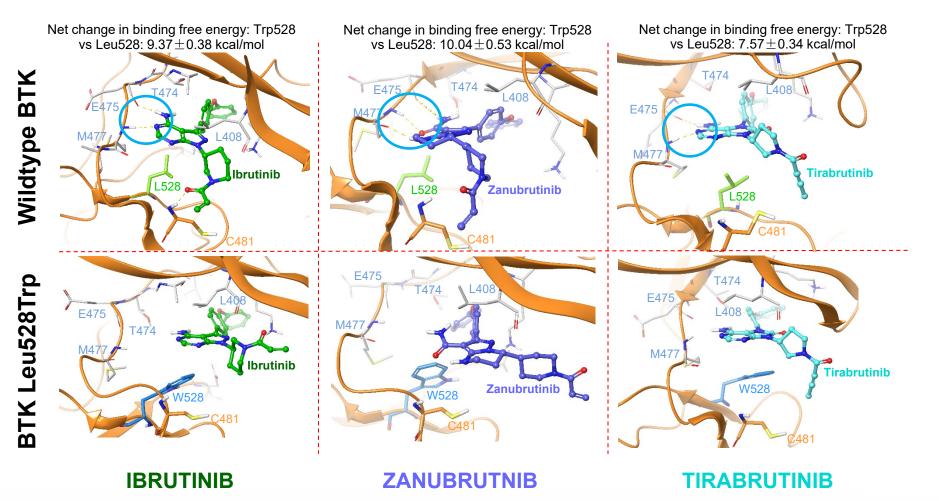
## BTK Leu528Trp is rarely observed in patients treated with ibrutinib

Author	BTK Leu528 codon assessed	Patients with progressive CLL	BTK Cys481 (and non-Leu528Trp)	BTK Leu528Trp
Woyach et al, NEJM 2014	Yes	6	5	0
Maddocks et al, JAMA 2015	Yes	19 (8 RT)	13	1
Sharma et al, Oncotarget 2016	Yes	1	1	0
Woyach et al, J Clin Oncol 2017	Yes	46	37	0
Gango et al, Int J Cancer 2019	Yes	20	8	0
Kanagal-Shamanna et al, Cancer 2019	Yes	29	19	0
	121	83 (68.5%)	1 (0.8%)	

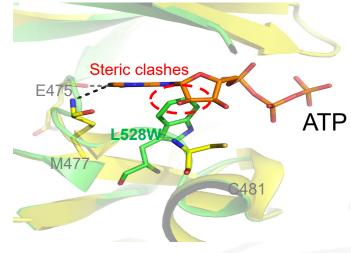
49 patients with progressive CLL on ibrutinib (European Research Initiative CLL [ERIC])

- Targeted next generation sequencing (Haloplex)
- 0/49 patients found to harbor the BTK Leu528Trp

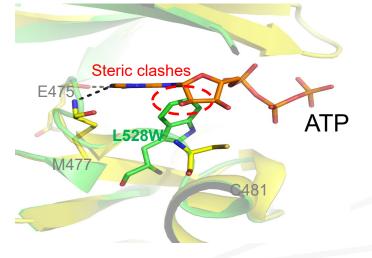
#### BTK Leu528Trp mutation disrupts the binding pose of ibrutinib, zanubrutinib and tirabrutinib

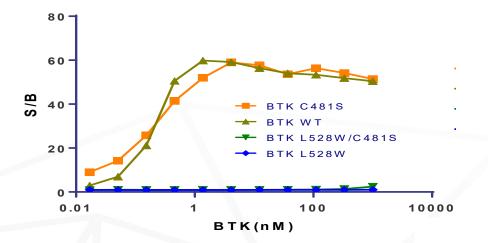


# BTK Leu528Trp leads to abrogated kinase function in biochemical and cellular models

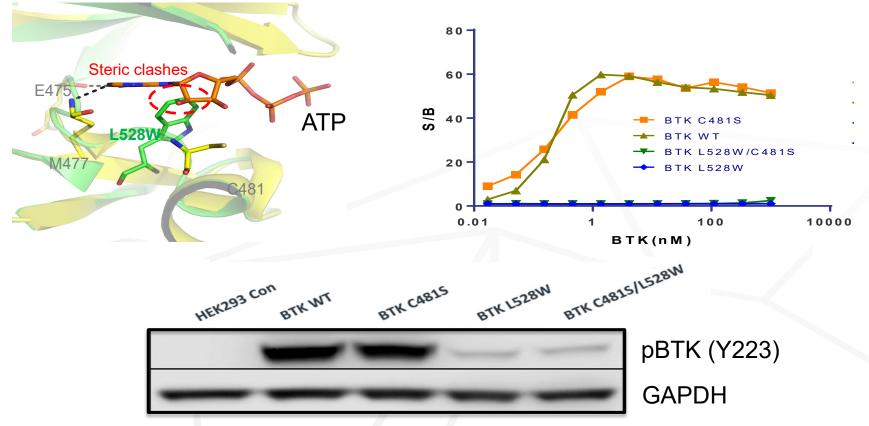


# BTK Leu528Trp leads to abrogated kinase function in biochemical and cellular models

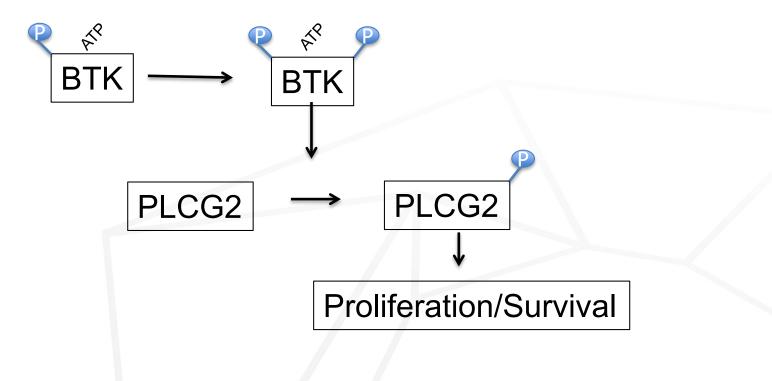




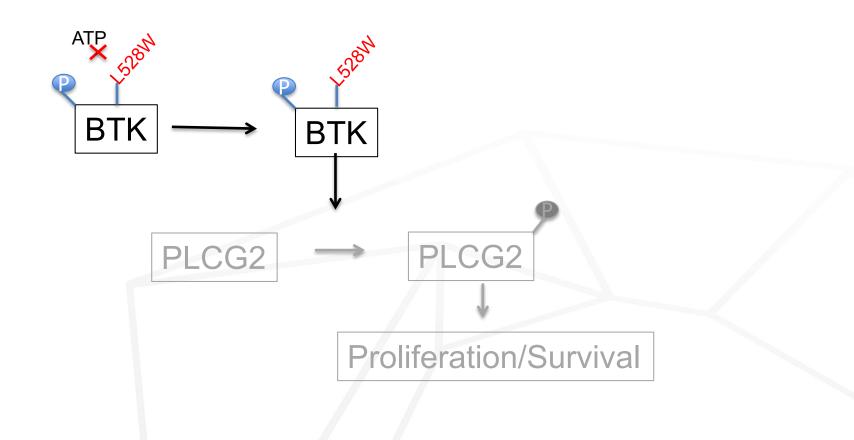
# BTK Leu528Trp leads to abrogated kinase function in biochemical and cellular models



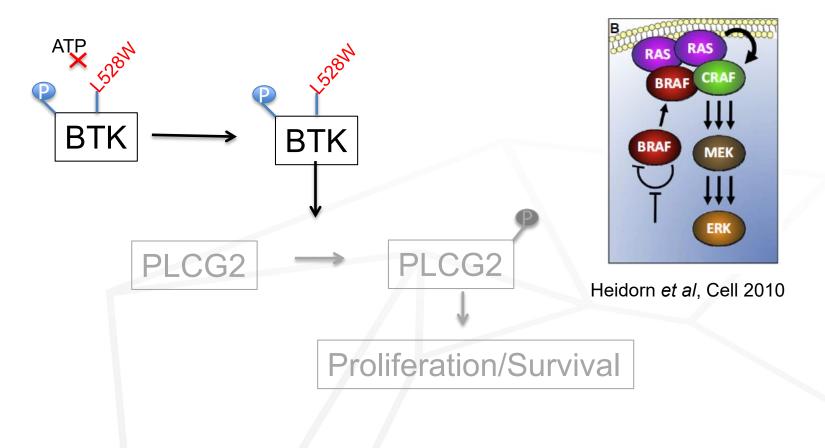
#### BTK Leu528Trp and downstream signalling

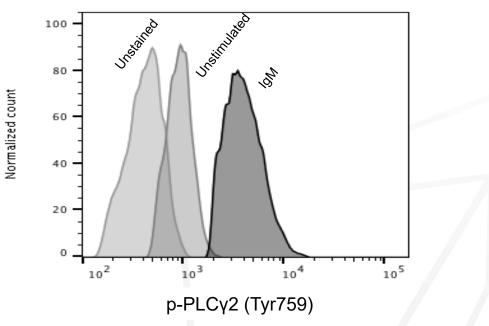


#### BTK Leu528Trp and downstream signalling

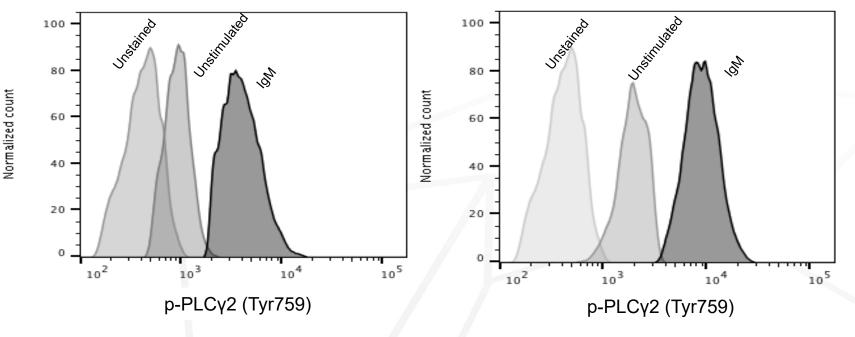


### BTK Leu528Trp and downstream signalling

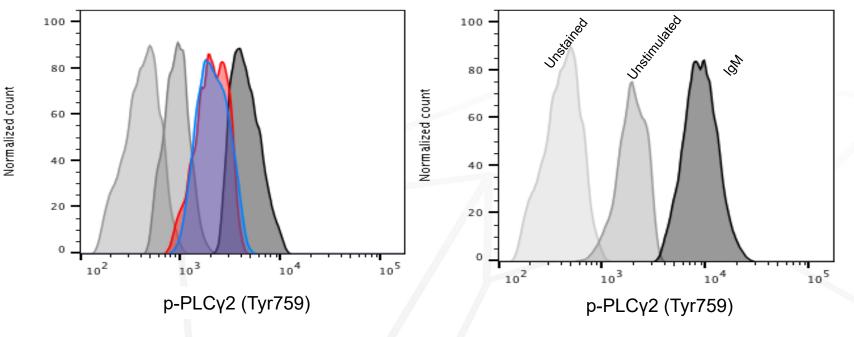




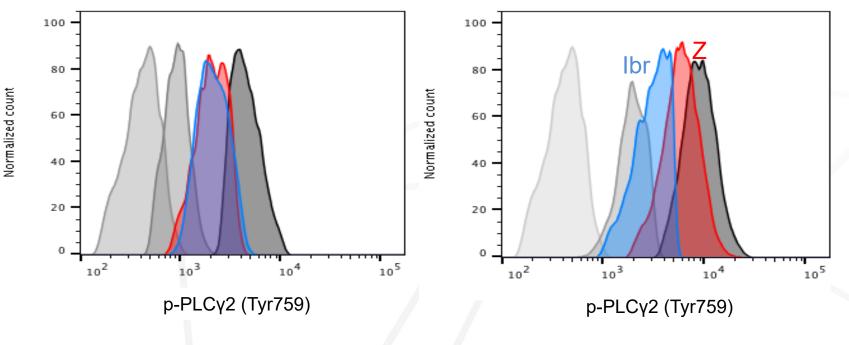












**BTK WILDTYPE** 

CD40L 100 100 CD40L CD40I + br IL21 80 80 Normalized count Normalized count 0 104 60 -104 60 Cell Trace Violet Cell Trace Violet 40 40 20 20 0 0 10<sup>3</sup> 105 10<sup>2</sup> 104 105 10<sup>2</sup> 103 10<sup>4</sup> p-PLCγ2 (Tyr759) p-PLCy2 (Tyr759)

BTK Leu528Trp

### Summary

- BTK Leu528Trp mutations are enriched in CLL progression on zanubrutinib compared to ibrutinib
- BTK Leu528Trp results in a marked impairment of binding of zanubrutinib (as well as ibrutinib and tirabrutinib) to BTK
- BTK Leu528Trp occurs with Cys481 mutations but is present in different CLL cells in the tumor compartment
- BTK Leu528Trp is associated with loss of native kinase function however downstream signalling pathways appear intact in patient CLL cells suggesting an alternative mechanism of PLCG2 phosphorylation



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