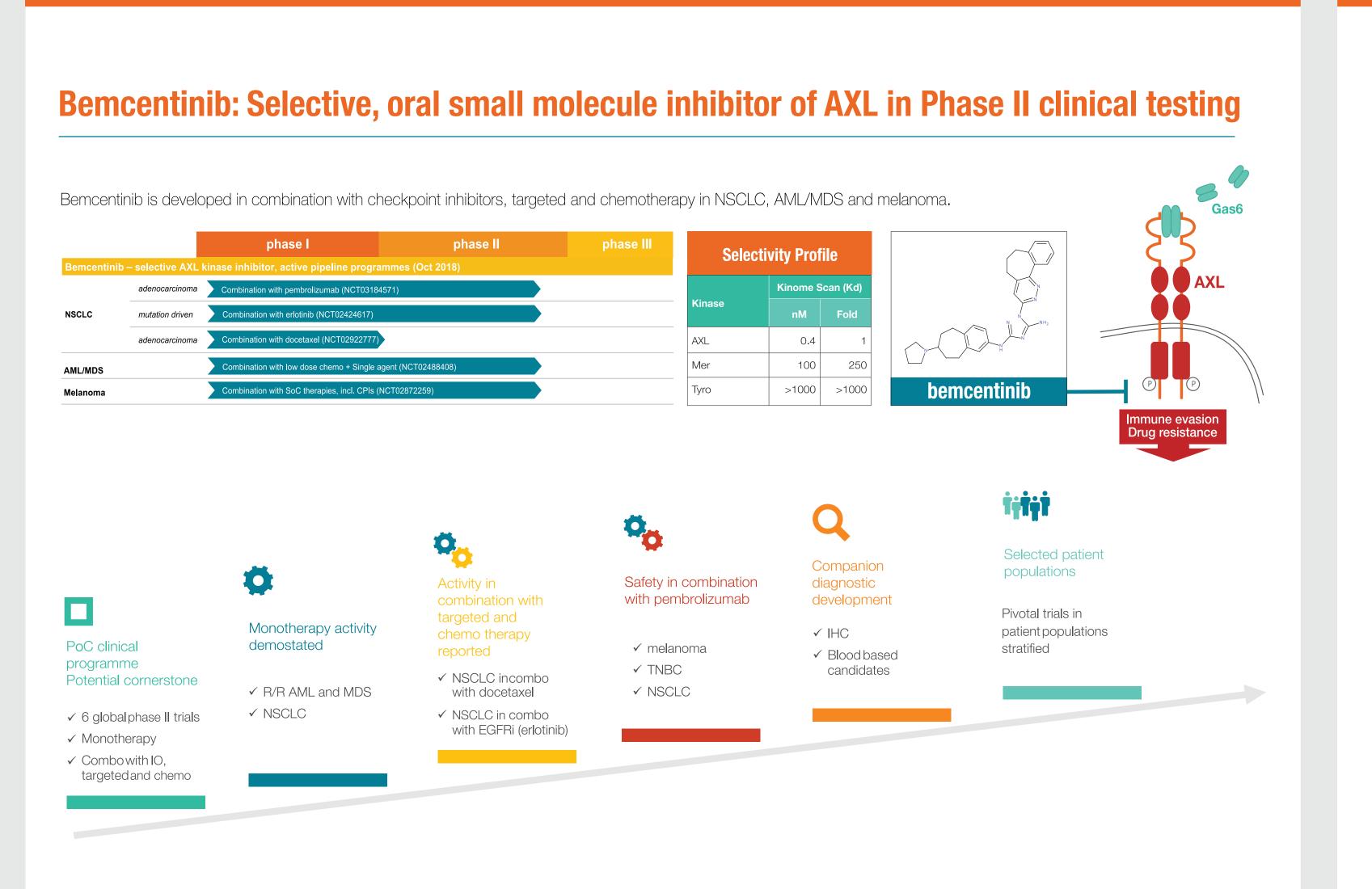
63PD: Predictive and Pharmacodynamic Biomarkers Associated with the Selective and Orally Bioavailable AXL Inhibitor Bemcentinib Across Multiple Clinical Trials



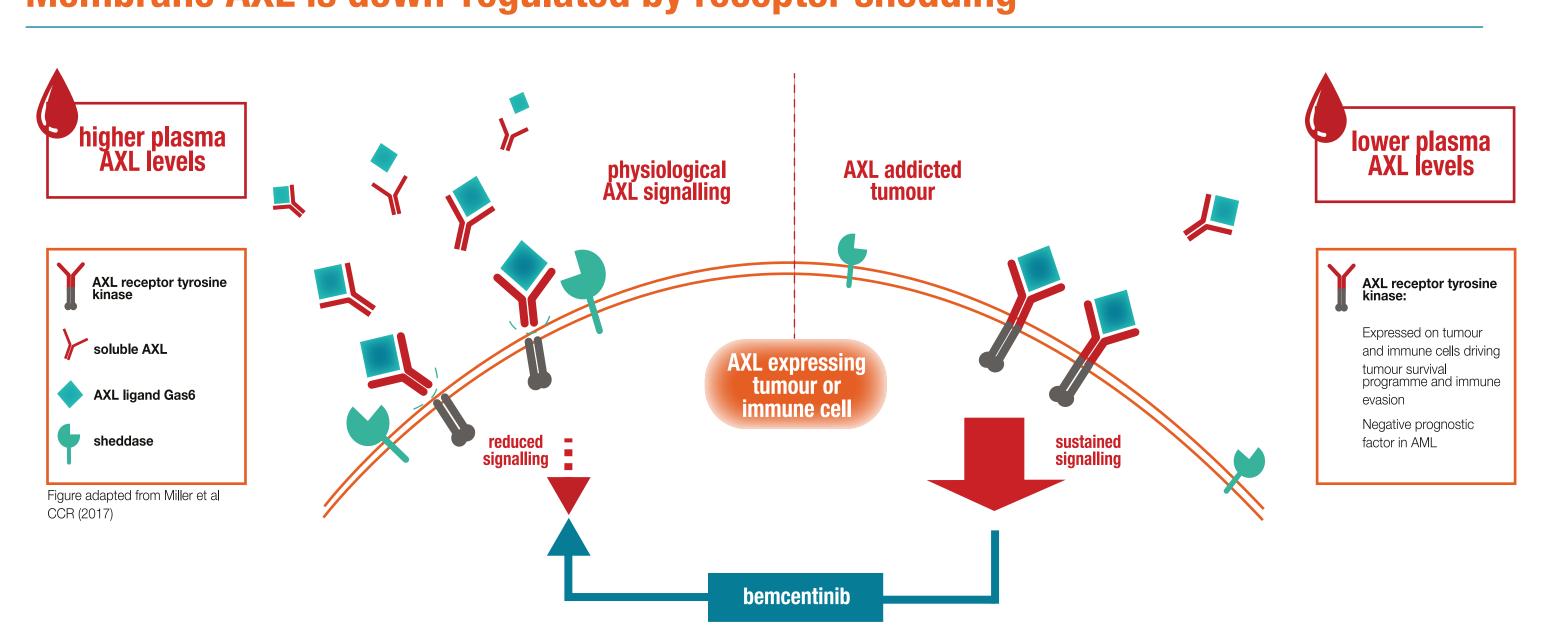
*BGBM014 is a transmembrane protein overexpressed in a wide variety of tumour types and induced by hypoxia.

Robert J Holt¹, David Micklem¹, Anthony Brown¹, Murray Yule¹, Oddbjørn Straume², James Lorens¹; (1) BerGenBio, (2) Department of Biomedicine, University of Bergen; Bergen, Norway

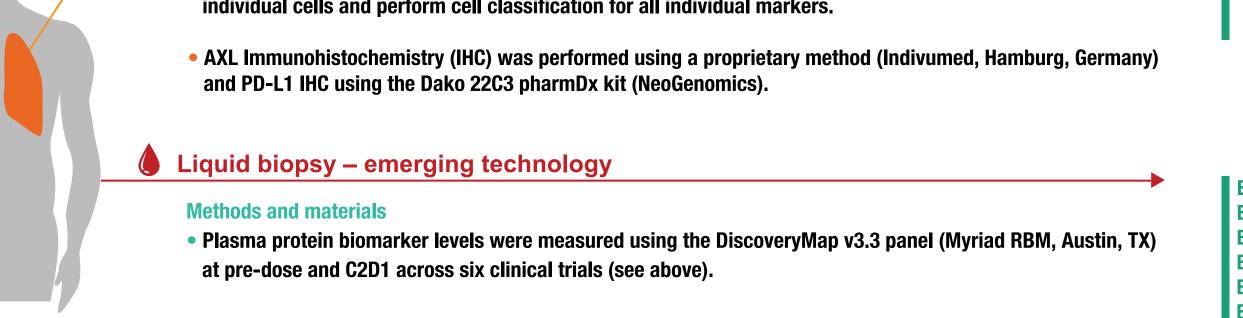
Bemcentinib & AXL Biology



Membrane AXL is down-regulated by receptor shedding



Tumour tissue biopsy - detect membrane AXL and other biomarker candidates A panel of biomarkers including AXL, CD68, and tumour segmentation marker PanCK was analysed using multiplexing methodology (MultiOmyx™, NeoGenomic Fort Myers, FL) coupled with deep learning to identify individual cells and perform cell classification for all individual markers AXL Immunohistochemistry (IHC) was performed using a proprietary method (Indivumed, Hamburg, Germany) and PD-L1 IHC using the Dako 22C3 pharmDx kit (NeoGenomics)



ormalised Myriad datasets: including (i) pre-dose (C1D1) Responders vs Non-Responders and (ii) Paired

of the lung (BGBC008). Data presented at WCL0

2018 (Lorens *et al*)

Materials & Methods

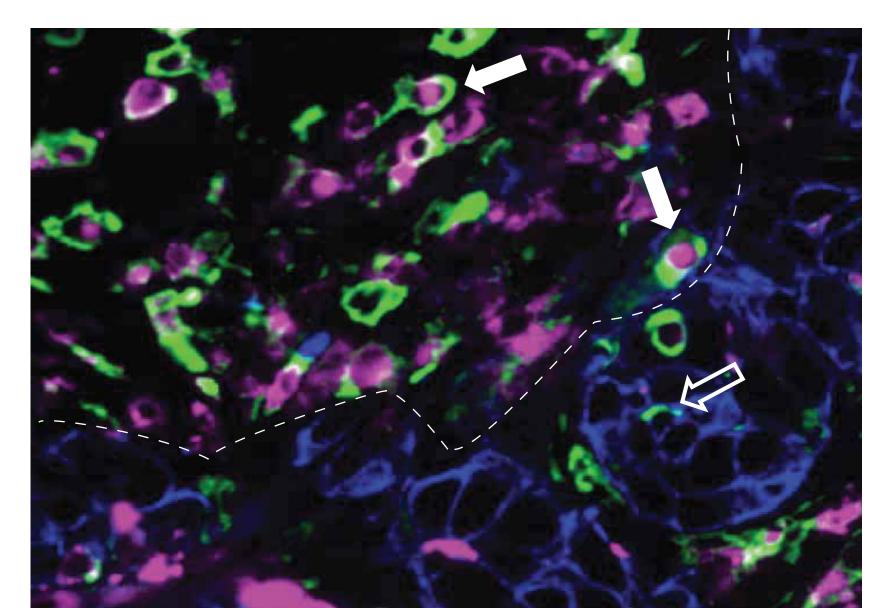
Timepoint comparison, C1D1 vs C2D1 for all patients.

Jonas Lies vei 91 1 Robert Robinson Ave Oxford, UK

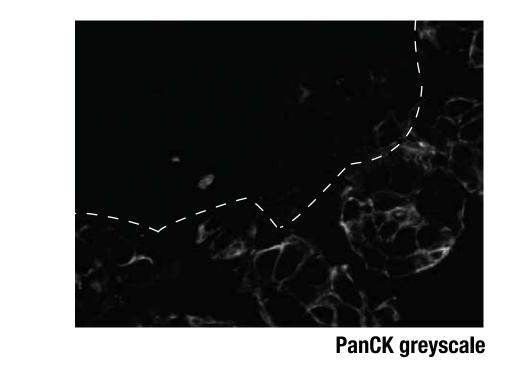
www.bergenbio.com @BGenBid post@bergenbio.com +47 559 61 159

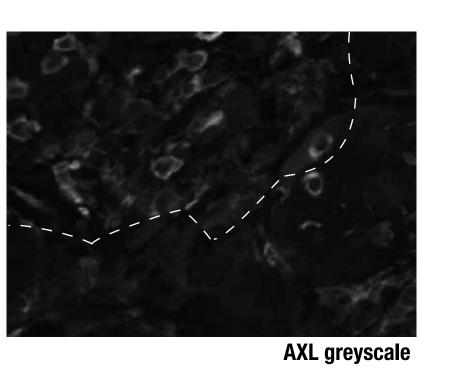
Predictive biomarker candidates to bemcentinib monotherapy and combination therapy with immuno- & targeted therapy have been identified in tissue and plasma

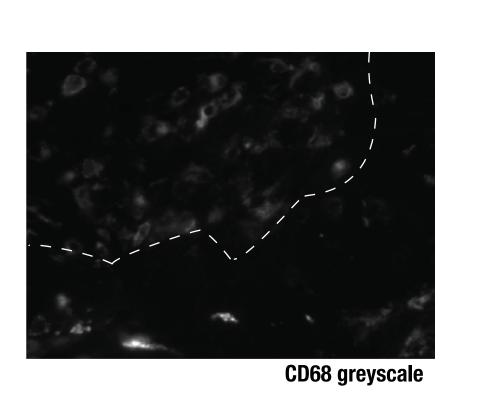
AXL is present in both tumour and tumour infiltrating immune cells



observed. Additionally a mainly weak cytoplasmic staining of stromal cells was seen (arrowheads).

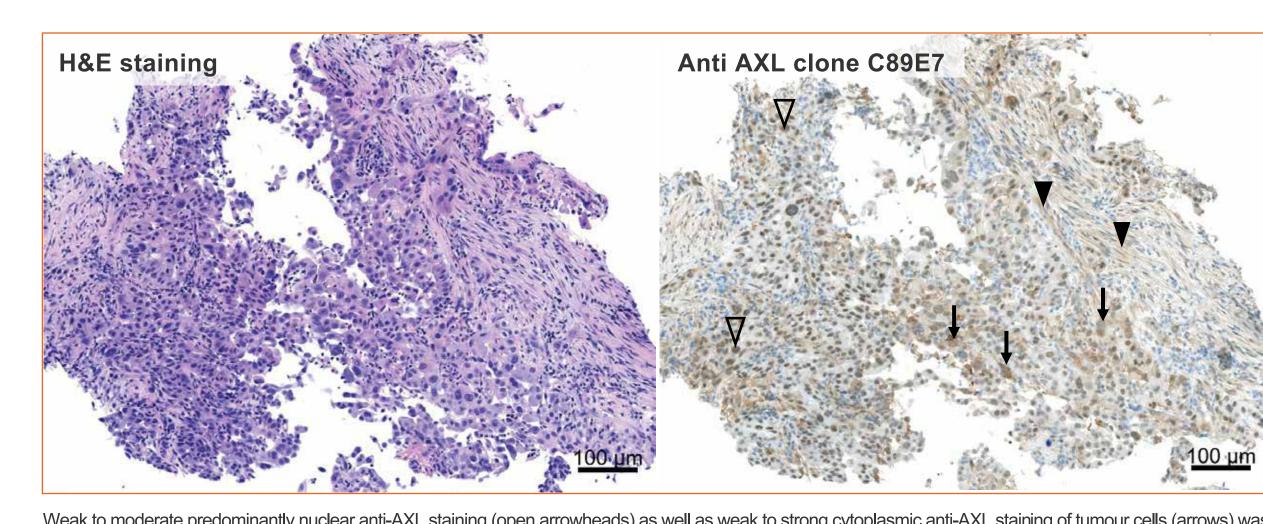


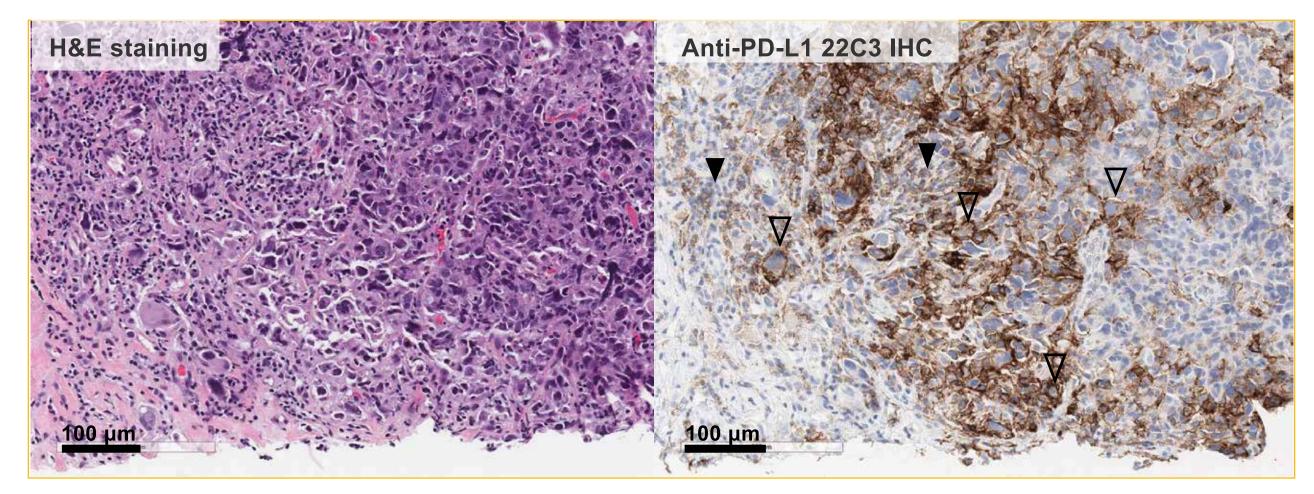




- A significant proportion of AXL expressing cells are macrophages and NK cells
- Investigation of correlation of AXL on immune cells with response is ongoing

AXL IHC identifies subset of NSCLC patients with increased response to bemcentinib + pembrolizumab (incl. PD-L1 negative patients)¹

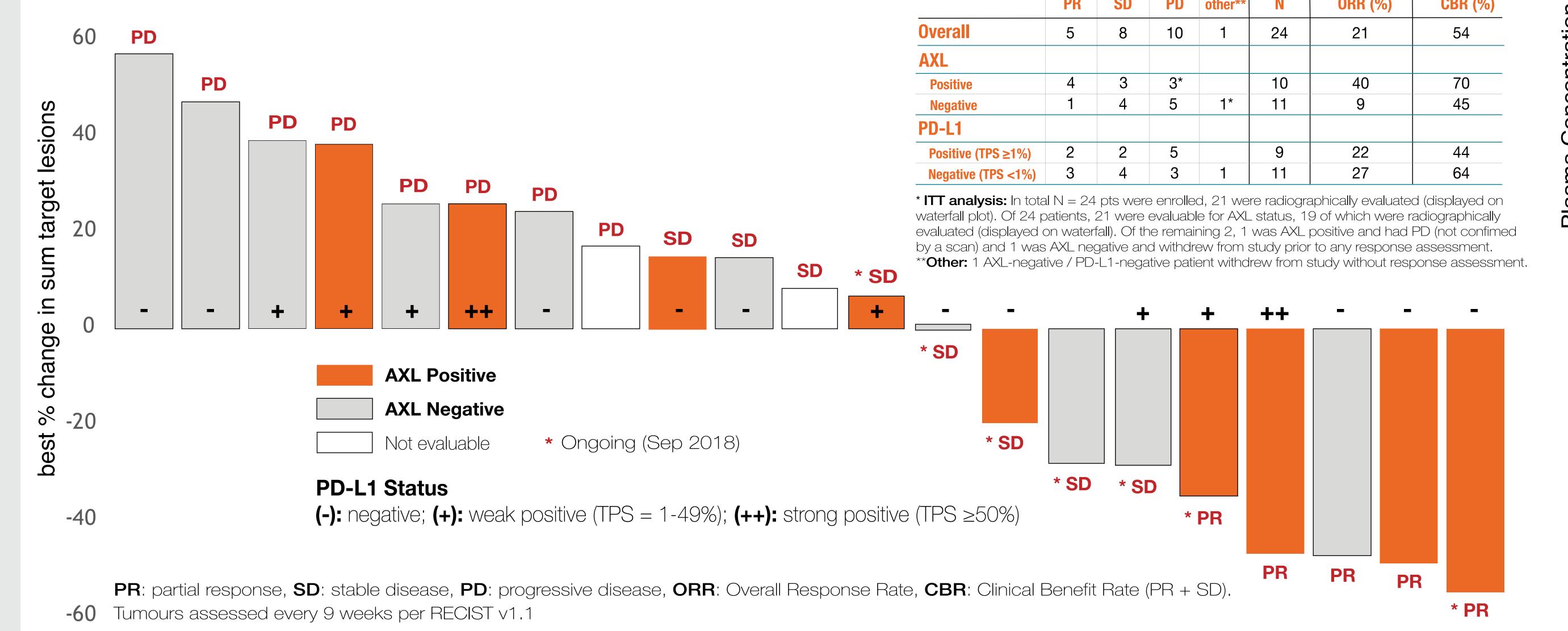




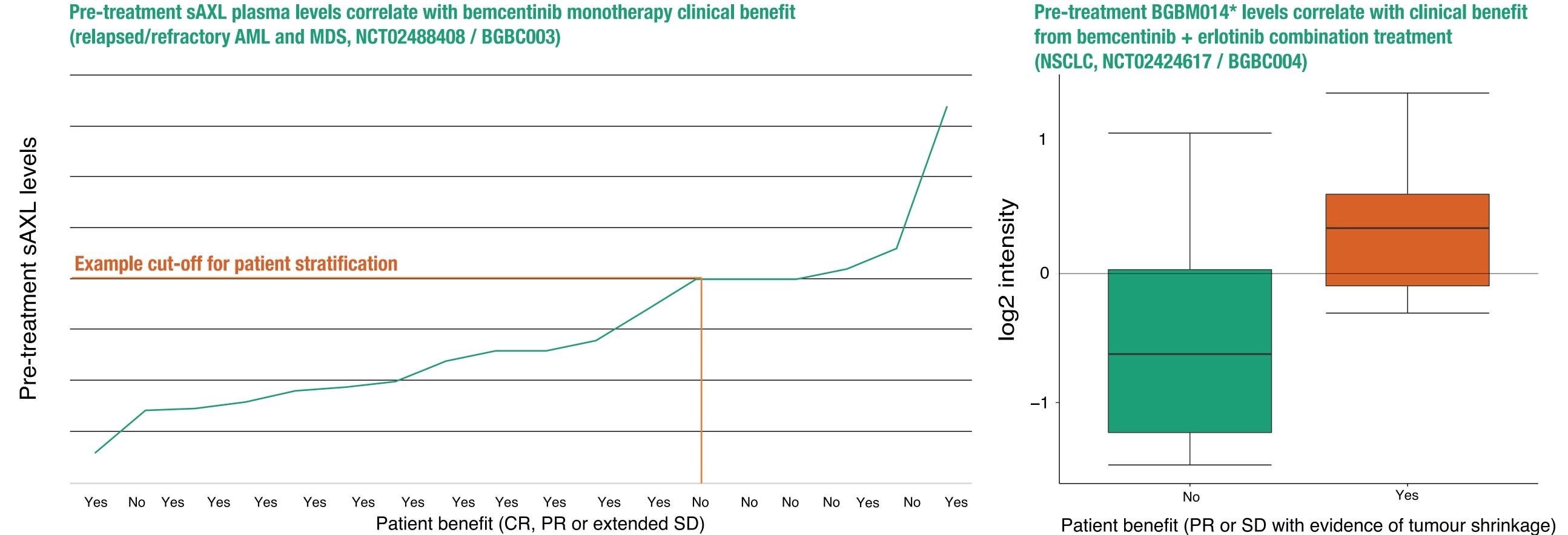
A predominantly moderate to strong membranous anti-PD-L1 22C3 IHC staining of tumour cells was observed (open arrowheads) as well as a weak to moderate membranous/cytoplasmic anti-PD-L1 22C3 IHC staining of immune cells (filled arrowheads).

Phase II trial exploring bemcentinib in combination with pembrolizumab in advanced NSCLC (NCT03184571): approximately half of patients found to be positive for tumour AXL staining, 7 of 10 AXL positive patients show clinical benefit incluing 4 PRs (ORR = 40%)¹

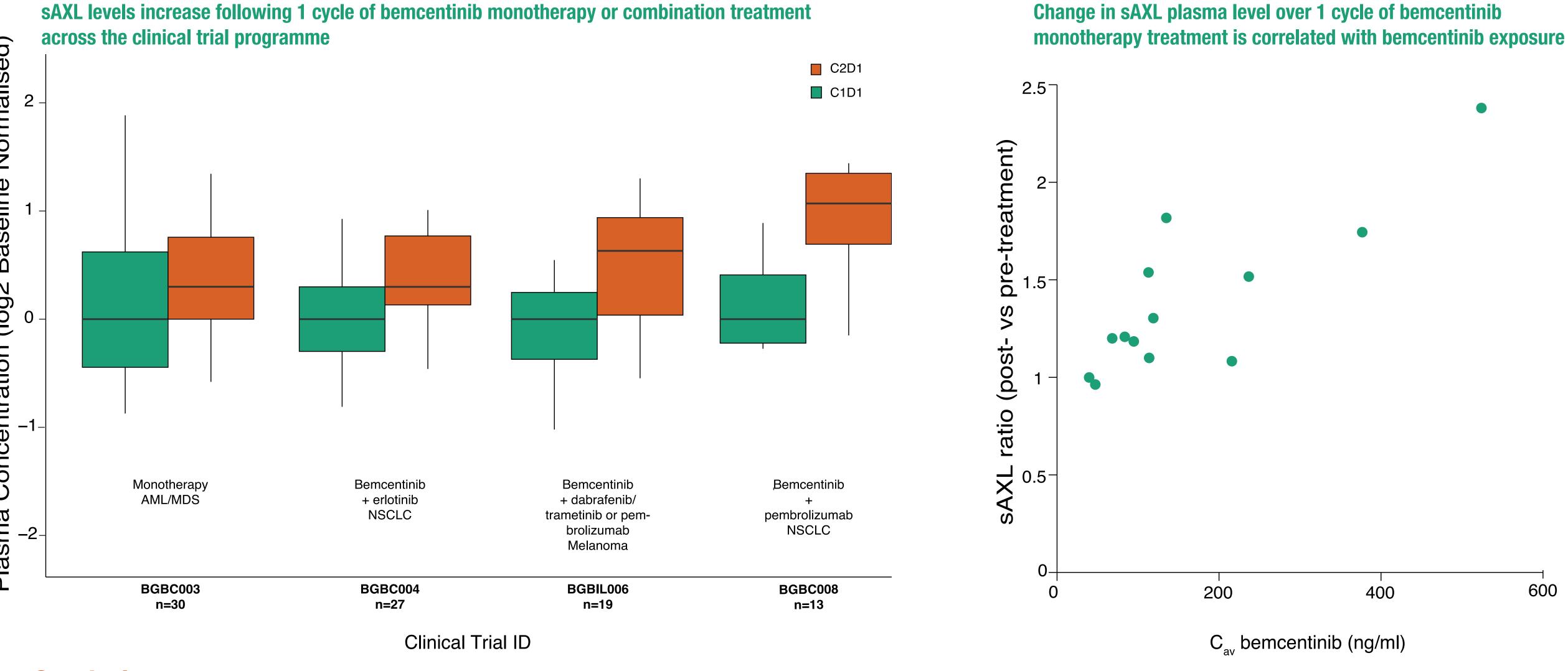
Waterfall plot: Best % change sum target lesions n = 21 pts with radiographic evaluation following treatment Best overall response: Intention-to-treat (ITT) & biomarker status analysis*



Pre-treatment levels of soluble AXL (sAXL) and other soluble biomarkers are predictive of patient benefit in AML/MDS and NSCLC



Pharmacodynamics: Soluble AXL (sAXL) levels in plasma samples increase following monotherapy and combination treatment



Conclusions

- Predictive biomarker candidates have been identified across multiple phase II monotherapy and combination trials:
- 1. Plasma sAXL is predictive of patient benefit from bemcentinib monotherapy in AML/MDS
- 2. Tissue AXL IHC identifies a subgroup of NSCLC patients with increased response to bemcentinib + pembrolizumab combination treatment
- 3. Plasma BGBM014 is predictive of NSCLC patient benefit from bemcentinib combination treatment with erlotinib
- Tissue AXL is found on tumour cells and tumour infiltrating immune cells (macrophages and NK cells)
- Plasma sAXL is also a pharmacodynamic marker across trials indicating bemcentinib on-target activity:
- 1. Plasma sAXL levels correlate with bemcentinib exposure
- 2. Plasma sAXL increases following bemcentinib treatment in all indications following monotherapy as well as combination treatment