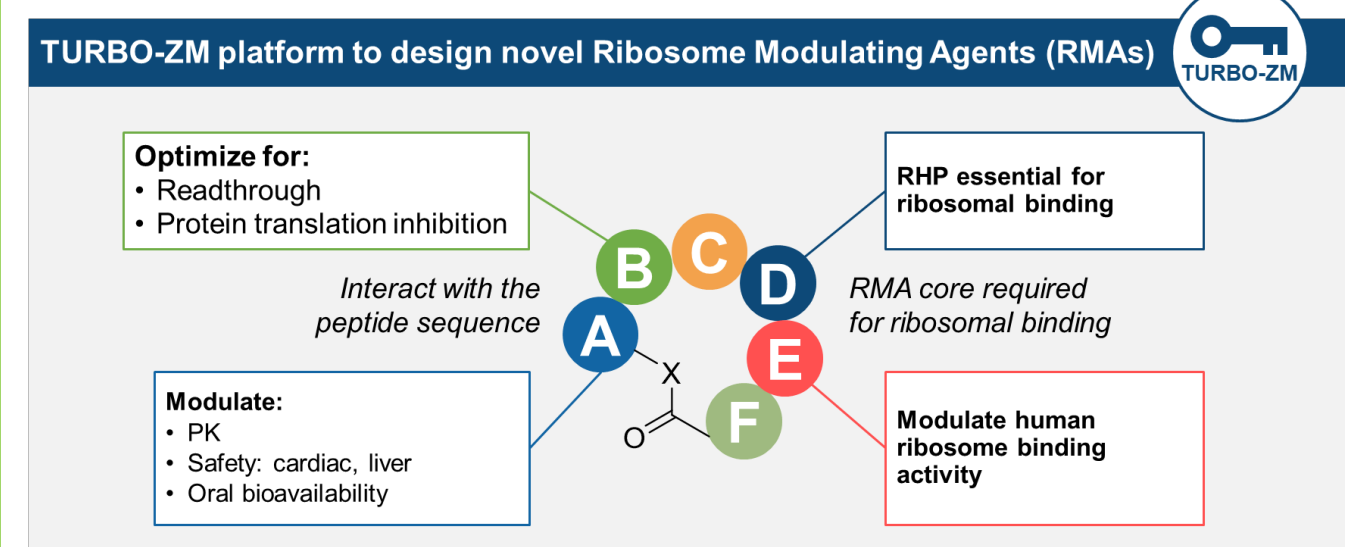


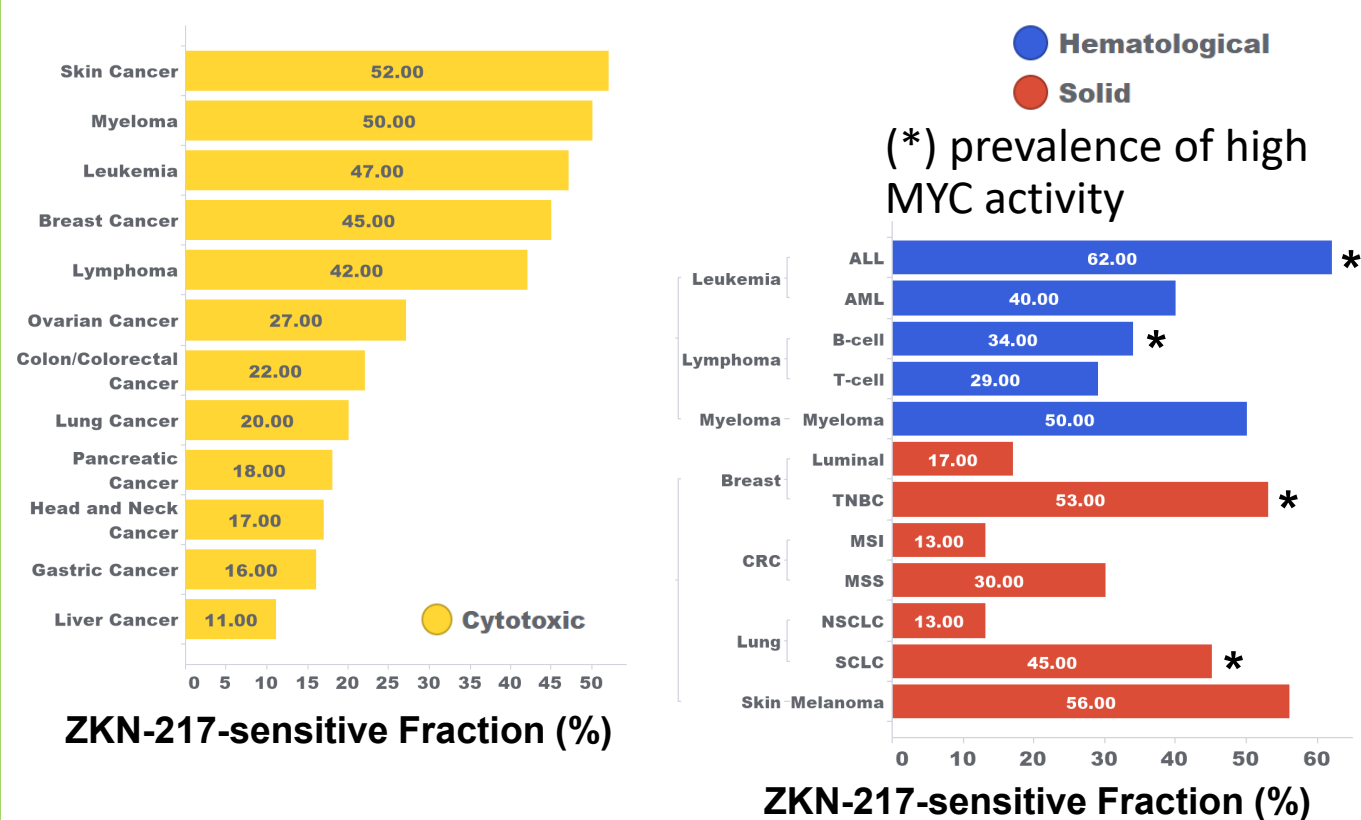
Introduction

Small cell lung cancer (SCLC) has a very poor prognosis and limited treatment options. Genomic studies in SCLC show that MYC isoforms are key oncogenic drivers in most of SCLC tumors. As a result, these tumors tend to have high proliferation and protein translation rates. We synthesized ribosome modulating agents (RMAs), a novel class of macrolides, capable of selectively inhibiting translation of a subset of molecules enriched for ribogenesis and translation-related proteins in sensitive SCLC cell lines. ZKN-217 selectively inhibits growth and induces apoptosis in a subset of SCLC cell lines characterized by high MYC activity and high ribogenesis rates. Furthermore, ZKN-217 is synergistic with current SCLC standard of care DNA intercalating drugs. The dependency of MYC-driven cancers on abnormally high protein translation rates can therefore be exploited by our RMAs presenting a novel therapeutic opportunity for this recalcitrant disease with high unmet needs.

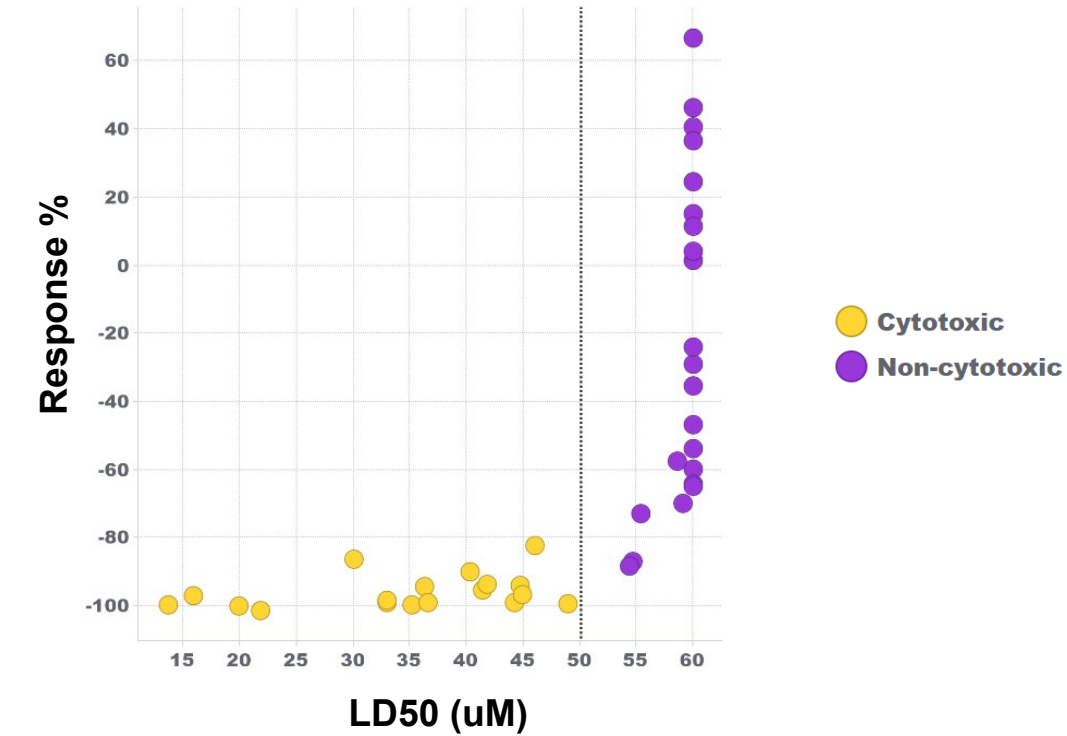
Novel synthetic macrolide-based allosteric modulators of the human ribosome



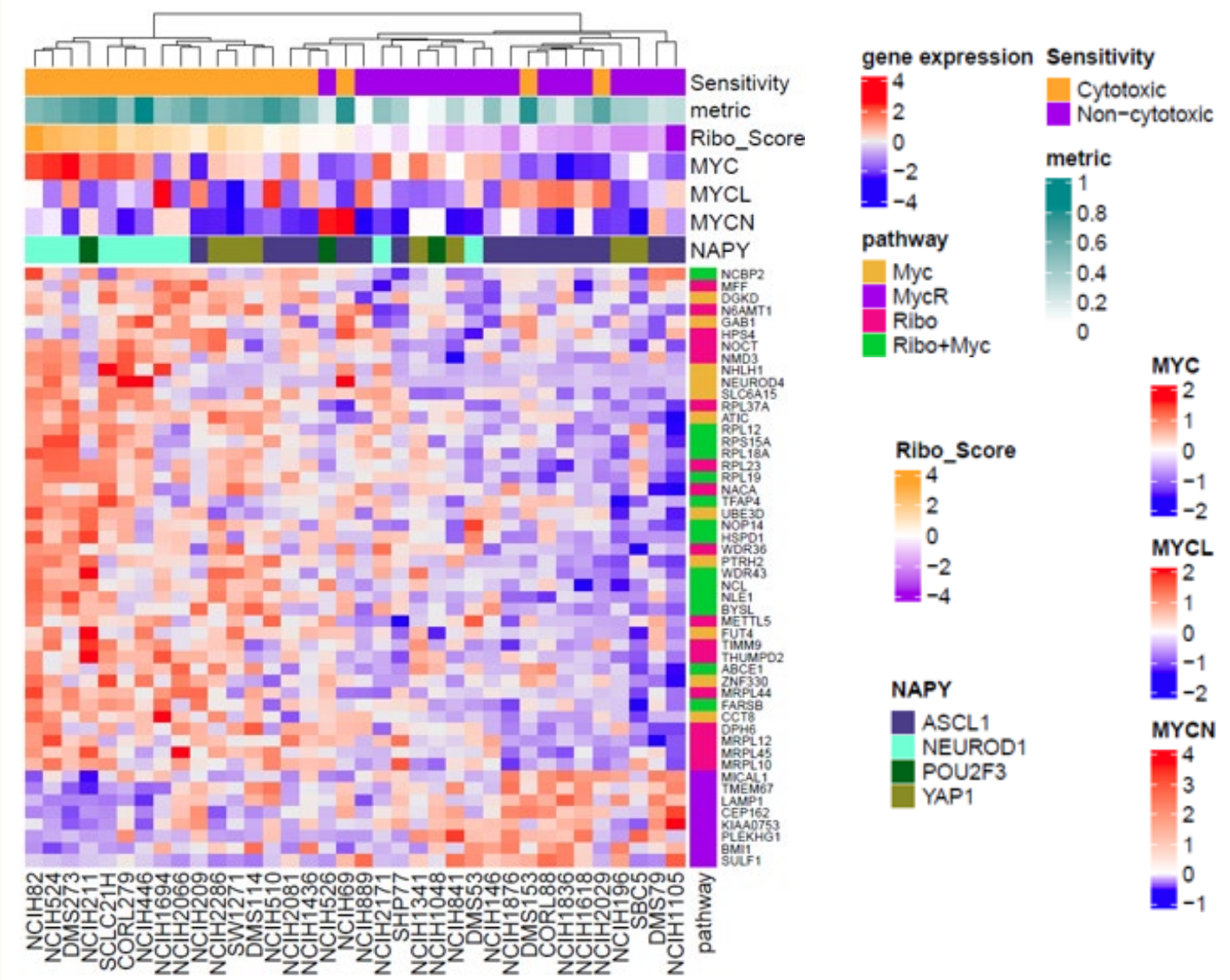
Several clinical subtypes in solid and hematological cancers are highly sensitive to ZKN-217



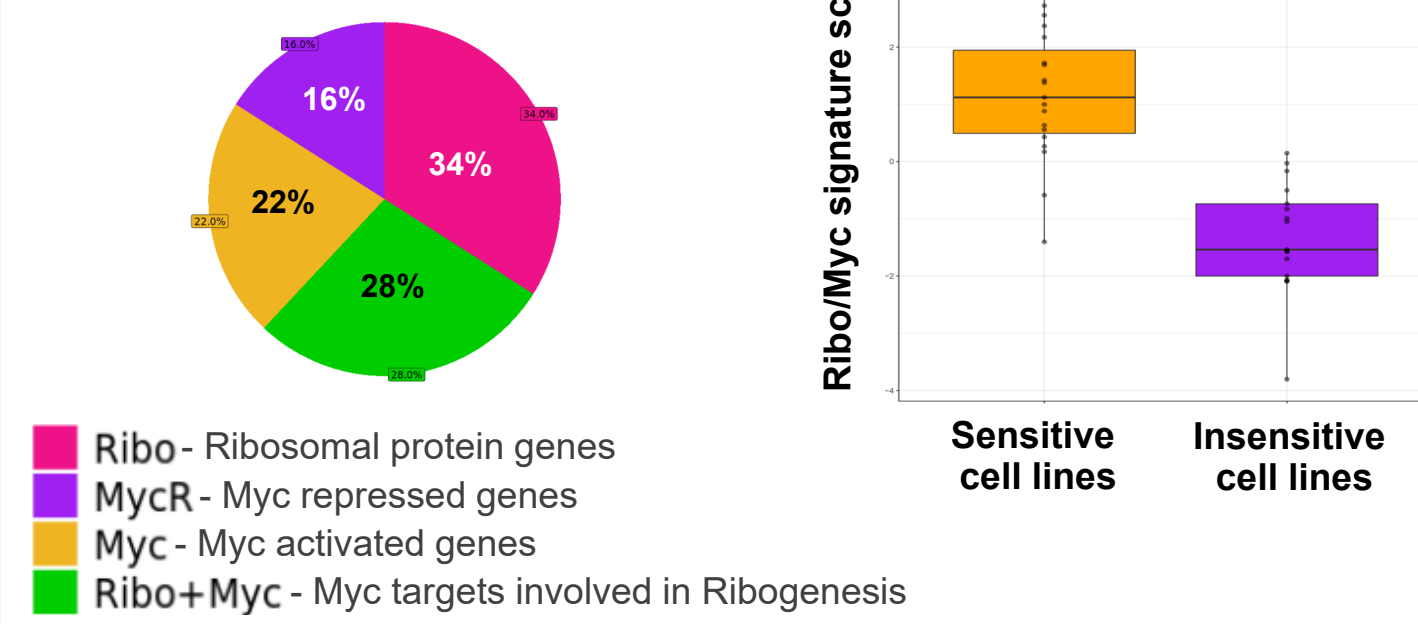
45% of SCLC cell lines are highly sensitive to ZKN-217



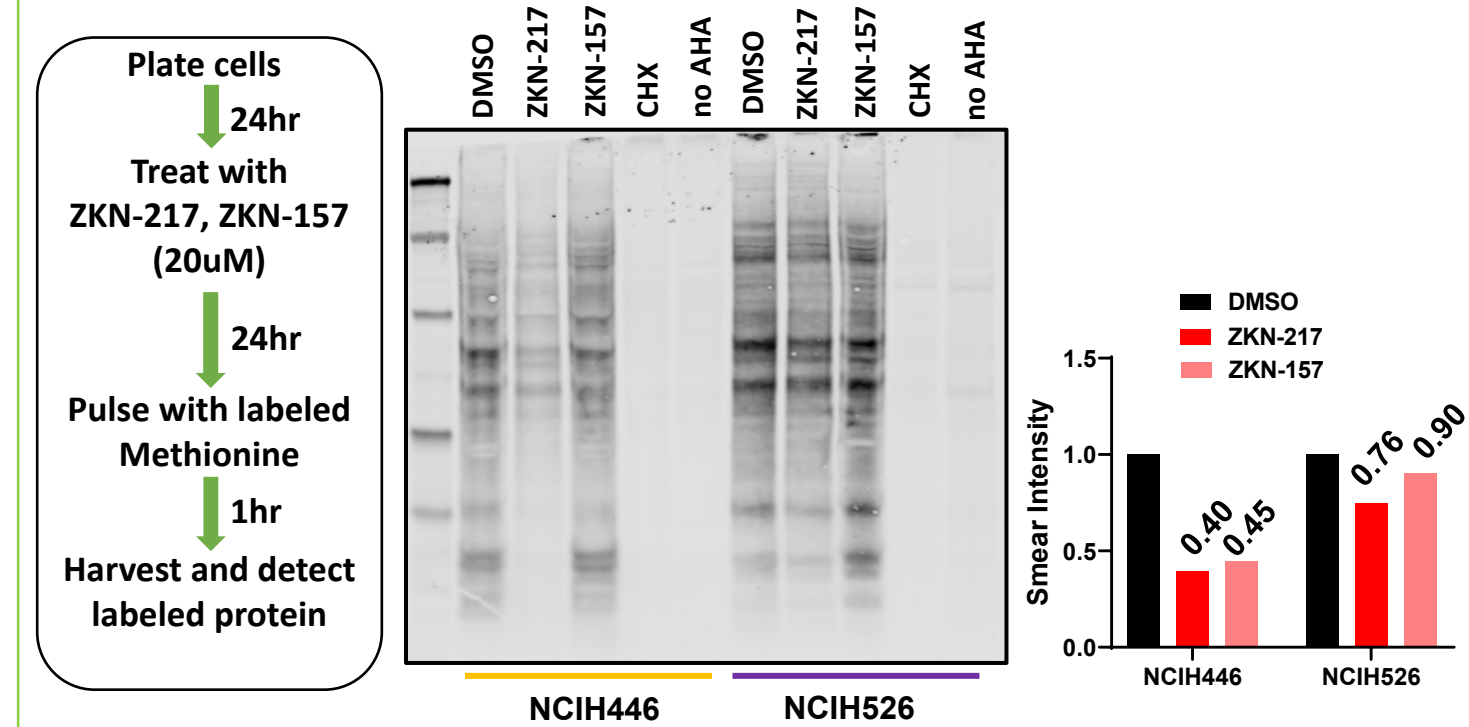
Sensitivity to ZKN-217 is associated with high expression of Ribogenesis, protein translation, and MYC pathway genes



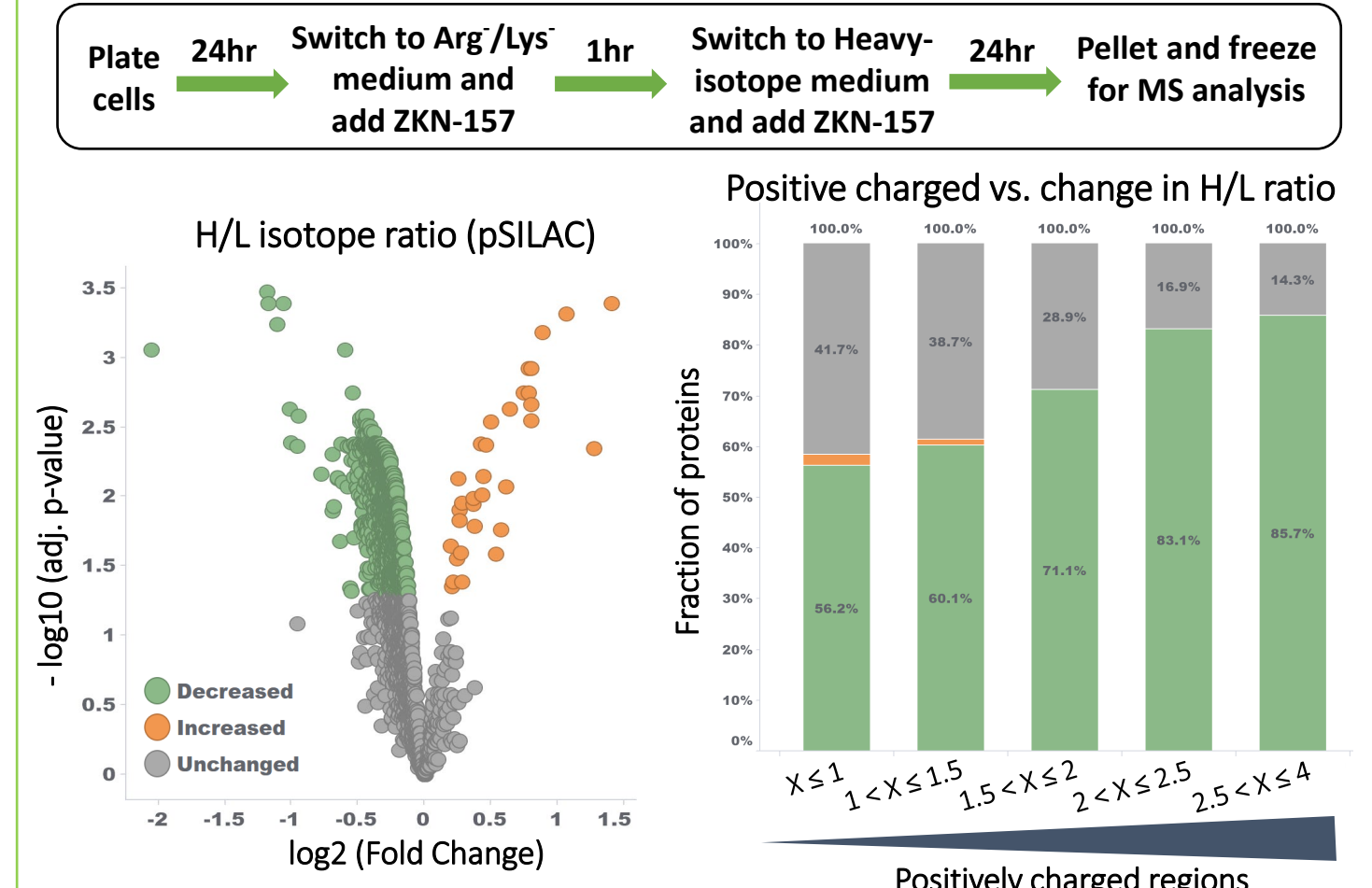
Top 50 Leading edge genes Classification



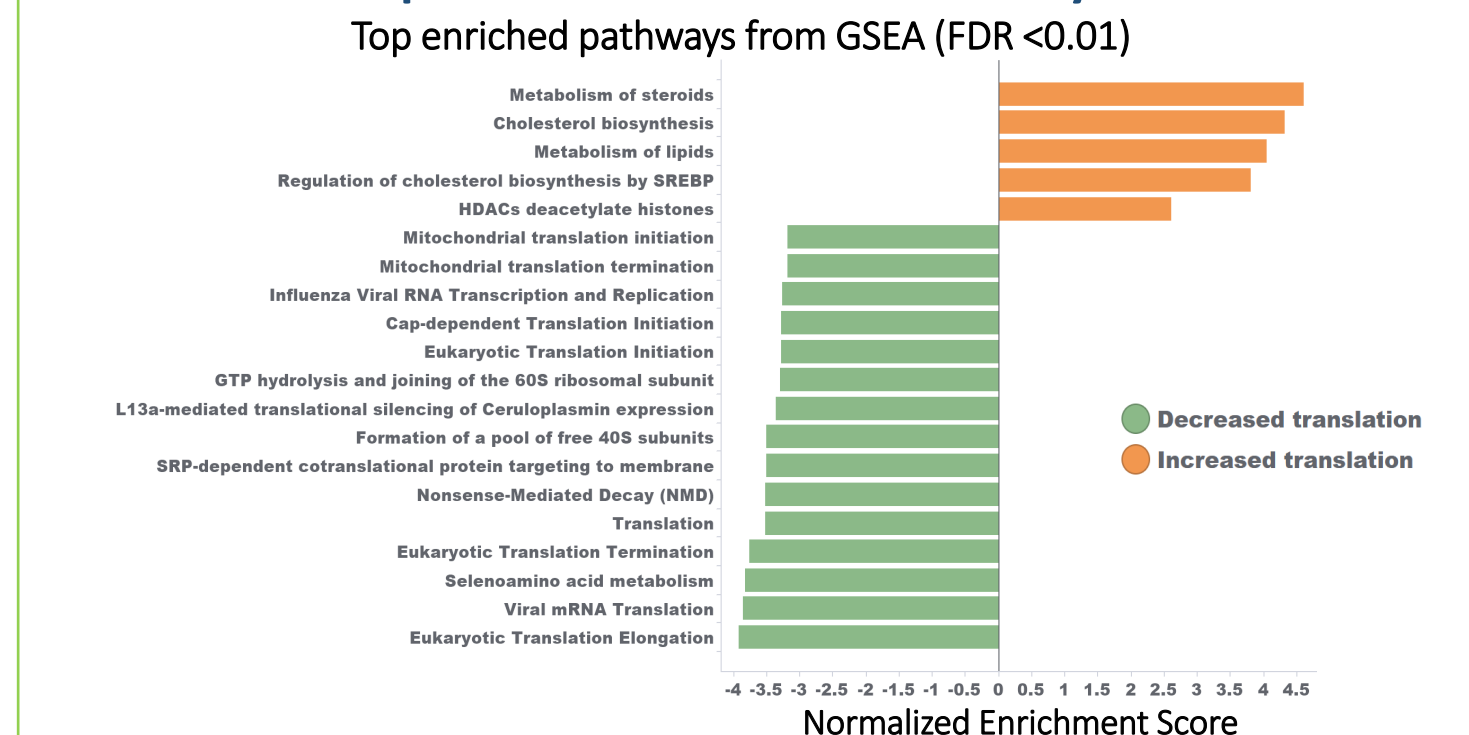
Anti-proliferative effect of RMAs is driven by selective inhibition of new protein synthesis



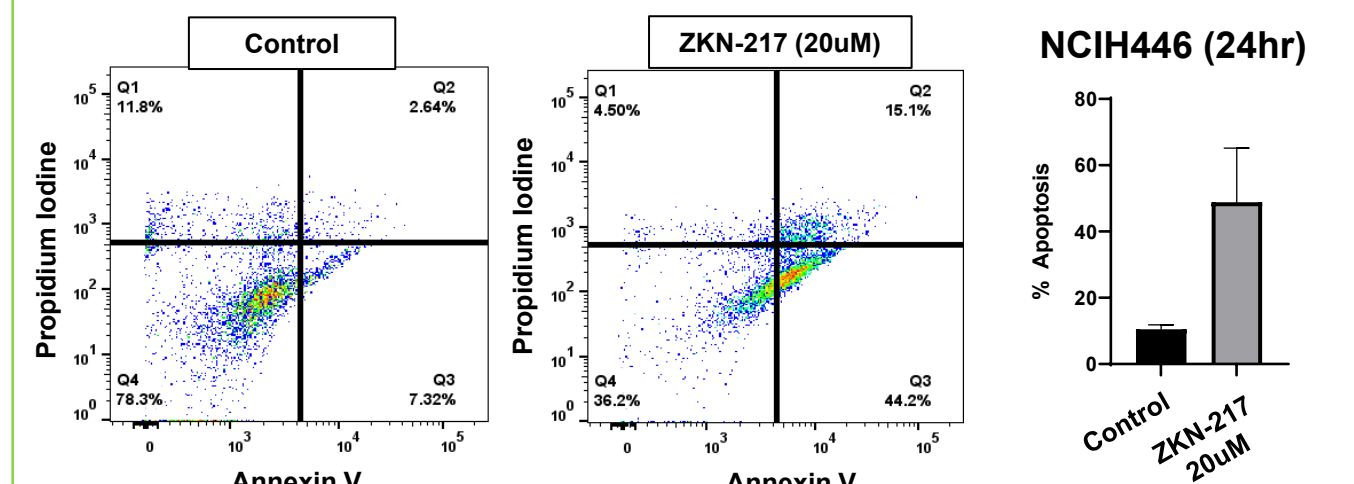
Proteins with higher positively charged regions are more sensitive to translation inhibition by RMAs



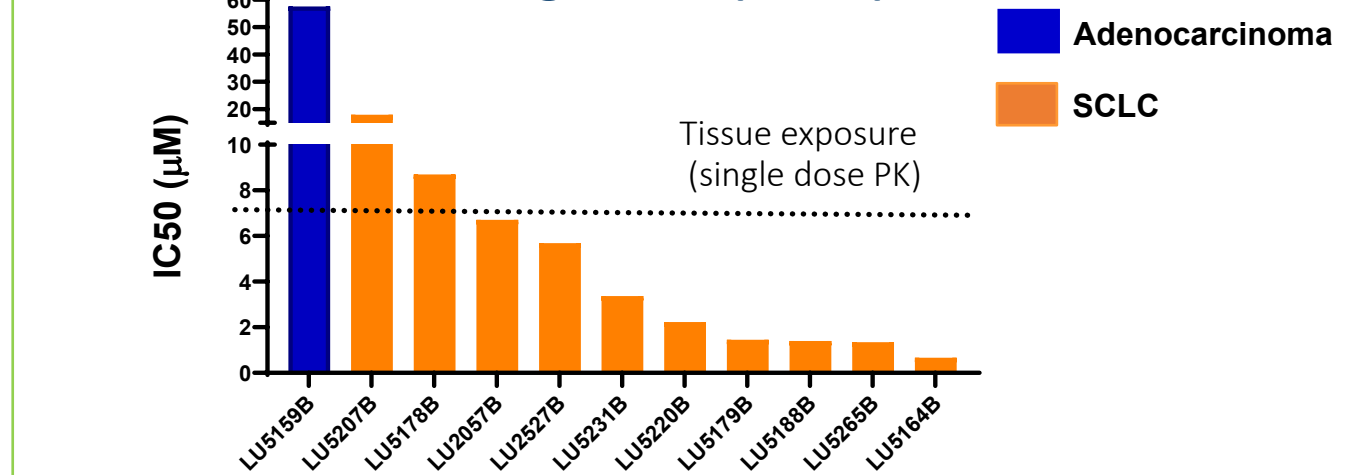
Decreased proteins show large impact on Ribogenesis and protein translation machinery



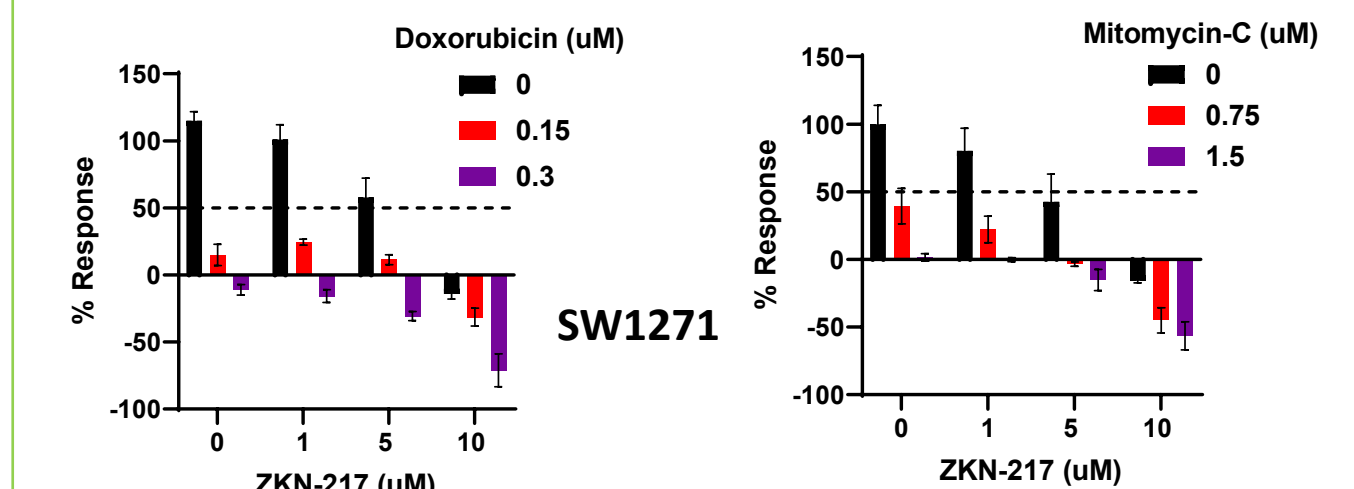
ZKN-217 selectively induces apoptosis in sensitive cells



ZKN-217 selectively inhibits SCLC patient-derived organoids (PDOs)



ZKN-217 shows combination synergy with DNA targeting chemotherapeutics known to inhibit rRNA synthesis



Multiple levels of selectivity of allosteric inhibition by RMAs

Novel Activity	Cause	Effect (of RMAs)
Ribosome selectivity	Ribosome heterogeneity	Target Ribosomes in a subset of cancers
Translation inhibition selectivity	Proteins containing regions with high density of positively charged amino acids	Inhibit translation of a subset of proteins

Conclusion

- Several cancer subtypes with prevalence of high MYC activity are sensitive to novel allosteric ribosome modulators
- Sensitivity in SCLC is associated with high expression of MYC target genes involved in Ribogenesis and protein translation
- Anti-proliferative effect is driven by selective inhibition of ribogenesis and new protein synthesis
- RMAs show additive and synergistic effects with major chemotherapy backbone agents