

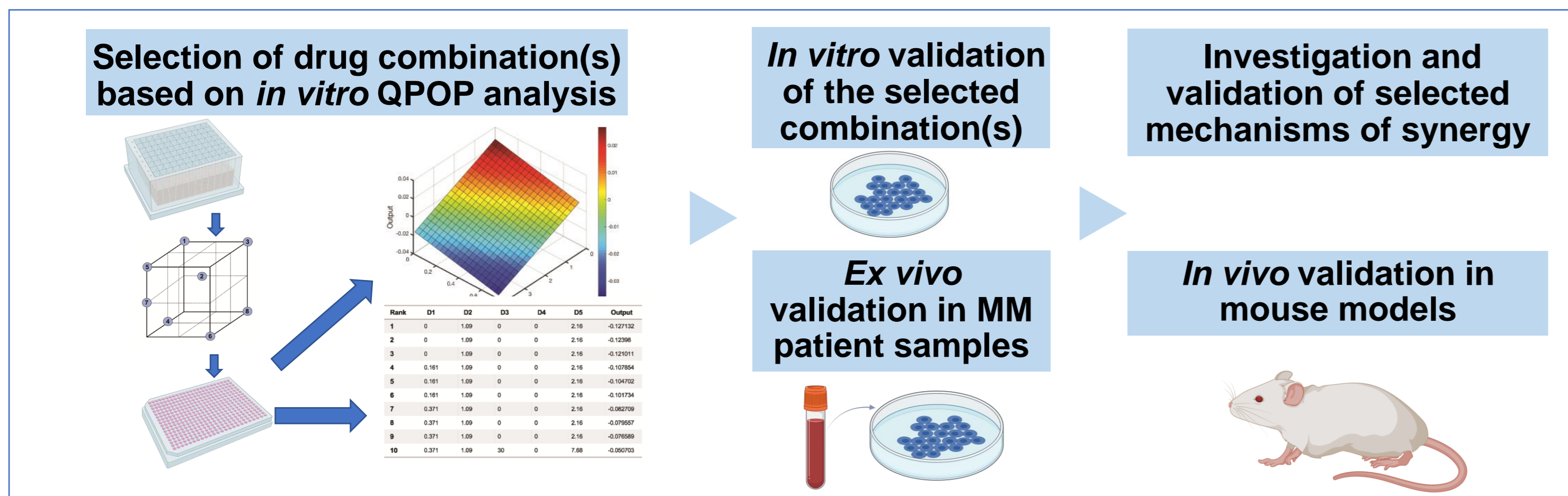
Overview

- We applied quadratic phenotypic optimisation platform (QPOP), a phenotypic-analytical hybrid multidrug interrogation platform developed by our lab to identify 8-Chloroadenosine and Pomalidomide as an effective novel drug combination against Multiple myeloma.
- The synergy of this combination has been validated *in vitro* and *in vivo*.
- This combination works potentially through:
 - Downregulation of c-Myc and its downstream targets
 - Alteration of the metabolic profiles, particularly glutamine metabolism
 - Stimulation of release of reactive oxygen species (ROS) and induction of apoptosis
- This study also reinforces the idea of targeting MM through inhibition of glutamine uptake, which has been implicated in several published literature

Introduction

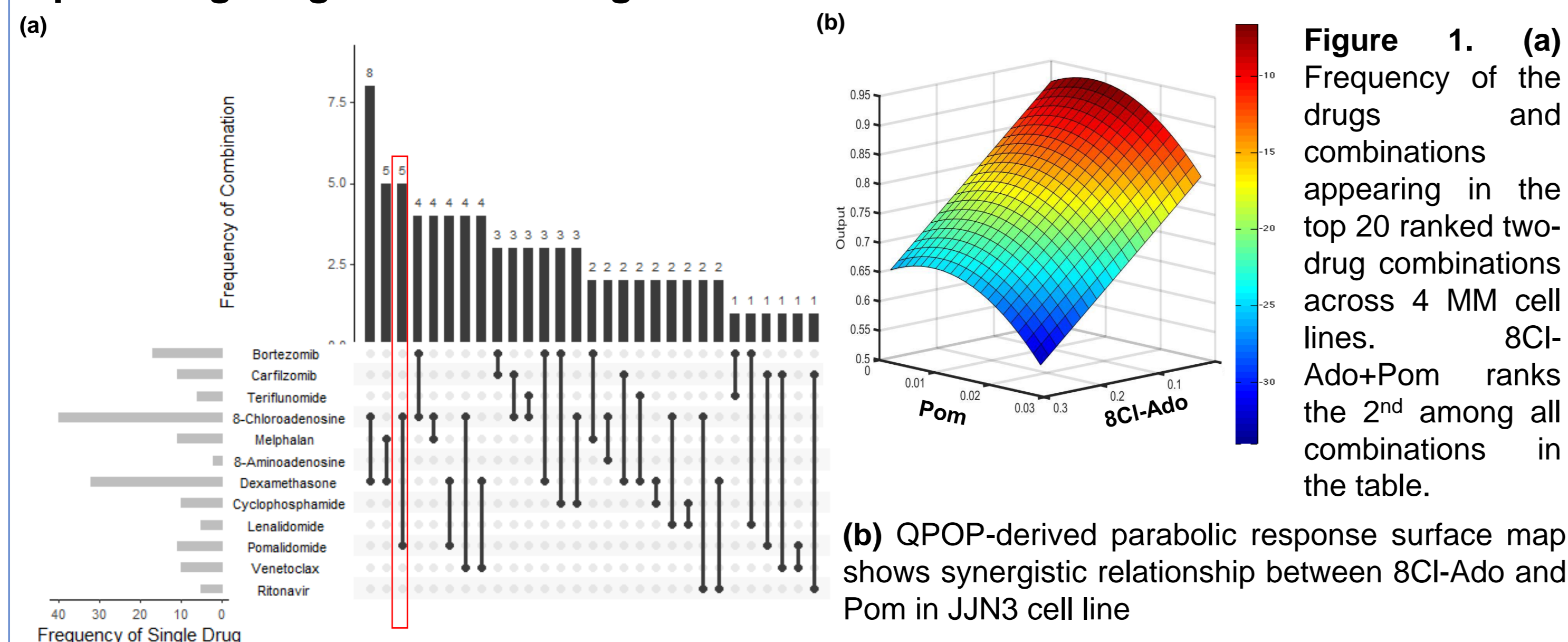
Multiple myeloma (MM) is a cancer disease characterised by malignant plasma cells, a type of white blood cells secreting immunoglobulins. It accounts for 10% of all haematological malignancies, ranked the second most common blood cancers only to Non-Hodgkin's Lymphoma worldwide.¹ Despite the advance in treatment strategies, MM is still an incurable disease, with no established curative treatments available. One main issue associated with this is the development of resistance to the current therapeutic approach. An effective strategy for overcoming drug resistance is to use combination therapy. Rapid advance in existing classes of treatments as well as the emergence of novel therapeutic approach has provided a large number of potential drug candidates to be used as combinatory therapies in second-line treatments.² Hence, our study aims to (1) apply a small dataset rational design approach (Quadratic Phenotypic Optimization Platform, QPOP)^{3,4} to identify effective drug-combinations in MM, (2) validate their efficacy *in vitro* and *in vivo*, and to (3) interrogate the molecular mechanisms underlying the selected drug combination in MM.

Methods

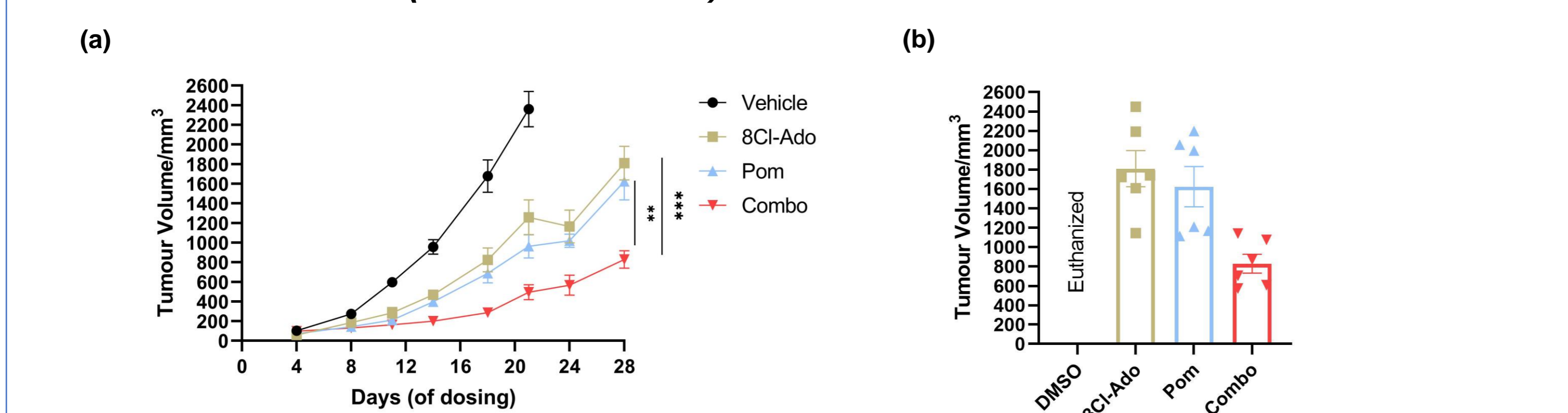


Results

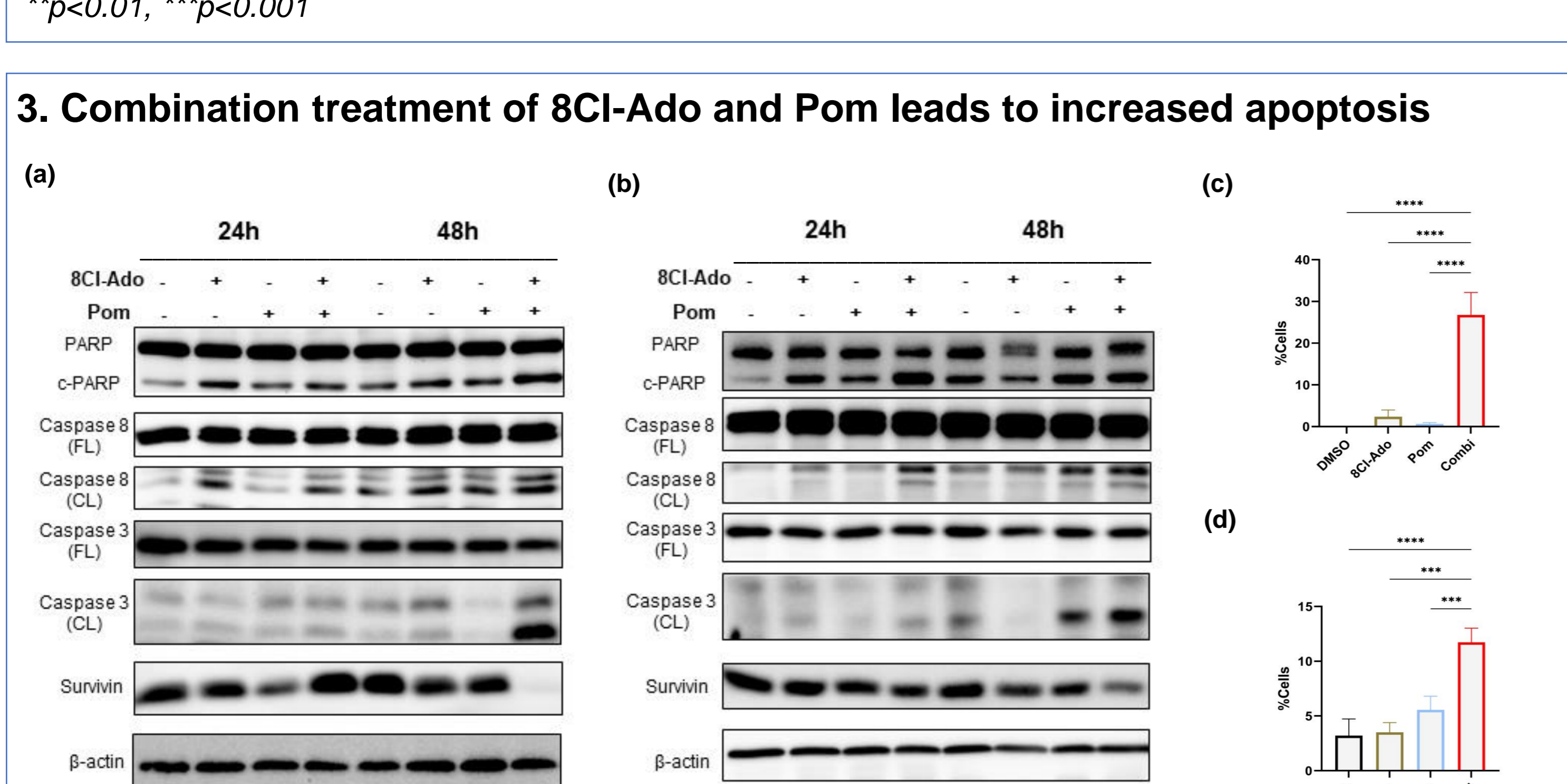
1. QPOP identifies 8-Chloroadenosine (8CI-Ado) and Pomalidomide (Pom) as a top-ranking drug combination against MM cells *in vitro*



2. 8CI-Ado and Pom demonstrates synergistic anti-tumour activity against MM tumour cells *in vitro* (data not shown) and *in vivo*



3. Combination treatment of 8CI-Ado and Pom leads to increased apoptosis



Results (continued)

4. Combination treatment of 8CI-Ado and Pom exerts their anti-myeloma effects through c-Myc downregulation

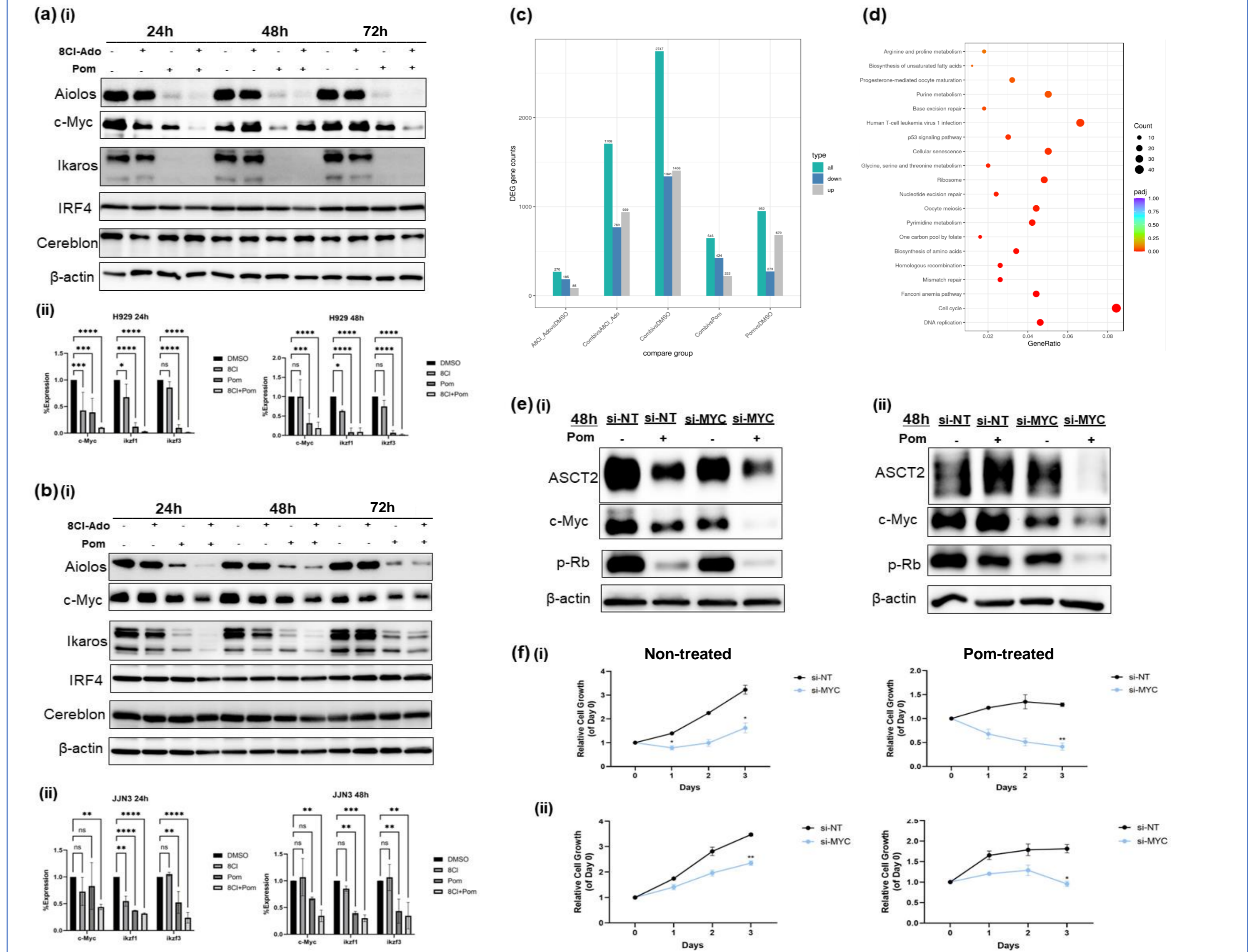


Figure 4. Immunoblots of (a)(i) H929 and (b)(i) JJJ3 and densitometric quantification of immunoblots (ii) and (iv) confirm the downregulation of the CRBN-IKZF signalling pathways, leading to significant downregulation c-Myc. (c) Combination treatment induces greater transcriptional change in terms of differentially-expressed genes. (d) KEGG enrichment analysis reveals the combination treatment induces transcriptional alterations in a c-Myc-regulated pathways. (e) Transient silencing of c-Myc in (i) H929 and (ii) JJJ3 enhances the effect of Pom, mimicking the synergistic effects of 8CI-Ado+Pom. (f) Transient silencing of c-Myc suppresses proliferation of (i) H929 (non-treated and Pom-treated) (ii) JJJ3 (non-treated and Pom-treated). **p*<0.05, ***p*<0.01, ****p*<0.001, *****p*<0.0001

5. Combination treatment of 8CI-Ado and Pom alters the metabolic profiles of MM cells

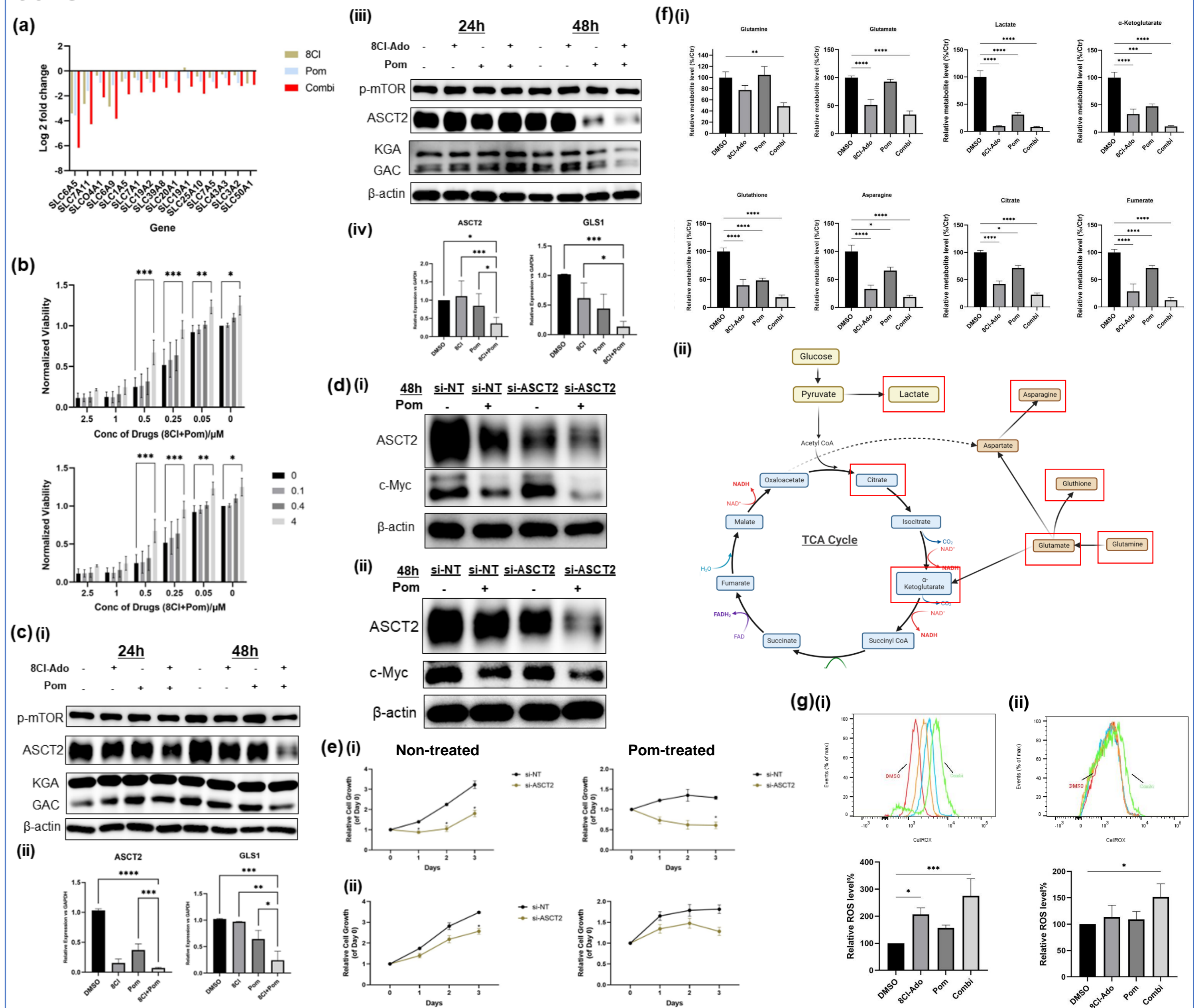


Figure 5. (a) The combination treatment induces significant downregulation of the solute-carrier (SLC) superfamily members. (b) Presence of extracellular glutamine partially rescues (i) H929 and (ii) JJJ3 from cytotoxic effects of the combination treatment. (c) Immunoblots and RT-qPCR of (i)(iii) H929 and (ii)(iv) JJJ3 show that combination treatment induces significant downregulation of the major glutamine transporter ASCT2 and the major glutaminase GLS1 in MM. (d) Transient silencing of ASCT2 in (i) H929 and (ii) JJJ3 enhances the effect of Pom, mimicking the synergistic effects of 8CI-Ado+Pom. (e) Transient silencing of ASCT2 suppresses proliferation of (i) H929, and Pom-treated H929, (ii) JJJ3 and Pom-treated JJJ3. (f)(i) LC-MS analysis reveals that the combination treatment reduces the cellular metabolic activities significantly in H929. (ii) Diagrammatic illustration of metabolites altered by the combination treatment in H929. (g) Combination treatment increases intracellular reactive oxygen species (ROS) level in (i) H929 and (ii) JJJ3, which could be the concomitant result of downregulation of the antioxidant glutathione. **p*<0.05, ***p*<0.01, ****p*<0.001, *****p*<0.0001

Conclusions

- QPOP analysis identifies that 8CI-Ado and Pom work synergistically against MM *in vitro* and *in vivo*
- This novel combination induces apoptosis and suppresses proliferation via c-Myc downregulation, leading to subsequent downregulation of pathways regulated by this transcription factor
- Combination treatment alters the metabolic profiles in MM cells, particularly suppressing glutamine uptake and metabolism. As ASCT2 and GLS1 are transcriptionally regulated by c-Myc, these alterations are potentially the concomitant results of upstream c-Myc degradation
- Future work : Comparison of expression of selective markers (e.g. Ki67, c-Myc, cleaved caspases, ASCT2) *in vivo*; Incorporation of more MM patient samples to identify specific biomarkers for this combination; Construction of Splice-switch oligonucleotide (SSO) to test *in vitro* and *in vivo*

References

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