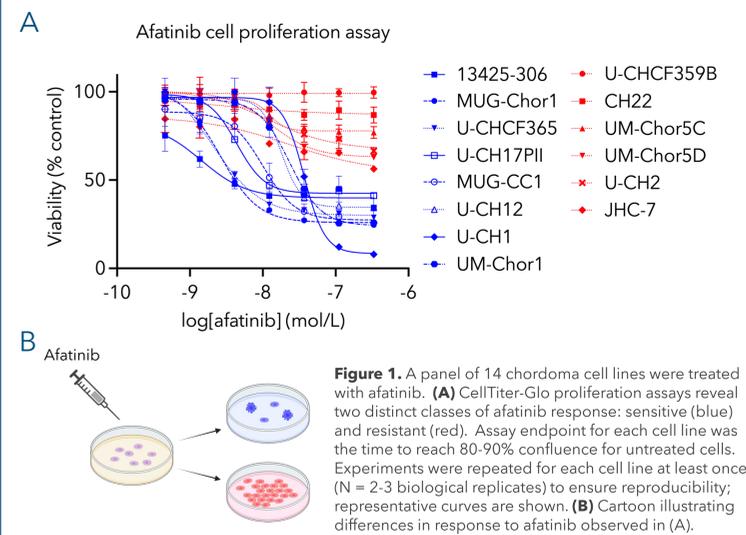


ABSTRACT

- Target discovery studies have identified EGFR as a promising therapeutic target in chordoma, motivating Phase II clinical trials with afatinib (NCT03083678) and cetuximab (NCT05041127).
- Recently-reported results from the afatinib trial include ORR = 9% with median DoR = 27.9 months (A. Lipplaa et al., *J Clin Oncol*, 2024), underscoring the need to identify therapeutic biomarkers. EGFR is not mutated in chordoma, leaving the mechanisms associated with sensitivity or resistance to EGFR inhibition unclear.
- We treated a panel of 14 chordoma cell lines with afatinib, an EGFR inhibitor (EGFRi) with potency against the wild-type receptor, and observed striking differential sensitivity. Sensitive cell lines have EC50 values < 50 nM, whereas resistant cell lines have EC50 values > 1 μM.
- Analysis of differentially-expressed genes in resistant versus sensitive cell lines revealed an association of interferon (IFN) signaling with afatinib resistance. Tumor cell-intrinsic IFN signaling has been associated with EGFRi resistance in lung cancer (K. Gong et al., *Nat Cancer*, 2020).
- In chordoma cells, modulation of IFN signaling does not reverse EGFRi sensitivity or resistance, suggesting the resistance driver may exist in an upstream IFN regulatory pathway. Studies are ongoing to identify the putative resistance driver.
- High IFN in chordoma may be linked to genomic instability and unresolved DNA damage, which leads to the activation of cytosolic nucleotide sensors. EGFR may play a role in DNA damage repair in chordoma.

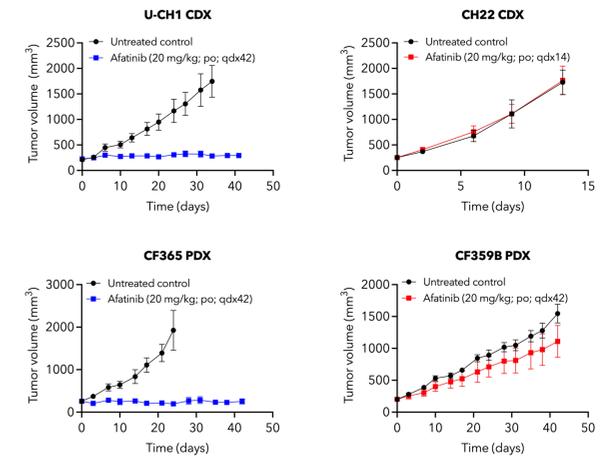
RESULTS

1. Chordoma preclinical models exhibit differential sensitivities to afatinib treatment

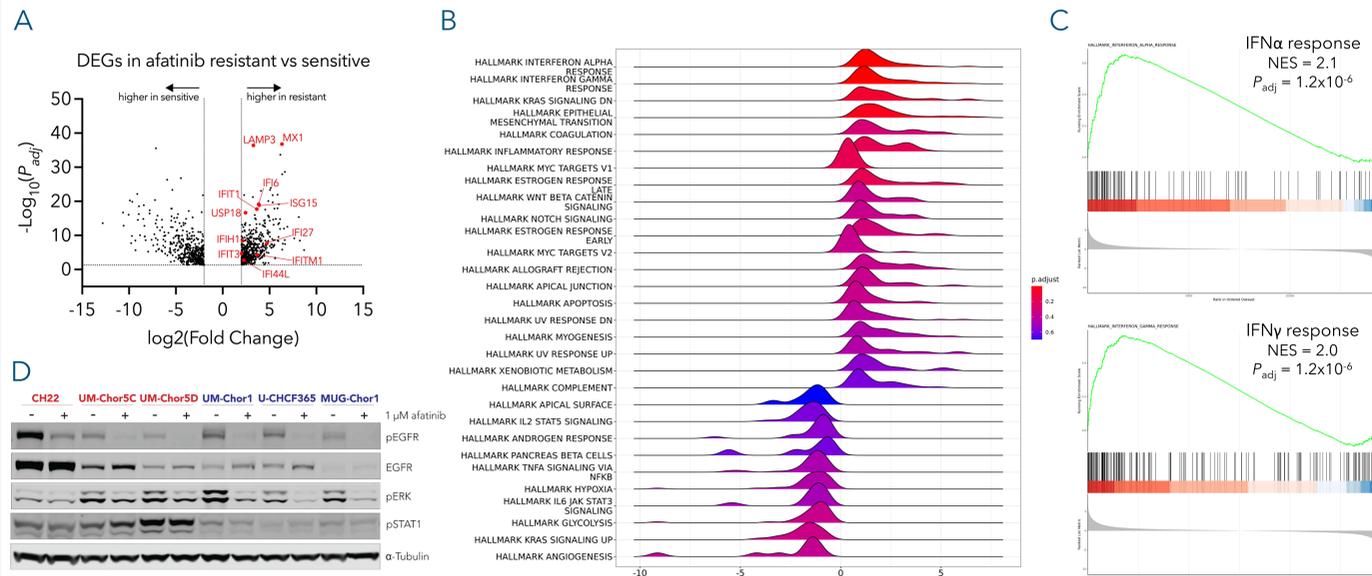


| Cell line | Abs EC50 (nM) | Response |
|------------|---------------|-----------|
| 13425-306 | 3 | Sensitive |
| U-CHCF365 | 5 | Sensitive |
| MUG-Chor1 | 8 | Sensitive |
| U-CH17PII | 11 | Sensitive |
| MUG-CC1 | 14 | Sensitive |
| U-CH12 | 27 | Sensitive |
| U-CH1 | 23 | Sensitive |
| UM-Chor1 | 33 | Sensitive |
| U-CHCF359B | > 1000 | Resistant |
| CH22 | > 1000 | Resistant |
| UM-Chor5C | > 1000 | Resistant |
| UM-Chor5D | > 1000 | Resistant |
| U-CH2 | > 1000 | Resistant |
| JHC-7 | > 1000 | Resistant |

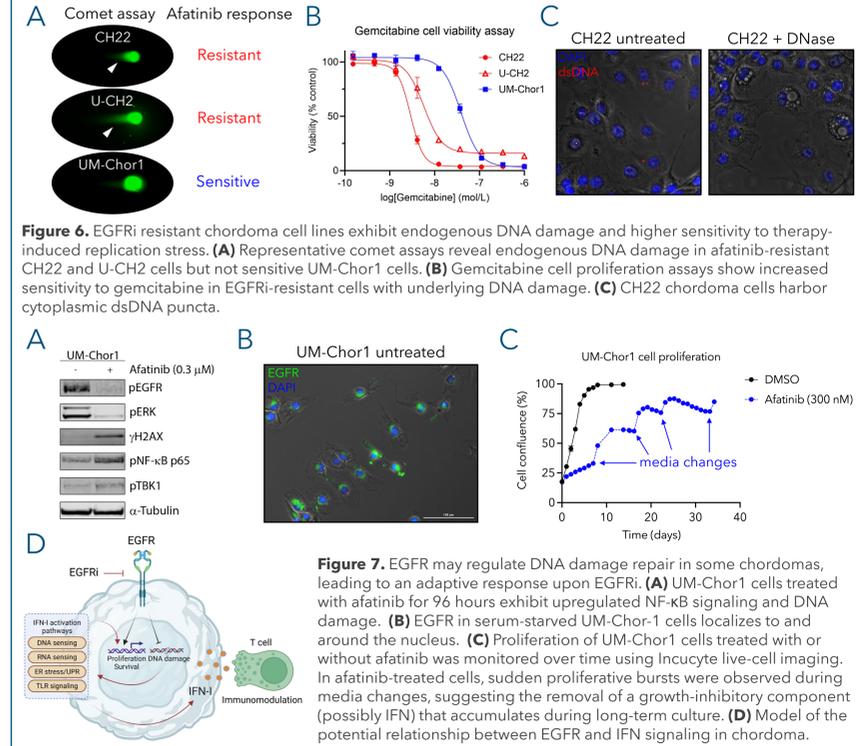
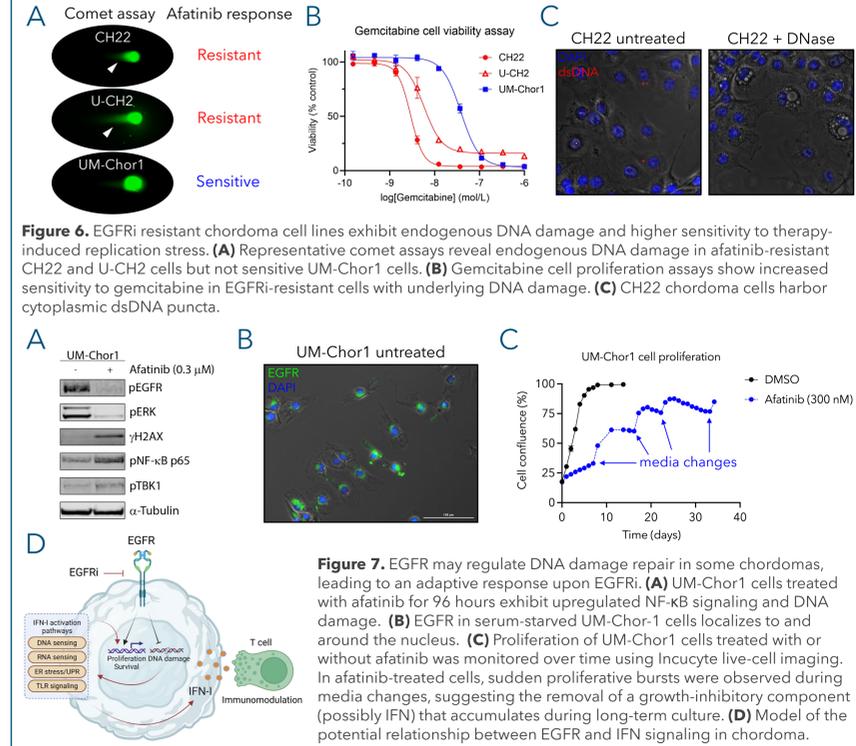
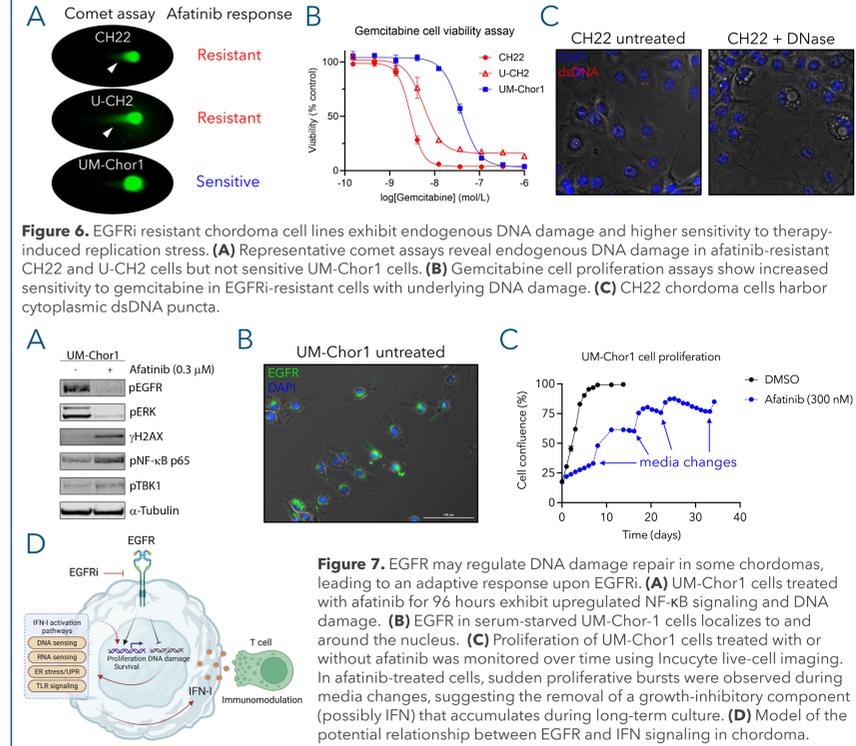
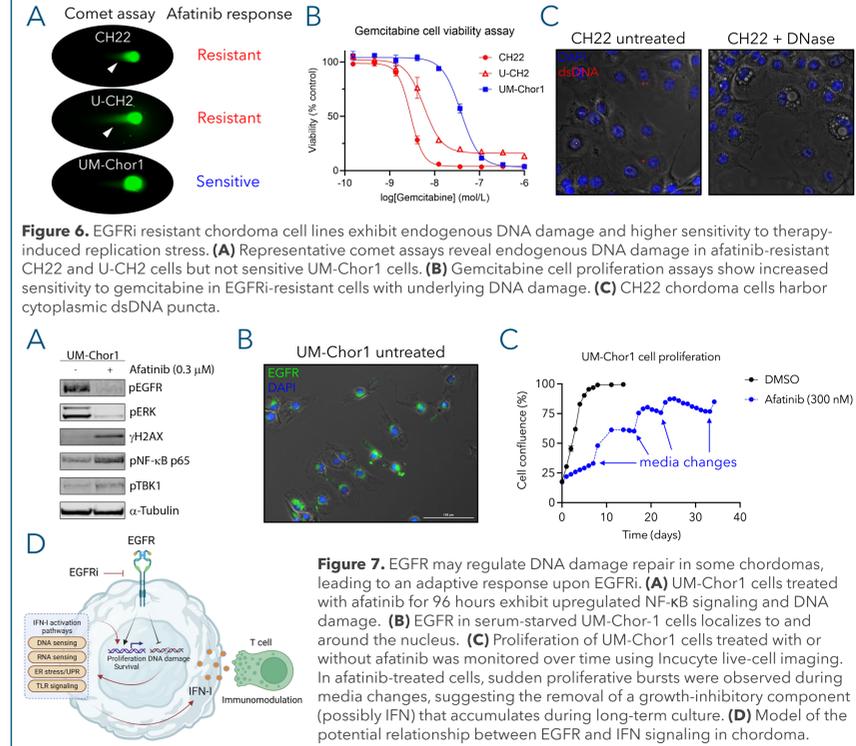
Table 1. Absolute EC50 values (in nanomolar) plotted for chordoma cell lines. The Abs EC50 is defined as the afatinib concentration that inhibits proliferation by 50% compared to control cells.



2. Differential gene expression analysis reveals an association between interferon signaling and afatinib resistance in chordoma



3. High IFN in chordoma may be linked to DNA damage



SUMMARY

- Chordoma cell lines exhibit striking differential sensitivity to afatinib, which is mirrored in xenograft mouse models.
- Combining *in vitro* afatinib sensitivity data with gene expression analysis identified an enrichment of interferon (IFN) signaling in resistant cell lines.
- IFN signaling per se does not promote resistance in chordoma cell lines. Instead, it may serve as a biomarker to guide patient selection for EGFR-targeted therapy.
- Studies are ongoing to identify the putative EGFRi resistance driver, and to characterize the causes and consequences of high IFN in chordoma.
- Acknowledgements:** We thank Kurt Bachman, Chris Moy, and Zayed Albertyn at Janssen R&D for providing in-kind bioinformatics support and expertise. Figures 1B, 5A, and 7D were created in BioRender.com.

