

E-cadherin loss imparts mitotic vulnerabilities rendering breast cancer cells synthetic lethal to crizotinib and up-regulation of Src signalling reverses this effect

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Lay Abstract

The E-cadherin molecule is often missing in lobular breast cancer cells. While its main job is to help cells stick together, it also plays a role in cell division. Previous work in our lab have found that a drug called crizotinib can be effective against cancers that do not have E-cadherin. Our current work has discovered that crizotinib causes problems in cell division, resulting in mistakes that can ultimately cause the cell to die. We identified a molecule called *CSK* that regulates this process and when *CSK* is turned off, the cancer cells become resistant to crizotinib. We also find that *CSK* may be involved in a pathway that controls the formation of important structures during cell division. This research could lead to new treatments that targets these vulnerabilities in lobular breast cancer.

Introduction

- Loss of E-cadherin (*CDH1*) is the pathognomonic alteration in lobular breast cancer.
- We have previously demonstrated synthetic lethality between the clinical *ROS1* tyrosine kinase inhibitor crizotinib and E-cadherin loss.
- Genetic perturbation screens of E-cadherin defective breast cancer cells exposed to crizotinib identified increased dependencies on genes controlling mitotic processes.
- E-cadherin has roles beyond cell adhesion and is known to regulate mitotic spindle orientation as well as centrosome clustering.
- Our top validated crizotinib resistance gene hit was *CSK* - the negative regulator of Src-family kinase (SFK) signalling.
- In this poster we describe the crizotinib synthetic lethal mechanism, linking E-cadherin loss with dysregulation of Aurora-A kinase localisation and provide a possible resistance pathway mediated through loss of *CSK*.

1. Uncovering novel mediators of crizotinib resistance

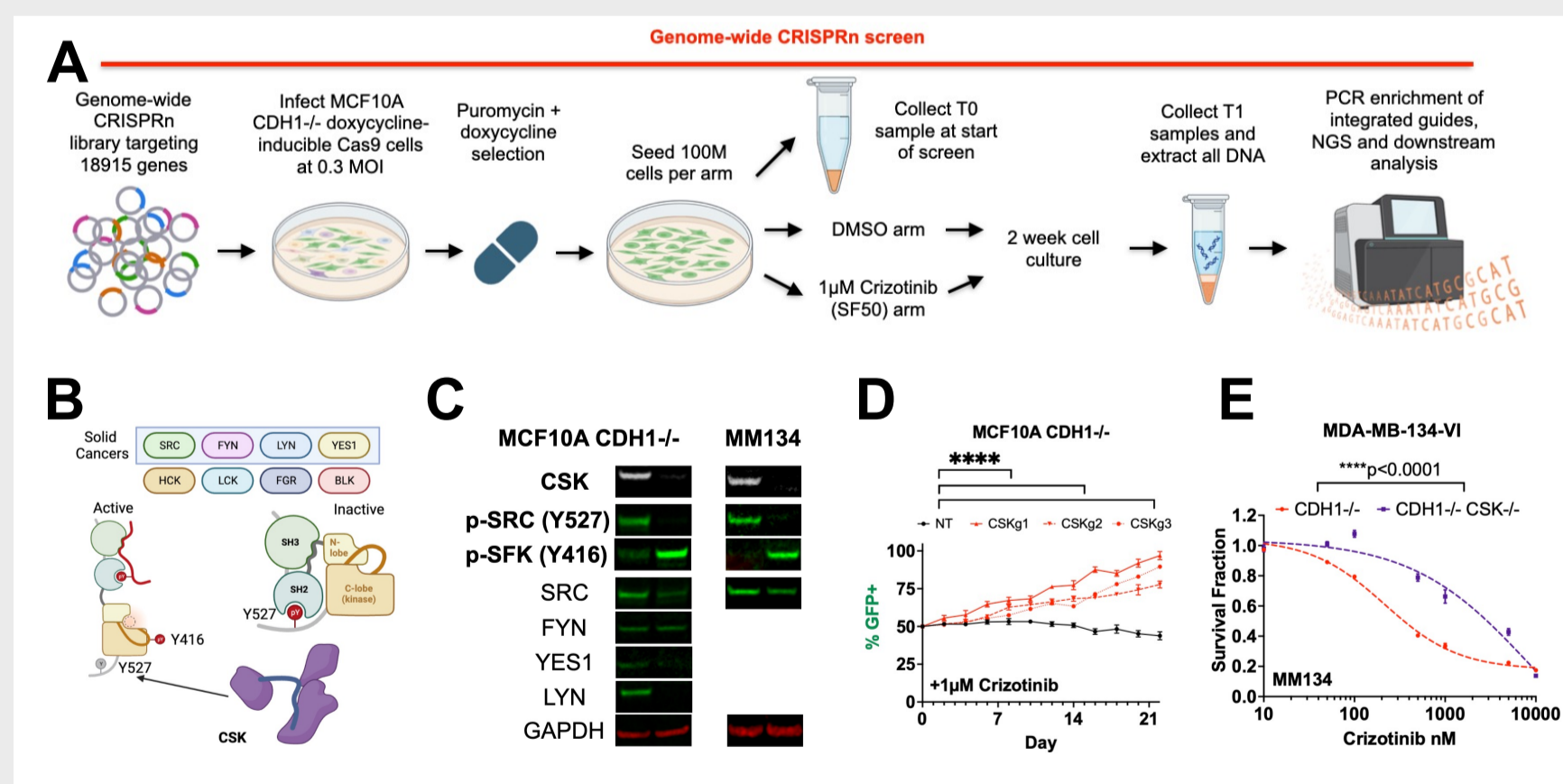


Figure 1. Genome-wide CRISPR screen uncovers novel mediators of crizotinib resistance in E-cadherin deficient cells.

A. Schematic of CRISPR screen. **B.** Knockout of *CSK*, negative regulator of Src family kinases (SFK) validated as a top resistance hit. **C.** *CSK*^{-/-} lines were generated and effects on SFK activity defined by western blot. **D.** Knockout of *CSK* using three different CRISPR guides validated crizotinib resistance in competition assays. **E.** ILC MM134 *CSK*^{-/-} cell line was crizotinib resistant.

2. E-cadherin loss weakens centrosome functions

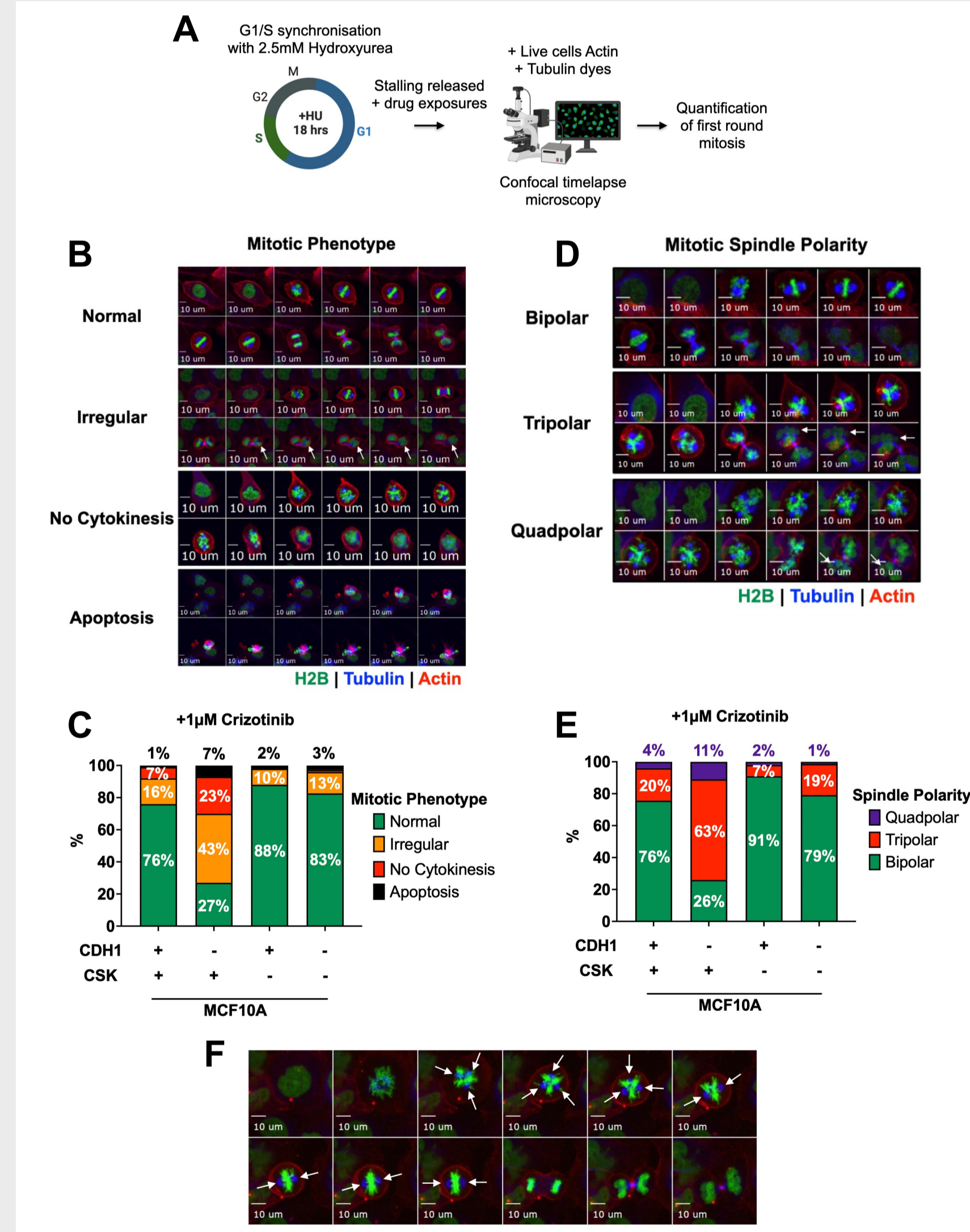


Figure 2. E-cadherin loss exacerbates formation of multipolar mitotic spindles after crizotinib exposure and up-regulation of SFK-activity rescues this defect.

A. Cells were synchronised with Hydroxyurea and confocal time-lapse microscopy carried out. **B+C.** Cells exposed to crizotinib were classified into four phenotypes. **D+E.** Quantification of spindle polarity prior to anaphase. **F.** *CSK*^{-/-} cells were able to cluster fragmented spindle poles, restoring bipolarity.

3. E-cadherin and SFKs modulates centrosomal Aurora-A

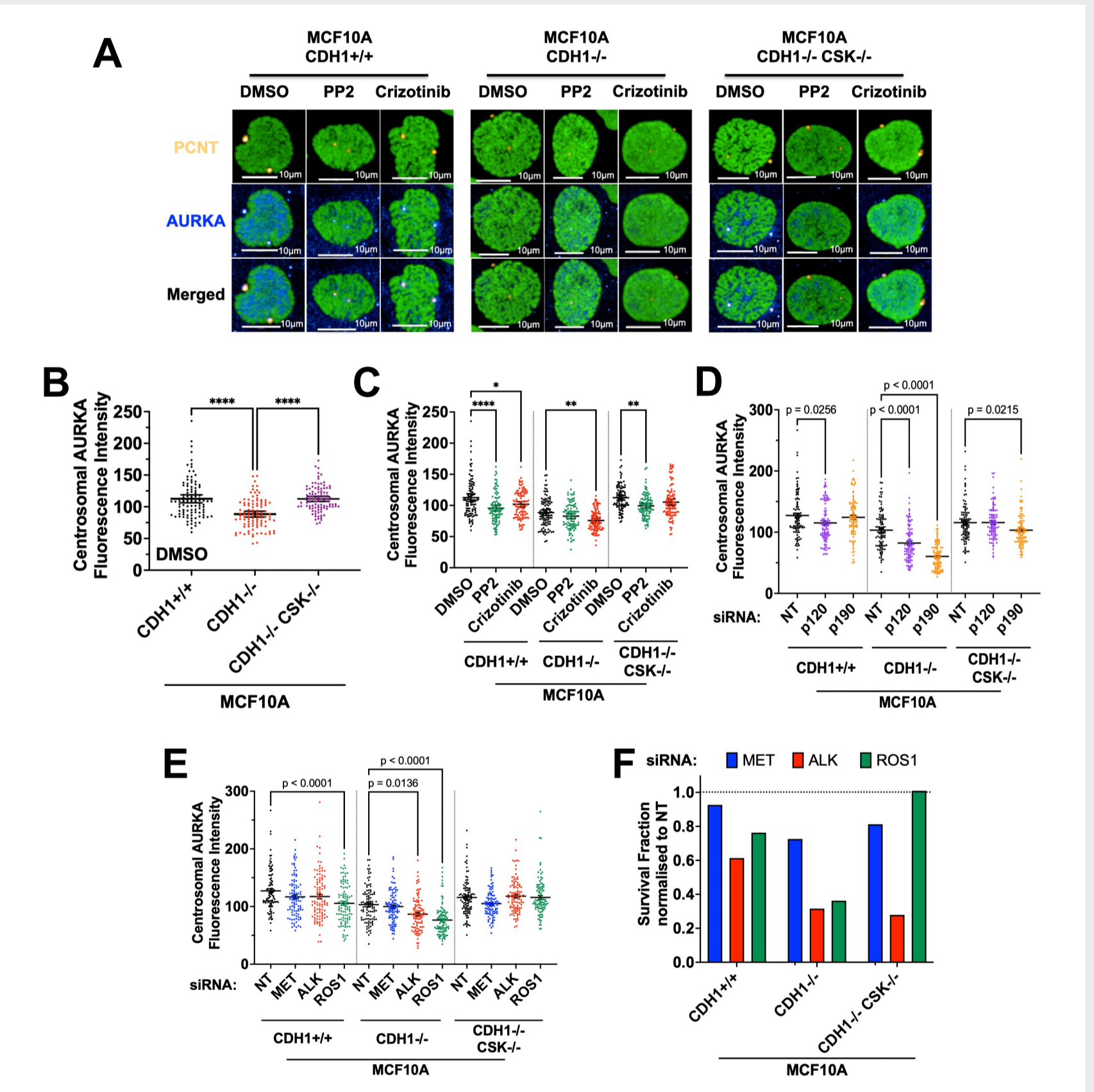


Figure 3. E-cadherin and SFK-activity modulates centrosome maturation by promoting Aurora-A kinase localisation during late G2/early M phase.

A. Cells expressing H2B-GFP were stained for Pericentrin (PCNT) and Aurora-A kinase (AURKA). **B.** Fluorescence intensity of AURKA at the centrosome was quantified. **C.** Centrosomal AURKA quantification following exposure to PP2 (pan SFK-inhibitor) or crizotinib. **D.** p120 or p190 RNAi reduced centrosomal AURKA in *CDH1*^{-/-} cells. **E.** *ROS1* or *ALK* RNAi reduced centrosomal AURKA in *CDH1*^{-/-} cells. **F.** *ROS1* RNAi synthetic lethality was reversed following *CSK*-loss.

Conclusions

- Loss of E-cadherin weakens centrosome maturation through reducing Aurora-A kinase localisation during late G2/early M phases of the cell cycle.
- This defect following E-cadherin loss can be targeted through *ROS1* and *ALK* inhibition with crizotinib.
- Up-regulation of Src-family kinase activity confers resistance through restoration of Aurora-A kinase expression.