

Table: 1879P

| Drugs                                     | H-1703                         | Calu-1<br>IC50 (nM)            | SK-Mes-1                       |
|---|--------------------------------|--------------------------------|--------------------------------|
| PF-05212384<br>(PI3K/mTOR inhibitor)      | 18                             | 26                             | 335                            |
| AZD-2014<br>(Dual mTORC1/C2<br>inhibitor) | 145                            | 217                            | 215                            |
| Everolimus<br>(mTORC1 inhibitor)          | Unable<br>to determine<br>IC50 | Unable<br>to determine<br>IC50 | Unable<br>to determine<br>IC50 |
| MK-2206<br>(pan-Akt inhibitor)            | Unable<br>to determine<br>IC50 | Unable<br>to determine<br>IC50 | Unable<br>to determine<br>IC50 |

**Conclusions:** Overall, the results of our study suggest the potential implication of PI3K/mTOR-Rictor pathway in SQLC oncogenesis, thus rendering it a promising target for a targeted approach. Among the mTOR components, RICTOR CNG seems to predict a higher sensitivity to PI3K/mTOR inhibition and might represent a potential biomarker to be explored as a stratification tool in clinical trials. Confirmatory RICTOR silencing experiments are currently ongoing.

**Legal entity responsible for the study:** Emilio Bria.

**Funding:** AIRC (Associazione Italiana per la Ricerca sul Cancro).

**Disclosure:** All authors have declared no conflicts of interest.

**1879P Potential role of RICTOR copy number gain (CNG) as a key biomarker of mTOR activity: A comprehensive preclinical analysis in squamous cell lung cancer (SQLC) models**

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**Background:** We previously performed a multi-step genomic study in almost 100 resected SQLC patients dichotomized according to the prognosis. Among the pathways with a biological impact on SQLC oncogenesis, PI3K/mTOR-Rictor emerged as a crucial axis [Pilotto WCLC 2016]. In order to explore the potentiality of mTOR inhibition, we present a set of in vitro experiments in RICTOR-aberrant SQLC preclinical models.

**Methods:** Next-generation sequencing (NGS), fluorescence in situ hybridization (FISH) and Western Blot were performed in 3 SQLC cell lines (H-1703, SK-Mes-1, Calu-1) for detecting CNG/protein profile of the PI3K/mTOR-Rictor components. The activity of PI3K/mTOR pathway targeted inhibitors in the SQLC cell lines was examined in short- (72 hours) and long-term (1 week) cell viability assays.

**Results:** NGS analysis revealed a different amount of RICTOR CNG among SQLC cell lines. FISH confirmed that H-1703 harbors the highest number of RICTOR copies (6) followed by SK-Mes-1 (4) and Calu-1 (3.5), suggesting polysomy of the short arm of chromosome 5 as the main mechanism of RICTOR gain. Although Rictor protein levels were similar among the cell lines, p-mTOR S2448 (active form of mTOR complexes) was higher in H-1703 than SK-Mes-1 and Calu-1. PI3K/mTOR inhibition proved more effective in H-1703, with lower IC50 values in short term treatment (Table). Similar findings were confirmed in long-term assays.