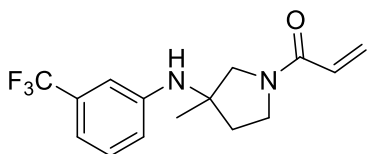
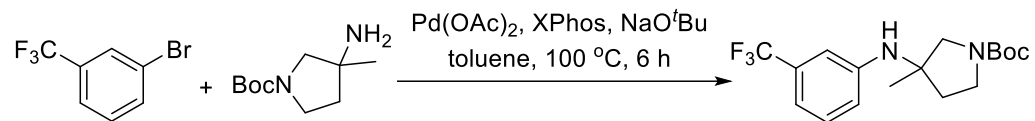


**MYF-01-037**Chemical Formula: C<sub>15</sub>H<sub>17</sub>F<sub>3</sub>N<sub>2</sub>O

Molecular Weight: 298.30921

| Category                  | Parameter                          | Description   |
|---------------------------|------------------------------------|---|
| Compound                  | Name                               | MYF-01-037  |
|                           | Citation                           | Kurppa, K.J., Liu, Y., To, C., Zhang, T., Fan, M., Vajdi, A., Knelson, E.H., Xie, Y., Lim, K., Cejas, P., et al. (2020). Treatment-Induced Tumor Dormancy through YAP-Mediated Transcriptional Reprogramming of the Apoptotic Pathway. <i>Cancer Cell</i> 37, 104-122.e112. |
|                           | Chemical descriptors               | O=C(C=C)N(C1)CCC1(C)NC2=CC(C(F)(F)F)=CC=C2  |
|                           | Chemical name                      | 1-(3-methyl-3-((3-(trifluoromethyl)phenyl)amino)pyrrolidin-1-yl)prop-2-en-1-one   |
|                           | Entries in chemical databases      | N/A   |
|                           | Availability                       | N/A   |
|                           | Additional comments                | Negative control MYF-01-104 also available  |
| <i>In vitro</i> profiling | Target (potency)                   | Gel-Based TEAD2 palmitoylation inhibition assay (IC <sub>50</sub> > 50uM)   |
|                           | Target (potency)                   | TEAD mCherry reporter assay on PC-9 cells (IC <sub>50</sub> ~ 5uM)  |
|                           | Selectivity                        | Chemoproteomics data available, R value = 2.5 (Chouchani lab)   |
|                           | Potential reactivity               | acrylamide  |
|                           | SAR                                | In progress, not published yet  |
|                           | Mechanism of inhibition            | Covalent binding to conserved Cys on TEAD palmitate pocket (for example C380 on TEAD2)  |
|                           | Structure of target-probe complex  | Docking only  |
| Cellular profiling        | Validation of cellular target      | Pull down by dethiobiotin probe MYF-01-108  |
|                           | Validation of cellular specificity | Chemoproteomics data available, R value = 2.5 (Chouchani lab)   |
|                           | Additional comments                |   |
| Pharmacodynamics          |                                    | N/A   |
| Pharmacokinetics          |                                    | N/A   |



Synthetic scheme

