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## Modulation of the aspartic acid scaffold to identify a new septin-4 covalent binder with anti-metastatic activity in a mouse model of melanoma

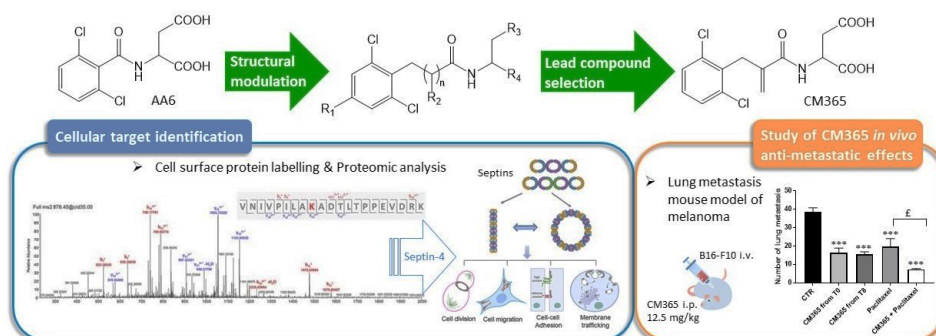
F. Boccato,<sup>a</sup> F. Blua,<sup>a</sup> A. A. Altomare,<sup>b</sup> S. Vittorio,<sup>b</sup> N. Clemente,<sup>c</sup> C. Monge,<sup>a</sup> E. Marini,<sup>a</sup> M. Bertinaria,<sup>a</sup>

<sup>a</sup> Department of Drug Science and Technology, University of Turin, Via P. Giuria 9, Turin, Italy

<sup>b</sup> Department of Pharmaceutical Sciences, University of Milan, Via L. Mangiagalli 25, Milan, Italy

<sup>c</sup> Settore Centri di Ricerca e Infrastrutture, University of Piemonte Orientale, Corso Trieste 15/A, Novara, Italy  
francesca.boccato@unito.it

Tumour metastases are still the leading cause of cancer-related death, being often responsible for the failure of current therapies.<sup>1</sup> In our previous work, we have developed a compound, AA6, that proved able to counteract the metastatic spread in a mouse model of breast cancer.<sup>2</sup> To identify the essential structural elements and the cellular target involved in the activity of our lead compound, a new series of AA6-derivatives was designed and synthesised. New compounds were tested for the inhibitory activity on tumour cell invasion, migration and adhesion in B16-F10 melanoma cells. From the *in vitro* results, compound CM365 was selected as the most promising derivative, showing good efficacy and most likely able to interact with the target through covalent binding. To pinpoint the cellular target, a proteomic analysis was carried out and this allowed to select septin-4 as the most likely protein involved. Septins are GTP-binding proteins, able to assemble into large filaments on the plasma membrane, modifying its rigidity in migrating cells and promoting tumour metastasis.<sup>3</sup> To study and compare the binding modes of the two derivatives, AA6 and CM365, with the molecular target, computational analyses were performed. Finally, the anti-metastatic properties of CM365 were evaluated *in vivo* in a murine model of metastatic melanoma. This compound proved capable of reducing metastases dissemination at different times of administration and of enhancing the antitumour effect of a known anticancer drug, paclitaxel, when administered in combination. The discovery of septin-4 as a new molecular target for the prevention of metastatic spread may encourage the future development of more specific inhibitors.



### References:

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