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Type I IFNs promote cancer cell stemness by triggering the epigenetic regulator KDM1B

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Cancer stem cells (CSCs) are a subpopulation of cancer cells endowed with high tumorigenic, chemoresistant and metastatic potential. Nongenetic mechanisms of acquired resistance are increasingly being discovered, but molecular insights into the evolutionary process of CSCs are limited. Here, we show that type I interferons (IFNs-I) function as molecular hubs of resistance during immunogenic chemotherapy, triggering the epigenetic regulator demethylase 1B (KDM1B) to promote an adaptive, yet reversible, transcriptional rewiring of cancer cells towards stemness and immune escape. Accordingly, KDM1B inhibition prevents the appearance of IFN-I-induced CSCs, both in vitro and in vivo. Notably, IFN-I-induced CSCs are heterogeneous in terms of multidrug resistance, plasticity, invasiveness and immunogenicity. Moreover, in breast cancer (BC) patients receiving anthracycline-based chemotherapy, KDM1B positively correlated with CSC signatures. Our study identifies an IFN-I → KDM1B axis as a potent engine of cancer cell reprogramming, supporting KDM1B targeting as an attractive adjunctive to immunogenic drugs to prevent CSC expansion and increase the long-term benefit of therapy.

SCs, also known as tumor-initiating or tumor-propagating cells, are a relatively rare stem-like cell subpopulation within the tumor capable of self-renewal and multilineage differentiation, and responsible for tumor initiation, progression, spreading and therapy resistance^{1,2}. Mounting evidence indicates that CSCs can evolve over space and time leading to a high degree of genotypic, phenotypic and functional heterogeneity^{2,3}. Along with this, it is emerging that non-CSC subsets can adapt to the changes in the tumor microenvironment (TME), undergoing cell reprogramming and (re)generating CSCs².

Epigenetic dysregulations critically affect cancer-immune cell interactions and coevolution during disease onset, progression and response to therapy by influencing cellular states and fates⁴. Not surprisingly given their role in normal stem cell maintenance, epigenetic mechanisms have also been involved in CSC preservation⁵. This feature, together with the inherent reversibility of epigenetic modifications, makes the use of epigenome-targeting drugs (epidrugs) a unique opportunity to rationally target CSCs in combination with conventional therapies^{4,6–9}.

One key concept in tumor immunology is that some chemotherapeutics, including (but not limited to) anthracyclines (for example, doxorubicin, DOX), oxaliplatin (OXP) and cyclophosphamide^{10,11} induce cancer immunogenic cell death (ICD), a form of regulated cell death that initiates adaptive immune responses by the emission

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1 Musella M, Guarracino A, Manduca N, Galassi C, Ruggiero E, Potenza A, Maccafeo E, Manic G, Mattiello L, Soliman Abdel Rehim S, Signore M, Pietrosanto M, Helmer-Citterich M, Pallocca M, Fanciulli M, Bruno T, De Nicola F, Corleone G, Di Benedetto A, Ercolani C, Pescarmona E, Pizzuti L, Guidi F, Sperati F, Vitale S, Macchia D, Spada M, Schiavoni G, Mattei F, De Ninno A, Businaro L, Lucarini V, Bracci L, Aricò E, Ziccheddu G, Facchiano F, Rossi S,

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