



RESEARCH TOPIC CLI4

Single cell dissection of the tumor microenvironment in patient with urothelial bladder cancer for the identification of predictive tools and new therapies

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Abstract

Advanced Bladder Cancer (BC) represents a challenge in oncology due to its high incidence and the scarcity of therapeutic options. At first diagnosis most of the patients present a non-muscle invasive cancer (NMIBC), that is treated with transurethral resection (TURBT). Intravesical instillation of Bacillus Calmette-Guérin (BCG) also shows strong efficacy in this context. Nevertheless, a percentage of cancers eventually recur or show resistance to MMC and/or BCG, thus evolving in a muscle-invasive disease (MIBC) associated with a high mortality. Radical cystectomy (RC) with or without neo-adjuvant chemotherapy is the standard of care, but morbidity and mortality remain high. There is thus an unmet clinical need for improving the rate of care and quality of life. Importantly, RC may be applied to treat BCG-resistant NMIBC but markers that can indicate rate of response and guide therapy decision making are missing. Thus, novel predictive tools are needed to guide surgery decision in this context. Also, novel therapeutics that may improve response to BCG or be applied as alternative option are strongly needed. On this regard, immunotherapies are under investigation and hold strong promise. NK adoptive cellular therapies have been tested in clinic and showed efficacy in solid tumors. Preclinical in vitro and in vivo efforts demonstrated that CAR-NK-92 cells efficiently kill urogenital cancer cells and inhibit BC growth in mice. Nevertheless, cancer exerts an inhibitory activity on infiltrating NK cells, thus partially hindering cell therapy efficacy. Here we will test the hypothesis that the composition of the TME can be used to as prognostic factor and that specific immune subsets can be targeted to prevent progression. Moreover, we will test NK cell therapies in BC preclinical models and elaborate strategies to improve efficacy. Preliminary data from the Consortium indicate that the TME strongly varies between MIBC and NMIBC. Also, we identified the CXCL12-CXCR4 axis as a strategy to improve the efficacy of NK-cell therapies in urogenital cancers. We propose a single cell-based approach at the RNA and protein level to unveil the dynamics of the TME in BC during progression and in resistance to therapy. We will determine new prognostic and predictive markers and we will test novel immunotherapies in preclinical settings.

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