

INTRODUCTION

Multiple myeloma (MM) is the second most frequent hematological malignancy. It is characterized by the progressive accumulation of plasma cells in the bone marrow (BM), which provides a favourable microenvironmental *niche* for uncontrolled growth, escape from apoptosis and genetic selection of increasingly biologically aggressive subclones. The introduction of immunomodulatory drugs, such as lenalidomide (Lena) and pomalidomide, has improved both quality of life and prognosis of patients, which however remain still poor. Novel therapeutic targets are thus urgently needed.

Recently it was demonstrated that in myelodysplastic syndromes and in acute myeloid leukemia, lenalidomide treatment induced a Cereblon-mediated ubiquitination and degradation of protein kinase CK1α [1].

CK1α is a serine/threonine kinase essential for the function of signaling pathways that could be involved in MM pathobiology: the canonical Wnt/β-catenin cascade, Hedgehog, NF-κB pathways and p53-driven response (here by stimulating the binding to Mdm2 and p53 inhibition). Recent evidence indicates that protein kinase CK1α may play a key role in myeloma cell growth and its inactivation determines MM cell death [2].

Therefore, CK1α could be a potential candidate for mediating the cytotoxic /cytostatic effects of lenalidomide also in MM. Here, we sought to investigate the therapeutic potential of the combination of lenalidomide and CK1α inactivation on malignant plasma cells. Inhibition of CK1α in association with lenalidomide may represent an original molecular target approach for MM.

OBJECTIVES

We have analyzed CK1 α expression and its contribution to MM survival. We investigated the therapeutic potential of the combination of lenalidomide and CK1α inactivation studying whether lenalidomide could cause CK1 α degradation and whether CK1 α could take part in lenalidomide-induced MM cell apoptosis and cell cycle arrest. We next studied whether blocking CK1α could influence pro-survival signalling pathways, like β-catenin and AKT accounting for resistance to lenalidomide.

METHODS

MM cell lines: H929, U-266, (ATCC, USA), INA-6 (gift from Dr. M. Gramatzki, University of Kiel, Kiel, Germany) and SaMMi (a patient derived MM cell line, newly generated in our laboratory).

• A model of BM microenvironment was established colturing MM cells with HS-5 stromal cells.

• RNA interference for CK1 α was obtained through electroporation of CK1 α directed double strand oligonucleotides (AMAXA system). CK1 inhibition was obtained with D4476 (abcam).

• Apoptosis and cell cycle were investigated with AnnexinV/PI and Ki-67/PI stainings and cytofluorimetric analysis. Lenalidomide- and CK1α-dependent signalling events were analyzed by WB.

CK1α INACTIVATION IN MULTIPLE MYELOMA EMPOWERS LENALIDOMIDE **INDUCED CYTOTOXICITY AND CELL CYCLE ARREST**

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