EphA4 receptor as a potential therapeutic target in colorectal cancer

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INTRODUÇÃO

Colorectal cancer (CRC) is the second most common malignancy diagnosed in women and third in men, being the fourth most common cause of cancer mortality worldwide (Jemal et al., 2011). Radiotherapy (RT) is widely used as a neoadjuvant treatment for advanced rectal tumors in order to reduce tumor size. This approach reduces the risk of local recurrence in comparison with surgery alone. However, this confers only a small impact on distant metastasis formation (Glynne-Jones et al., 2013). Tumor recurrence tends to be more aggressive with invasive metastatic conditions and shorter survival expectancy after preoperative RT (Vicini et al., 2003; Sauer et al., 2004). In previous studies, we demonstrated that the progeny derived from HT-29 CRC cell line submitted to 5 Gy irradiation displayed an EMT-like phenotype characterized by downregulation of E-cadherin and increased migratory and invasive potentials (Fig. 1) (Bastos et al., 2014; Marcondes et al., 2016). These cellular events could be responsible for the high rates of therapeutic failure promoting local invasion and metastasis in rectal cancer after RT.







Fig. 1. Illustration of the signalling pathways induced in the progeny of colorectal cancer. Unirradiated (left) and irradiated (right) cells (Marcondes et al., 2016).



Fig. 3. HT-29 cells were preteated for 1 hr with kyl (30 μ M) and after with doxasozin (23 μ M) the cells were maintained for 24h. (A) Cell morphology analyzed using phase-contrast microscopy. Scale bar 50 μ m. (B) Western blot analysis of p-EphA4 (Tyr-602) and EphA4 protein levels. GAPDH protein was used as a loading control. Numbers below the figure represent the ratio of band's optical density as fold change of protein expression normalized by GAPDH. Doxa: doxazosin



Analyze the use of EphA4 as a therapeutic target for CRC. For this, we will use a small peptide designated KYL that competitively inhibit Ephrin binding to EphA4 receptor.

METHODS



PRELIMINARY RESULTS

Table 1. Microarray analysis of the main genes involved in the EMT process and the Ephrin genes that bind to the EphA4 receptor.

Gene Symbol	Gene Assignment	Fold-Change (IR vs. Control)
SNAI2	Snail homolog 2 (Drosophila)	2,25
STAT3	Signal Transducer and Activator of Transcription 3	1,39
EFNA4	Ephrin - A4	1,35
SIP1	Survival of Motor Neuron Protein Interacting Protein 1	1,31
VIM	Vimentin	1,22
SNAI1	Snail Homolog 1 (Drosophila)	1,17
TGFB1	Transforming Growth Factor Beta 1	1,17
CDH1	E-cadherin (epithelial)	1,15
EFNB1	Ephrin-B1	1,14
EFNA2	Ephrin-A2	1,10
EFNA5	Ephrin-A5	1,09
CDH2	N-cadherin (neuronal)	1,05
TGFBR1	Transforming Growth Factor Beta Receptor 1	1,04
EFNB2	Ephrin-B2	1,00
EPHA4	EPH receptor A4	-1,06
ZEB1	Zinc Finger E-box Binding Homeobox 1	-1,07
ZEB2	Zinc Finger E-box Binding Homeobox 2	-1,08
TWIST1	Twist Homolog 1 (Drosophila)	-1,24





Fig. 4. HT-29 cells were preteated for 1 hr with kyl (30 μM) and after with doxasozin (23 μM) the cells were maintained for 24h. Cells were grown in glass coverlips until colony formation and subjected to immunofluorescence analysis. The nucleus was stained with DAPI. Doxa: doxazosin





Fig. 5. HT-29 F1 cells, the group Irradiated + kyl were pretreated for 1 hr with 30 μ M of Kyl and submeted to irradiation. (A) Cells morphology of the HT-29 F1 cells analyzed using phase-contrast microscopy. Scale bar 50 μ m. (B) Western blot analysis of p-EphA4 (Tyr-602) and EphA4 protein levels. GAPDH protein was used as a loading control. Numbers below the figure represent the ratio of band's optical density as fold change of protein expression normalized by GAPDH.



Fig. 2. Western blot analysis of p-EphA4 (Tyr-602) and EphA4 protein levels in lysates derived from HT-29. The cells were pretreated for 1 hr with different concentrations of Kyl (5, 15, 30 and 60 μ M) and after the cells were treated with doxazosin (23 μ M) for 15 min. GAPDH protein was used as a loading control. Numbers below the figure represent the ratio of band's optical density as fold change of protein expression normalized by GAPDH. Data presented as the mean ± SEM of three independents experiments. Doxa: doxazosin

Fig. 6. Illustration of the signaling pathways induced in the progeny of colorectal cancer and the possible mechanism of action of Kyl. Ephrins as well as others factors can activated the EphA4 phosphorylation which leads to activation of downstream molecules such as ERK and the internalization of E-cadherin/EphA4 complex that culminates in migration and invasion (left). Kyl is a molecule that competitively inhibits Ephrin binding to EphA4 receptor and theoretically inhibits downstream pathways activated by EphA4 and allows the maintenance of cell-cell contacts through the retention of E-cadherin in the membrane (right).



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