

Cannabinoid Receptor Profiling Reveals a Low CB1 and High CB2 signature in Clear Cell Renal Cell Carcinoma

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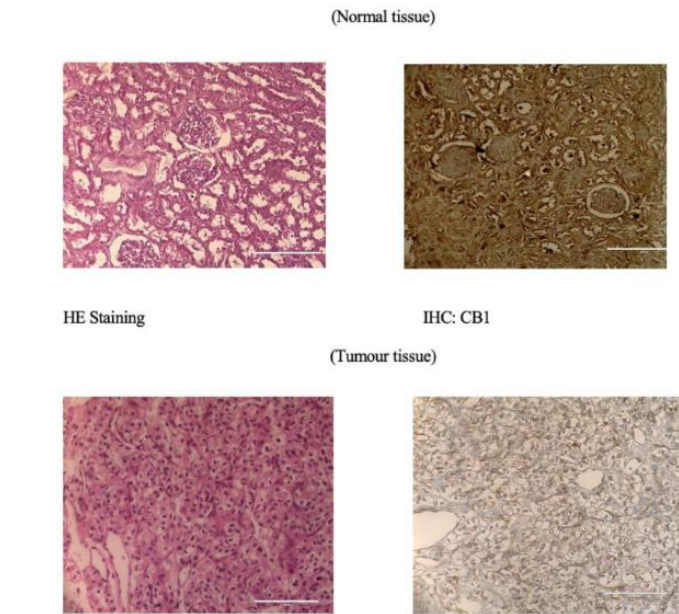
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Background: The endocannabinoid system has emerged as a potential regulator of tumour biology; however, the expression and clinical significance of cannabinoid receptors in clear cell renal cell carcinoma (ccRCC) remain poorly understood. We aimed to characterize the expression profile of cannabinoid receptor 1 (CB1) and cannabinoid receptor 2 (CB2) in ccRCC and evaluate their association with clinicopathological features and clinical outcomes.

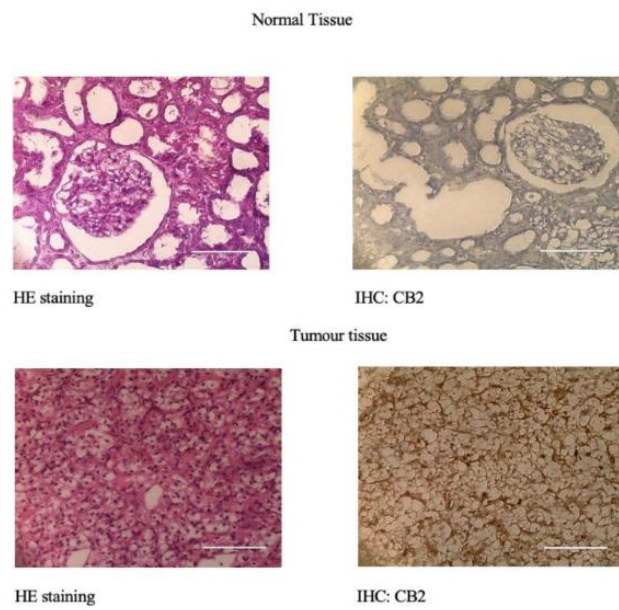
Methods: A total of 90 patients with histologically confirmed ccRCC were included, comprising a retrospective cohort (n=60) and a prospective cohort (n=30). Immunohistochemistry was performed to evaluate CB1 and CB2 expression in tumour and adjacent normal renal tissues using staining intensity, percentage positivity, and immune-reactive score (IRS). Protein and transcript-level validation was performed in the prospective cohort using Western blotting and quantitative real-time PCR. Associations with clinicopathological parameters and survival outcomes were analysed using Fisher's exact test and Kaplan-Meier survival analysis.

Results: Tumour tissues demonstrated a distinct cannabinoid receptor signature characterised by reduced CB1 expression and increased CB2 expression compared with adjacent normal kidney tissue (Figure 1). CB1 IRS was significantly lower in tumour samples ($P=0.001$), whereas CB2 IRS was significantly elevated ($P<0.001$). Western blot analysis confirmed reduced CB1 protein expression ($P=0.0466$) and increased CB2 protein expression ($P=0.0283$) in tumour tissues. At the transcriptional level, CB2 mRNA expression was significantly upregulated in tumours ($P=0.0070$), while CB1 mRNA expression showed no significant difference ($P=0.8830$), suggesting post-transcriptional regulation of CB1. High CB2 expression was associated with male sex ($P=0.010$) and higher tumour grade ($P=0.040$). No significant associations were observed between CB1 expression and clinicopathological variables. Neither receptor demonstrated independent prognostic significance for overall survival.

Conclusions: ccRCC exhibits a reproducible low CB1/high CB2 molecular phenotype. The selective overexpression of CB2 in tumour tissue highlights its potential as a biomarker and candidate therapeutic target in renal cell carcinoma.



a)



b)

Figure 1. Differential expression of cannabinoid receptors in clear cell RCC: a) CB1 expression. b) CB2 expression in normal kidney and tumour tissue.