CLINICAL IMPORTANCE OF GENETIC VARIANTS IN CAPECITABINE BIOACTIVATION PATHWAY FOR THE PREDICTION OF RESPONSE IN COLORECTAL CANCER PATIENTS

Y. CURA¹, C. PÉREZ-RAMÍREZ², A. SÁNCHEZ-MARTÍN¹, M.R. CANTUDO CUENCA¹, A. JIMENEZ-MORALES¹

¹UNIVERSITY HOSPITAL VIRGEN DE LAS NIEVES, PHARMACOGENETICS UNIT. HOSPITAL PHARMACY SERVICE, GRANADA, SPAIN. ²INSTITUTE OF NUTRITION AND FOOD TECHNOLOGY "JOSÉ MATAIX"- CENTER OF BIOMEDICAL RESEARCH- UNIVERSITY OF GRANADA, BIOCHEMISTRY AND MOLECULAR BIOLOGY II, GRANADA, SPAIN.





Colorectal cancer (CRC) is one of the most prevalent neoplasms worldwide. Capecitabine (Xeloda®), an oral prodrug of 5-Fluorouracil, is one of the standard treatments for patients with advanced CRC (Stages III-IV). In clinical practice, capecitabine response shows high interindividual variability. This variability may be due to the presence of polymorphisms in genes related to the bioactivation of capecitabine to fluorouracil (CES1, CES2, CDA, TYMP), that may alter drug bioavailability.

To assess treatment response and evaluate the influence of genetic polymorphisms in *CES1* (rs71647871, rs71647871), *CES1P1* (rs rs7187684, rs11861118), *CES2* (rs11075646), *CDA* (rs532545, rs602950, rs2072671), *TYMP* (rs11479) as predictive biomarkers in CRC patients treated with capecitabine.

MATERIAL AND METHODS



Prospective cohort study in CRC patients under adjuvant capecitabine treatment



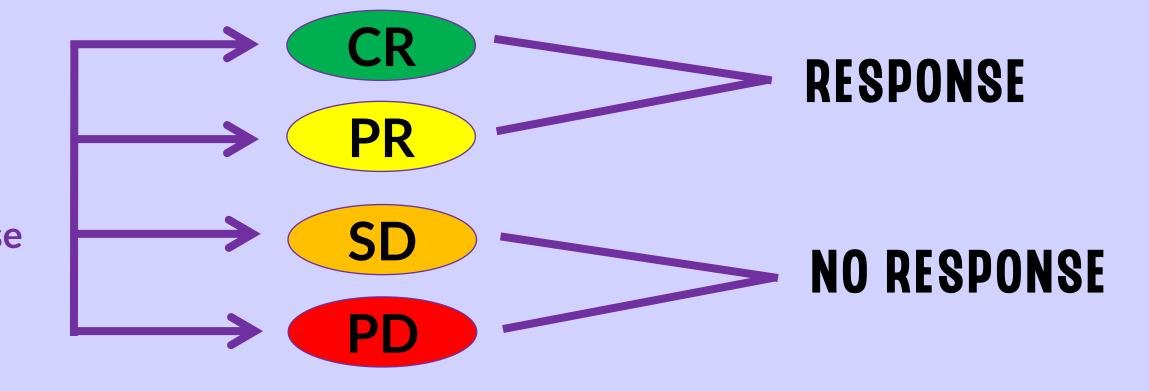
DNA extracted from buccal swabs

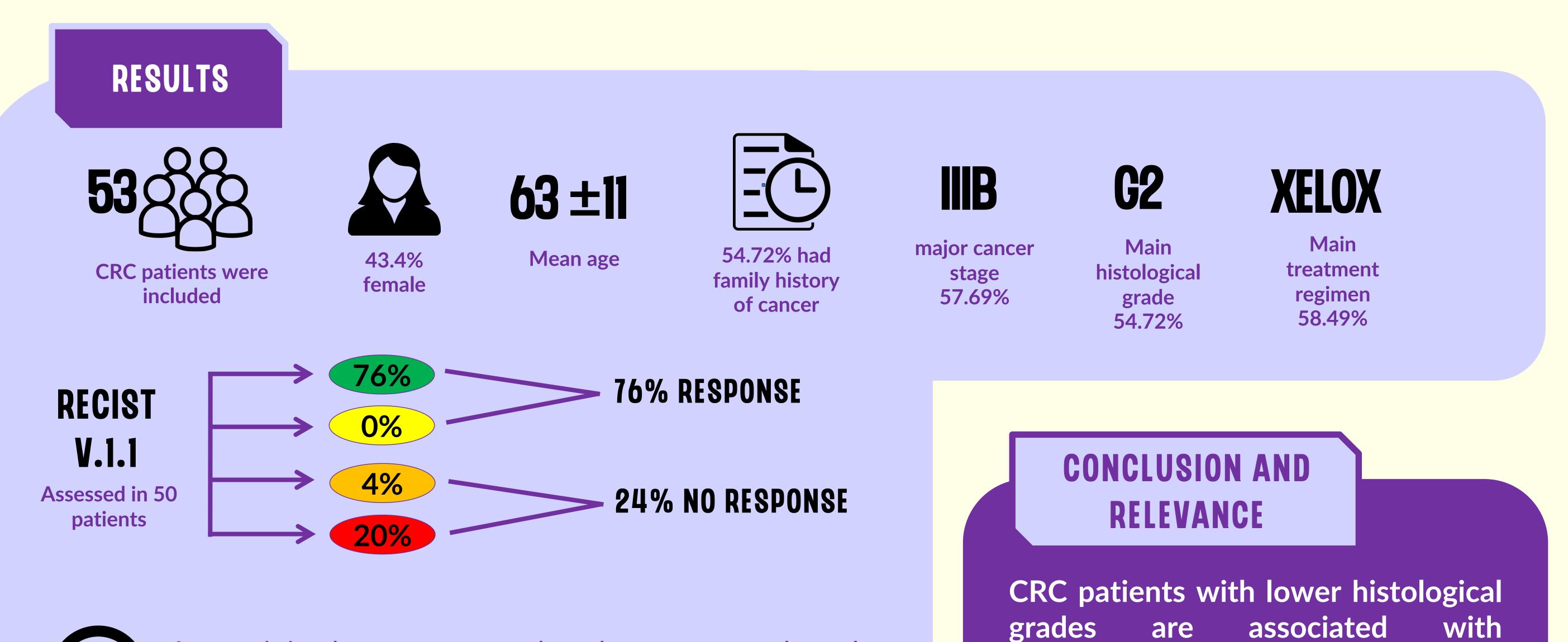


Genetic variants determined by RT-PCR with TaqMan® probes



Treatment response was assessed using RECIST criteria v1.1







An association between tumor grade and response was observed (p=0.03), OR = 2,71; Cl95% [1.82-189.39] for G1 vs G3 and OR = 2.17; Cl95% [1.35-78.39] for G2 vs G3.



No significant association was found between treatment response and the analyzed polymorphisms (p>0.05).

capecitabine positive response. No significant association was found between response and genetic variants in CES1, CES2, CDA, and TYMP.

Keywords: Colon cancer, Capecitabine, Bioactivation, Polymorphisms, Response







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