

## HUMAN GENOTYPING- CEGEN UNIT

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### OVERVIEW

In the Unit we implement high-throughput methods for detection of genetic variation (single nucleotide variants, indels, structural variants) and methylation analysis using DNA microarray and next-generation DNA sequencing technologies. Complementarily, research focused on identifying predictive biomarkers for precision medicine is undertaken.

**“Our aim is to identify predictive biomarkers in cancer patients in order to implement precision medicine in clinical practice.”**

#### • PUBLICATIONS

- Fachal L *et al.* (incl. Benítez J, González-Neira A) (2020). Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. *Nat Genet* 52, 56-73.
- Zhang H *et al.* (incl. Benítez J, González-Neira A) (2020). Genome-wide association study identifies 32 novel breast cancer susceptibility loci from overall and subtype-specific analyses. *Nat Genet* 52, 572-581.
- Escala-García M *et al.* (incl. Benítez J, González-Neira A) (2020). A network analysis to identify mediators of germline-driven differences in breast cancer prognosis. *Nat Commun* 11, 312.
- Peña-Chilet M *et al.* (incl. González-Neira A) (2020). CSVS, a crowdsourcing database of the Spanish population genetic variability. *Nucleic Acids Res.* PMID: 32990755.
- Kramer I *et al.* (incl. Benítez J, González-Neira A) (2020). Breast cancer polygenic risk score and contralateral breast cancer risk. *Am J Hum Genet* 107,837-848.
- Barnes DR *et al.* (incl. Benítez J, González-Neira A) (2020). Polygenic risk scores and breast and epithelial ovarian cancer risks for carriers of BRCA1 and BRCA2 pathogenic variants. *Genet Med* 22, 1653-1666.
- Núñez-Torres R, Martín M, García-Sáenz JA, Rodrigo-Faus M, Del Monte-Millán M, Tejera-Pérez H, Pita G, de la Torre-Montero JC, Pinilla K, Herráez B, Peiró-Chova L, Bermejo B, Lluch A, González-Neira A (2020). Association between ABCB1 genetic variants and persistent chemotherapy-induced alopecia in women with breast cancer. *JAMA Dermatol* 156, 987-991.
- Ruiz-Pinto S, Pita G, Martín M, Nuñez-Torres R, Cuadrado A, Shahbazi MN, Caronia D, Kojic A, Moreno LT, de la Torre-Montero JC, Lozano M, López-Fernández LA, Ribelles N, García-Saenz JA, Alba E, Milne RL, Losada A, Pérez-Moreno M, Benítez J, González-Neira A (2020). Regulatory CDH4 genetic variants associate with risk to develop capecitabine-induced hand-foot syndrome. *Clin Pharmacol Ther.* PMID: 32757270.
- Triviño JC *et al.* (incl. Pita G, González-Neira, Benítez J) (2020). Combination of phenotype and polygenic risk score in breast cancer risk evaluation in the Spanish population: a case-control study. *BMC Cancer* 20, 1079.
- Lin SH *et al.* (incl. González-Neira) (2020). Low-frequency variation near common germline susceptibility loci are associated with risk of Ewing sarcoma. *PLoS One* 15, e0237792.
- Feng H *et al.* (incl. Benítez J, González-Neira A, Osorio A) (2020). Transcriptome-wide association study of breast cancer risk by estrogen-receptor status. *Genet Epidemiol* 44, 442-468.

### RESEARCH HIGHLIGHTS

*Novel predictive genetic markers for adverse drug reactions in breast cancer (BC) patients.* Persistent chemotherapy-induced alopecia (pCIA) and capecitabine-induced hand-foot syndrome (CiHFS) are 2 common adverse drug reactions in cancer treatment. pCIA occurs in its most severe form in up to 10% of BC patients treated with docetaxel-based therapies, having a profound psychological impact on them. CiHFS is a dermatological toxicity affecting around 30% of patients, and the main cause of dose reductions and chemotherapy delays. By GWAS, we identified a regulatory variant associated with pCIA appearance in patients; this finding was validated in the replication cohort (ORcombined 4.05; 95% IQR, 2.46-6.67; P=3.946 x 10<sup>-8</sup>). This variant affects ABCB1 mRNA expression, being the risk allele associated with decreased expression. The ABCB1 gene encodes P-glycoprotein, an efflux pump responsible for the elimination of docetaxel, and lower expression could cause decreased drug elimination and thus its intracellular accumulation. Carriers of the risk allele would experience high drug exposure in the hair follicle and alopecia may become permanent, owing to the destruction of hair follicle stem cells. In addition, we discovered and replicated a cluster of 4 variants associated with decreased levels of CDH4 mRNA and the protein it encodes, R-cadherin, which localises in the granular layer of the epidermis. This resulted in reduced expression of involucrin, a protein of the cornified envelope, an essential structure for skin barrier function.

*Identifying variants of pharmacogenomic interest using CSVS, a crowdsourcing database of the Spanish population genetic variability.* Genetic differences between human populations are becoming increasingly recognised as important factors accounting for interindividual variations in drug responsiveness. Using data from the CSVS repository, we addressed how population-specific differences in genes involved in drug absorption, distribution, metabolism, excretion and toxicity (ADMET) could affect the rates and risks of drug inefficacy and/or adverse drug reactions in the Spanish population. We studied the Spanish genetic variability in a total of 421 pharmacogenes and, interestingly, a non-negligible percentage of private variation was observed in genes encoding proteins involved in drug metabolism, transport, and response.

*Detection of mutations in liquid biopsies from paediatric CNS tumours.* Paediatric CNS tumours are the most fatal cancer diseases in childhood. Due to their localisation and infiltrative nature, some tumour resections or biopsies are not feasible. We conducted the first study to compare different sources of liquid biopsies in paediatric cancers, an unmet need for clinical practice. We found serum to be more promising than plasma for BRAF V600E by dPCR detection in liquid biopsy of CNS paediatric cancers. ■