

Paraskevi Kosmidou, MSc

Scientific Technical Personnel

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Key skills Extensive practical experience in using laboratory equipment and computer software, ability to work both independently and cooperatively in a team environment and good communication skills

Education

1998-1999: Master of Science (MSc) in Medical Molecular Genetics, University of Aberdeen, UK

1995-1998: Bachelor of Science (BSc Honours) in Genetics, University of Wales, Cardiff, UK

Work experience

2006-present: Scientific Technical Personnel in Laboratory of Signal-Mediated Gene Expression, Institute of Chemical Biology, National Hellenic Research Foundation, Athens

Key techniques: DNA, RNA and protein extraction, PCR, RT, real-time PCR, NGS, cell culture, Western Blotting, Flow Cytometry, Confocal Microscopy

2001-2006: Research Assistant in Team 78 (Cancer Genome Project), Wellcome Trust Genome Campus, Hinxton, Cambridge, UK

Key techniques: Cell culture, DNA extraction, PCR, Sanger Sequencing, Sequencing analysis

2000-2001: Assistant Scientific Classifier in Medicines Control Agency, London, UK

Key area of the job: Data entry onto the Agency's Product License User System

Participation in EU research consortia:

- 2009-2013: "EpiDiaCan" Development of sensitive methodologies for exploitation of early epigenetic marker diagnosis in major types of cancer. FP7 EU-Cooperation – Theme "Health".
- 2006-2010: "Oncodeath" Resistant determinants and sensitization of solid tumour cells to death receptor related therapies: combination of TRAIL with other therapeutic molecules. EU-Combating Cancer.

Participation in National funded grants:

- 2018-2021: HNPM "Hellenic Network for Precision Medicine", a national network for precision oncology.
- 2017-2020: "STHENOS-b", Targeted therapeutic approaches against degenerative diseases with special focus on cancer and ageing-optimisation of the targeted bioactive molecules. National Strategic Reference Framework. "Competitiveness, entrepreneurship and innovation".
- 2013-2015: "STHENOS", Targeted therapeutic approaches against degenerative diseases, with special focus on cancer and ageing. National Strategic Reference Framework, Action "Developmental Projects of Research Organisations- Kripis".
- 2012-2015: "THERACAN", Exploiting molecular pathways of apoptotic cell death for the rational design of therapeutic strategies for colon cancer. National Strategic Reference Framework, Action "Co-operation II".
- 2010-2015: "POM", PIK3CA Oncogenic Mutations in Breast and Colon Cancers: Development of Targeted Anticancer Drugs and Diagnostics. National Strategic Reference Framework, Action "Co-operation".

Publications

1. Koumaki K et al. BRAF paradox breakers PLX8394, PLX7904 are more effective against BRAFV600E CRC cells compared with the BRAF inhibitor PLX4720 and shown by detailed pathway analysis. *Biochim Biophys Acta Mol Basis Dis*, 1867, (4): 166061. doi:10.1016/j.bbadis.2020.166061 (2021).
2. Kosmidou V et al. Noxa upregulation and a 5-gene apoptotic biomarker panel in colorectal cancer. *Eur J Clin Invest*, 51, (1):e13353. doi:10.1111/eci.13353 (2021).
3. Devetzi M et al. Death receptor 5 (DR5) and a 5-gene apoptotic biomarker panel with significant differential diagnostic potential in colorectal cancer. *Sci Rep*, 6: doi:10.1038/srep36532 (2016).
4. Kosmidou V et al. Tumor heterogeneity revealed by KRAS, BRAF and PIK3CA pyrosequencing: KRAS and PIK3CA intratumor mutation profile differences and their therapeutic implications. *Hum Mutat*, 35, 329-340 (2014).

5. Ferraro A et al. Epigenetic regulation of miR-21 in colorectal cancer: ITGB4 as a novel miR-21 target and a three-gene network (miR-21-ITGB4-PCDC4) as predictor of metastatic tumor potential. *Epigenetics* 9, 129-141 (2014).
6. Ferraro A et al. EZH2 is regulated by ERK/AKT and targets integrin alpha2 gene to control Epithelial-Mesenchymal Transition and anoikis in colon cancer cells. *Int J Biochem Cell Biol*, 45, 243-254 (2013).
7. Oikonomou E et al. TRAIL receptor upregulation and the implication of KRAS/BRAF mutations in human colon cancer tumours. *Int. J. Cancer*, 125, 2127-2135 (2009).
8. Hunter C et al. A hypermutation phenotype and somatic MSH6 mutations in recurrent human malignant gliomas after alkylator chemotherapy. *Can Res.*, 66, 3987-3991 (2006).
9. Bignell G et al. Sequence analysis of the protein kinase gene family in human testicular germ-cell tumours of adolescents and adults. *Genes, Chrom and Cancer*, 45, 42-46 (2006).
10. Ikediobi ON et al. Mutation analysis of 24 known cancer genes in the NCI-60 cell line set. *Mol Cancer Ther* 5(11):2606-12 (2006).
11. Davies H et al. Somatic mutations of the protein kinase gene family in human lung cancer. *Cancer Research*, 65(17), 7591-7595 (2005).
12. Stephens P et al. A screen of the complete protein kinase gene family reveals diverse patterns of somatic mutations in human breast cancer. *Nature Genetics*, 37, 590-592 (2005).
13. Stephens P et al. Intragenic ERBB2 kinase mutations in tumours. *Nature*, 431, 525-526 (2004).
14. Davies H et al. Mutations of the BRAF gene in human cancer. *Nature*, 417, 949-954 (2002).