## SEARCH FOR NOVEL COMMON VARIANTS INFLUENCING DIFFERENTIATED THYROID CANCER

<u>Gisella Figlioli</u><sup>1§</sup>, Aleksandra Köhler<sup>2§</sup>, Bowang Chen<sup>2</sup>, Stefano Landi<sup>1</sup>, Rossella Elisei<sup>3</sup>, Cristina Romei<sup>3</sup>, Monica Cipollini<sup>1</sup>, Alfonso Cristaudo<sup>3</sup>, Franco Bambi<sup>4</sup>, Per Hoffmann<sup>5,6,7</sup>, Stefan Herms<sup>5,6,7</sup>, Michał Kalemba<sup>8</sup>, Dorota Kula<sup>8</sup>, Shelley Harris<sup>9</sup>, Peter Broderick<sup>9</sup>, Richard Houlston<sup>9</sup>, Susana Pastor<sup>10,11</sup>, Ricard Marcos<sup>10,11</sup>, Antonia Velázquez<sup>10,11</sup>, Barbara Jarząb<sup>8</sup>, Kari Hemminki<sup>2,12</sup>, Asta Försti<sup>2,12\*</sup>, Federica Gemignani<sup>1\*</sup>

<sup>1</sup>Department of Biology, University of Pisa, Pisa, Italy; <sup>2</sup>Molecular Genetic Epidemiology, German Cancer Research Center (DKFZ), Heidelberg, Germany; <sup>3</sup>Department of Endocrinology and Metabolism, University of Pisa, Pisa, Italy; <sup>4</sup>Blood Centre, Azienda Ospedaliero Universitaria A. Meyer, Firenze, Italy; <sup>5</sup>Department of Genomics, Life and Brain Center, University of Bonn, Bonn, Germany; <sup>6</sup>Institute of Human Genetics, University of Bonn, Bonn, Germany; <sup>7</sup>Division of Medical Genetics, University Hospital Basel; Department of Biomedicine, University of Basel, CH-4058 Basel, Switzerland; <sup>8</sup>Department of Nuclear Medicine and Endocrine Oncology, Maria Sklodowska-Curie Memorial Cancer Center and Institute of Oncology, Gliwice Branch, 44-101 Gliwice, Poland; <sup>9</sup>Molecular and Population Genetics, Division of Genetics and Epidemiology, Institute of Cancer Research, Sutton, Surrey SM2 5NG, United Kingdom; <sup>10</sup>Grup de Mutagènesi, Departament de Genètica i de Microbiologia, Facultat de Biociències, Universitat Autònoma de Barcelona, 08193 Cerdanyola del Vallés, Barcelona, Spain; <sup>11</sup>CIBER Epidemiologia y Salud Pública, ISCIII, 28029 Madrid, Spain; <sup>12</sup>Primary Health Care Research, Clinical Research Center, Lund University, 205 02 Malmö, Sweden

§G.F. and A.K. contributed equally to this work

Thyroid cancer is a common endocrine malignancy with a rapidly increasing global incidence in the recent decades. Differentiated thyroid cancer (DTC), arising from follicular cells, includes the most common histological subtypes, papillary and follicular thyroid cancer, representing 80% and 15% of all thyroid cancers, respectively. Genome-wide association studies (GWASs) have identified robust associations with polymorphisms at 9q22.33 (*FOXE1*) and 14q13.2 (*NKX2-1*) and the disease. However, most of the inherited genetic risk factors of DTC remain to be discovered.

To search for new DTC risk variants we performed a GWAS in the high incidence Italian population and followed up the most significant associations in the lower incidence populations from Poland, UK and Spain. After excluding previously identified *loci*, the strongest association was observed for *DIRC3* at 2q35 ( $P=6.4\times10^{-10}$ ). Additionally promising associations were attained for *IMMP2L* at 7q31 ( $P=4.1\times10^{-6}$  and  $P=5.7\times10^{-6}$ ), *RARRES1* at 3q25.32 ( $P=4.6\times10^{-5}$ ) and *SNAPC4/CARD9* at 9q34.3 ( $P=3.5\times10^{-5}$ ).

Our findings provide insights into the genetic and biological basis of inherited genetic susceptibility to DTC. To further improve our knowledge on the disease, new *loci*, selected on the basis of association signals in our GWAS, will be analyzed.

<sup>\*</sup>A.F. and F.G. contributed equally to this work