

# Clinical and genetic characteristics of global cohort of 132 individuals with Janus kinase-3 deficiency

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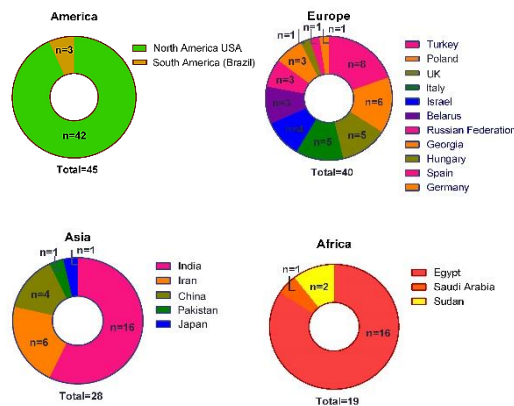
## Background and aims

Janus kinase-3(JAK3) deficiency, first described in 1995, is an autosomal recessive inborn error of immunity that mostly results in variants of severe combined immunodeficiency (SCID). The frequency is estimated to account for 7-14% of heritable SCID, with sporadic cases in the Western world. Neither preferential "hot-spots" nor founder effects have yet been documented. Hereby, we aim to describe the global experience of JAK3-related diseases regarding clinical spectrum, genetic landscape, including founder variants and treatment strategies.

## Materials and methods

We extracted clinical, genetic, immunological data from published cases on patients with CID/SCID phenotype caused by defects in the JAK3 gene. The literature search included unpublished cases from collaborators, reports from meetings of the ESID, of the CIS, published data in the PubMed from 1995 to 2024.

Our cohort includes 132 patients with 47 unique genetic defects, including 35 novel variants (18-homozygous, 17-heterozygous)



## Results

Forty five (35%) of patients from cohort were born to consanguineous parents in 31 families from Georgia (n=3,1-family), Russian Federation (n=3/2-family), Sudan (n=1), Turkey (n=8,5-family), Israel (n=4, 2 family), India (n=3, 2 family), Egypt (n=13, 8 family), UK (n=5, 4 family), China(n=1), Brazil(n=1), Spain(n=1), Italy (n=1), Pakistan(n=1), Saudi Arabia (n=1).

Title		JAK3(n=132)
Zygosity genetic defect	Homozygote	90
	Compound Heterozygote	40
	Germline heterozygous gain of function	2
Omenn syndrome		6
Treatment	HSCT	61
Outcome	Died	17
	Alive	44

## Conclusions

We describe for the first time a global cohort of JAK3 with 132 patients with founder effects in a subgroups. Patients were identified in four continents but is most common in countries with high rate of consanguinity. Founder effect was identified in 14 regions.

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## Background and aims

Severe combined immunodeficiency with Janus kinase 3 (*JAK3*) deficiency has immunologic phenotype T-B+NK-, frequency~10% of SCID cases.

## Materials and methods

We report on 10 patients (5 females and 5 males) from different ethnic groups with the SCID/CID phenotype (Fig1.).

In order to obtain a diagnosis, immunophenotyping of lymphocytes and molecular analysis using the Sanger sequencing and NGS were performed.

The *JAK3*-cohort included 10 patients from: the Republic of Belarus (2 - Eastern Slavs, 1- Arabs (Sudan origin), 1-Romany), the Russian Federation (3-Turkic), Georgia (3-Azerbaijani Georgians). (Fig1.). Consanguineous marriages were confirmed in 5 out of 7 families.

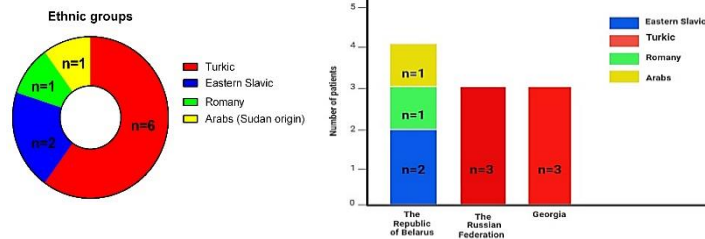


Fig. 1. The *JAK3*-cohort

## Results

Genetic variants in the *JAK3* gene were identified in all families

	Title	JAK3(n=10)
Immuno phenotype	T-B+NK-	8
	T-low B+NK-	1
	(CD3+CD8-)B+NK-	1
Type of mutation	Missense	2
	Frame shift	6
	Splicing site	2
Zygosity	Homozygote	8
	Compound	1
	Heterozygote	1
Pathogenicity (ACMG)	Pathogenic	2
	VUS	7
	Gain of function	1

1 patient underwent HSCT, 8 died at the age of 9(9-18) months, 2 are alive (1-preparing for HSCT).

## Conclusions

Our study expands the mutational spectrum of *JAK3* deficiency. Pathogenic variants in the *JAK3* gene leading to combined immunodeficiency are accompanied by heterogeneity of immunological phenotypes, which creates the need for detailed immunological and molecular studies, especially in closed ethnic groups.

