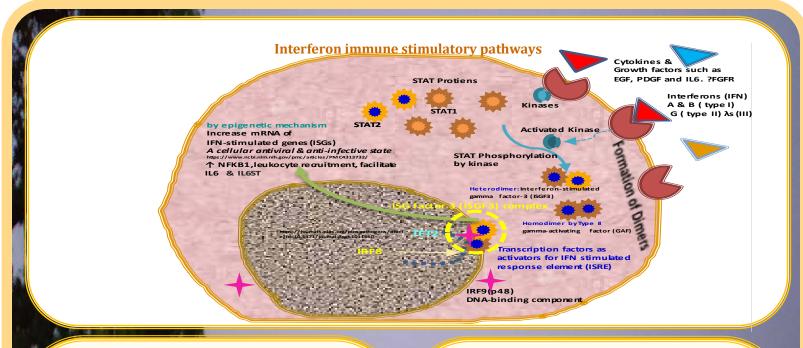


Insight:

- 1. What is the impact of recurrent inflammation on the innate immune system at the cellular level?
- 2. What are genes responsible for bronchiectasis related to immune regulatory pathways?
- 3. How do the interferon immune stimulatory pathways act at the cellular level?
- 4. What is the possible phenotype for case IV: 2?



Continue...

Genes related to immune regulation & responsible for bronchiectasis due to Recurrent Respiratory tract infections +/- Multisystem inflammatory disease

Gene/locatio n /MOI	Gene function	Phenotypes / Miscellaneous
STAT1/ 2q32.2/ AD, & AR IRF8/16q24.1	Signal transducer and transcription activator, for interferon-alpha/beta (IFNA/B) and interferon- gamma (IFNG) signalling pathways Regulate expression of genes	Immunodeficiency (ID) 31 A, 31 C (AD) 31 B (AR) Severe recurrent respiratory tract infections of the, including mycobacterial disease ID 32A (AD)- Mycobacteriosis
/AD & AR	stimulated by type I interferon	ID 32B (AR)- Recurrent viral disease
IGHM/14q32. 33/AR	The IGHM gene encodes the Constant region of immunoglobulin heavy chains	Agammaglobulinemia 1 - Infantile onset & neutropenia
TPP2/ <u>13q33.1</u> /AR	A dynamic supramolecular structure, proteasome like function, epitope generation for MHC class I antigen presentation	ID 78 with autoimmunity(Al) & developmental delay Ectodermal changes: Paronychia. Alopecia, dermo-hypodermitis, & dermatitis
IRF9/14q12/ AR	Mediates signalling by type I IFNs	ID 65 Pivot role in anti-viral immunity
NFKB1/4q24/ AD	NFKB is a transcription regulator as activators or repressors	ID, common variable (CV), 12 Type I diabetes mellitus, Pernicious anemia, Hepatitis, & Chronic enteropathy

Gene/locatio n /MOI	Gene function	Phenotypes / Miscellaneous
LRBA/4q31.3/ AR	Involved in intracellular signalling by regulates CTLA4 & post bacterial lipopolysaccharides (LPS) exposure	ID, CV, 8, with AI Early childhood onset, ITP, IBD, Asthma, & recurrent Conjunctivitis
TAP2/6p21.3 2/AR	Intricate in antigen presentation	Bare lymphocyte syndrome, type I Emphysema, Localized cutaneous necrobiosis lipoidica
IL6ST/5q11.2 /AD & AR	Initiating signal transmission for CNTF, IL11, IL6, LIF, CTF1, OSM and BSF3 as a part of the cytokine receptor complex	 I.ID 94 with autoinflammation and dysmorphic facies (AD) Hyper-IgE syndrome 4A, with recurrent infections (AD) Hyper-IgE syndrome 4B, with recurrent infections (AR) Stuve-Wiedemann syndrome 2 (AR) Macrocephaly, Prognathia, Hypertelorism, Downslanting palpebral fissures, Small nose Granulomatous interstitial lung disease, & organomegaly
LIG1/19q13.3 3/AR	DNA ligase, role in DNA replication, recombination, & the repair process	ID 96 Early childhood onset, sensitive to DNA- damaging agents
TET2/4q24/ AR	Activate the genes by DNA demethylation Convert methylcytosine to 5- hydroxymethylcytosine	ID 75 Myelodysplastic syndrome, somatic Reported with various myeloproliferative disorders

Possible phenotype for case IV: 2- Overall expected milder phenotype because of low penetrance, & have long term good prognosis.

Thought Riveting:

- What is the possible interaction of TPP2 & LRBA genes with IFN stimulated response element (ISRE)?
- How does IFN signaling act on CTLA4 (Cytotoxic T-Lymphocyte Associated Protein 4)?
- What is the predictable interaction of chloroquine with the LRBA gene in the pathological cellular stage?
- Does any overactivity of the IL-17 pathway with STAT1 loss of function mutations protect from candida infection?
- Do somatic mutations in the immune regulatory genes & their pathways lead to various autoimmune & inflammatory disorders?