The expanding spectrum of rare monogenic autoinflammatory diseases

Isabelle Touitou1,2,3*, Caroline Galeotti6, Linda Rossi-Semerano4, Véronique Hentgen5, Maryam Piram4, Isabelle Koné-Paut4 and for the CeRéMAI, French reference center for autoinflammatory diseases

Abstract

Monogenic autoinflammatory diseases are a group of hereditary disorders characterized by a clinical and biological inflammatory syndrome in which there is little or no evidence of autoimmunity. The discovery of the first causative gene in 1997 was rapidly followed by the identification of many others from the same group. The mutated proteins can be directly or indirectly involved in the regulation of inflammation. The available literature includes numerous reviews, which address the principle diseases, but we wanted to focus on the most recent rare syndromes. A comprehensive review is thus provided, including taxonomic, genetic, and epidemiological data, along with characteristics defining positive and differential diagnoses and treatment. We believe that this update will assist physicians in correctly naming their patient’s illness. This is an essential step for the effective and targeted management of an orphan disease.

Keywords: Autoinflammatory diseases, Histiocytosis and lymphadenopathy syndrome (H Syndrome), NLRP12 Associated Periodic Syndrome (NAPS12), Deficiency of Interleukin 1 Receptor Antagonist (DIRA), Deficiency of interleukin 36 receptor antagonist (DITRA), Pityriasis rubra pilaris (PRP), Disseminated Superficial Actinic Porokeratosis (DSAP), Autoinflammation LipoDystrophy and Dermatosis syndrome (ALDD)

Abstract in french

Les maladies autoinflammatoires monogéniques sont un groupe de pathologies héréditaires caractérisées par un syndrome inflammatoire clinique et biologique, dans lesquelles on ne retrouve pas ou peu de signe d’autoimmunité. La découverte du premier gène en 1997 a été rapidement suivie par l’identification de nombreux autres appartenant au même groupe. Les protéines mutées participent directement ou indirectement à la régulation de l’inflammation. De nombreuses revues balayant les principales entités sont disponibles dans la littérature, mais nous avons voulu mettre la lumière sur les formes rares les plus récentes. Nous proposons une revue exhaustive incluant des données taxonomiques, génétiques, épidémiologiques, ainsi que des éléments cliniques de certitude et différentiels, et de traitement. Nous espérons que cette mise à jour aidera le médecin à mettre un nom sur la maladie dont souffrent leurs patients atteints de maladie autoinflammatoire, une étape indispensable pour une prise en charge efficace et ciblée.

* Correspondence: isabelle.touitou@inserm.fr
1Laboratoire de génétique des maladies rares et auto-inflammatoires, CHRU de Montpellier, Montpellier, France
2INSERM U844, Montpellier, France
Full list of author information is available at the end of the article

© 2013 Touitou et al.; licensee BioMed Central Ltd. This is an open access article distributed under the terms of the Creative Commons Attribution License (http://creativecommons.org/licenses/by/2.0), which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited.
Introduction
Autoinflammatory diseases arise from disorders of the innate immune system [1]. The prototype disease is familial Mediterranean fever (FMF), which belongs to the group of hereditary recurrent fevers. Periods of inflammation are predominantly triggered following unnoticed proinflammatory signals. The levels of serum markers of inflammation such as C-reactive protein are always increased during attacks, however autoantibodies remain mostly undetectable. Since the discovery in 1997 of the gene responsible for FMF [2,3], dramatic insights have been gained through establishing the central role of the inflammasomes by unraveling their components and interacting cellular partners [4-6]. Subsequent to the identification of causal mutations in monogenic autoinflammatory diseases, several sequence variants in the causative genes were found to act as susceptibility factors in the more common multifactorial diseases such as rheumatoid arthritis or multiple sclerosis [7]. More recently, monogenic autoinflammatory disorders expanded to include the so far-called inflammatory skin diseases without systemic involvement. Enhanced knowledge relating to the critical physiological mechanisms altered in these diseases fuelled the development of novel, dramatically effective targeted biological drugs such as Interleukin-1 inhibitors [8]. As autoinflammatory diseases are rare, a national referral center nominated by the French ministry of health was set-up (Ce Ré MAI; [http://asso.orpha.net/CEREMAI/EN/index.html]) in order to improve care for patients. This referral center created Infevers [9] [fmf.igh.cnrs.fr/infevers/], a dedicated online database for autoinflammatory mutations [9].

This review focuses on monogenic autoinflammatory diseases identified within the last five years, including epidemiological and genetic data (Table 1), an exhaustive description of the corresponding disease name and synonyms where available, OMIM and Orphocode numbers, main clinical signs and therapeutic approaches (Table 2).

Rare recently recognized monogenic autoinflammatory diseases
The correct diagnosis of monogenic autoinflammatory diseases relies on the physicians’ awareness. Due to the rarity of these disorders, a very low number of cases are reported. For example, less than five patients with clear autoinflammatory features and causative mutations have been published for each of H syndrome [10-13], Autoinflammation and PLCG2-associated antibody deficiency and Immune Dysregulation syndrome (APLAID) [14], and RANBP-Type and C3HC4-Type Zinc Finger-Containing 1 (RBCK1)-related disease [15]. These three conditions are thus cited in Tables 1 and 2 only. Moreover, as mutations in these rare conditions have apparently systemic and pleiotropic effects, the patients’ clinical spectrum is heterogeneous and not yet completely delineated. Molecular analysis of the candidate genes may also fail to detect mutations. Consequently, the frequency of these diseases is unknown and too little feedback and data exist on the clinical significance and penetrance of the variants to ensure adequate genetic counseling for these patients.

NLRP12 Associated Periodic Syndrome (NAPS12)
This disease is characterized by recurrent periods of fever, combined with various systemic manifestations such as myalgia, arthralgia, headaches and skin urticarial-like rash although some patients may display arthralgia or myalgia only [16-18]. The most severely affected patients suffer from neurosensory signs (headache, deafness) [18]. The inflammatory attacks last 5-10 days and are triggered by physical exertion or cold exposure. These attacks can be very disabling in everyday life and can have a major impact on school attendance or work. The disease appears to be more active in children and teenagers than in adults.

Treatment is not standardized and no long-term medication has proven to be effective. Patients with a mild phenotype may be treated symptomatically with non-steroidal anti-inflammatory drugs. There is currently no therapeutic option available for patients with the most severe phenotype, and long-term treatment with interleukin (IL)-1 inhibitors (Anakinra, Kineret®) failed in the two cases tested [19].

The genetic defect of this disease was identified in 2008 [18] as a mutation in the gene encoding NLR pyrin domain-containing protein 12 (NLRP12) belonging to the intracellular Nod-like receptor (NLR) family. This protein is an intracellular sensor of the innate immune system, that regulates inflammatory processes through inhibition of nuclear factor kappa B (NF-kB) [18] and IL1B [16] signaling. NLRP12 mutations are likely responsible for a loss of protein function resulting in upregulation of this pro-inflammatory pathway. However, the exact role of NLRP12 remains unknown. It may participate in the formation of an NLRP3-independent inflammasome and be crucial for the recognition of intracellular pathogens. NLRP12 could also play an important role in the homeostasis of the gut by controlling the genesis of inflammatory diseases of the gastrointestinal tract and/or colorectal cancer [20,21]. Two unambiguous mutations, c.850C>T, p.Arg284*; c.2072+2dupT, p.Val635Thrfs*12 [18] as well as two missense substitutions c.1054C>T, p.Arg352Cys [16] and c.882C>G, p.Asp294Glu [17] were proven pathogenic through functional studies.

Deficiency of Interleukin 1 Receptor Antagonist (DIRA)
Sterile multifocal osteomyelitis with periostitis and pustulosis (OMPP) is more commonly referred to as
| Year of gene discovery | Disease* | N patients** | M/F ratio | Ethnicity*** | Ref | Mode of inheritance | Gene**** | Chromosome location | Most frequent mutation |
|------------------------|----------|--------------|-----------|--------------|-----|---------------------|----------|---------------------|------------------------|
| 2008                   | H Syndrome | Histiocytosis lymphadenopathy plus syndrome | 4 | 2/2 | 1 Indian case, 1 Pakistani case, 1 Moroccan case, 1 Tunisian case | Recessive | SLC29A3 | 10q22.1 | ND |
| 2008                   | NAPS12 | NLRP12 Associated Periodic Syndrome | 19 | 13/6 | 2 Guadeloupean families, 1 Italian family, 1 Armenian case, 1 Italian case, 1 Guadeloupean case, 1 Irish case | Dominant | NLRP12 | 19q43.42 | c.1054C>T, p.Arg352Cys |
| 2009                   | DIRA | Deficiency of Interleukin 1 Receptor Antagonist | 17 | 9/8 | 1 Turkish family, 1 Lebanese family, 2 Dutch families, 1 Dutch case, 1 Canadian case, 2 Puerto-Rican cases, 2 Brazilian cases, 2 Unknown, 3 Dutch, 1 Lebanese, 1 Turkish, 1 from unknown ancestry | Recessive | IL1RN | 2q13 | c.64_169del, p.IL1F9_IL1RNdel, c.213_227del, p.Asp72_Ile76del |
| Year | Disease | Description | Patients | Location(s) | Mutation Details |
|------|---------|-------------|----------|-------------|-----------------|
| 2011 | DITRA   | Deficiency of Interleukin 36 | 66       | 26/37       | Recessive IL36RN c.80C>T, p.Leu27Pro |
|      |         |             |          |             | 3 Tunisian families, 1 Algerian family |
|      |         | 36 Receptor Antagonist |         |             | 19 European cases, 10 Asian cases, 6 Tunisian cases, 6 German cases, 3 Japanese cases |
|      |         |             |          |             | 33,34 |
|      |         |             |          |             | 32,35,37 |
|      |         |             |          |             | c.338C>T, p.Ser113Leu |
|      |         |             |          |             | c.28C>T, p.Arg10* |
|      |         |             |          |             | 1 Algerian family, 10 Asian cases, 3 Japanese cases, 1 Tunisian case, 1 German case, 1 Algerian family |
|      |         |             |          |             | 6 Tunisian cases, 1 Japanese case, 3 Nigerian cases, 1 Spanish case, 1 Turkish case, 1 Iraqi case, 1 Russian case |
|      |         |             |          |             | 33,34 |
|      |         |             |          |             | 32,35,37 |
|      |         |             |          |             | c.338C>T, p.Ser113Leu |
|      |         |             |          |             | c.28C>T, p.Arg10* |
| 2012 | PSORS2  | Psoriasis susceptibility 2 | 45      | 24/21       | Dominant CARD14 c.390G>A, p.Gly117Ser |
|      |         |             |          |             | 1 European family, 1 Taiwanese family, 1 Tunisian family |
|      |         |             |          |             | 42 |
|      |         |             |          |             | 42 |
|      |         |             |          |             | 42 |
|      |         |             |          |             | c.390G>A, p.Gly117Ser |
|      |         |             |          |             | 1 Chinese family, 1 Chinese family |
|      |         |             |          |             | 42 |
|      |         |             |          |             | 42 |
|      |         |             |          |             | c.390G>A, p.Gly117Ser |
|      |         |             |          |             | 1 Chinese family, 1 Chinese family |
|      |         |             |          |             | 42 |
|      |         |             |          |             | 42 |
|      |         |             |          |             | c.390G>A, p.Gly117Ser |
| 2012 | DSAP    | Disseminated superficial Actinic Porokeratosis | >45     | 21/16/ND | Dominant MVK C.604G>A, p.Gly202Arg |
|      |         |             |          |             | 20 Chinese families, 6 Chinese cases |
|      |         |             |          |             | 48.49 |
|      |         |             |          |             | 12q24 |
|      |         |             |          |             | C.604G>A, p.Gly202Arg |
| 2010 | ALDD (JMP, NNS, CANDLE) | Autoinflammation, lipodystrophy, and dermatosis syndrome | 40      | 26/14       | Recessive PSMB8 c.224C>T, p.Thr75Met |
|      |         |             |          |             | 22 Japanese families |
|      |         |             |          |             | 53,57 |
|      |         |             |          |             | 6p21.32 |
|      |         |             |          |             | c.224C>T, p.Thr75Met |
|      |         |             |          |             | c.602G>T, p.Gly201Val |
|      |         |             |          |             | 56 | 1 Portuguese family, 1 Mexican family, 3 Hispanic cases, 3 Spanish cases, 2 American cases |
|      |         |             |          |             | 56 |
|      |         |             |          |             | 56 |
|      |         |             |          |             | 56 |
|      |         |             |          |             | 56 |
|      |         |             |          |             | 56 |
|      |         |             |          |             | 56 |
|      |         |             |          |             | 56 |
|      |         |             |          |             | 56 |
|      |         |             |          |             | 56 |
|      |         |             |          |             | 56 |
|      |         |             |          |             | 56 |
| Year | Disease Name | Number of Families | Family Origin | Mode of Inheritance | Gene | Chromosome Location | Notes |
|------|--------------|--------------------|---------------|---------------------|------|---------------------|-------|
| 2012 | APLAID       | 2                  | 1 family of unknown origin | Dominant | PLCG2 | 16q23.3 | ND |
| 2012 | NA           | 3                  | 1 French family 1 Italian case | Recessive | RBCK1 | 20p13 | ND |

* Commonly used disease names as defined in Table 2 ** Genetically confirmed. *** “Family” refers to multiplex families; “Case” refers to sporadic patients **** Approved HUGO. NA: Not assigned yet. ND: Not determined.
### Table 2 Clinical features of rare recently recognized monogenic autoinflammatory diseases

| OMIM   | ORPHA- CODE | Disease acronym | Disease extended name* | Synonyms | Age at onset | Key symptoms | Differential diagnosis | Treatment |
|--------|-------------|----------------|------------------------|----------|--------------|--------------|------------------------|-----------|
| 612373 | 168569      | HS y n d r o m e | Histiocytosis-lymphadenopathy plus syndrome | PHID | Infancy, Childhood | Urticaria, fever, myalgia, arthralgia | FHC, FCAS3 | NSAID (IL-1 and TNF blockades not effective) |
|        | 254707      |                |                        | FHC      |              |              | FCAS1, FCAS2            |           |
|        | 254712      |                |                        | SHML     |              |              | ALDD Other causes of insulin-dependent diabetes |           |
|        | 254723      |                |                        | Sinus Histiocytosis with Massive Lymphadenopathy |              |              |                         |           |
| 611762 | 247868      | FCAS2          | Familial Cold Autoinflammatory Syndrome 2 | NAP512, NLRP12 | Infancy | Urticaria, fever, myalgia, arthralgia | FHC, FCAS3 | NSAID (IL-1 blockades not effective) |
|        |              |                |                        | NLRP12-associated disorder | Childhood |              | FCAS1, FCAS3            |           |
|        |              |                |                        |          | Adulthood    |              | ALDD Other causes of insulin-dependent diabetes |           |
| 612852 | 210115      | OMPP           | Osteomyelitis, sterile Multifocal, with Periostitis and Pustulosis | DIRA | Neonatal | Neutrophilic pustular dermatosis, periostitis, aseptic multifocal osteomyelitis |          | PSORP Other dermatologic and infectious conditions |
|        |              |                |                        |          |              |              | OMPP Other autoinflammatory diseases |           |
| 614204 | 247353      | PSORP          | Pustular Psoriasis, Generalized | DITRA | Infancy, Childhood, Adulthood | Diffuse erythematous pustular rash, fever, malaise and diffuse pain, systemic inflammation |          | NSAID, Vitamin D3, Acitretin, TNF and IL-1 blockades |
|        |              |                |                        | GPP      |              |              | OMPP Other autoinflammatory diseases |           |
| 602723 | NA          | PSORS2         | Psoriasis susceptibility 2 |          | Variable | Round, well circumscribed erythematous plaques covered by a thick silver scale with a predilection for elbows, knees, scalp, lumbosacral and anogenital regions |          | Corticosteroids, calcineurin inhibitor, calcipotriene, emollients, keratolytic agents, ultraviolet light, retinoids, methotrexate, cyclosporine, anti-TNF agents |

Toutou et al. Orphanet Journal of Rare Diseases 2013, 8:162

http://www.ojrd.com/content/8/1/162
| OMIM 1 | OMIM 2 | Disease | Genes | Neutrophilic skin lesions, dysregulated cytokines, autoimmune phenotypes, inflammatory bowel disease, recurrent infections, neutrophilic skin lesions, IBD, PLAID |
|--------|--------|---------|-------|----------------------------------------------------------------------------------|
| 173200 | 2897   | PRP     | Pityriasis Rubra Pilars           | Neonatal Early childhood, Small keratotic follicular papules, disseminated salmon-colored scaly plaques surrounding islands of normal skin, diffuse red-orange palmoplantar keratoderma |
| 175900 | 79152  | POROK3  | Porokeratosis 3, Disseminated Superficial Actinic Type | Adult, UV sensitive, Epidermal cornification, round and brownish lesions |
| 256040 | 2615   | ALDD    | Autoinflammation, LipoDystrophy, and Dermatosis syndrome | Neonatal Fever, skin rash, panniculitis, lipoatrophy |
| 614878 | 324530 | APLAID  | Autoinflammation, antibody deficiency, and immune dysregulation syndrome | Neutrophilic skin lesions, IBD, recurrent sino pulmonary infections |

*Approved OMIM; NA Not assigned yet; ND: too few patients to delineate clear criteria. Commonly used disease names are in bold text.*

**Notes:**
- NSAID: nonsteroidal anti-inflammatory drugs.
- CINCA: Chronic, Infantile, Neurologic, Cutaneous and Articular syndrome, IBD: inflammatory bowel disease, PLAID: PLCg2-Associated antibody Deficiency.
DIRA [22]. DIRA is an early-onset auto-inflammatory disease that presents with neutrophilic pustular dermatosis, periostitis, aseptic multifocal osteomyelitis, and high acute-phase reactants. Patients develop pustular skin rashes, gastrointestinal reflux, and multifocal osteomyelitis during the neonatal period. One patient presented with early onset intrauterine disease and intrauterine fetal demise [23]. Massive systemic inflammatory syndrome resulting in patient death has been described [22]. Other manifestations include osteopenia, interstitial pneumonia often causing hypoxemia and dyspnea or localized ground-glass opacities and areas of atelectasis at chest computed tomography, vesicular stomatitis, mouth ulcers, ribs widening, periosteal reaction, joint swelling, cervical vertebral fusion, hepatosplenomegaly, thrombosis and vasculopathy [22,24-26]. Skin biopsies show neutrophilic epidermal and dermal infiltration, concomitant superficial folliculitis, with subcorneal localization of the pustules. Differential diagnosis includes other autoinflammatory conditions especially deficiency of the Interleukin 36 receptor antagonist (DITRA), dermatologic conditions (infantile pustular psoriasis, tinea, bacterial folliculitis, scabies, eosinophilic pustular folliculitis, erythema toxicum neonatorum, transient neonatal pustular melanosis…), and infectious diseases (arthritis, osteomyelitis).

Antibiotics and conventional disease-modifying anti-rheumatic drugs including steroids are of limited benefit, however blocking IL-1 signaling with anakinra dramatically improved clinical symptoms within days, resolving osteolytic lesions, normalizing acute-phase reactants, and permitting appropriate growth [22,24-29]. Patients with DIRA need to remain on lifelong IL-1 inhibitory therapy. Previous attempts at stopping or weaning anakinra have resulted in disease flares.

The disease is caused by mutations of the gene encoding the protein IL1RA. This protein is a natural antagonist of the potent proinflammatory interleukins-1 cytokines and mutations result in over activity of IL-1α as well as IL-1β is expected in case of IL-1RA deficiency. Two founder deletions were reported, one Puerto Rican: c.-64_1696del, p.IL1F9_IL1RNdel [22,24,27], and one Brazilian: c.213_227del, p.Asp72_Ile76del [25].

Deficiency of IL-36 receptor antagonist (DITRA)

Interleukin-36-receptor antagonist deficiency is an hereditary auto-inflammatory disease characterized by repeated flares of generalized pustular psoriasis (GPP). The flares include sudden diffuse pustular erythematous rash associated with high fever, general malaise, systemic inflammation and sometimes geographic tongue' and nail dystrophy [30]. Organ inflammation such as cholangitis can occur during the disease course. In most patients, symptoms develop during childhood though the age of onset can vary considerably. Neonatal onset has also been reported. Failure to thrive may occur. Elevated C-reactive protein serum levels and marked leukocytosis are constant during disease attacks. Differential diagnosis includes other causes of GPP, mainly DIRA [22], in cases where symptoms appear during the first weeks of life.

There is no established treatment. Topical and oral steroids, vitamin D3, acitretin and anti-TNF-α, are effective. Our group has recently successfully treated an infant with DITRA not responding to topical and oral steroids or to acitretin with anakinra [31].

IL-36 receptor antagonist (IL-36RA) is a natural antagonist of IL-36α, IL-36β and IL-36γ, three cytokines belonging to the IL-1 family. Mutations result in decreased IL-36RA antagonistic effects due to a reduction of both IL-36RA protein stability and affinity for its receptor, the consequence of which is uncontrolled inflammation [30,32-37]. The high resulting expression of IL-36R and of the three agonists in epithelial tissues like skin likely causes GPP, the main clinical feature in all DITRA patients. In a recent study, Setta-Kafetzii et al. identified IL36RN variant alleles in palmarplantar pustulosis and acrodermatitis continua of hallopeau, two forms of GPP genetically distinct from psoriasis vulgaris [35].

Among the most frequent genetic alterations are one Tunisian founder missense mutation: c.80C > T, p.Leu27Pro1; one European recurrent missense mutation: c.338C > T, p.Ser113Leu2; and one Japanese founder nonsense mutation: c.28C > T, p.Arg10*3.

Psoriasis susceptibility (PSORS)2 and pityriasis rubra pilaris (PRP)

Psoriasis is a common and chronic immune-mediated inflammatory disease of the skin affecting approximately 2% of individuals of European descent with a strong genetic susceptibility [38]. Classic lesions consist of round, well-circumscribed erythematous plaques covered by a thick silver scale with a predilection for elbows, knees, scalp, lumbosacral and anogenital regions. Fingernail plate involvement includes pitting and distal onycholysis. Inverse psoriasis (flexural distribution), guttate psoriasis, pustular or erythrodermic psoriasis are other clinical variants. Pityriasis rubra pilaris (PRP) is a chronic skin disorder of abnormal keratinization classified by Griffiths into five types based on the age of onset and the clinical features [39]. Familial PRP belongs to the type V group (atypical juvenile) and accounts for about 5% of all cases. This chronic skin disorder usually presents at birth or appears during early childhood. It is characterized by small keratotic follicular papules forming disseminated salmon-colored scaly plaques that surround islands of normal skin, and diffuse red-orange palmoplantar keratoderma. Lesions typically progress rostral to caudal. While nails can be dystrophic, pitting of nails is not a...
feature of this disease. Skin biopsies can be performed if the clinical features or the responses to treatment are unusual. Histopathological analysis shows psoriasiform hyperplasia presenting as acanthosis with elongation of the rete ridges, parakeratosis and mononuclear cell infiltrate. Compared to psoriasis, the epidermal hyperplasia in PRP is more irregular, the granular layer is preserved, follicular plugging is characteristic and neutrophils in the stratum corneum are not essential. Differential diagnosis of psoriasis includes other papulosquamous disorders. PRP shows clinical features similar to phrynoderma (vitamin A deficiency), psoriasis, erythrodermatodermia and other causes of cornification.

Management of psoriasis includes patient education, topical therapy (corticosteroids, calcineurin inhibitor, calcipotriene, emollients, keratolytic agents), ultraviolet light, and systemic therapy (retinoids, methotrexate, cyclosporine and anti-TNF agents). The treatment of PRP is often disappointing. Milder cases of the disease can be treated with emollients, topical corticosteroids, tazarotene, keratolytic agents or calcineurin inhibitor whereas systemic retinoids and/or TNFs blocking agents are required in severe cases.

The prognosis of both disorders is that of most chronic diseases, with spontaneous remissions and exacerbations. Arthritis is present in about one fourth of psoriasis patients, and there is an increased risk of metabolic syndromes and cardiovascular diseases.

Both phenotypes have been associated to CASpase Recruitment Domain-containing protein 14 (CARD14) mutations [36,41-43]. CARD14 is an epidermal regulator of NF-$\kappa$B transcription factor. Mutations in CARD14 initiate a process that includes inflammatory cell recruitment by keratinocytes and perpetuate a vicious cycle of epidermal inflammation and regeneration that cause the abnormal keratinization. More specifically, c.349G > A, p.Gly117Ser (the most prevalent mutation), and c.413A > C, p.Glu138Ala (a de novo mutation) were shown to lead to enhanced NF-$\kappa$B activation and upregulation of a subset of psoriasis-associated genes in keratinocytes [43].

Disseminated Superficial Actinic Porokeratosis (DSAP)
This disease described in 1966 by Chernosky [44] represents the most common form of porokeratosis, a group of diseases characterized by epidermal cornification with epidermal cell hyperproliferation and premature apoptosis of keratinocytes. Triggers include ultraviolet light exposure and immunosuppression. Multiple small (0.5-1 cm) round and brownish lesions surrounded by a hyperkeratotic rim develop on sun-exposed areas of the skin particularly on the lower legs and forearms. Lesions usually appear in summer and improve or disappear during winter. The disease usually starts during the third or fourth decades of life, and rarely affects children. While it is usually benign, squamous cell carcinoma or Bowen's disease may occasionally develop within the lesions.

Histopathological examination of the skin sampled from the edge of the peripheral rim characteristically shows an angulated cornoid lamella and a parakeratotic column overlying an area of epidermis with an absent or reduced granular layer [45]. Differential diagnosis includes neoplastic or hyperplastic squamous skin lesions.

Cryotherapy, topical reagents (retinoids, imiquimod, 5-fluorouracil, keratolytic), electrodessication, laser ablation or photodynamic therapy are used to destroy the abnormal keratinocyte clones [46].

The mode of inheritance is dominant however many sporadic cases of porokeratosis have been described [47]. Mutations in the MeValonate Kinase (MVK) gene were found in DSAP patients [48,49]. The MVK enzyme is involved in the biosynthesis of cholesterol and isoprenoid. It is also thought to play a role in regulating calcium-induced keratinocyte differentiation and could protect keratinocytes from apoptosis induced by type A ultraviolet radiation. All genetically confirmed patients are of Chinese origin. Fourteen different mutations have been found, among which, 10 accounted for one third of the familial cases. Mutations in this gene were initially demonstrated as responsible for recessive complete (mevalonic aciduria [50], OMIM 610377) and incomplete (hyperIgD syndrome [51,52], OMIM 260920) mevalonate kinase deficiency.

Autoinflammation, LipoDystrophy, and Dermatosis syndrome (ALDD)
ALDD is a systemic inflammatory condition starting within the first months of life, and comprising elevated fever, panniculitis with lipatrophy, purplish and swollen eyelids, arthralgia, developmental retardation, and increased acute phase reactants [53,54]. Key symptoms include a persistent fever of >38.5°C, steroid-sensitive erythema nodosum-like (edematous and purpuric) plaques, long clubbed fingers, hyperhidrosis, myositis, hepatosplenomegaly, macroglossia, facial and limbs lipatrophy, and developmental (height, weight and IQ) retardation. Skin biopsies show immature myeloid-lineage cells with mitoses. Some patients may have joint contracture, auricular and nasal chondritis, and calcification of the basal ganglia. Acute cardiovascular event is the leading cause of death in these patients for whom life expectancy is notably reduced. Differential diagnosis includes other autoinflammatory conditions, in particular Sweet's syndrome, Still's disease and Chronic, Infantile, Neurological, Cutaneous and Articular (CINCA) syndrome, storage diseases (mucopolysaccharidoses), lupus, dermatomyositis, laminopathies (progeroid syndromes), and other interferonopathies like Aicardi Goutieres syndrome, an interferon type I related disease. Biological findings include microcytic or normocytic
anemia, thrombopenia (rare), mild increase in levels of amino transferase enzymes, perturbation of glucose and lipid metabolism (hyperglycemia and glucose intolerance, increased cholesterol and triglycerides), increased erythrocyte sedimentation rate and γ globulins levels, and some markers of autoimmunity (antinuclear and anti-neutrophilic cytoplasmic auto-antibodies, lupus anticoagulant, positive Coombs test).

Management of these patients is by palliative care. The need for steroids is very high even in combination with anti-IL-1, anti-IL-6 or anti-TNF treatments. Drugs targeting interferon γ are warranted. JAK kinase inhibitors are currently under clinical trials at the National Institute of Health, Bethesda, USA [55].

Most (N = 28) of the reported patients carried homozygous mutations in the Proteasome Subunit, Beta-Type, 8 (PSMB8) gene [53,54,56,57]. This gene encodes β5i, a catalytic subunit of the immunoproteasome, an intracellular protease complex specialized for degradation of polyubiquitinated proteins. This ubiquitin-proteasome system degrades unnecessary proteins and regulates the cell cycle, gene repair and nuclear factor NF-κB activation [53]. The mutated protein impairs the assembly of this immunoproteasome. One nonsense (c.405C > A, p.Cys135*) [54] and five missense mutations of which two are recurrent (c.224C > T, p.Thr75Met and c.602G > T, p.Gly201Val) have been reported [9]. As the prognosis is so severe, prenatal diagnosis can be discussed.

Conclusions

Hereditary recurrent fevers, as well as Majeed syndrome, Blau syndrome, and Pyogenic sterile Arthritis, Pyoderma gangrenosum and Acne (PAPA) syndrome, have been extensively reviewed since the concept of autoinflammation was created in 1999 [58]. However, very little information has been assembled regarding the phenotype, genetics and epidemiology of the more recently discovered autoinflammatory disorders. Here in this review, we aimed to offer novel diagnostic options to those patients for whom a genetic link could not be established. More than 25 autoinflammatory genes were listed during the last international conference on autoinflammation, Lausanne, 23-26 May, 2013, including at least five to six new ones in the pipeline. Both phenotype and genotype heterogeneity have been described which further hinders correct diagnosis and highlights the necessity for a gene and disease nomenclature consensus. A systematic in-depth description of the clinical features of these patients has been undertaken thanks to the Eurofever initiative [59,60]. Massively parallel sequencing approaches should help in identifying the causative gene among those already known or novel genes through whole exome or genome sequencing. Development of such innovative genetic diagnostic tools is in progress through an International network [I Touitou, personal communication].

The correct diagnosis of these diseases is essential for the relevant management of patients affected by rare autoinflammatory diseases as remarkably effective therapies have been developed recently which have the potential to avoid the risk of fatal complications such as amyloidosis, or sensorineural impairment. Rarely emphasized yet equally important is being able, even after many years as in some cases, to name the patient’s disease, which can be extremely useful from a psychological point of view.

Competing interests

The authors declare that they have no competing interests.

Authors’ contributions

IT and IKP conceived and designed the manuscript. IKP coordinated and assigned the writing of the different disease items: CG, DIRA; LC, DIISRA; JKP, ALDD; VH: NALP12, MP: DSAP and PRP. IT collected the data, assembled the first draft, and polished the manuscript. All authors read the final manuscript and gave approval for the version to be published. All authors read and approved the final manuscript.

Authors’ information

All authors are members of the French reference centre for autoinflammatory diseases (CeRéMAI, [http://asso.orpha.net/CEREMAI/EN/indexhtml]) coordinated by IKP. IT developed infeners [http://fmfligh.cnrs.fr/ISSAID/infeners/], an online registry devoted to autoinflammatory disease mutations.

Acknowledgements

This work was supported by the French Ministry of Health, and the Kremlin Bicêtre, Versailles and Montpellier hospitals. We thank Angloscribe for editing the manuscript for English language.

Author details

1Laboratoire de génétique des maladies rares et auto-inflammatoires, CHRU de Montpellier, Montpellier, France. 2INSERM U844, Montpellier, France.
3Université Montpellier 1, UM1, Montpellier, France. 4Service de pédiatrie générale et rhumatologie pédiatrique, APHP, CHU de Bicêtre, Université de Paris Sud, Le Kremlin-Bicêtre, France.
5Service de pédiatrie, CH de Versailles, Versailles, France.

Received: 16 August 2013 Accepted: 5 October 2013
Published: 16 October 2013

References

1. Kastner DL, Akersentjevich I, Goldbach-Mansky R: Autoinflammatory disease reloaded: a clinical perspective. Cell 2010, 140:784–790.
2. Consortium TFF: A candidate gene for familial Mediterranean fever. Nat Genet 1997, 17:25–31.
3. TheInternationalMFCConsortium: Ancient missense mutations in a new member of the Ro/RO gene family are likely to cause familial Mediterranean fever. Cell 1997, 90:797–807.
4. Conforti-Andreoni C, Riccardi-Castagnoli P, Montefaro A: The inflammasomes in health and disease: from genetics to molecular mechanisms of autoinflammation and beyond. Cell Mol Immunol 2011, 8:135–145.
5. Schroder K, Tschopp J: The inflammasomes. Cell 2010, 140:821–832.
6. Jeru I, Amselem S: Inflammasomes and interleukin 1. Rev Med Interne 2011, 32(4):218–224. doi:10.1016/j.revmed.2010.02.013. Epub 2010 Jun 11. French.
7. Touitou I: Inheritance of autoinflammatory diseases: shifting paradigms and nomenclature. J Med Genet 2013, 50:349–359.
8. Ter Haar N, Lachmann H, Czen S, Woo P, Uzel Y, Modesto C, Kone-Paut I, Cantarini L, Insalaco A, Neven B, et al.: Treatment of autoinflammatory diseases: results from the Eurofever Registry and a literature review. Ann Rheum Dis 2013, 72(5):678–685. doi:10.1136/annrheumdis-2011-201268. Epub 2012 Jun 29.
25. Jesus AA, Osman M, Silva CA, Kim PW, Pham TH, Gadina M, Yang B, Bertola et al. Inflammatory bowel disease. In: Slanetz BJ, ed. Inflammatory bowel disease. 2013. 39: 311–319.

26. Altiok E, Aksoy F, Perk Y, Taylan F, Kim PW, Ilikkan B, Asal GT, Goldbach-Mansky R. Noncanonical NF-kappaB signaling. In: Slanetz BJ, ed. Inflammatory bowel disease. 2013. 39: 320–329.

27. Reddy S, Aja S, Geoffrey R, Lorier R, Suchi M, Broeckel U, Hessner MJ, Verbsky et al. Autoinflammatory disease due to homozygous deletion of the IL1RN locus. N Engl J Med 2009, 360:2426–2437.

28. Chen Y, Xiao Y, Shi H, Li Q, Wang F, Hu Y, et al. Variations in IL36RN in Severe Adverse Drug Reactions Manifesting as Acute Generalized Exanthematous Pustulosis. J Invest Dermatol 2013, 133:1904–1907.

29. Griffiths WA. Pityriasis rubra pilaris: the problem of its classification. J Am Acad Dermatol 1992, 26:140–142.

30. ToCi LC, Spain SL, Knight J, Ellinghaus E, Stuart PE, Caron F, Ding J, Li Y, Tiejavs I, Gudjonsson JE et al. Identification of fifteen new psoriasis susceptibility loci highlights the role of innate immunity. Nat Genet 2012, 44:1341–1348.

31. Ammar M, Bouchka-Cousins S, Zaraa I, Helms C, Doss N, Bolz F, Dhaoui SF, Bezares et al. Familial pityriasis rubra pilaris is caused by mutations in CARD14. J Investig Dermatol 2000, 115:163–170.

32. Jordan CT, Cao L, Roberson ED, Pierson KC, Yang CF, Joyce CE, Ryan C, Duan S, Helms CA, Liu Y, et al. PSORS2 is due to mutations in CARD14. J Am Acad Dermatol 2012, 67:352–359.

33. Chenosky ME. Porokeratosis: report of twelve patients with multiple superficial lesions. South Med J 1966, 59:289–294.

34. Rouhani P, Fischer M, Mehegan S, Pomeranz MK. Disseminated superficial actinic porokeratosis. Dermatol Online J 2012, 18:24.

35. Skupsky H, Skupsky J, Goldenberg D. Disseminated superficial actinic porokeratosis: a review. J Dermatolog Treat 2012, 23:52–56.

36. Kaur S, Thami GP, Mohan H, Kanwar AJ. Coexistence of variants of porokeratosis: a case report and a review of the literature. J Dermatolog Treat 2002, 23:305–309.
48. Zhou Y, Liu J, Fu X, Yu Y, Shi B, Yu G, Shi Z, Wu W, Pan F, Tian H, et al: Identification of three novel frameshift mutations of the MVK gene in four Chinese families with disseminated superficial actinic porokeratosis. Br J Dermatol 2013, 169:193–195.

49. Zhang SQ, Jiang T, Li M, Zhang X, Ren YQ, Wei SC, Sun LD, Cheng H, Li Y, Yin XY, et al: Exome sequencing identifies MVK mutations in disseminated superficial actinic porokeratosis. Nat Genet 2012, 44:1156–1160.

50. Schafer BL, Bishop RW, Kratunis VJ, Kalinowski SS, Mosley ST, Gibson KM, Tanaka RD: Molecular cloning of human mevalonate kinase and identification of a missense mutation in the genetic disease mevalonic aciduria. J Biol Chem 1992, 267:13229–13238.

51. Drentth JP, Cuisser L, Grateau G, Vasseur C, van de Velde-Visser SD, De Jong JG, Beckmann JS, van der Meer JW, Delpech M: Mutations in the gene encoding mevalonate kinase cause hyper-IgD and periodic fever syndrome. International Hyper-IgD Study Group. Nat Genet 1999, 22:178–181.

52. Houten SM, Kuis W, Duran M, De Koning TJ, Van Royen-Kerkhof A, Romeijn GJ, Frenkel J, Dorland MM, Huijbers WA, et al: Mutations in MVK, encoding mevalonate kinase, cause hyperimmunoglobulinaemia D and periodic fever syndrome. Nat Genet 1999, 22:175–177.

53. Arima K, Kinoshita A, Mishima H, Kanazawa N, Kaneko T, Mizushima T, Ichinose K, Nakamura H, Tsujino A, Kawakami A, et al: Proteasome assembly defect due to a proteasome subunit beta type 8 (PSMB8) mutation causes the autoinflammatory disorder, Nakajo-Nishimura syndrome. Proc Natl Acad Sci U S A 2011, 108:14914–14919.

54. Liu Y, Ramot Y, Torrelo A, Paller AS, Si N, Babay S, Kim PW, Sheikh A, Lee CC, Chen Y, et al: Mutations in proteasome subunit beta type 8 cause chronic atypical neutrophilic dermatosis with lipodystrophy and elevated temperature with evidence of genetic and phenotypic heterogeneity. Arthritis Rheum 2012, 64:895–907.

55. Compassionate Use Treatment Protocol IV-MC-JAGA: Treatment of Autoinflammatory Syndromes Expected to Benefit from JAK Inhibition. http://clinicalstudies.info.nih.gov/cgi/wais/bold032001.pl?A_12-AR-8001.html@CANDLE@@@@.

56. Agarwal AK, Xing C, DeMartino GN, Mizrachi D, Hernandez MD, Sousa AB, Martinez DeVillarreal L, DosSantos HG, Garg A: PSMB8 encoding the beta5i proteasome subunit is mutated in joint contractures, muscle atrophy, microcytic anemia, and panniculitis-induced lipodystrophy syndrome. Am J Hum Genet 2010, 87:866–872.

57. Kanazawa N: Nakajo-Nishimura syndrome: an autoinflammatory disorder showing pernio-like rashes and progressive partial lipodystrophy. Allergol Int 2012, 61:197–206.

Submit your next manuscript to BioMed Central and take full advantage of:

- Convenient online submission
- Thorough peer review
- No space constraints or color figure charges
- Immediate publication on acceptance
- Inclusion in PubMed, CAS, Scopus and Google Scholar
- Research which is freely available for redistribution

Submit your manuscript at www.biomedcentral.com/submit