INVESTIGATIVE REPORT

Genotype–Phenotype Correlations in a Prospective Cohort Study of Paediatric Plaque Psoriasis: Lack of Correlation Between HLA-C*06 and Family History of Psoriasis

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This study aims to investigate associations between observed clinical parameters and known genetic risk factors of psoriasis in a well-defined prospective cohort of paediatric patients with plaque psoriasis (n = 151). Significant associations were found for paediatric-onset psoriasis with ERAP1 (p = 0.002), IL23R (p = 0.01), LCE3C, LCE3B-de (p = 0.00049) and HLA-C*06 (p = 3.15 × 10−30). Psoriasis severity was associated with the single nucleotide polymorphisms tagging IFIH1 and ERAP1 (p < 0.05). An onset before 10 years of age was associated with IL12B (p = 0.02). Nail psoriasis was more often seen in HLA-C*06-negative patients (p = 0.008). Remarkably, family history is clearly not associated with HLA-C*06 in this specific group. The large proportion of patients with a positive family history in HLA-C*06 negative patients (and the lack of correlation between the two) indicates that other genes, either alone or interaction between two or more genes, may have significant effects on heritability. Key words: genotype-phenotype correlations; paediatric psoriasis; genetic risk factors; clinical parameters.

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Psoriasis is a clinically heterogeneous skin disease with a complex genetic background (1–3). It affects around 2% of the population and in approximately 30% of these patients the disease first appears during childhood (1, 4). By means of genome-wide association studies, the number of genome regions identified to be associated with psoriasis increased recently to 36 (5). Different pathways evidently play a role in the pathophysiology of psoriasis, as the innate and adaptive immune system, the Th17 pathway and the skin barrier function are genetically linked with psoriasis (2, 3, 6–13). Some studies have investigated correlations between genetic risk factors of psoriasis and clinical parameters, and these were mainly focused on major histocompatibility (MHC) gene HLA-C*06 (14–23). In a cohort of adult psoriasis patients a strong association was demonstrated between psoriasis severity and single nucleotide polymorphisms (SNPs) tagging HLA-C*06 and a deletion of LCE3B and LCE3C (22, 34). Several studies have shown that early age at onset may be associated with distinct genetic factors, such as ERAP1 and HLA-C*06 (14, 17, 23, 24). Previous studies detected an association between HLA-C*06 and early onset psoriasis (onset < 30 or 40 years), and a positive family history of psoriasis and early onset of psoriasis (14, 16, 17, 20), suggesting an association between HLA-C*06 and family history of psoriasis in patients with early onset psoriasis. This association has however, never been investigated. Data about associations between clinical parameters and genetic risk factors in paediatric psoriasis are lacking. The current study aims to investigate associations between observed clinical parameters and known genetic risk factors of psoriasis in a well-defined cohort of paediatric patients with plaque psoriasis.

MATERIALS AND METHODS

Sample collection
DNA samples were obtained from children with psoriasis referred to the outpatient clinic of the Department of Dermatology of the Radboud University Medical Center. Only patients of European descent with a primary diagnosis of plaque-type psoriasis, before the age of 18 years, were included in this study. Patients with guttate psoriasis were excluded. The phenotype classification is based on phenotype at examination and not at onset of disease. We obtained clinical data from our prospective observational paediatric psoriasis registry in daily clinical practice, called Child-CAPTURE (Continuous Assessment of Psoriasis Treatment Use Registry). In this registry, patient characteristics including age at onset, family history of psoriasis (up to the third-degree relative) and Koebner phenomenon are recorded at the first visit. Psoriasis severity and presence of observed nail psoriasis by the physician are recorded every visit.

The characteristics of the control group of European descent were previously described (24). Only self-reported data were available for all controls, and individuals reported to have psoriasis were excluded from this study. All participants/parents gave written informed consent.

Genotyping
Genotyping was executed as previously described (24). In short, SNP genotyping was performed using Taqman® SNP
typing 5% of the samples were analysed in duplicate; all ge-
homozygotes and heterozygotes. As quality control on the SNP
(2), which does not allow distinction between
primers for
C*06
tagging SNP) was determined by PCR with sequence specific
previous described PCR (2). itself (rather than a
HLA-C*06:02
power to detect an association
between paediatric-onset psoriasis and healthy
control study. Age at onset was analysed comparing 2 groups;
before 10 years and at or after 10 years. Clinical severity of
psoriasis was assessed by a clinician using 3 different severity
scales: Psoriasis Area and Severity Index (PASI; range 0–72),
Physician Global Assessment (PGA; range 0–5) and Body
Surface Area (BSA; range 0–100) (25). The highest reported
psoriasis severity scores for each individual were used. Patients
were divided in 2 groups based on psoriasis severity scores;
mild-to-moderate and severe psoriasis. Mild-to-moderate pso-
riasis was defined as those patients that have never reached
PASI ≥ 10, PGA ≥ 3 or BSA ≥ 10. Severe psoriasis was defined
as those patients that have ever reached PASI ≥ 10, PGA ≥ 3 or
BSA ≥ 10. Nail psoriasis was scored by using the Nail Psoriasis
Severity Index (NAPSI; range 0–80) (26). Koebner phenomenon
was assessed based on a 4-point scale, how often a psoriasis
plaque appeared after skin damage of their non-involved skin:
never, rarely, often, or very common. The individuals who re-
sponded with “often” or “very common” were considered as
Koebner-positive patients and the others as Koebner-negative
patients, which is in line with our previous study (27). To test
for the association with psoriasis clinical variables, chi-square
tests were executed from the allele frequency table (2 × 2 tables).
Logistic regression analyses were performed to calculate ORs
and 95% CIs. The level of significance was considered to be
0.05. Power calculations were performed using the Genetic
Power calculator (28).

RESULTS
Cohort characteristics
The investigated cohort for the replication of the asso-
ciations between paediatric-onset psoriasis and healthy
controls consisted of 151 cases and 450 controls, which
is a doubling of the number of cases compared to our
previous study (24). The power to detect an association
ranged between 18% and 100%. Patient cohort charac-
teristics are reported in Table I. A female preponderance
was found (59%) in the cases, which was also present
in the control group (60.9%) (24). For a total of 139
psoriasis patients clinical data were available from our
Child-CAPTURE registry (see below). We found no significant phenotype differences between patients
with an age at onset < 10 years and ≥ 10 years.

Replication of genetic associations with paediatric-
onset psoriasis
Logistic regression analysis of our extended cohort
demonstrated the same associations with 4 of the 7
tested loci as in our previous study (24), albeit this
time with increased significance. The 4 genes were
IL23R (OR 2.42, 95% CI 1.23–4.74), ERAP1 (OR 1.55,
95% CI 1.18–2.03), LCE3B_LCE3C-del (OR 1.67,
95% CI 1.25–2.22, and HLA-C*06 (OR 17.1, 95% CI
10.5–27.9) (Table II).

Associations with clinical psoriasis characteristics
Age at onset of psoriasis. The mean age ± SD at onset of
psoriasis in our cohort was 8.2 ± 4.1 years. The group
of children with an early onset (< 10 years; n = 86) had a
mean age at onset of 5.5 ± 2.2 years, and the other group
(≥ 10 years; n = 53) had a mean age at onset of 12.7 ± 2.0
years. For IL12B, analysed using SNP rs3213094, we
demonstrated a significant association between age at
onset < 10 years and the risk (T) allele, with an OR of
2.59 (95% CI 1.14–5.88, p = 0.02); 17% of patients
with an age at onset < 10 years carried this risk allele,
compared to 7.5% of the patients with an age at onset
≥ 10 years (Table III). None of the other analysed loci
showed significant associations.

Family history of psoriasis. In 38.1% (n = 53) of the
patients a first-degree relative stated to have psoriasis.
Two thirds of the patients (67.6%, n = 94) reported a
positive family history of psoriasis up to third degree
relatives. However, we found no significant associations
between family history of psoriasis (first and up to third
degree relatives) and the allele frequency of the psoriasis

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Table I. Patient cohort characteristics

<table>
<thead>
<tr>
<th>Cohort characteristics</th>
<th>Number of cases</th>
<th>Male, n (%)</th>
<th>Age, years, mean ± SD (range)</th>
<th>Age at onset of psoriasis, years, mean ± SD (range)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Family history of psoriasis, n (%)</td>
<td>151</td>
<td>57 (41.0)</td>
<td>13.0 ± 4.1 (4–18)</td>
<td>8.2 ± 4.1 (0–17)</td>
</tr>
<tr>
<td>First-degree relatives</td>
<td>53 (38.1)</td>
<td></td>
<td></td>
<td>32.9 ± 35.9 (1–154)</td>
</tr>
<tr>
<td>Up to the third degree relatives</td>
<td>94 (67.6)</td>
<td></td>
<td></td>
<td>17.8 ± 15.4 (0–54)</td>
</tr>
</tbody>
</table>

*Data were available for 139 subjects.

SD: standard deviation.
risk factors. Remarkably, even HLA-C*06 showed no significant association with family history of psoriasis (first degree relatives \( p = 0.92 \) and up to third degree relatives \( p = 0.76 \)). In patients with a positive family history up to third degree relatives 76.1% were HLA-C*06 positive and in patients with only first-degree relatives with psoriasis, 76.5% were HLA-C*06 positive. In patients with a negative family history up to third degree relatives, 78.6% were HLA-C*06-positive compared to 77.2% in patients with no first-degree relatives with psoriasis (Table IV).

**Psoriasis severity.** In our cohort 55.4% (\( n = 77 \)) of the patients were classified as having severe psoriasis. Significant associations were demonstrated between severe psoriasis and the risk (T) allele of SNP rs17716942 (IFIH1) with an OR of 2.41 (95% CI 1.14–5.12, \( p = 0.019 \)) and the risk (A) allele of IFIH1 SNP rs17716942 (\( p = 0.002 \)) with an OR of 2.41 (95% CI 1.14–5.12, \( p = 0.019 \)) and the risk (T) allele of IL12B SNP rs3213094 (\( p = 0.047 \)) (Table III).

**Nail involvement.** In HLA-C*06 negative patients we found a significant increase of nail involvement (OR 0.32, 95% CI 0.14–0.76, \( p = 0.008 \), Table III). None of the other psoriasis risk alleles showed a positive or negative association with this clinical parameter.

**Koebner-phenomenon.** Forty-three patients (30.9%) were Koebner-positive. None of the investigated risk factors showed a significant association with Koebnerization in our cohort.

### DISCUSSION

This is the first study to report associations between clinical parameters and genetic risk factors in a prospective paediatric psoriasis cohort. We could confirm and strengthen, in a larger patient group, our previous findings that paediatric-onset psoriasis is associated with HLA-C*06, LCE3C_LCE3B deletion and SNPs in the ERAP1 and IL23R loci (24). We performed additional analyses based on clinical data from these children and demonstrated that age at onset, psoriasis severity and nail psoriasis are associated with different genetic risk factors of psoriasis. Remarkably, family history of psoriasis is clearly not associated with HLA-C*06 in this specific group.

For analysis of a possible effect of age at onset, we divided the patients into 2 groups with a cut-off point at 10 years of age, which is in line with data previously published by Lysell et al. (23). They demonstrated, in a Swedish cohort, that ERAP1 showed an association, albeit weak, with a psoriasis onset before 10 and 20 years. Also the strongest association with HLA-C*06 was found for this age group. We did not, however, find associations between age at onset and ERAP1 and HLA-C*06 in the Dutch paediatric psoriasis patients, which may be due to ethnic variation. In our cohort, an age at onset before 10 years was demonstrated to be associated with IL12B, which encodes the p40 subunit of interleukin (IL)-23 and IL-12 and is involved in both the IL12/Th1 and IL23/Th17 pathway of psoriasis (29, 30).

### Table II. Genetic associations with paediatric-onset psoriasis (stratified by age of onset adjusted for sex)

<table>
<thead>
<tr>
<th>Single nucleotide polymorphism</th>
<th>Gene</th>
<th>Allele</th>
<th>Frequency of risk alleles</th>
<th>Odds ratio (95% confidence interval)</th>
</tr>
</thead>
<tbody>
<tr>
<td>rs11209026</td>
<td>IL23R</td>
<td>G/A</td>
<td>0.97</td>
<td>0.010 2.42 (1.23 to 4.74)</td>
</tr>
<tr>
<td>rs3213094</td>
<td>IL12B</td>
<td>T/C</td>
<td>0.13</td>
<td>0.088 0.72 (0.50 to 1.05)</td>
</tr>
<tr>
<td>rs27524</td>
<td>ERAP1</td>
<td>A/G</td>
<td>0.43</td>
<td>0.002 1.55 (1.18 to 2.03)</td>
</tr>
<tr>
<td>rs17716942</td>
<td>IFIH1</td>
<td>T/C</td>
<td>0.88</td>
<td>0.445 1.16 (0.79 to 1.72)</td>
</tr>
<tr>
<td>rs240993</td>
<td>TRAF3IP2</td>
<td>T/C</td>
<td>0.29</td>
<td>0.132 1.09 (0.82 to 1.45)</td>
</tr>
<tr>
<td>direct PCR</td>
<td>LCE3C_LCE3B</td>
<td>DEL/WT</td>
<td>0.72</td>
<td>0.00049 1.67 (1.25 to 2.22)</td>
</tr>
<tr>
<td>direct PCR</td>
<td>HLA-C*06</td>
<td>POS/NEG</td>
<td>0.78</td>
<td>3.15E-30 17.1 (10.5 to 27.9)</td>
</tr>
</tbody>
</table>

*The first allele is the allele associated with psoriasis. DEL: deletion; WT: wild type; POS: positive; NEG: negative. Significant \( p \)-values are shown in bold.

### Table III. Significant associations between clinical characteristics and psoriasis risk factors

<table>
<thead>
<tr>
<th>Gene/locus</th>
<th>Clinical characteristics</th>
<th>Frequency of risk allele</th>
<th>Odds ratio (95% confidence interval)</th>
<th>( p )-value</th>
</tr>
</thead>
<tbody>
<tr>
<td>IL12B/rs3213094</td>
<td>Age at onset</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td>&lt; 10 years</td>
<td>0.17</td>
<td>2.59 (1.14–5.88)</td>
<td>0.020</td>
</tr>
<tr>
<td></td>
<td>( \geq 10 ) years</td>
<td>0.08</td>
<td></td>
<td></td>
</tr>
<tr>
<td>ERAP1/rs27524</td>
<td>Psoriasis severity</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td>Mild-to-moderate psoriasis</td>
<td>0.36</td>
<td>1.64 (1.01–2.67)</td>
<td>0.047</td>
</tr>
<tr>
<td></td>
<td>Severe psoriasis</td>
<td>0.48</td>
<td></td>
<td></td>
</tr>
<tr>
<td>IFIH1/rs17716942</td>
<td>Psoriasis severity</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td>Mild-to-moderate psoriasis</td>
<td>0.83</td>
<td>2.41 (1.14–5.12)</td>
<td>0.019</td>
</tr>
<tr>
<td></td>
<td>Severe psoriasis</td>
<td>0.92</td>
<td></td>
<td></td>
</tr>
<tr>
<td>HLA-C*06/direct PCR</td>
<td>Nail involvement</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td></td>
<td>Yes</td>
<td>0.61</td>
<td>0.32 (0.14–0.76)</td>
<td>0.008</td>
</tr>
<tr>
<td></td>
<td>No</td>
<td>0.83</td>
<td></td>
<td></td>
</tr>
</tbody>
</table>
and a positive family history. Clearly this cannot be a meaningful classifier, the distinction between paediatric (<18 years) and adult psoriasis (≥18 years) is probably more informative than the cut-off point of age 30 or 40 years, which is used in early and late onset psoriasis.

The most striking clinical association in our paediatric cohort with any of the genetic risk factors was the observation that nail psoriasis was found more often in HLA-C*06 negative patients ($p<0.008$). This association has been previously reported for a larger cohort (unstratified for age) by Gudjonsson et al. (19).

Psoriasis severity in adults was previously demonstrated to be associated with HLA-C*06 and LCE3C_LCE3B deletion (17, 19, 20, 22). In our pediatric cohort we did not find these associations. We did, however, identify an association between psoriasis severity and IFIH1 and ERAP1. IFIH1 encodes the interferon-induced helicase C domain 1 (innate immune system), which triggers type I interferon in response to microbial infection (31), and variants are associated with type 1 diabetes (32). ERAP1 encodes an amino peptidase, which regulates the quality of peptides bound to MHC class I molecules, such as HLA-C*06 (33).

A limitation of this study is the modest sample size for genetic studies. Considering that only children were included, it is, however, the largest cohort described with clinical features. In our cohort more than 50% of the patients was defined as severe psoriasis which could introduce a selection bias.

In conclusion, our findings suggest that genetic polymorphisms in both innate and adaptive immunity play a role in paediatric plaque psoriasis severity and age at onset of psoriasis. We confirm earlier associations found in adult psoriasis between HLA-C*06 with respect to nail involvement. The large proportion of patients with a positive family history in HLA-C*06 negative patients (and the lack of correlation between the two) indicates that other genes, either alone or interaction between two or more genes (34), may have significant effects on heritability.

ACKNOWLEDGEMENTS

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The authors declare no conflict of interest.

REFERENCES


3. Strange A, Capon F, Spencer CC, Knight J, Weale ME, Allen MH, et al. A genome-wide association study identifies new psoriasis susceptibility loci and an interaction between...


