

ORIGINAL ARTICLE

Identification of infants with increased type 1 diabetes genetic risk for enrollment into Primary Prevention Trials—GPPAD-02 study design and first results

Christiane Winkler^{1,2}  | Florian Haupt^{1,2} | Martin Heigermoser¹ | Jose Zapardiel-Gonzalo¹ | Jasmin Ohli¹ | Theresa Faure¹ | Evdokia Kalideri¹ | Angela Hommel³ | Petrina Delivani³ | Reinhard Berner⁴ | Olga Kordonouri⁵ | Frank Roloff⁵ | Thekla von dem Berge⁵ | Karin Lange⁶ | Mariusz Oltarzewski⁷ | Ryszard Glab⁷ | Agnieszka Szybowska⁸ | Matthew D. Snape⁹ | Manu Vatish¹⁰ | John A. Todd¹¹ | Helena E. Larsson^{12,13}  | Anita Ramelius¹⁴ | Jeanette Å. Kördel¹⁴ | Kristina Casteels^{15,16} | Jasmin Paulus¹⁵ | Anette G. Ziegler^{1,2,17} | Ezio Bonifacio³ | the GPPAD Study Group

¹Institute of Diabetes Research, Helmholtz Zentrum München, German Research Center for Environmental Health, Munich, Germany

²Forschergruppe Diabetes e.V. at Helmholtz Zentrum München, German Research Center for Environmental Health, Munich, Germany

³Center for Regenerative Therapies Dresden (CRTD), Faculty of Medicine, Technische Universität Dresden, Dresden, Germany

⁴Department of Pediatrics, University Hospital Carl Gustav Carus, Technische Universität Dresden, Dresden, Germany

⁵Hannoversche Kinderheilstalt, Kinder- und Jugendkrankenhaus AUF DER BULT, Hannover, Germany

⁶Department of Medical Psychology, Hannover Medical School, Hannover, Germany

⁷Institute of Mother and Child, Warsaw, Poland

⁸Department of Paediatrics, Medical University of Warsaw, Warsaw, Poland

⁹Department of Paediatrics, NIHR Oxford Biomedical Research Centre, University of Oxford, Oxford, UK

¹⁰Nuffield Department of Women's & Reproductive Health, University of Oxford, Oxford, UK

¹¹Wellcome Centre for Human Genetics, Nuffield Department of Medicine, University of Oxford, Oxford, UK

¹²Unit for Pediatric Endocrinology, Department of Clinical Sciences Malmö, Lund University, Lund, Sweden

¹³Department of Paediatrics, Skåne University Hospital, Malmö, Sweden

¹⁴Lund University, Lund, Sweden

¹⁵Department of Pediatrics, University Hospitals Leuven, Leuven, Belgium

¹⁶Department of Development and Regeneration, KU Leuven, Leuven, Belgium

¹⁷Forschergruppe Diabetes, Technical University Munich, Klinikum rechts der Isar, Munich, Germany

Correspondence

Christiane Winkler, Institute of Diabetes Research, Helmholtz Zentrum München, Ingolstaedter Landstr. 1, 85764 Neuherberg, Germany.
Email: christiane.winkler@helmholtz-muenchen.de

Abstract

Primary prevention of type 1 diabetes (T1D) requires intervention in genetically at-risk infants. The Global Platform for the Prevention of Autoimmune Diabetes (GPPAD) has established a screening program, GPPAD-02, that identifies infants with a genetic high risk of T1D, enrolls these into primary prevention trials, and follows

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the children for beta-cell autoantibodies and diabetes. Genetic testing is offered either at delivery, together with the regular newborn testing, or at a newborn health care visits before the age of 5 months in regions of Germany (Bavaria, Saxony, Lower Saxony), UK (Oxford), Poland (Warsaw), Belgium (Leuven), and Sweden (Region Skåne). Seven clinical centers will screen around 330 000 infants. Using a genetic score based on 46 T1D susceptibility single-nucleotide polymorphisms (SNPs) or three SNPs and a first-degree family history for T1D, infants with a high (>10%) genetic risk for developing multiple beta-cell autoantibodies by the age of 6 years are identified. Screening from October 2017 to December 2018 was performed in 50 669 infants. The prevalence of high genetic risk for T1D in these infants was 1.1%. Infants with high genetic risk for T1D are followed up and offered to participate in a randomized controlled trial aiming to prevent beta-cell autoimmunity and T1D by tolerance induction with oral insulin. The GPPAD-02 study provides a unique path to primary prevention of beta-cell autoimmunity in the general population. The eventual benefit to the community, if successful, will be a reduction in the number of children developing beta-cell autoimmunity and T1D.

KEYWORDS

beta-cell autoantibodies, genetic risk for type 1 diabetes, type 1 diabetes

1 | BACKGROUND

The primary prevention of common chronic disease is a major public health goal. Type 1 diabetes (T1D) is among the more frequent chronic diseases in childhood, and is increasing in incidence.¹ T1D results from an immune-mediated destruction of pancreatic islet beta-cells resulting in insulin deficiency. This process is identified by circulating autoantibodies to beta-cell antigens, indicative of a break in immunological self-tolerance.^{2,3} Neonates or infants who are at increased genetic risk for multiple beta-cell autoantibodies and T1D can be identified using genetic markers and T1D family history.⁴ Risk stratification by such genetic markers is now sufficiently advanced to bring testing to a population-based level and, thereby, facilitates the primary prevention of beta-cell autoimmunity and T1D.⁵

Primary prevention of T1D has a strong rationale. There is a clear peak incidence period of beta-cell autoantibody seroconversion between age 9 months and 3 years demonstrated in German,⁶ Finnish,⁷ and The Environmental Determinants of Diabetes in the young (TEDDY) studies.⁸ This provides a convenient study follow-up time in order to execute controlled clinical trials. Insulin is a first autoantibody target.^{6,8,9} It is widely held that if infant tolerance to beta-cell autoantigens could be enhanced, this could prevent or delay the onset of pre- or asymptomatic T1D (defined as loss of tolerance and multiple beta-cell autoantibodies), and hence prevent or delay the clinical onset of diabetes.¹⁰ As a primary prevention, we have laid the foundation for autoantigen-specific tolerance induction initiated prior to an autoimmune memory response. We have identified a dose of insulin that when administered orally on a daily basis to beta-cell autoantibody negative children at increased genetic risk is safe (neither affecting

plasma glucose levels nor causing an allergic reaction) and appears to engage the immune system in a manner that is consistent with immune-mediated, tolerogenic protection.¹¹ Therefore, the task is now to identify neonates or infants at increased genetic risk for T1D to execute the first insulin-based primary prevention trial. The objective of the Global Platform for the Prevention of Autoimmune Diabetes (GPPAD)-02 study protocol is to identify neonates or infants who have >10% risk for developing multiple beta-cell autoantibodies by the age of 6 years, which is the risk we have defined as the requirement for participation in a primary prevention randomized controlled trial that tests the effects of daily oral insulin vs placebo on the incidence of beta-cell autoantibody seroconversion. GPPAD will continue and likely improve screening for future primary prevention clinical trials.

2 | STUDY ORGANIZATION

GPPAD is a network of collaborating investigators from seven clinical trial centers located in five European countries: Germany (Bavaria, Saxony, Lower Saxony); UK (Oxford); Poland (Warsaw); Belgium (Leuven), and Sweden (Region Skåne). GPPAD was created to allow a multidisciplinary approach to prevent T1D and is coordinated by a GPPAD coordinating center in Germany (Munich).

2.1 | Identification of high genetic risk for T1D

T1D has a multifactorial etiology, which is determined by genetic and environmental factors.³ Lifetime risk in a European population is around 0.4%. A first-degree family history of T1D is associated with a

5% risk for T1D.⁴ There are at least 50 regions of the genome where genetic variation is associated with T1D risk,¹² the most important of these being in the HLA DR-DQ region of chromosome 6. Certain HLA DR-DQ genotypes confer markedly elevated risk for T1D. Notably, infants who are first-degree relatives of patients and who have the HLA DR4-DQ8 haplotype, and infants who have either the HLA DR3/DR4-DQ8 or the DR4-DQ8/DR4-DQ8 genotype have a risk of around 5%.^{4,13} Typing at additional T1D susceptibility regions can identify neonates or infants with risks that are 10% or more.⁵ We have recently developed a T1D genetic score that identified neonates or infants without family history of T1D who had a greater than 10% risk for pre-symptomatic T1D, and a nearly 2-fold higher risk than children identified by high-risk HLA genotypes alone. Thus, family history and genetic markers can be used to identify neonates or infants with 25-fold increased risk for T1D.⁵ Genetic risk in GPPAD-02 is based on these newly developed risk scores derived from a total of 46 single-nucleotide polymorphisms (SNPs) including SNPs that define HLA DR3, HLA DR4, and HLA DQ8 alleles as well as SNPs from HLA class I and non-HLA T1D susceptibility genes, and from HLA class II protective alleles (Supplemental Table 1).

3 | STUDY PROCEDURES

3.1 | Testing for T1D risk

Newborn screening for genetic, endocrine, and metabolic disorders is routinely done within the first days after birth at obstetric clinics or pediatrician offices, using a few drops of blood from the heel onto filter paper cards, or venous blood taken from the back of the hand. Testing for T1D risk is offered to families together with the regular newborn screening as a supplemental test with separate consent (Belgium, Germany, Poland, UK). Alternatively, it is offered at delivery using cord blood (Sweden) or at a pediatric visit before the age of 5 months (Sweden, Germany). At least one and a maximum of two blood spots are collected for testing of T1D risk using a separate GPPAD-02 filter paper card (Belgium, Germany, Sweden) or using the newborn screening filter paper card (UK, Poland). The dried blood spot filter paper cards are sent to the specialized newborn screening laboratories or the local clinical study center for subsequent retrieval. The screening target is 330 000 newborns or infants prior to age of 5 months.

3.2 | Consent procedure

Parents or legal guardians of neonates or infants are asked whether they wish to participate in the study. A qualified physician, midwife, or study nurse performs the informed consent in accordance with country-specific guidelines and ethical review board requirements, providing written and verbal information explaining the objectives of the GPPAD-02 study and the opportunities of primary prevention and follow-up for infants with high genetic risk. The families are told that if their child is found to have a high risk for T1D, they will be contacted and offered the possibility of participation in the prevention study with further informed consent. In Sweden, additional

information on a clinical trial aiming to prevent celiac disease is given to children with intermediate risk for T1D and risk for celiac disease is also provided.

Parents or guardians are given sufficient time to read the informed consent material and have any questions answered. It is explained that participation in the project is voluntary and that consent can be withdrawn at any time without providing a reason and without disadvantages by doing so. Confidentiality and protection of the data of every participant is maintained throughout the entire study. Only the local study personnel have access to personal identifiers. The informed consent form must be signed and dated by at least one parent/guardian. The parent or guardian is given a copy of the informed consent.

3.3 | Demographic data

Name, contact information of the parents, child's date of birth, gender, weight, date of blood collection, mother's date of birth, first-degree family history of T1D are collected either by a separate questionnaire or by extracting the information from the regular newborn screening card. Only pseudonymized data are entered into a central database held at the GPPAD coordinating center, Helmholtz Zentrum Muenchen, Germany.

3.4 | Genetic testing

DNA is extracted from a 3.2-mm punch of one dried blood spot and tested for the 46 SNPs (Supplemental Table 1). DNA extraction and SNP typing are performed in a central genotyping lab (LGC group, TEDDINGTON, Middlesex, UK [<https://www.lgcgroup.com>]; Figure 1A).

For sites where extra blood spots are collected for the GPPAD study (Germany, Sweden, Belgium), the remainder of the card is kept with the local GPPAD-02 team for at least 3 months in case repeat testing is required. Thereafter, the card is destroyed or, if there is consent for it to be kept for further research, stored at -20°C or -80°C .

The genetic risk score is calculated as previously described,⁷ by multiplying the number of risk alleles (ie, 0, 1 or 2 for each single SNP) with the weight assigned to each SNP (Supplemental Table 1). The weighted contributions of all SNPs plus an additive constant for each of the two HLA class II categories: 3.15 for infants who have the HLA DR4-DQ8/DR4-DQ8 genotype, 3.98 for infants who have the HLA DR3/DR4-DQ8 genotype, 3.12 for infants who have the HLA DR3/DR3 genotype, 2.08 for infants who have the HLA DR4-DQ8/DRX genotype, 1.55 for infants who have the HLA DR3/DRX genotype are summed.

3.5 | Definition of high genetic risk

For neonates or infants without a first-degree family history of T1D, high genetic risk is defined as a DR3/DR4-DQ8 or DR4-DQ8/DR4-DQ8 genotype, and a high genetic risk score >14.4 . For neonates or infants with a first-degree family history of T1D, high genetic risk is defined as having HLA DR4 and DQ8, and none of the protective

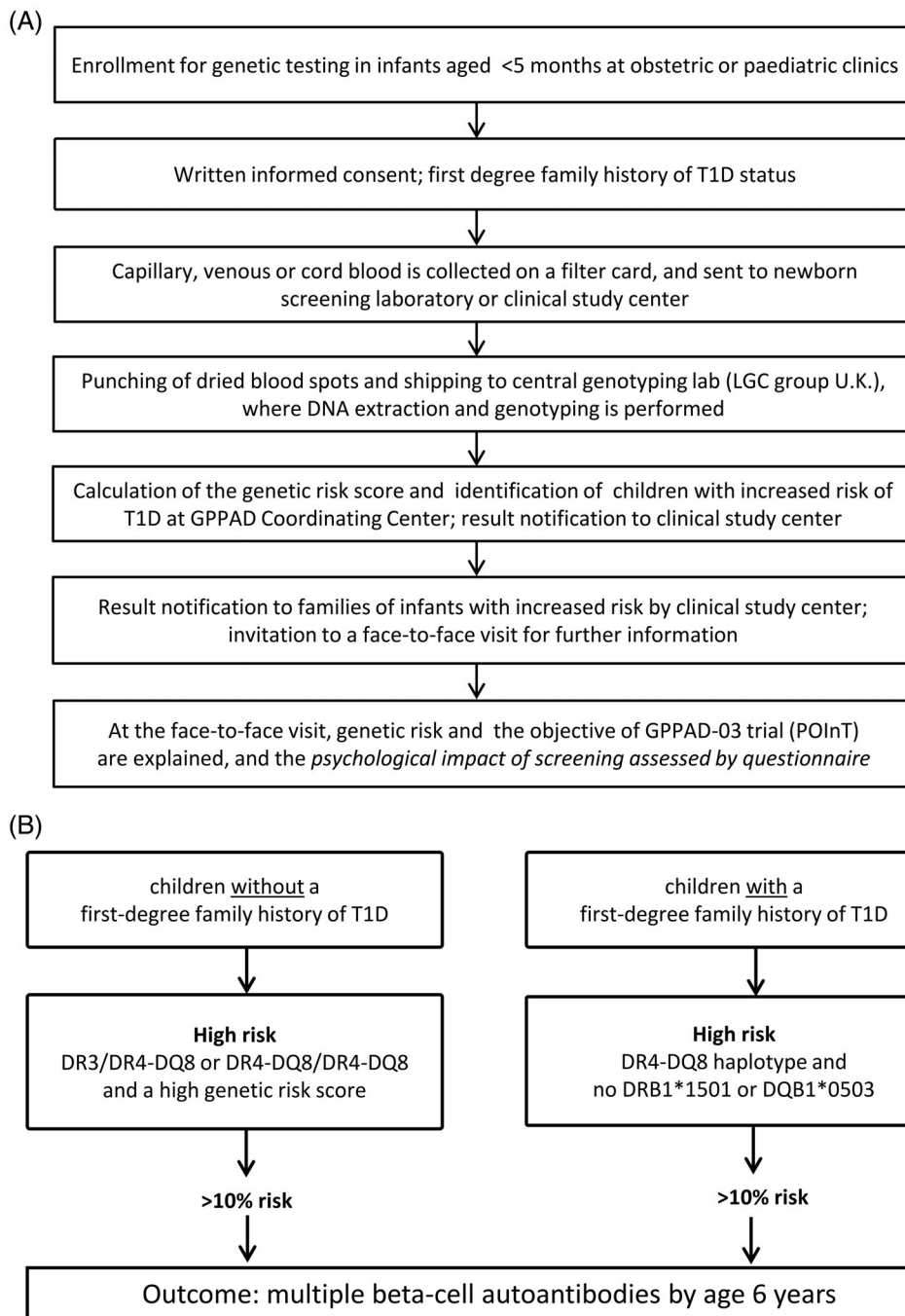


FIGURE 1 (A) GPPAD-02 study design and (B) Definition of increased genetic risk

alleles DRB1*1501 (SNP tag rs3129889) or DQB1*0503 (SNP tag rs1794265). Neonates or infants identified with high genetic risk are expected to have >10% risk for developing multiple beta-cell autoantibodies by the age of 6 years.⁹ The estimated prevalence of neonates or infants with this high risk for T1D is ~1% (Figure 1B).

3.6 | Result notification

Families of infants with a high risk for T1D are contacted by phone by an experienced physician/pediatrician or trained nurse from the local GPPAD-02 team. The family is invited to a face-to-face consultation

at the clinical study center, or alternatively at the primary care pediatrician's office. Dependent on the site-specific arrangements, the family may receive a result letter confirming the date of the consultation, and a brochure, which explains the meaning of T1D risk to the family, describes T1D symptoms, and informs about trials to prevent disease initiation and progression. The face-to-face consultation visit is planned to take place before 5 months of age and includes all parents/guardians. The content of the brochure may also be discussed during the consultation (Figure 1A).

Families of children with an estimated risk that is less than 10% are not contacted unless there is a country-specific follow-up of

children with intermediate risk of T1D or who have a high risk for celiac disease (Sweden), but are provided with a contact phone number if they want to inquire about the result.

3.7 | Invitation to participate in prevention trial

Infants with high genetic risk for T1D are asked to participate in a randomized controlled trial aiming to prevent the initiation of beta-cell autoimmunity and T1D (GPPAD-03 POInT Trial; ClinicalTrials.gov Identifier: NCT03364868). Families who decide not to participate in GPPAD-03 (POInT) may be asked to participate in existing natural follow-up studies aiming to prevent metabolic complications such as diabetic ketoacidosis through an early diagnosis of T1D program, and to investigate immunological alterations in the pathogenesis of the disease (ie, Fr1da in Bavaria, Fr1dolin in Lower Saxony). Participation in GPPAD-03 (POInT) and Fr1da/Fr1dolin requires a separate consent (Figure 1). In Sweden, subjects at intermediate risk for T1D and high risk for celiac disease are asked to participate in a prevention study of celiac disease (ClinicalTrials.gov Identifier: NCT03562221).

3.8 | Psychological assessment

The psychological impact of the consultation in which the increased genetic risk for developing T1D is communicated is subsequently monitored using a standardized questionnaire (Patient Health Questionnaire) in Germany and Sweden. The questionnaire looks for evidence of depression, anxiety and burden, and is handed to the parents at the face-to-face consultation visit. A trained and experienced psychologist of the GPPAD-02 team checks the questionnaire within 5 days. Families with elevated anxiety and depression levels (depression score > 5) are referred for consultation to a psychologist. In general, the parents are offered to contact the free telephone GPPAD-02-hotline with any questions or concerns.

3.9 | Statistical analyses

Analyses that are planned as part of the study will be limited to descriptive statistics. These will include the number of infants consented and tested per month per region and country, the frequency of infants with and without a first-degree relative with T1D, the proportion of children with high genetic risk, the proportion of families attending the consultation visit, the proportion of families with increased depression or anxiety and the distribution of scores from the Patient Health Questionnaire, and the proportion of families consenting to participate in the GPPAD-03 prevention trial.

Comparisons of the genetic risk score between the study centers were performed using Kruskal-Wallis test, one-way analysis of variance with *P*-values corrected by the Bonferroni adjustment and the comparison between boys and girls was performed using the Mann-Whitney *U* test.

4 | DATA SHARING

GPPAD is committed to open data sharing in compliance with all applicable European and GPPAD Consortium Member State, Data Protection and Privacy Protection laws, rules and regulations to facilitate scientific progress and patient benefit. GPPAD makes anonymized data of GPPAD-02 available to the scientific community on a regular basis (<https://www.gppad.org/en/data-sharing/>).

5 | PRELIMINARY RESULTS

From October 2017 to December 2018, 50 669 infants, including 891 (1.76%) with a first-degree family history of T1D (FDR), have been screened for increased genetic risk of T1D (Figure 2). The T1D-associated HLA risk genotypes DR3/DR4-DQ8 or DR4-DQ8/DR4-DQ8 were present in 1263 (2.5%) children from the general population and the HLA DR4-DQ8 haplotype was present in 250 (28.1%) FDR children (Table 1). Among the general population children with HLA DR3/DR4-DQ8 or DR4-DQ8/DR4-DQ8 genotypes, 327 (0.7% of all screened) children had a genetic risk score >14.4 which is the threshold above which there is a >10% risk for developing multiple beta-cell autoantibodies by the age of 6 years. Of the remainder, 631 (1.3% of all screened) children had a genetic risk score between 13.1 and 14.4, and 305 (0.6% of all screened) a genetic risk score below 13.1. In FDR children, 250 (28.1%) of 891 had a HLA DR4-DQ8 haplotype and no protective allele associated with >10% risk to develop multiple beta-cell autoantibodies by the age of 6 years (Table 1). The minor allele frequency of each SNP and the median genetic risk score are provided in Table 2. The median genetic risk score of all children was 10.59 (IQR: 9.76-11.62) and of the eligible infants was 14.55 (IQR: 12.98-15.02). The genetic risk score of screened children was higher in Sweden (Region Skåne) (median: 10.96, IQR: 9.86-11.99, *P* < .01) and lower in Leuven, Belgium (median: 10.22, IQR: 9.59-11.18, *P* < .01) when compared to the other

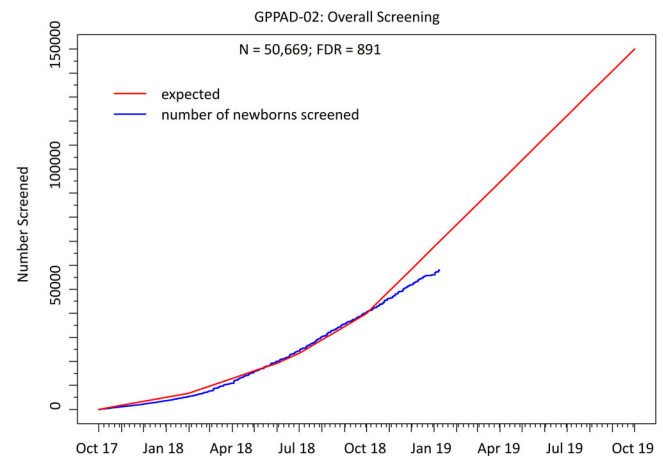


FIGURE 2 Cumulative number of newborns/infants screened in the GPPAD-02 study. Numbers are shown from the start of the study in October 2017. The red line represents the expected numbers and the blue line shows the actual number of screened infants

TABLE 1 First results from population-based screening for type 1 diabetes genetic risk (GPPAD-02)

	General population					
	Total	Germany	Poland	Sweden	Belgium	UK
Total screened	49 778	34 677	12 971	448	344	1338
HLA DR3/DR4-DQ8 or DR4-DQ8/DR4-DQ8	1263 (2.5%)	885 (2.6%)	300 (2.3%)	29 (6.5%)	3 (0.9%)	46 (3.4%)
Risk score in HLA DR3/DR4-DQ8 or DR4-DQ8/DR4-DQ8						
<13.1	305 (0.6%)	214 (0.6%)	73 (0.6%)	5 (1.1%)	1 (0.3%)	12 (0.9%)
13.1-14.4	631 (1.3%)	447 (1.3%)	146 (1.1%)	15 (3.4%)	2 (0.6%)	21 (1.6%)
>14.4 ^a	327 (0.7%)	224 (0.7%)	81 (0.6%)	9 (2.0%)	0 (0%)	13 (1.0%)
HLA DR3/DR3 ^b	557 (1.1%)	354 (1.0%)	160 (1.2%)	11 (2.5%)	3 (0.9%)	29 (2.2%)
	First-degree relatives					
	Total	Germany	Poland	Sweden	Belgium	UK
Total screened	891	709	149	20	5	8
HLA DR4-DQ8 and no DRB1*1501 or DQB1*0503 ^a	250 (28.1%)	195 (27.5%)	38 (25.5%)	8 (40.0%)	1 (20.0%)	8 (100.0%)

^aEligible for POInT Trial (GPPAD-03).^bHigh risk for celiac disease.**TABLE 2** Minor allele frequency of each SNP and the median genetic risk score

SNP	Gene	Minor allele frequency						
		Dresden	Hannover	Leuven	Malmö	Munich	Oxford	Warsaw
rs17426593	DR4	0.135	0.151	0.132	0.216	0.137	0.175	0.112
rs2187668	DR3	0.100	0.115	0.102	0.12	0.101	0.131	0.106
rs7454108	DQ8	0.090	0.099	0.075	0.14	0.090	0.091	0.079
rs3129889	DRB1*1501	0.125	0.121	0.091	0.127	0.121	0.114	0.121
rs1794265	DQB1*0503	0.025	0.034	0.031	0.024	0.032	0.030	0.021
rs1264813	HLA A 24	0.120	0.124	0.110	0.108	0.126	0.101	0.116
rs2395029	HLA B 5701	0.035	0.033	0.035	0.032	0.032	0.035	0.035
rs2476601	PTPN22	0.108	0.093	0.068	0.112	0.092	0.088	0.130
rs2816316	RGS1	0.178	0.181	0.195	0.198	0.177	0.180	0.182
rs3024505	IL10	0.164	0.166	0.140	0.154	0.161	0.145	0.168
rs1990760	IFIH1	0.385	0.405	0.399	0.376	0.402	0.424	0.354
rs3087243	CTLA4	0.425	0.444	0.451	0.412	0.437	0.459	0.369
rs10517086	C4orf52	0.283	0.287	0.293	0.278	0.286	0.279	0.285
rs2069763	IL2	0.360	0.350	0.305	0.397	0.338	0.318	0.357
rs6897932	IL7R	0.246	0.250	0.277	0.290	0.243	0.248	0.254
rs3757247	BACH2	0.447	0.450	0.425	0.412	0.456	0.457	0.447
rs9388489	C6orf173	0.466	0.479	0.483	0.461	0.481	0.486	0.444
rs6920220	TNFAIP3	0.187	0.195	0.163	0.229	0.183	0.204	0.165
rs1738074	TAGAP	0.383	0.389	0.451	0.425	0.393	0.445	0.344
rs7804356	SCAP2	0.225	0.234	0.244	0.231	0.230	0.231	0.189
rs4948088	COBL	0.042	0.040	0.038	0.044	0.038	0.042	0.038
rs7020673	GLIS3	0.469	0.468	0.439	0.496	0.465	0.471	0.468
rs12722495	IL2RA	0.100	0.094	0.074	0.099	0.099	0.084	0.092
rs947474	PRKCQ	0.194	0.199	0.194	0.168	0.195	0.197	0.191
rs10509540	RNLS/C10orf59	0.267	0.267	0.267	0.302	0.267	0.259	0.266

TABLE 2 (Continued)

SNP	Gene	Minor allele frequency						
		Dresden	Hannover	Leuven	Malmö	Munich	Oxford	Warsaw
rs1004446	INS	0.357	0.358	0.385	0.332	0.350	0.380	0.351
rs4763879	CD69	0.381	0.364	0.332	0.370	0.368	0.355	0.380
rs2292239	ERBB3	0.309	0.305	0.275	0.341	0.309	0.324	0.300
rs3184504	SH2B3	0.496	0.477	0.442	0.448	0.483	0.403	0.496
rs1465788	ZFP36L1	0.265	0.272	0.244	0.258	0.267	0.282	0.257
rs17574546	RASGRP1	0.180	0.177	0.166	0.198	0.180	0.179	0.176
rs3825932	CTSH	0.327	0.327	0.373	0.326	0.332	0.362	0.312
rs12708716	CLEC16A	0.357	0.345	0.387	0.324	0.367	0.364	0.345
rs4788084	IL27	0.442	0.417	0.383	0.435	0.411	0.390	0.472
rs7202877	CTRB2	0.116	0.100	0.093	0.126	0.106	0.091	0.127
rs2290400	ORMDL3	0.494	0.489	0.448	0.463	0.488	0.491	0.468
rs7221109	CCR7	0.406	0.384	0.368	0.387	0.405	0.354	0.434
rs45450798	PTPN2	0.149	0.147	0.159	0.145	0.144	0.157	0.138
rs763361	CD226	0.482	0.477	0.500	0.480	0.488	0.499	0.444
rs425105	PRKD2	0.160	0.156	0.136	0.150	0.159	0.173	0.172
rs2281808	SIRPG	0.327	0.313	0.316	0.324	0.319	0.318	0.333
rs3788013	UBASH3a	0.410	0.409	0.386	0.418	0.412	0.423	0.388
rs5753037	RPS3AP51	0.353	0.361	0.366	0.361	0.353	0.339	0.324
rs229541	IL2B	0.423	0.425	0.430	0.413	0.432	0.451	0.400
rs5979785	TLR8	0.266	0.287	0.296	0.263	0.283	0.299	0.232
rs2664170	GAB3	0.307	0.318	0.346	0.307	0.307	0.339	0.307
Genetic risk score (median)		10.58	10.65	10.22	10.96	10.54	10.56	10.57

Abbreviation: SNPs, single-nucleotide polymorphisms.

study centers and was not significantly different between male (median: 10.59, IQR: 9.76-11.64) and females (median: 10.58, IQR: 9.75-11.61; $P = .43$).

6 | SIGNIFICANCE AND NOVELTY

The GPPAD-02 study will provide a unique population-based path to primary prevention of beta-cell autoimmunity. Participation in GPPAD-02 also provides a platform to enable assessment of interventions for secondary prevention of T1D if children develop beta-cell autoantibodies in the course of the prevention or during natural follow-up. GPPAD and the GPPAD-02 study will, therefore, closely collaborate with INNODIA (www.innodia.eu), TrialNet (www.diabetestrialnet.org), and others for such opportunities. The eventual benefit to the community, if successful, will be a reduction in the number of children developing T1D.

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ORCID

Christiane Winkler <https://orcid.org/0000-0002-7606-3687>

Helena E. Larsson <https://orcid.org/0000-0003-3306-1742>

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SUPPORTING INFORMATION

Additional supporting information may be found online in the Supporting Information section at the end of this article.

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APPENDIX A

GPPAD STUDY GROUP

GPPAD-Coordinating Center (CC)

Melanie Gündert¹, Stefanie Arnolds¹, Robin Assfalg¹, Corinna Barz¹, Karina Blasius¹, Cigdem Gezginci¹, Cordula Falk¹, Joerg Hasford², Florian Haupt¹, Martin Heigermoser¹, Bianca Höfelschweiger¹, Verena Hoffmann¹, Manja Jolink¹, Nana Kwarteng¹, Ramona Lickert¹, Claudia Matzke¹, Rebecca Niewöhner¹, Michaela Ott¹, Peter Ruile¹, Marlon Scholz¹, Katharina Schütte-Borkovec¹, Mira Taulien¹, Lorena Wendel¹, Katharina Wystub-Lis¹, José Maria Zapardiel Gonzalo¹

¹Institute of Diabetes Research, Helmholtz Zentrum München, Neuherberg, Germany

²Institut für Medizinische Informationsverarbeitung, Biometrie und Epidemiologie, Ludwig-Maximilians-Universität München, Munich, Germany

Belgium Clinical Center

Kristina Casteels, Goele Smeets, Hilde Morobé, Jasmin Paulus, Renka Van Heyste, Sophie Achten, Emma Larivière, Janne Houben, Lionel Marcelis, Luc Regal

Germany, Dresden Clinical Center

Ezio Bonifacio, Reinhard Berner, Uta Ceglarek, Petrina Delivani, Sevina Dietz, Yannick Fuchs, Gita Gemulla, Manja Gottschalk, Sophie Heinke, Angela Hommel, Anne Karasinsky, Susann Kowal, Fabian Lander, Robert Morgenstern, Katharina Nitzsche, Bianca Schlee, Marina Stopsack, Marc Weigelt, Pauline Wimberger, Marie-Luise Zielmann, Nicole Zubizarreta

Germany, Hannover Clinical Center

Olga Kordonouri, Torben Biester, Thomas Danne, Nils Janzen, Ute Holtkamp, Karin Lange, Erika Marquardt, Frank Roloff, Kerstin Semler, Thekla von dem Berge

Germany, Munich Clinical Center

Anette G. Ziegler^{1,2}, Peter Achenbach¹, Melanie Bunk¹, Anita Gavrisan¹, Katharina Gestrich¹, Willi Grätz¹, Pascale Heim-Ohmayer¹, Melanie Herbst¹, Julia Hirte¹, Theresa Faure¹, Anna Hofelich¹, Evdokia Kalideri¹, Cornelia Kraus¹, Yvonne Kriesen¹, Karin Lange¹, Jasmin Ohli¹, Claudia Ramminger¹, Jennifer Schairer¹, Christiane Winkler¹, Susanne Wittich¹, Stephanie Zillmer¹

¹Institute of Diabetes Research, Helmholtz Zentrum München, Neuherberg, Germany

²Forscherguppe Diabetes, Klinikum rechts der Isar, Technische Universität München, Medical faculty, Munich, Germany

Poland Clinical Center

Agnieszka Szypowska, Mariusz Ołtarzewski, Sylwia Dybkowska, Katarzyna Dzygała, Lidia Groele, Karolina Dłużniak-Gońska, Dorota Owczarek, Katarzyna Popko, Agnieszka Skrobot, Anna Taczanowska, Beata Zduńczyk

Sweden Clinical Center

Helena Elding Larsson, Markus Lundgren, Åke Lernmark, Daniel Agardh, Jeanette Åkerström Kördel, Carin Andrén Aronsson, Rasmus Bennet, Charlotte Brundin, Annika Fors, Lina Fransson, Berglind Jónsdóttir, Ida Jönsson, Zeliha Mestan, Anita Ramelius, Evelyn Tekum Amboh, Carina Törn

UK Clinical Center

Matthew Snape, John A Todd, Owen Bendor-Samuel, James Bland, Edward Choi, Rachel Craik, Kimberly Davis, Arancha de la Horra, Yama Farooq, Clare Scudder, Ian Smith, Manu Vatish, Louise Willis, Tabitha Wishlade