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Autoimmune diabetes mellitus after COVID-19 vaccination in adult population: a systematic review of case reports

Ali S Alsudais^{1,2}, Raghad S Alkanani^{1,2}, Abdulaziz B Fathi^{1,2}, Saleh S Almuntashiri^{1,2}, Jafar N Jamjoom^{1,2}, Mustafa A Alzhrani^{1,2}, Alaa Althubaiti^{1*} and Suhaib Radi^{1,2,3}

Abstract

Background Autoimmune/type 1 diabetes mellitus (T1DM) is a recently described rare occurrence following the administration of adjuvants such as coronavirus disease 2019 (COVID-19) vaccines. This systematic review aimed to review all available literature on the potential association between COVID-19 vaccines and T1DM.

Methods The Directory of Open Access Journals, MEDLINE, Google Scholar, and Scopus were systematically searched for all published studies from inception to July 2022. Articles reporting T1DM development within 8 weeks of administration of COVID-19 vaccine were included. Two reviewers independently performed the risk of bias assessment following the Joanna Briggs Institute (JBI) Critical Appraisal Checklist for Case Reports.

Results Eight eligible studies were retrieved, comprising 12 patients diagnosed with T1DM after being vaccinated with a COVID-19 vaccine. Six patients (50%) reported T1DM after receiving the second dose. Five patients (41.7%) presented with diabetic ketoacidosis, of which four presented within the first eight days after vaccination. Five patients (41.7%) had genetic susceptibility, with RNA binding motif protein 45 (*RBM45/DRB1*) and major histocompatibility complex, class II, DQ beta 1 (*HLA-DQB1*) mutations being prominent.

Interpretation In this review, we have shown a small number of new-onset diabetes cases coincidently occurring soon after the COVID-19 vaccine, especially in those with genetic susceptibility. Despite being older, these patients had a similar phenotype to T1DM. While there might be a causal relationship between COVID-19 vaccines and T1DM development, this should not influence decisions regarding vaccination since the overall benefit outweighs the risk. Further larger prospective trials are needed to assess causal relationship and to clarify the potential roles of COVID-19 vaccine-derived antigens in autoimmune disease development.

Protocol registration PROSPERO-CRD42022342093.

Significance statement

With the rapid and wide-spread rate of COVID-19 vaccines developed, many complications have appeared, including autoimmune diseases. In this review, we have shown a small number of new onset diabetes cases coincidently occurring soon after the COVID-19 vaccine. Despite being older, these patients will have a similar

*Correspondence: Alaa Althubaiti thubaitia@ksau-hs.edu.sa

Full list of author information is available at the end of the article



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phenotype to type 1 diabetes and risk is higher in those with genetic susceptibility. Most will present with diabetic ketoacidosis and require insulin therapy. However, incidence remains rare, and the overall benefit outweighs the risk. Such a possible complication would be important to recognize to appropriately screen and treat affected patients.

Keywords COVID-19 vaccine, Autoimmune diabetes mellitus, Type 1 diabetes mellitus

Background

Since December 2019, the coronavirus disease 2019 (COVID-19) pandemic has imposed devastating health, social, and economic burdens on health systems worldwide [1, 2]. Many strategies and precautionary measures (e.g., social isolation, personal hygiene, wearing face masks, and frequent hand washing) have been implemented globally to limit COVID-19's spread. However, a protective vaccine is required to achieve sufficient herd immunity to prevent the progression of the COVID-19 pandemic. Many vaccines have eventually been approved for clinical use, and around 12 billion doses have been administered globally [3, 4]. The most globally administered COVID-19 vaccines are based on messenger ribonucleic acids (e.g. mRNA-1273 [Moderna] and BNT162b2 [Pfizer-BioNTech]), viral vectors (e.g. ChAdOx1-S [Vaxzervria]), or inactivated viruses (e.g. CoronaVac [Sinovac Biotech]) [3, 4].

While the virus' unprecedented toll caused the accelerated development of multiple vaccines by many biopharmaceutical companies, the breakneck pace of their development has raised safety concerns from the general population. A study on understanding the causes of COVID-19 vaccine hesitancy reported that concerns about vaccine side effects increased the odds of becoming vaccine hesitant by 31% [5]. Most of the reported adverse effects of COVID-19 vaccines are non-specific, including fever, fatigue, and headache [6]. However, emerging evidence indicates an intricate relationship between the COVID-19 vaccine and autoimmune/type 1 diabetes (T1DM) [7]. The administration of adjuvants such as COVID-19 vaccines, potentially containing virusderived proteins, to genetically predisposed individuals could activate autoimmune cascades such as T1DM [7]. Despite vague evidence, the literature suggests additional autoimmune manifestations such as vaccine-induced thrombotic thrombocytopenia, autoimmune liver disease, immunoglobulin A (IgA) nephropathy, and Guillian-Barre syndrome [8]. In this context, T1DM is of great public concern. Therefore, this study aimed to review the literature and the evidence on the potential association between COVID-19 vaccines and T1DM. It describes the clinical presentation and outcomes of COVID-19-associated diabetes and assesses the quality of reports.

Methods

Protocol, registration, and search strategy

This systematic review of case reports was performed according to the Preferred Reporting Items for Systematic Reviews and Meta-Analysis (PRISMA) guidelines [9] and registered with the PROSPERO online database (identifier: CRD42022342093).

A systematic literature search was performed to identify all published studies from inception without language or country restrictions. The search was performed in July 2022 in the following databases: The Directory of Open Access Journal, MEDLINE, Google Scholar, and Scopus. It was conducted using the search terms: ('COVID-19' OR 'SARS-CoV-2') AND ('Autoimmune Diseases' OR 'Type 1 Diabetes Mellitus' OR 'Autoimmune Disorder' OR 'Autoimmunity') AND ('Vaccines' OR 'COVID-19 Vaccines' OR 'Immunization'). Detailed queries are provided in Supplement Table 1. The reference lists of relevant articles were searched for additional studies in October 2022.

Eligibility criteria

A search of case reports and case series of patients receiving a COVID-19 vaccine and developing T1DM within eight weeks of any vaccine dose was conducted. The T1DM diagnosis must have been established by either the presence of any positive antibodies known to cause the disease (anti-glutamic acid decarboxylase [anti-GAD] antibodies, anti-tyrosine phosphatase [anti-IA2] antibodies, islet cell antibodies, or insulin autoantibodies), or in the case of negative antibodies, and low C-peptide levels following the acute presentation. However, this study excluded articles with overlapping patient data, reporting diabetes development>3 months after a vaccine injection, reporting cases where the diabetes diagnosis was insufficiently documented or unclear, and reporting cases known to have T1DM before vaccination.

Study selection

A detailed literature search was performed in the four databases mentioned above, identifying 1264 articles. Access to the predefined databases were granted by the Saudi Digital Library. All articles were exported to Microsoft Excel through which duplication removal and screening process were executed manually. Two authors (AsA and AbF) independently scrutinized all 1264 papers according to specific eligibility criteria. After review, 1254

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articles were excluded, and ten were initially retained. An author raised conflicts about two articles, and a third author (SR) was involved in resolving the inter-author conflict, resulting in the exclusion of both articles. The remaining eight articles met the inclusion criteria and were included in the analysis (Fig. 1). The eligible studies were published between October 2021 and July 2022.

Data extraction and quality assessment

Data were extracted in Microsoft Office Excel using a predefined template. The extracted data included the following information: author details, publication year, study characteristics, patient demographics, patients' family and history of autoimmune diseases or diabetes mellitus, vaccine type, symptom onset (after which dose), signs and symptoms, diagnosis day after vaccination, diabetic ketoacidosis (DKA), antibody presence, fasting C-peptide levels, stimulated C-peptide levels, hemoglobin A1C (HbA1C) levels before vaccination, HbA1C levels at diagnosis, genetic susceptibility, and other T1DM triggers. Two reviewers independently performed the risk of bias assessment. Studies were scored as "yes," "no," or "not applicable," following the Joanna Briggs Institute (JBI) Critical Appraisal Checklist for Case Reports based on an eight-item scale [10].

Data analysis

Since this review is descriptive, we used descriptive statistics to describe the data with median (range) for continuous variables and frequencies and percentages for categorical variables. Microsoft Excel software was used for all calculations.



Table 1 Clinical features and	l baseline la	boratory resui	ts of the cases	s at presentati	uc							
	Case 1	Case 2	Case 3	Case 4	Case 5	Case 6	Case 7	Case 8	Case 9	Case 10	Case 11	Case 12
First Author (reference)	Yano et al. 2022 [11]	Bleve et al. 2022 [12]	Bleve et al. 2022 [12]	Sasaki et al. 2022 [13]	Sakurai et al. 2022 [14]	Sasaki et al. 2022 [15]	Tang et al. 2022 [16]	Patrizio et al. 2021 [17]	Aydoğan et al. 2022 [18]			
Country	Japan	Italy	Italy	Japan	Japan	Japan	China	Italy	Turkey	Turkey	Turkey	Turkey
Age (years)	51	57	61	73	36	45	50	52	56	48	27	36
Sex	ш	ц	ш	щ	Ц	ц	M	X	M	Σ	ш	M
BMI (kg/m ²)	18.3	N/A	N/A	N/A	N/A	20.6	18.1	N/A	27.4	21.9	20	22.8
Medical history	None	A / N	Acquired hypothyroid- ism	- Osteoporsis - Non- tuberculous mycobacte- rial infection - Diet-con- trolled type 2 diabetes	N en en	Bronchial asthma	None	- Vitiligo - T2DM for 8 years on oral agents	Vitiligo and Hashimoto thyroiditis	None	None	None
Family history	T2DM (father)	T2DM, vitiligo, Hashimoto's thyroiditis	N/A	None	None	N/A	T2DM (mother)	N/A	N/A	None	None	None
Type of vaccination	Moderna mRNA- 1273	ChAdOx1-S	Pfizer-BioN- Tech	Moderna mRNA-1273	Pfizer-BioN- Tech	Pfizer-BioNTech	CoronaVac	Pfizer- BioN- Tech	Pfizer- BioNTec	Pfizer- BioN- Tec	Pfizer- BioN- Tec	Pfizer- BioN- Tec and Coro- naVa
Duration from vaccination to symptoms onset (weeks)	4 (28 days)	1 (8 days)	Since receiv- ing the 2nd dose of the vaccine	4	<1 (3 days)	1 (6 days)	1 (6 days)	4	2 (15 days)	œ	m	m
Dose after which the symptoms appeared	1st	- s	2nd	2nd	-s	1 ⁵	1 ^s	2nd	2nd	2nd	2nd	4th
DKA at presentation	Yes	N/A	N/A	N/A	Yes	Yes	Yes	N/A	No	N/A	N/A	Yes
Triggers (e.g. steroids use, pan- creatic disease, viruses) Laboratory findings	No infection	No infection	N/A	N/A	No infection	No infection	N/A	N/A	N/A	N/A	N/A	N/A
HbA1c (%) before vaccination (reference range)	5.6 (4.6–6.2)	N/A	N/A	> 7%	N/A	N/A	N/A	7%	5.9 (4.0-5.6)	5.6 (4.0-5.6)	N/A	N/A
HbA1c (%) at diagnosis (refer- ence range)	10.3 (4.6–6.2)	10.4	11.5	9.3 (4.9–5.9)	7 (4.6–6.1)	7.6 (4.6–6.2)	"Near normal"	10.1% (4.6–6.1)	8.2 (4.0-5.6)	10.1 (4.0-5.6)	12.5 (4.0-5.6)	12.6 (4.0-5.6)
Anti-GAD antibodies presence	No	Yes	Yes	Yes	No	No	No	Yes	Yes	Yes	Yes	Yes

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	Case 1	Case 2	Case 3	Case 4	Case 5	Case 6	Case 7	Case 8	Case 9	Case	Case 11	Case
Other antibodies presence	- Insulin autoanti- bodies - TPOAh*	- Anti-IA2 - Anti-Trans- Glut IgA	TPOAb	Insulin auto- antibodies	None	None	None	- TRAb^ - TgAb - TPOAb	N/A	N/A	N/A	N/A
2-peptide (ng/mL) (refrence ange)	0.4 (0.80–2.50)	N/A	N/A	0.54 (0.74–3.18)	0.35 (0.8–2.3)	0.33 (0.80–2.5)	Undetectable/ Low levels	1.0 (1.0-3.5)	1.5 (1.1–4.4)	0.97 (1.1– 4.4)	0.87 (1.1– 4.4)	0.39 (1.1– 4.4)
HLA-DNA typing (genetic usceptibility)		N/A	N/A	DRB1 *04:05:01-	DRB1*0405- DQB1*0401	DRB1*04:05:01/ *13:02:01 and	DQB1*02:03/ 03:03 and	N/A	N/A	N/A	N/A	A/N

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Results

Characteristics of the eligible studies and patients' characteristics

This review included 12 cases from all eight included articles [11-18], summarized in Table 1. All patients were diagnosed with T1DM after being vaccinated with a COVID-19 vaccine. Seven of the 12 patients were female (58.3%). A summary of patients' characteristics is provided in Table 2. Their median age was 49 years. Body mass index (BMI) was reported for seven patients, of which four were normal ($18.5-24.9 \text{ kg/m}^2$), two were underweight, and one was overweight. A family history of type 2 diabetes was recorded for three patients (25%). One patient (8.3%) had a family history of autoimmune diseases, including vitiligo and Hashimoto's thyroiditis. Three patients (25%) had a personal history of other autoimmune diseases (vitiligo and Hashimoto's thyroiditis). Moreover, two patients (16.7%) had a history of type 2 diabetes mellitus for >7 years that was effectively controlled with diet modifications and oral antidiabetic agents. The Pfizer-BioNTech vaccine was given to seven patients (58.3%), the Moderna mRNA-1273 vaccine to two (16.7%), the CoronaVac to one (8.3%), and the ChAdox1-s vaccine to one (8.3%). The last patient (8.3%) received four doses of two different vaccines. They first received two CoronaVac doses, followed by two Pfizer-BioNTech doses, after which symptoms appeared.

Six patients (50%) reported T1DM after receiving their second dose, and symptoms appeared within eight weeks of administration. Additionally, five patients (41.7%) had T1DM after their first dose, and symptoms appeared within four weeks of administration. One patient (8.3%) had T1DM after their fourth dose, with symptoms appearing within three weeks. The shortest duration from vaccination to symptom onset was three days, while the longest was approximately eight weeks. DKA was recorded in five (41.7%) patients, one (8.3%) presented before developing DKA, while the rest (n=6; 50%) had no available data on DKA at the time of T1DM presentation. C-peptide levels were quantified and documented for ten patients (83.3%), and antibodies were documented for nine (75%). Six of the 12 patients had their T1DM diagnosis established by low C-peptide levels and positive antibodies, three by low C-peptide levels only, and three by positive antibodies only. The most common antibody present was the anti-GAD (n=8; 66.7%). Only five patients had valid data for HbA1c levels before vaccination; two had high values (HbA1c \geq 7%), one had a prediabetic HbA1c value (HbA1c=5.9%), and two had normal values (HbA1c \leq 5.6%). Post-vaccine data showed that all patients had a very high HbA1c level indicative of T1DM, except one with no HbA1c data reported. The estimated mean HbA1c level at diagnosis was 9.96%. Triggers that might cause T1DM development including steroid use,

Table 2 Summary of patients' characteristics

Variable		Descriptive
		statistics
Age, <i>median (range)</i> years		50.5 (27–73)
Sex, n (%)	Female	7 (58.3)
	Male	5 (41.7)
Type of vaccine received, n (%)	Moderna mRNA-1273	2 (16.7)
	ChAdOx1-S	1 (8.3)
	Pfizer-BioNTec	7 (58.3)
	CoronaVac	1 (8.3)
	Pfizer-BioNTec and CoronaVa	1 (8.3)
Duration from vaccination to sym (range) weeks	ptoms onset, <i>median</i>	2.5 (0–8)
Dose after which symptoms ap-	1st dose	5 (41.7)
peared, <i>n (%)</i>	2nd dose	6 (50)
	4th dose	1 (8.3)
Past medical history of autoim- mune disease, <i>n</i> (%)		3 (25)
Diabetic ketoacidosis at presentat	tion (DKA), <i>n (%)</i>	5 (41.7)

Table 3 Characteristics of articles included

First author [reference number]	Quality score*	Overall quality
Yano et al. 2022 [11]	б	Good
Bleve et al. 2022 [12]	4	Intermediate
Sasaki et al. 2022 [13]	7	Good
Sakurai et al. 2022 [14]	6	Good
Sasaki et al. 2022 [15]	6	Good
Tang et al. 2022 [16]	7	Good
Patrizio et al. 2021 [17]	4	Intermediate
Aydoğan et al. 2022 [18]	7	Good

*JBI risk assessment tool was used

pancreatic diseases, and viruses; have been reported nor established. Five patients (41.7%) had genetic susceptibility, with RNA binding motif protein 45 (*RBM45/DRB1*) and major histocompatibility complex, class II, DQ beta 1 (*HLA-DQB1*) mutations being prominent.

A basal-bolus insulin therapy regimen was used as the treatment mode in all but one patient to correct the acute presenting condition. The final patient refused insulin and was treated with diet modification, showing a good response. Eight of the 11 treated patients achieved sufficient glycemic control with insulin therapy only; the remaining three patients had missing data on their post-intervention status. Two of the seven patients who had data regarding their post-discharge condition stopped insulin and were being managed with diet modification only, showing good glycemic control (Supplementary Table 2).

The current study embodies systematic review of case reports, wherein the original articles have thoroughly complied with the ethical obligation of obtaining informed consent from the subjects under study.

Quality assessment

The overall quality of the included studies was intermediate to good. Two studies were classified as being of intermediate quality [12, 17], while the remaining studies as good quality (Table 3). Question #7 from the JBI tool was not applicable to this systematic review because no intervention was studied. Details of each study's quality assessment are presented in the Appendix (Supplementary Table 3).

Interpretation

To our knowledge, this is the first and largest systematic review of reported cases of COVID-19 vaccine-associated T1DM. This systematic review of eight articles identified 12 case reports with T1DM, defined by either the presence of autoantibodies or low C-peptide levels at diagnosis. Four different vaccines were administered to the 12 patients. They were all mRNA, recombinant DNA, viral vector, or inactivated virus vaccines, with most cases (83.3%) given an mRNA-based vaccine. Of the 12 patients, 41.7% developed T1DM symptoms after their first dose, of which 80% presented with symptoms within the first 10 days. However, 50% developed symptoms after their second dose. The average time to symptom onset varied from days to weeks, with the shortest duration being three days and the longest being eight weeks after vaccination. Five patients (41.7%) presented with DKA, an uncontrolled diabetes complication, of which four presented within the first eight days after vaccination.

Upon assessment of the patients in this study, it became evident that aside from genetic susceptibility and a history of autoimmune diseases, there were no risk factors for developing T1DM. There were no triggers for developing autoimmune diseases, such as infections, steroid use, or pancreatic diseases. BMIs were mostly normal or underweight. All patients were far from T1DM's peak incidence age, usually within childhood (Table 4). Genetic susceptibility was only examined for five patients. One patient had a family history of autoimmune diseases. Three patients were already diagnosed with autoimmune diseases, such as Hashimoto's thyroiditis and vitiligo. While it is likely for those with autoimmune diseases to develop other autoimmune diseases, a trigger is usually required. Two patients with elevated HbA1c levels before vaccination were known to have type 2 diabetes. However, their HbA1c level increased significantly after vaccination, and they likely converted into T1DM triggered by the vaccine with autoantibody development. Although long-standing type 2 diabetes can lead to beta cell failure and low C-peptide, the fact that their diabetes was well controlled on oral agents, and they developed autoantibodies makes them less likely to be type 2 diabetes. Two patients had no data of C-peptide presented with a

Table 4Characteristics similarities & differences between typesof autoimmune diabetes [19, 20]

Characteristics	Usual Type 1 Diabetes Mellitus	Late Autoim- mune Diabetes in Adults (LADA)	Autoimmune Diabetes fol- Iowing COVID- 19 Vaccine
Age at onset	Most commonly in Childhood	Adult age (30–50 years of age); but may occur at any age	Adult age (25–75 years of age)
Typical weight	Normal – underweight	Normal – overweight	Normal – underweight
Symptoms at onset	Acute	Insidious	Acute
Time to requir- ing insulin	At onset	Months-years	At onset
Presence of Autoantibodies	Yes	Yes	Yes
C-peptide levels	Low-undetect- able	Low-normal	Low-undetect- able
Personal or family history of other autoim- mune disease	Yes	Yes	Yes
Genetic susceptibility	Yes	Yes	Yes
DKA at presentation	Common	Rare	Common
Honeymoon phase	Yes (usually after 6–18 months)	No	Possible (after 4–8 weeks)

significantly elevated HgA1c (>10%). However, both had either a personal or family history of autoimmune disease and were not diabetics before. In addition to the significant elevation in HgA1C (>10%), the presence of type 1 diabetes-related antibodies might support the hypothesis that these 2 cases are type 1 rather than type 2 diabetes. Furthermore, three patients presented with a rapid development of DKA within one week of receiving COVID-19 vaccines. Although this might be a coincidence, the fact that their HbA1C levels was around 7% at time of diagnosis indicates that their severe hyperglycemia was of very recent onset. Moreover, these patients had genetic susceptibility and these genes have been linked to fulminant type 1 diabetes, which might be the reason for this very quick development of ketonemia.

Following T1DM presentation and diagnosis, all but one patient (n=11) were managed with an intensive basal-bolus insulin therapy regimen. Two patients steadily decreased their insulin doses and discontinued insulin treatment due to recurrent hypoglycemic episodes. Those who discontinued insulin therapy were proposed to have entered the honeymoon phase (or partial remission). This is a phase within some diabetic patients' courses involving increased pancreatic B-cell activity and insulin sensitivity, resulting in a progressive decrease in insulin dependence to the point where they no longer require insulin therapy [8]. This phase was reported to occur in 3-61% of newly diagnosed T1DM patients, with the highest incidence within the first six months to one year after diagnosis, but its pathophysiology remains unclear [8]. Many patients with partial remission require low-dose insulin therapy (<0.5 U/kg/day) or some oral antidiabetic medications. Those who completely refrain from medication use may be categorized as having complete remission, provided they maintain HbA1c levels < 6% and normal glycemic levels without any medications [8, 21, 22]. Interestingly, two of the four patients with post-discharge follow-up data (50%) were off insulin and showed good glycemic control, indicating a possible honeymoon phase. This transition occurred around three months after diagnosis, which is relatively shorter than in usual T1DM patients. The lack of long-term follow-up information for most of the included cases precludes us from reaching a definitive conclusion regarding the rate and duration of remission in such cases.

COVID-19 vaccines have been proposed to act as triggers for autoimmune diseases via multiple pathways, including immune system hyperstimulation, autoantibody formation, and molecular mimicry [23]. The severe acute respiratory syndrome coronavirus 2 (SARS-COV-2) vaccine has been suggested to cause a "cytokine storm" involving the release of multiple cytokines to induce an inflammatory state in those with mild to severe infections. Hyperstimulation is associated with autoimmune disease development since those with severe COVID-19 developed anti-phospholipid and antinuclear antibodies, interferon-neutralizing autoantibodies, and antineutrophil cytoplasmic antibodies against myeloperoxidase (MPO/p-ANCA) and proteinase 3 (PRTN3/c-ANCA) [4].

Furthermore, T1DM is not the only autoimmune disease associated with COVID-19. Multiple case reports have suggested a relationship between the SARS-CoV-2 vaccine and Guillain–Barre syndrome [24], Graves' disease [6, 25], warm and cold autoimmune hemolytic anemia [26], and Kawasaki disease [27]. The basis for developing various autoimmunity forms following COVID-19 vaccination is thought to sequence homology between SARS-CoV-2 and human proteins. New antibodies created to attack the virus have the potential to cross-react with the host's cells. Many human proteins shared with SARS-COV-2 have been identified as potentially pathogenic when perturbed (e.g. mutation, alteration, and improper function) [23].

As indicated by autoimmunity development, mRNA vaccines result in more post-vaccination T1DM cases than all other vaccine types. The mRNA-based COVID-19 vaccines have a higher pooled risk ratio for developing adverse events and local adverse reactions following immunization than all other vaccine types [6].

Furthermore, while rare, developing autoimmunity following vaccination is a known risk and is categorized under the term autoimmune/inflammatory syndrome by adjuvants (ASIA syndrome), covering vaccinationinduced side effects and other complications. This syndrome is thought to develop in those with pre-existing genetic risk factors after exposure to an adjuvant in the vaccine by activating autoimmune pathways.

Though not explicitly explained, higher adverse reaction rates following mRNA vaccination are important since they greatly influence vaccine hesitancy and compliance in populations. Sallam's [28] systematic review of vaccine acceptance rates showed a large variance in SARS-CoV-2 vaccine acceptance among countries, ranging from as low as 23.4% and 28.3% in Kuwait and Jordan, respectively, to as high as 97.0% and 94.3% in Ecuador and Malaysia, respectively. The most common causes of hesitancy were low disease risk perception, lack of trust in vaccination safety and effectiveness, and vaccine affordability and delivery [28]. Therefore, when presenting information like that offered by this study, it is important to place the findings in the context of the entire vaccination program. Over 12.85 billion doses of COVID-19 vaccines [29] have been administered, with a minority of individuals experiencing severe adverse effects due to vaccination. The benefit of vaccinating the population and preventing disease propagation far outweighs the individual vaccination risks, as shown by the 6.6 million deaths worldwide [29] and severe hospitalizations that strained healthcare systems globally, resulting in the inability to provide care to many individuals.

Limitations

Our review had multiple limitations. First, given that included articles were case reports and series, missing information is a significant issue. For example, multiple parameters were expected to act as baseline measurements for the participants in this study, including BMI, HbA1c level, genetic susceptibility, and a history of exposure to autoimmune disease triggers such as infection. They are used to assess whether the phenotype of vaccine-induced diabetes is similar to T1DM. Unfortunately, many of these measurements were missing in the assessed case reports. In two of the case reports an intermediate overall quality score was seen. This low score was attributed to important missing variables in terms of diagnosis and laboratory work up. Despite the lack of major risk factors for developing T1DM in patients with complete data, these sets of information or their lack thereof limit our ability to conclude with certainty that vaccination is the cause of T1DM development. Secondly, there might be an underdiagnosis and underappreciation of T1DM development triggered by COVID-19 vaccination, especially in low-income countries where testing for antibodies and C-peptides might be limited. In such countries, most of these patients might be labelled as type 2 diabetes, given their older age at diagnosis.

Conclusions

We have shown in this review a small number of new onset diabetes cases coincidently happening soon after COVID-19 vaccine, especially in those with genetic susceptibility. Despite being older, these patients had a similar phenotype to T1DM. Most presented with DKA and required insulin therapy but needed to be monitored closely since their insulin requirement might rapidly decline. Moreover, sudden unexplained severe hyperglycemia and increased HbA1c with development of diabetes-related autoantibodies in otherwise well-controlled type 2 diabetes should alert physicians about the possibility of conversion to T1DM. There might be an underappreciation and underdiagnosing of this COVID-19-induced T1DM. Nevertheless, its incidence remains rare compared to the huge number of vaccine doses given worldwide. While there might be an association in a few case reports between COVID-19 vaccines and T1DM development, this should not influence decisions regarding vaccination since the overall benefit outweighs the risk. Further larger prospective trials are needed to assess causal relationship and to clarify the potential roles of COVID-19 vaccine-derived antigens in autoimmune disease development.

Supplementary Information

The online version contains supplementary material available at https://doi.org/10.1186/s12902-023-01424-0.

Supplementary Material 1

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None.

Authors' contributions

A.S.A., A.B.F., and S.R. contributed to screening. A.S.A., R.S.A., and S.R. have extracted the data and assessed the quality of the eligible studies. A.B.F., J.N.J., S.S.A., and M.A.A. wrote the manuscript. A.S.A., R.S.A., A.B.F., J.N.J., S.S.A., M.A.A., A.A., S.R. contributed to development of the study design, reviewed, and revised the manuscript critically for important intellectual content, gave final approval of the version to be published and agreed to be accountable for all aspects of the work. All the authors revised it critically for important intellectual content.

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None.

Data Availability

The details of extracted data, and quality assessment are in the appendices. Further data may be available upon a reasonable request to the corresponding author.

Declarations

Ethical approval and consent to participate

No ethical approval was required. The present investigation represents a systematic review of case reports, in which the original articles have adhered to the ethical principle of obtaining informed consent.

Consent for publication

N/A.

Approval date of registry and the registration no. of the study/trial

Protocol registration: PROSPERO-CRD42022342093, Date: 04/07/2022.

Competing interests

The authors declare no competing interests.

Author details

¹College of Medicine, King Saud bin Abdulaziz University for Health Sciences, Jeddah, Saudi Arabia

²King Abdullah International Medical Research Centre, Jeddah, Saudi Arabia

³Department of Internal Medicine, Division of Endocrinology, Ministry of the National Guard-Health Affairs, Jeddah, Saudi Arabia

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