

Citation: Navas-Madroñal M, Valero-Mut A, Martínez-Zapata MJ, Simón-Talero MJ, Figueroa S, Vidal-Fernández N, et al. (2017) Absence of antibodies against KIR4.1 in multiple sclerosis: A three-technique approach and systematic review. PLoS ONE 12(4): e0175538. https://doi.org/ 10.1371/journal.pone.0175538

**Editor:** Markus Reindl, Medizinische Universitat Innsbruck, AUSTRIA

Received: September 5, 2016 Accepted: March 27, 2017 Published: April 17, 2017

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**Data Availability Statement:** All relevant data are within the paper and its Supporting Information files.

**Funding:** The experimental study has been funded with an unrestricted grant from Novartis Spain. LQ is supported by the JR13/00014 of the Instituto de Salud Carlos III, Ministry of Economy and Innovation, Spain. The systematic review was performed independently and was not funded by the leading funding source.

RESEARCH ARTICLE

# Absence of antibodies against KIR4.1 in multiple sclerosis: A three-technique approach and systematic review

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# **Abstract**

#### Introduction

Antibodies targeting the inward-rectifying potassium channel KIR4.1 have been associated with multiple sclerosis (MS) but studies using diverse techniques have failed to replicate this association. The detection of these antibodies is challenging; KIR4.1 glycosylation patterns and the use of diverse technical approaches may account for the disparity of results. We aimed to replicate the association using three different approaches to overcome the technical limitations of a single technique. We also performed a systematic review to examine the association of anti-KIR4.1 antibodies with MS.

#### Methods

Serum samples from patients with MS (n = 108) and controls (n = 77) were tested for the presence of anti-KIR4.1 antibodies using three methods: 1) by ELISA with the low-glycosylated fraction of recombinant KIR4.1 purified from transfected HEK293 cells according to original protocols; 2) by immunocytochemistry using KIR4.1-transfected HEK293 cells; and 3) by immunocytochemistry using the KIR4.1-transfected MO3.13 oligodendrocyte cell line. We developed a systematic review and meta-analysis of the association of anti-KIR4.1 antibodies with MS according to the Preferred Reporting Items for Systematic Reviews and Meta-Analyses (PRISMA) guidelines.

#### Results

We did not detect anti-KIR4.1 antibodies in the MS patients or in controls using ELISA. Neither did we detect any significant reactivity against the antigen on the cell surface using the KIR4.1-transfected HEK293 cells or the KIR4.1-transfected MO3.13 cells. We included 13 prospective controlled studies in the systematic review. Only three studies showed a positive association between anti-KIR4.1 and MS. Clinical and statistical heterogeneity between studies precluded meta-analysis of their results.



Competing interests: The experimental study was funded through an unrestricted grant from Novartis Spain. LQ is supported by a grant (JR13/00014) from the Instituto de Salud Carlos III, Ministry of Economy and Innovation, Spain, and from research grants from the GBS-CIDP Foundation. The other authors declare no conflict of interest. The corresponding author had full access to all data and final responsibility for the decision to submit the article for publication. This does not alter our adherence to PLOS ONE policies on sharing data and materials.

#### Conclusion

We found no association between anti-KIR4.1 antibody positivity and MS. Although this lack of replication may be due to technical limitations, evidence from our study and others is mounting against the role of KIR4.1 as a relevant MS autoantigen.

## Introduction

Multiple sclerosis (MS) is an autoimmune and neurodegenerative disease of the central nervous system that arises in genetically susceptible individuals exposed to environmental risk factors[1]. Clinical, genetic and experimental data in MS suggest a primary autoimmune response against oligodendrocytes is followed by a secondary neurodegenerative process that leads to disability[2]. Although MS is generally considered a T-cell mediated disease, B-cells populate the demyelinating lesions and the cerebrospinal fluid (CSF) and form lymphoid follicles in the brain parenchyma and the meninges[3]. Importantly, the presence of oligoclonal bands in the CSF, the most frequently observed lab abnormality, is useful for diagnosis and MS conversion risk stratification [4,5]. Moreover, B-cell depletion has proven an effective treatment for MS, suggesting B-cells play a central role in the pathogenesis of MS [6]. The search for the target antigen(s) of the immune response is an active field of MS research[7]. The discovery of antibodies targeting aquaporin-4 provided a biological marker to classify these patients as a specific disease subset (neuromyelitis optica) and the proof of principle that humoral factors could also mediate MS pathogenesis[8]. Despite exhaustive research using diverse approaches, the target antigen (or antigens) of the MS immune response remains largely unknown. Thus, MS lacks a disease-specific diagnostic biomarker and diagnosis relies on clinical and brain imaging criteria[9]. In 2012, Srivastava et al published a landmark article describing the presence of antibodies against the inward-rectifying potassium channel 4.1 (KIR4.1) in approximately half of the MS patients they studied [10]. A similar study in children with MS by the same group yielded similar results[11]. Antibodies against KIR4.1 were originally described using ELISA assays in which either whole protein KIR4.1 or KIR4.1 extracellular loop peptides served as substrate antigens, but subsequent studies failed to replicate the peptide ELISA results[12-15]. Lennon et al failed to replicate the original report using a cellbased assay with KIR4.1-transfected human embryonic kidney (HEK) cells[16]. KIR4.1 is present in diverse tissues and cell-types, including oligodendrocytes, where it is expressed as KIR4.1 homotetramers, and astrocytes, where it forms heterotetramers with KIR5.1[17]. Furthermore, KIR4.1 glycosylation varies depending on the cell type and the whole protein ELISA results are heavily influenced by the glycosylation status of the protein[18]. These KIR4.1 features could account for the conflicting results, prompting replication studies using the original ELISA system and addressing the influence of glycosylation patterns. Using the original ELISA system, two independent groups recently published reports that fail to replicate KIR4.1 as a frequent autoantigen in MS[19,20].

Our study aims to solve the controversy using a four-fold approach. Experimentally, we used a whole-KIR4.1 ELISA to try to replicate the results reported in a former study and two cell-based assays. One assay used KIR4.1-transfected HEK cells to display KIR4.1 in non-denaturating conditions on the surface of cells used to express and purify the protein for the ELISA test. The other used the KIR4.1-transfected MO3.13 cell line to display the KIR4.1 on a cell type that would replicate the natural glycosylation patterns of KIR4.1 better than HEK cells. Finally, we systematically reviewed the evidence on the association of MS with anti-KIR4.1 antibodies and attempted a meta-analysis to help solve this controversy.



## **Methods**

# Subjects, informed consent and protocol approvals

Consecutive patients fulfilling the McDonald diagnostic criteria for MS[9] and visited in our Unit from June 2014 to June 2016 were included in the study. Controls (n=77) consisted of sera from healthy subjects (NC; n=13) and from subjects with other neuroinflammatory and degenerative disorders: multifocal motor neuropathy (NMM; n=16); chronic inflammatory demyelinating polyneuropathy (CIDP; n=24); myasthenia gravis (MG; n=16); amyotrophic lateral sclerosis (ALS; n=8). Serum samples were aliquoted and stored at -80°C until needed. All patients gave written informed consent to participate in the study under a protocol approved by the Ethics Committee at Hospital de la Santa Creu i Sant Pau (protocol IIBSP-KIR-2014-12). Sera from patients and controls were collected in two different sample collections following Instituto de Salud Carlos III protocols for sample collection (C.0003085, for patients; C.0002365, courtesy of Professor Isabel Illa, for controls)

# Enzyme-linked immunosorbent assay (ELISA)

HEK293 cells were cultured in 100 mm culture dishes in 5% fetal bovine serum-supplemented DMEM medium plus penicillin-streptavidin, sodium-pyruvate and L-glutamine. At 70% cell confluence, the mammalian expression vector pcDNA3.1 encoding full-length human KIR4.1 with a His-tag in the C-terminus)[20] was transfected with Lipofectamine 2000 (Thermo-Fisher, Spain) and incubated at 37°C. After 24 hours, cells were washed with PBS, centrifuged and KIR4.1 purified following the original Srivastava et al[10] protocol. Briefly, cells were lysed, centrifuged and KIR4.1 protein purified using the HisPur Cobalt resin (Thermo Fisher, Spain). ELISA was performed as described by Srivastava et al. using a final concentration of 7ug per milliliter of elution fraction 3 (EF3), in which low-glycosylated KIR4.1 oligomers are found. Elution fraction 1 reactivity was used as the background control for each sample. Results were read using a Beckman AD340 plate reader (Beckman-Coulter, Indianapolis, IN). Results were represented as the difference of the elution fraction 3 minus elution fraction 1 (EF1) optical density (OD). Samples were considered positive by ELISA when they had a  $\Delta$ OD higher than mean healthy control ΔOD plus five standard deviations. All samples were tested at the same time and only analyzed if the ELISA fulfilled the quality criteria proposed by Srivastava[18] and Marnetto et al[21]: ELISA signal of the rat monoclonal anti-KIR4.1 antibody 20F9 (courtesy of Srivastava and Hemmer) with an OD lower than 0,36, EF3 OD higher than 1,361 and assay diluent OD lower than 0,1 were the quality criteria used to proceed with ELISA analysis. Positive control samples were used to help set-up the assay (kindly provided by Rajneesh Srivastava)

Sensitivity and specificity of the KIR4.1 ELISA and their respective confidence intervals were calculated. The non-parametric Kruskal Wallis test with post-hoc multiple comparisons test was used to compare average OD between groups. All statistical analyses were performed with GraphPad Prism v5.0.

# Immunocytochemistry (ICC)

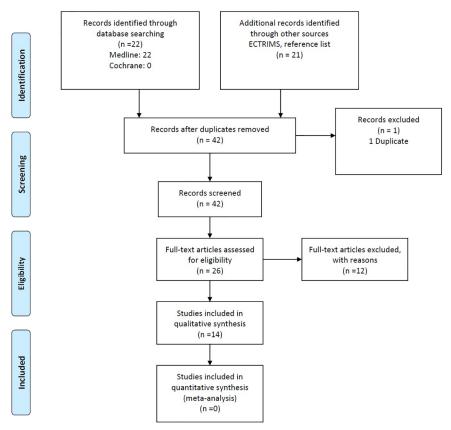
KIR4.1-transfected MO3.13 oligodendrocyte cell line (Cedarlane Labs, Canada) and HEK293 cells were used for ICC experiments. Briefly, HEK293 cells or the MO3.13 oligodendrocyte cell line were cultured in 60mm culture plates with poly-D-Lysine-coated coverslips and transfected as previously described. After 24 hours, cells were washed with PBS, fixed with 4% paraformaldehyde and blocked with 5% goat serum in PBS for 1 hour. Coverslips were then incubated with patients' sera at 1:100 in blocking buffer and with a 1:500 commercial anti-



KIR4.1 antibody (Millipore AB8518) at room temperature. After 1 hour, cells were washed with PBS and incubated with Alexa-Fluor goat-anti-human 594nm and goat anti-rabbit 488nm (Thermo-Scientific, Spain) secondary antibodies diluted 1:500. Finally, coverslips were washed and mounted using Vectashield with DAPI (Vector Labs, UK). Evaluators (LQ, MN) were blinded for the diagnosis. Demonstration of KIR4.1 expression on MO3.13 cells was performed in an independent experiment described in Supplementary Methods.

# Systematic review

We aimed to assess the association of anti-KIR4.1 antibodies with MS. We proceeded according to the Preferred Reporting Items for Systematic Reviews and Meta-Analyses (PRISMA) guidelines [22]. We included studies published in English or Spanish, addressing anti-Kir4.1 antibody positivity in patients with MS diagnosed according to the 2010 McDonald criteria compared to healthy controls. Exclusion criteria included lack of outcomes of interest in the report (Fig.1). The search strategy included the Cochrane Central Register of Controlled Trials, Ovid Medline (1946 to August 2016) and the European Committee for Treatment and Research in MS (ECTRIMS) abstract databases. The medical subject heading (MeSH) terms and key words used in the search included the terms: "multiple sclerosis AND Kir4.1" and ("multiple sclerosis" [MeSH Terms] OR ("multiple" [All Fields] AND "sclerosis" [All Fields]) OR "multiple sclerosis" [All Fields]) AND kir4.1 [All Fields]. For data extraction, one author (AV) selected the studies, and two authors (LQ, AV) independently assessed each study to check its eligibility. References meeting the inclusion criteria were retrieved in full and further assessed



**Fig 1. Systematic review flowchart.** Thirteen studies addressing the prevalence of anti-KIR4.1 antibodies in MS and controls were identified. These 13 studies plus our own results were used for the systematic review.

https://doi.org/10.1371/journal.pone.0175538.g001



independently by the same two review authors. Data in each study were extracted by two independent reviewers (LQ and AV) using the same standardized method. We contacted study authors by e-mail to obtain additional information when data were missing or unclear. The methodological quality was assessed by the Effective Public Health Practice Project's Quality Assessment Tool (http://www.ephpp.ca/tools.html), which has six components: selection bias, study design, confounders, blinding, data collection methods, and withdrawals/drop-outs. Each component was classified as strong, moderate or weak. A global methodological quality assessment was obtained according to the number of components rated as weak (0, 1, or > 1)as strong, moderate or weak[23]. The information extracted was: publication data, study design, patients' characteristics, and patients with positivity against anti-KIR4.1 antibodies. When two or more different assays yielded different frequencies of positive anti-KIR4.1 antibody results in the same report, the standard test (ELISA) was chosen for analysis. The primary outcome was percentage of study participants with positive anti-KIR4.1 antibodies. The comparison was MS patients versus controls. The odds ratios (OR) and 95% confidence interval (CI) were calculated to compare groups. We used a random effect model [24] for all analyses, as we expected variation in effects due to differences in study populations and methods to measure anti-KIR4.1 antibodies. Heterogeneity between studies was evaluated using the I<sup>2</sup> statistic, categorized as follows: <30% not important; 30%-50% moderate; 50%-75% substantial; and 75%-100% considerable [25]. The data were not pooled if I<sup>2</sup> was over 75%. Statistical software used was Review Manager 5.3[26].

#### Results

One hundred and eight MS patients were included in the study and tested by ELISA and ICC with both KIR4.1-transfected HEK293 cells and the KIR4.1-transfected MO3.13 cell line. <u>S1</u> Table shows basic demographic features of patients and controls.

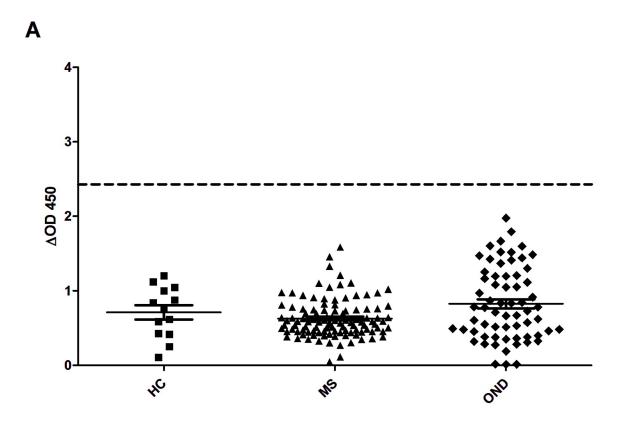
## **ELISA**

None of the 108 MS patients or the healthy and disease controls reached a  $\Delta$ OD higher than average healthy control  $\Delta$ OD plus 5 standard deviations and, thus, failed to be considered positive by ELISA according to the pre-established criteria (Fig 2A). The sensitivity of the assay to detect MS patients in comparison with any type of control was 0.000 (0.000 to 3.535; 95% confidence interval) and specificity was 1 (0.952 to 1, 95% confidence interval). Comparison of the average  $\Delta$ OD signal between groups did not differ between MS patients and healthy participants or disease controls (p = 0.08, Kruskal-Wallis). The average  $\Delta$ OD signal from patients with MMN was significantly higher than in any other group except CIDP (p<0.0001, Kruskal-Wallis). The CIDP average  $\Delta$ OD was higher than that in the ALS and MG groups (Fig 2B; p<0.0001, Kruskal-Wallis).

## **Immunocytochemistry**

We performed ICC experiments to further explore the presence of anti-KIR4.1 antibodies in MS patients. All patients and disease controls tested negative by immunocytochemistry over KIR4.1-transfected HEK293 cells despite abundant surface expression of KIR4.1 (Fig 3A–3F). Considering the possibility that glycosylation patterns could differ between HEK293 and oligodendrocytes, we repeated the ICC experiments using KIR4.1-transfected MO3.13 cells[18]. No MS patient or disease control testing was positive for surface staining on MO3.13 cells, despite efficient KIR4.1 transfection and cell surface expression (Fig 3G–3L). Fig 3 shows patients and disease controls with the highest  $\Delta$ OD signals by ELISA.





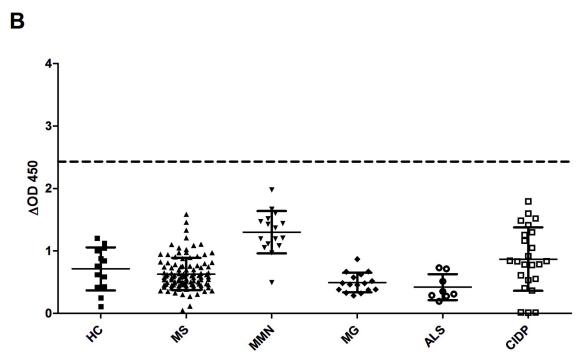


Fig 2. IgG reactivity against KIR4.1 does not differ in MS and controls by ELISA. Low-glycosylated fraction of the KIR4.1 whole protein complex was purified and used to detect anti-KIR4.1 antibodies by ELISA. None of the 108 MS patients



or disease controls reached a  $\Delta$ OD signal above healthy control average  $\Delta$ OD plus 5 SD. The average  $\Delta$ OD signal did not show significant differences between groups (A). The average  $\Delta$ OD signal from patients with MMN was significantly higher (p<0.0001, Kruskal-Wallis) than that from any other group except CIDP. The CIDP average  $\Delta$ OD was higher than the average in the ALS and MG groups (p<0.0001, Kruskal-Wallis; Fig 2B).

https://doi.org/10.1371/journal.pone.0175538.g002

# Systematic review

Thirteen studies were included, 12 in adult subjects and one in children (Fig 1 and S1 Fig). The studies were developed in the USA (3)[15,16,20], Germany (2)[10,11], France (2)[12,27], Japan (2)[28,29], Israel (1)[14], Italy (1)[21], Spain (1), Switzerland (1)[19], and Turkey (1) [13]. They showed substantial variability in the technical determination of anti-KIR4.1 and control group populations. The methodological quality was qualified as week in five studies [13,21,27–29], moderate in four studies[11,16,20,30] plus our own, and strong in three studies [10,14,15]. One study[19] was not included due to lack of normal controls (S1 Fig). Compared with studies in healthy controls, three studies in MS patients showed an association of positivity of anti-KIR4.1[10,11,14] and seven studies, including our own, showed no significant differences[12,13,16,20,21,27–29] (Fig 4). The statistical heterogeneity between studies was over 75% and did not allow data pooling to perform meta-analysis.

## **Discussion**

We found no differences in the proportion of MS patients harboring autoantibodies against KIR4.1 compared to disease and healthy controls using the same ELISA-based detection assay described in the original report and using two different cell-based assays[10,19,20].

The description for clinically useful biomarkers is of special relevance in MS[31].MS is a chronic, heterogeneous and disabling disease that mainly affects young adults. It is diagnosed based on clinical and radiological criteria but diagnostic uncertainty often persists in the initial phases. Moreover, there is clinical, pathological, experimental and genetic evidence that B-cells and autoantibodies play a role in MS pathogenesis [32]. For these reasons, the quest for autoantibodies is a topic of great relevance in MS research. The description of anti-KIR4.1 antibodies was a potential landmark discovery[33], first, because KIR4.1 was observed to co-localize with aquaporin-4 in the brain and represented a plausible antigen[34] and, second, because it provided the first solid evidence that disease-specific autoantibodies could be found in a substantial proportion of adult MS patients [10]. This first report was followed by a second study by the same group describing a similar proportion of positive patients among children with MS [11]. Importantly, these studies did not identify clinical, radiological or therapeutic features that were specifically associated with the presence of anti-KIR4.1 antibodies in MS patients. The results from early replication studies using the KIR4.1 extracellular loop (residues 83-120) peptide to test anti-KIR4.1 antibodies by ELISA were insufficiently consistent to demonstrate a clear association of anti-KIR4.1 and MS even though the same detection assay was used [12,27]. A thorough replication study including a multimodal approach also failed to confirm the association of anti-KIR4.1 antibodies with MS[16]. These reports did not use the fulllength KIR4.1 protein ELISA for anti-KIR4.1 antibody detection, and they did not take into consideration the existence of KIR4.1 oligomers with different degrees of glycosylation that were reported to influence the ELISA results[18]. At the 2015 ECTRIMS Congress, Marnetto et al presented a small study in which they addressed the technical difficulties associated with anti-KIR4.1 testing and partially replicated the results of the original report[21]. However, this presentation highlighted "session-specific" success in detecting the antibodies, further highlighting the inconsistent results. Recently, two replication studies by Chastre et al and Probstel et al, using the full KIR4.1 complex and addressing the technical caveats of the assay,



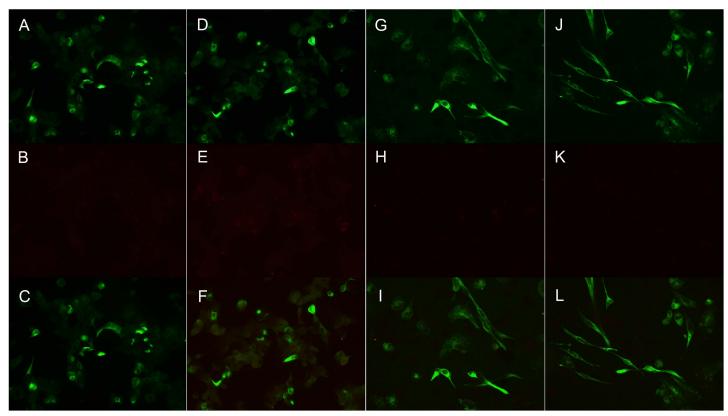


Fig 3. Lack of identification of KIR4.1 antibodies in serum samples using ICC. MS patients or disease controls did not show antibodies against KIR4.1 tested by ICC in KIR4.1 transfected HEK293 (A-F) or MO3.13 (G-L). Anti-KIR4.1 commercial antibody reactivity is shown in the upper row (A, D, G, J), serum IgG reactivity from an MS patient (B and H) and from a disease control (E and K) in the middle row, and merge images (C, F, I, L) in the lower row. The MS patient and disease control showing highest ΔOD were chosen to assemble this Figure.

https://doi.org/10.1371/journal.pone.0175538.g003

did not show differences in anti-KIR4.1 antibodies between patients and healthy or disease controls[19,20]. Other authors, using unbiased approaches to detect autoantibodies with protein arrays, and not specifically aimed to detect anti-KIR4.1 antibodies, also failed to identify KIR4.1 as a candidate antigen, but the expression systems, experimental designs and antigen sources on the arrays for these studies are significantly different to the original description [35,36].

In this context, we developed the present study to specifically address the technical differences that could lead to inconsistent results in the association of anti-KIR4.1 antibodies with MS. First, we used the full-length KIR4.1 protein in an ELISA using methods as described in the original report and in two recent studies[19,20], and we found no differences in the proportion of patients or controls testing positive for these antibodies. On the contrary, the average ELISA signal tended to be higher in specific types of neuroimmune controls, such as MMN and CIDP. This observation is in agreement with findings in a recent study[19] in which anti-KIR4.1 reactivity correlated significantly with reactivity against sham antigens, suggesting that higher nonspecific antibody promiscuity, independently of the disorder, could result in higher anti-KIR4.1 reactivity[19]. Second, we transfected KIR4.1 in HEK293 cells using the same mammalian expression vector (pDNA3.1+) that Srivastava et al, Chastre et al and ourselves had used for protein purification. ELISA allowed us to detect the low-glycosy-lated isoforms using the Srivastava and Marnetto criteria with the rat 20F9 antibody but we were unable to identify surface staining in KIR4.1-transfected HEK293 cells even in patients



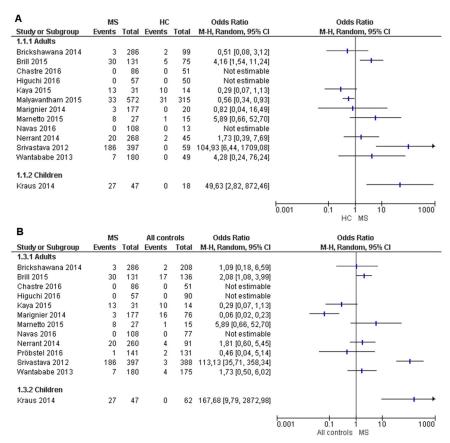


Fig 4. Forest plot. Thirteen studies were included in the systematic review. Three of them showed evidence of an association of KIR4.1 with MS patients compared to healthy controls (A) or all controls (B). One showed a weak association of KIR4.1 antibodies with healthy controls. None of the other reports found any association of KIR4.1 antibodies with MS, healthy or disease controls. The study by Probstel[19] was excluded from Fig 4A forest plot because it lacked comparison with healthy controls, as specified in the inclusion criteria (A). The study by Malyavantham[15] was excluded from Fig 4B forest plot because it lacked comparison with disease controls (B).

https://doi.org/10.1371/journal.pone.0175538.g004

and controls showing the highest ELISA signal. If low-glycosylated fractions of KIR4.1 are expressed in HEK293 cells and are detectable by ELISA, sera from MS patients should react against the surface of these cells, but we did not find any hint of reactivity. Brickshawana et al attempted to demonstrate surface staining of sera from MS patients against KIR4.1-transfected HEK293 cells [16]. However, although they demonstrated functional activity of the potassium channel and correct expression at the surface, KIR4.1 was encoded in the pEGFP-C2 expression vector. Expression of the fusion protein (KIR4.1-GFP) could, in theory, lead to epitope modifications and ultimately be the source of inconsistency with the original report. Finally, we performed ICC using patient sera over the MO3.13 human cell line, transfected with KIR4.1, and here again we failed to detect any relevant anti-KIR4.1 reactivity. The MO3.13 cell line was originated from the fusion of a human rhabdomyosarcoma line and adult human oligodendrocytes obtained from a surgical specimen. MO3.13 cells show features of immature oligodendrocytes and, under certain conditions, they can differentiate and express myelin proteins[37]. In fact, they are used as an in vitro model for human oligodendrocytes[38] and have been used for autoantibody screening in MS patients [39]. MO3.13 cells presumably recapitulate mature oligodendrocyte glycosylation patterns better than HEK293 cells. Therefore, the



KIR4.1 protein expressed on the MO3.13 surface should be a better antigen to detect relevant anti-KIR4.1 antibodies. Again, we did not find any surface staining in any patient or control using this system, ruling out a significant influence of the cell system in our KIR4.1-transfected HEK293 assay. There remains the possibility that the hybrid origin of the MO3.13 cells influences the glycosylation patterns of KIR4.1 and, thus the ability to detect anti-KIR4.1 antibodies in this system.

Finally, we developed a systematic review with the ultimate objective of performing a metaanalysis on the usefulness of anti-KIR4.1 antibodies. We included 13 studies with diverse methodological quality. The main methodological weakness of studies is that confounders were not considered (e.g. matching case and controls by age and sex) in the design or analysis. A publication bias was detected, with smaller studies showing no-association with anti-KIR4.1 and MS ( $\underline{S2}$  Fig). The meta-analysis was not performed, however, because the calculation of an  $\underline{I^2}$  test over the 75% limit showed high heterogeneity among the studies included.

In summary, we failed to identify anti-KIR4.1 antibodies in patients with MS using the full-length, low-glycosylated KIR4.1 ELISA assay and HEK293 and oligodendrocyte cell-based assays. Although technical caveats are still a matter of concern to detect antibodies targeting this complicated antigen, using very diverse approaches, evidence is accumulating against the confirmation of KIR4.1 as a relevant MS autoantigen.

# **Supporting information**

S1 PRISMA Checklist. PRISMA checklist disclosing information necessary for the standardization of the systematic review.

(DOC)

S1 Fig. This table summarizes all studies included in the systematic review and displays information about date and country of publication, n of patients and controls, technique used to detect anti-KIR4.1 antibodies, the study design and their methodological quality according to systematic review guidelines. MS: multiple sclerosis, OND: other neurological diseases, HC; healthy controls. (PDF)

**S2** Fig. Funnel plot comparison: Multiple sclerosis vs healthy people, outcome: Positive anti-KIR4.1. The funnel plot detects a publication bias, with smaller studies showing no-association with anti-KIR4.1 and MS. (PDF)

S1 Supplementary Methods. Description of additional immunocytochemistry experiments to demonstrate surface expression of recombinant Kir4.1 in transfected MO3.13 cells. (PDF)

S1 Table. Basic demographic features of patients with MS and controls included in the study. Epidemiological data other than gender were unavailable for 15 disease controls and average age reflects that of the ones in which information was available. MS: multiple sclerosis; HC: Healthy controls; OND: Other neurological diseases; RR: relapsing-remitting; SP: secondary progressive; PP: primary progressive.

(DOCX)

## **Acknowledgments**

We would like to thank Kevin O'Connor (Yale University, New Haven, CT), Rajneesh Srivastava and Bernhard Hemmer (Technische Universität, Munich, Germany) for kindly providing



antibodies and positive controls (RS, BH), KIR4.1 expression vectors (KOC) and technical advice (RS and KOC).

We also thank Novartis Spain for providing funds to complete this study.

## **Author Contributions**

Conceptualization: LQ AE.

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Formal analysis: LQ MN MST AVM MJMZ MLG.

Funding acquisition: LQ AE.

Investigation: LQ MN AVM MST MJMZ.

Methodology: LQ MN AVM MJMZ AE.

Project administration: LQ MN NV AE.

Resources: AE SF MST NV MLG.

Supervision: LQ AE MJMZ.

Visualization: LQ MN AVM MJMZ.

Writing - original draft: LQ MN AE AVM MJMZ.

Writing - review & editing: LQ MN AE AVM MJMZ.

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