



Original article

Q1 The impact of pathogen reduction on ABO isoagglutinin Q2 titers in apheresis platelets

Q3 Mikayel Yeghiazaryan^a, Yembur Ahmad^{ID a,b}, Jessie Singer^{ID a},
Vaanush Nazaryan^a, Craig Fletcher^{ID a,b}, Yamac Akgun^{ID a,b,*}

^a Department of Pathology and Laboratory Medicine, Children's Hospital Los Angeles, Los Angeles, CA, USA

^b Department of Pathology, Keck School of Medicine of USC, Los Angeles, California, USA

ARTICLE INFO

Article history:

Received 19 September 2024

Accepted 20 December 2024

Available online xxx

Keywords:

Pathogen reduction

Apheresis platelets

Isoagglutinin titers

Solid phase

ABSTRACT

Background: Platelet transfusions are a cornerstone of modern medical care, used across various clinical contexts. Ensuring the compatibility of blood products, especially regarding ABO isoagglutinins, is critical to minimize adverse reactions. Pathogen reduction technologies have been widely adopted to enhance the safety of blood products, however, the impact of such treatments on ABO isoagglutinin titers in platelet products remains unclear. **Methods:** This study analyzed 60 apheresis platelet donations, including type O, A, and B donors, using the INTERCEPT® Blood System for pathogen reduction. Samples were collected both from donor whole blood at the time of apheresis (Retention) and from the final pathogen-reduced platelet product after it had passed through the compound adsorption device (Post-CAD). ABO isoagglutinin titers, including both IgM and IgG classes, were measured using solid-phase technology on the NEO Iris platform.

Results: This study found a significant reduction in IgM isoagglutinin titers in Post-CAD samples, with 99% of Retention titers being greater than or equal to their Post-CAD counterparts. IgG titers exhibited more variability, with 9% of Post-CAD samples displaying higher titers than Retention samples. Statistical analysis confirmed differences between Retention and Post-CAD samples for both IgM and IgG titers, with p-values <0.05 in most comparisons.

Conclusion: Pathogen reduction using the INTERCEPT® Blood System effectively reduces ABO isoagglutinin titers in apheresis platelets, potentially lowering the risk of hemolytic transfusion reactions. This reduction is beneficial for safer out-of-group platelet transfusions, especially in vulnerable populations such as pediatric patients. These findings support the continued use of pathogen-reduced platelets in transfusion medicine to enhance both safety and availability of blood products.

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* Corresponding author. Yamac Akgun, Department of Pathology and Laboratory Medicine, Children's Hospital Los Angeles, Los Angeles, CA, USA.

E-mail address: akgunyamac@gmail.com (Y. Akgun).

<https://doi.org/10.1016/j.htct.2025.103840>

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1 Introduction

2 Platelet transfusions are a vital component of modern medi-
 3 cal care, utilized across various clinical settings ranging from
 4 trauma and surgery to the management of hematological dis-
 5 orders. The compatibility of blood products, including plate-
 6 lets, is paramount to ensure patient safety and the efficacy of
 7 treatment [1]. The ABO blood group system, characterized by
 8 the presence or absence of A and B antigens on red blood
 9 cells, plays a central role in blood transfusion compatibility
 10 [2]. In addition to these antigens, individuals also possess nat-
 11 urally occurring antibodies known as isoagglutinins, which
 12 are directed against the ABO antigens absent from their own
 13 blood [3]. ABO antibodies develop in individuals at 3–6
 14 months of age and reach adult levels at 5–10 years [4]. They
 15 are primarily IgM, and IgG antibodies often belonging to the
 16 IgG₂ subclass. Historically, ABO matching has been consid-
 17 ered less critical for platelet transfusions compared to red
 18 blood cell transfusions due to the lower expression of ABO
 19 antigens on platelets and the shorter lifespan in circulation
 20 [5]. As a result, out-of-group platelet transfusions, where the
 21 donor and recipient have different ABO blood groups, have
 22 been commonly practiced, especially in situations where
 23 ABO-matched platelets are unavailable or in high demand [6].

24 Despite the prevailing acceptance of out-of-group platelet
 25 transfusions, concerns regarding the potential risks associ-
 26 ated with ABO-incompatible platelet transfusions have
 27 prompted a reevaluation of transfusion practices [7]. One sig-
 28 nificant development in this regard is the recognition of ABO
 29 titers as a valuable tool in assessing the suitability of out-of-
 30 group platelet transfusions [8]. The semi-quantitative assess-
 31 ment of ABO isoagglutinin titers is valuable for evaluating the
 32 compatibility of blood products with the recipient and for
 33 minimizing the risk of adverse reactions such as hemolytic
 34 transfusion reactions [9]. Traditional methods for titration
 35 involve labor-intensive techniques such as tube agglutination
 36 or gel centrifugation. However, advancements in technology
 37 have introduced automated methodologies, offering high-
 38 throughput and standardized approaches for ABO isoaggluti-
 39 nin titration [10].

40 In addition, pathogen reduction of platelets represents an
 41 important advancement in blood safety technology and is
 42 increasingly being adopted by blood centers and hospitals
 43 worldwide [11]. Psoralen is a photosensitive compound that,
 44 when activated by ultraviolet (UV) light, forms cross-links
 45 between nucleic acids, thereby preventing replication and
 46 transcription of DNA and RNA in pathogens such as bacteria,
 47 viruses, and parasites [12]. Residual psoralen and byproducts
 48 are removed by adsorption via the compound adsorption
 49 device (CAD) to reduce toxicity [13].

50 Psoralen treatment of platelets is an effective method of
 51 pathogen reduction, yet its impact on the levels of ABO isoag-
 52 glutinins in the plasma of the final treated product is
 53 unknown. This study aims to evaluate ABO isoagglutinin
 54 titers (both IgM and IgG) in platelet donations using auto-
 55 mated solid-phase technology on the NEO Iris platform (Wer-
 56 fen, previously Immucor, Inc). By comparing titers between
 57 donor whole blood samples collected at the time of apheresis
 58 and final pathogen-reduced platelet product samples, this

study seeks to identify any changes in isoagglutinin levels 59
 that may occur during the manufacturing process. 60

Material and methods

61 Sixty apheresis platelet donations from 30 type O, 15 type A, 62
 and 15 type B donors collected using a Trima Accel Auto- 63
 mated Blood Collection System were analyzed. Each donation 64
 provided two samples: donor whole blood retention samples 65
 (Retention) collected in EDTA tubes at the time of apheresis, 66
 and final platelet product post-CAD samples (Post-CAD) col- 67
 lected after processing. 68

69 All platelets were treated with the pathogen reduction 69
 INTERCEPT® Blood System for Platelets System (Cerus 70
 Corp.). In the manufacturing process, platelets are ster- 71
 ilely transferred into a single-use processing set contain- 72
 ing amotosalen solution. The platelets are placed in an 73
 illumination device which delivers a controlled dose of 74
 ultraviolet A (UVA) light for each treatment, lasting 75
 approximately four minutes. After illumination, platelets 76
 are transferred to the bag containing the CAD and agi- 77
 tated for 6–24 h at room temperature. At completion of 78
 the CAD incubation, the platelets are transferred by grav- 79
 ity flow to the storage container in their final state as 80
 INTERCEPT platelets. The Post-CAD sample is then col- 81
 lected for testing. 82

83 Samples were tested for IgM and IgG classes of anti-A, 83
 anti-B, and anti-A/B isoagglutinins, totaling 360 individual 84
 tests split evenly between Retention and Post-CAD samples. 85
 ABO Isoagglutinin titers of both IgM and IgG classes were 86
 determined using solid-phase technology on the NEO Iris 87
 platform (Werfen, previously Immucor, Inc.) [14]. Initial IgM 88
 and IgG results were measured up to a dilution of 1:128. For 89
 IgG isoagglutinin titer results exceeding 128, reflex testing 90
 was conducted to establish titers up to a dilution of 1:2048. 91
 Automated protocols were unavailable for IgM titers above 92
 128. In the event of invalid automation results, the test was 93
 repeated up to two times and excluded from the study upon 94
 the third invalid result. 95

96 In the automated IgM protocol, 50 μ L of sample was seri- 96
 ally diluted to a dilution of 1:128, then incubated with 15 μ L of 97
 pooled A or B cells (2–4%, Immucor) for 10 min at 20 °C. The 98
 IgG protocol utilizes Capture-R® technology to detect IgG red 99
 blood cell antibodies. Pooled A or B cells (2–4%) were added 100
 to each well of Capture-R Select strips, followed by mixing 101
 with 50 μ L of system fluid. After centrifugation and washing 102
 steps, 50 μ L of system fluid and 100 μ L of sample were added 103
 and serially diluted up to 1:128. Subsequently, 100 μ L of Low 104
 Ionic Strength Saline (LISS) was added and incubated for 105
 15 min at 39 °C, followed by washing. Capture-R® Indicator 106
 cells (55 μ L) were added, and the plate was centrifuged, and 107
 read. Reflex testing employed the same IgG protocol at higher 108
 dilutions. 109

Statistical analysis

110 Since ABO titers represent non-continuous data, the results 111
 were transformed into titer steps using a logarithmic (log₂) 112

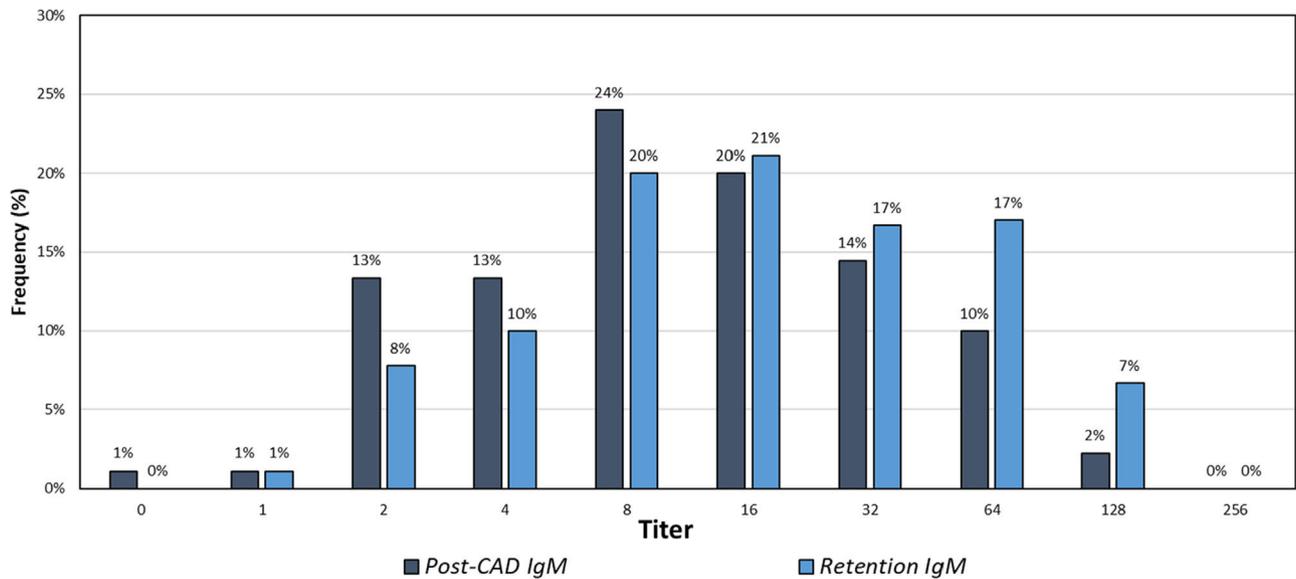


Figure 1 – Distribution of ABO IgM isoagglutinin titers for retention and post-CAD samples.

113 transformation. Specifically, a titer of 0 corresponded to -1, a
 114 titer of 1 corresponded to 0, and so forth, with each doubling
 115 dilution represented by a single digit increase in titer steps.
 116 The differences in titer steps between Retention and Post-
 117 CAD samples were then assessed to demonstrate if one sam-
 118 ple type produced higher or lower results. Meanwhile, the
 119 absolute difference was used to determine the percentage of
 120 concordance within ± 1 and ± 2 titer steps. To evaluate the sta-
 121 tistical significance between Retention and Post-CAD IgM and
 122 IgG results, a non-parametric Wilcoxon paired signed-rank
 123 test (using Z distribution) was employed, with a significance
 124 level set at 0.05.

Results

125

The distribution of titer results for IgM class isoagglutinin
 126 titers was close to symmetrical (Figure 1): Post-CAD results
 127 had a median titer of 8, mode of 8, and a sample coefficient of
 128 variation (CV) of 0.50 (in titer steps), and the Retention results
 129 had a median titer of 16, mode of 16, and CV of 0.42.
 130

The distribution for IgG class isoagglutinin titers (Figure 2)
 131 was asymmetrical: Post-CAD results had a median titer of 16,
 132 mode of 128, and CV of 0.65, and Retention results had a
 133 median of 32, mode of 128, and CV of 0.61. The data for IgG
 134

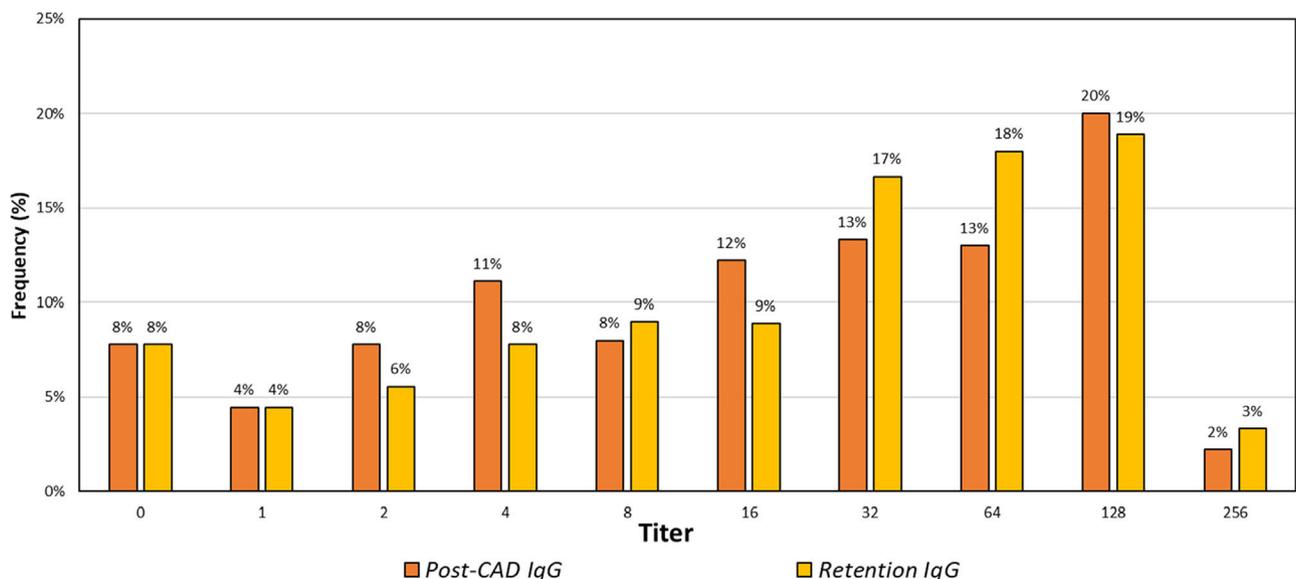


Figure 2 – Distribution of ABO IgG isoagglutinin titers for retention and post-CAD samples.

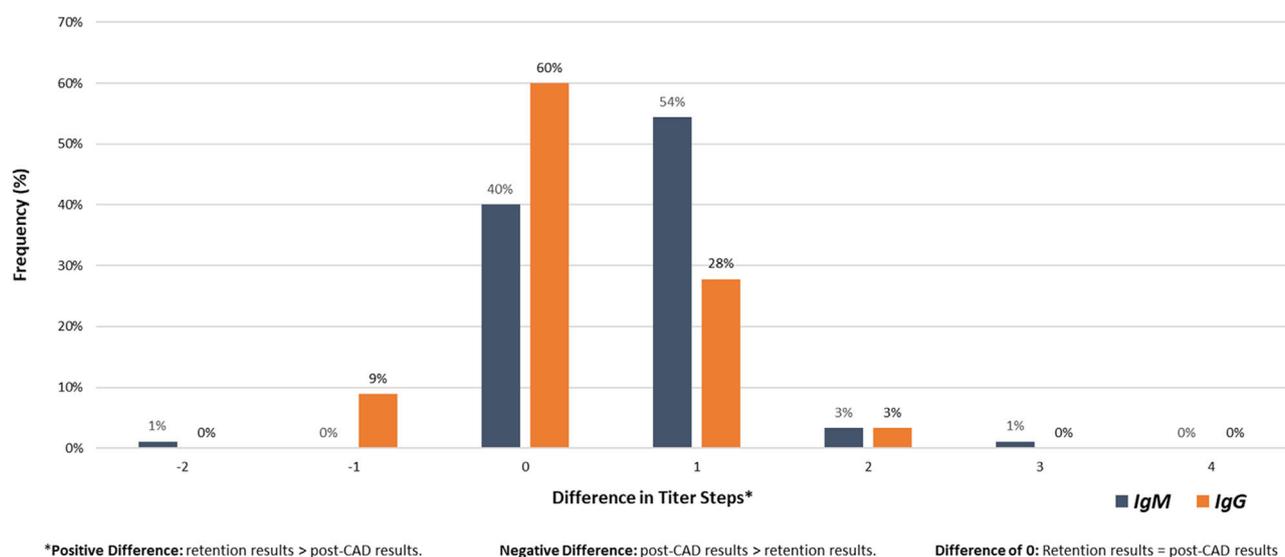


Figure 3 – Distribution of titer step differences (%) (retention – Post-CAD).

135 titers was more evenly distributed between titers 0 to 256, and
136 skewed towards higher results.

137 Comparing IgM Post-CAD and Retention samples, 99 % of
138 all Retention titer results were greater than or equal to their
139 Post-CAD counterparts (Figure 3). An exception to this was a
140 single outlier (1%) where the Post-CAD IgM result was two
141 titer steps greater than the corresponding Retention result. In
142 contrast to IgM results, IgG data had 9% of its Post-CAD sam-
143 ples yield higher titer results than the paired Retention sam-
144 ples.

145 Retention and Post-CAD samples were compared accord-
146 ing to type and isoagglutinin class (IgM or IgG) using Wilcoxon
147 paired signed-rank test. Overall, the p-values for both IgM and
148 IgG titers were <0.05, indicating significant differences
149 between the Post-CAD and Retention samples. Upon breaking
150 down the antibody classes by blood type, the p-values for
151 most comparisons indicated statistically significant differen-
152 ces between Post-CAD and Retention results (Table 1).

153 There were three cases in which the p-values exceeded
154 0.05: anti-A IgG for group O, anti-B IgG for group A, and anti-A
155 IgM for group B. In the last case, the high p-value was attrib-
156 uted to a single outlier which, when excluded ($n = 14$ for type
157 B) from the statistical test, yielded a p-value of less than 0.05
158 (p -value = 0.006). The percentage concordance between Reten-
159 tion and Post-CAD samples was calculated using titer steps.
160 All comparisons had 100% concordance within ± 2 titer steps.

161 Distribution of titers from all blood group results utilizing
162 maximum titer per sample was determined (Figure 4). Since
163 blood type O contains antibodies of both anti-A and anti-B
164 specificity, the higher of the two results was used to deter-
165 mine the maximum titer. For IgM, 83% of all Post-CAD sam-
166 ples and 69% of all Retention samples yielded titer results
167 ≤ 32 . For IgG, 65% of Post-CAD and 62% of Retention samples
168 yielded titer results ≤ 32 .

169 The distribution of the differences between IgG and IgM
170 (IgG-IgM) results was similar when comparing Post-CAD and
171 Retention samples. However, in both cases IgG isoagglutinin

172 testing generally yielded higher titer results than the IgM
173 counterparts. Refer to Table 1 to compare median titer results
174 between IgM and IgG class isoagglutinin titers.

175 Discussion

176 The ABO blood group system plays a pivotal role in transfu-
177 sion medicine, dictating the compatibility between donor and
178 recipient blood types [15]. In vulnerable patient populations
179 such as pediatric patients, ensuring compatibility is essential
180 to mitigating the risk of adverse reactions during platelet
181 transfusions [16]. This study investigated the quantitative
182 assessment of ABO isoagglutinin titers, focusing on both IgM
183 and IgG classes, in platelet donations using solid-phase tech-
184 nology [17]. By comparing titers between donor Retention
185 samples and final platelet Post-CAD samples, this study
186 aimed to evaluate the impact of the pathogen reduction
187 manufacturing process on isoagglutinin levels.

188 The results revealed notable differences in the distribution
189 of isoagglutinin titers between Retention and Post-CAD sam-
190 ples. Specifically, the median and mode titers for both IgM
191 and IgG classes differed between the two sample types. For
192 IgM isoagglutinin titers, Retention samples exhibited higher
193 median and mode titers compared to Post-CAD samples.
194 These differences suggest that the manufacturing process
195 influences isoagglutinin levels in platelet products. Specifi-
196 cally, 99% of IgM Retention titers were greater than or equal
197 to their Post-CAD counterparts, with only a single outlier. In
198 contrast, IgG results showed a more variable pattern, with 9%
199 of Post-CAD samples exhibiting higher titers than Retention
200 samples, and a greater percentage of equivalent results
201 between Retention and Post-CAD titers compared to IgM.

202 The observed reduction in isoagglutinin titers in Post-CAD
203 samples can be attributed to the pathogen reduction process
204 using the INTERCEPT® Blood System, which includes psoralen
205 treatment and UV light activation [18]. This process not only

Table 1 – Statistical results summary table.

Blood type	Sample n	Antibody specificity, class	Specimen source	Median	Mode	Min	Max	CV ^a	p value ^b	Concordance (%) ^c +/-1	Concordance (%) ^c +/-2
O	30	Anti-A, IgM	Retention	32	64	2	128	0.37	<0.05	93 %	100 %
			Post-CAD	16	64	2	64	0.41			
O	30	Anti-B, IgM	Retention	16	16	1	64	0.45	<0.05	100 %	100 %
			Post-CAD	8	8	0	64	0.49			
O	30	Anti-A, IgG	Retention	64	128	1	256	0.28	0.066	100 %	100 %
			Post-CAD	64	128	1	256	0.31			
O	30	Anti-B, IgG	Retention	32	128	2	128	0.36	<0.05	100 %	100 %
			Post-CAD	32	32	2	128	0.40			
A	15	Anti-B, IgM	Retention	8	8	2	128	0.50	<0.05	87 %	93 %
			Post-CAD	4	2	1	128	0.73			
A	15	Anti-B, IgG	Retention	1	0	0	8	3.32	0.484	100 %	100 %
			Post-CAD	1	0	0	8	2.50			
B	15 ^d	Anti-A, IgM	Retention	16	32	2	128	0.38	0.095 ^d	93 %	100 %
			Post-CAD	8	8	2	128	0.43			
B	15	Anti-A, IgG	Retention	8	32	0	64	0.60	<0.05	80 %	100 %
			Post-CAD	4	2	0	32	0.82			
All	60	Anti-A & Anti-B, IgM	Retention	16	16	1	128	0.42	<0.05	94 %	99 %
			Post-CAD	8	8	0	128	0.50			
All	60	Anti-A & Anti-B, IgG	Retention	32	128	0	256	0.65	<0.05	97 %	100 %
			Post-CAD	16	128	0	256	0.61			

^a Coefficient of variation was calculated using titer steps.

^b The results were compared using non-parametric Wilcoxon signed ranked test (using Z distribution) with a significance level at 0.05.

^c Concordance is calculated using absolute difference in titers steps between retention and post-CAD samples.

^d One of 15 Type B, IgM samples was an outlier with a post-CAD result two titer steps above its corresponding retention sample. This was the only pair of samples out of 60 pairs that had a higher IgM titer on the post-CAD sample than the retention sample. When the outlier is excluded (n = 14, for type B) from the analysis the p value is below 0.05 (p value of 0.006).

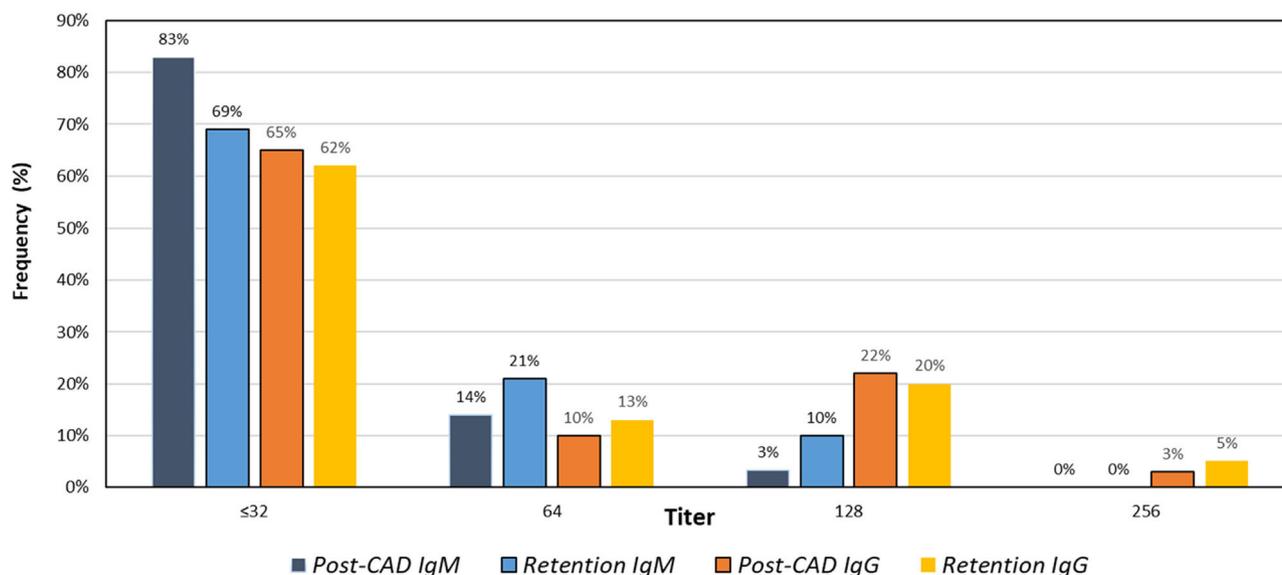


Figure 4 – Distribution of maximum titer results (anti-A or anti-B) per sample tested.

206 inactivates pathogens but may also impact the isoagglutinin
207 levels by binding and possibly denaturing these antibodies,
208 thereby reducing their effective concentration in the final
209 platelet product. This reduction is beneficial in minimizing
210 the risk of hemolytic transfusion reactions, particularly in
211 pediatric patients who are more susceptible to such adverse
212 events [19].

213 The statistical analysis, using the Wilcoxon paired signed-
214 rank test, indicated significant differences between Retention
215 and Post-CAD samples for both IgM and IgG titers, with p -
216 values <0.05 in most comparisons. This finding underscores the
217 consistent impact of the manufacturing process on reducing
218 isoagglutinin titers across different blood types and antibody
219 classes.

220 The percentage concordance within ± 2 titer steps was
221 100% for all comparisons, indicating a high degree of consis-
222 tency between Retention and Post-CAD samples, even with
223 the observed reductions in isoagglutinin titers. This high con-
224 cordance rate supports the reliability of the pathogen reduc-
225 tion process in maintaining relative titer levels while
226 reducing absolute concentrations.

227 Notably, three comparisons did not reach statistical signif-
228 icance: anti-A IgG for group O, anti-B IgG for group A, and
229 anti-A IgM for group B. The high p -value for anti-B IgG for
230 group A is likely due to the predominance of negative results
231 (mode of 0), which diminishes the potential for detecting sig-
232 nificant differences. After exclusion of the outlier in the anti-
233 A IgM for group B, the p -value reached statistical significance.
234 Anti-A IgG for group O had a p -value of 0.066, where a minor
235 increase in the sample size could make it significant.

236 The reduction in isoagglutinin titers after pathogen reduc-
237 tion highlights the potential for safer out-of-group platelet
238 transfusions with pathogen-reduced platelets, particularly in
239 settings where ABO-matched platelets are scarce [20]. By low-
240 ering the risk of hemolytic reactions, pathogen-reduced plate-
241 lets can be more safely used across different patient
242 populations, including vulnerable groups such as pediatric

243 patients. This aligns with current trends in transfusion medi- 243
244 cine that emphasize both safety and availability of blood 244
245 products [21]. Moreover, the use of solid-phase technology on 245
246 the NEO Iris platform for titer measurement offers a robust 246
247 and standardized approach for assessing isoagglutinin levels 247
248 [22]. This technological advancement facilitates high- 248
249 throughput, automated testing, enhancing the efficiency 249
250 and accuracy of compatibility assessments in transfusion 250
251 services. 251

252 While this study provides significant insights, it has limi- 252
253 tations that should be addressed in future research. The sam- 253
254 ple size, although adequate for demonstrating significant 254
255 differences, could be expanded to include a more diverse 255
256 range of donor demographics. Additionally, the impact of 256
257 other variables such as storage duration and donor health 257
258 status on isoagglutinin titers could be further investigated. 258

259 In conclusion, this study highlights the significant reduc- 259
260 tion in ABO isoagglutinin titers achieved through the patho- 260
261 gen reduction process, enhancing the safety profile of out-of- 261
262 group platelet transfusions. The use of advanced solid-phase 262
263 technology for titer assessment ensures precise and reliable 263
264 measurements, supporting informed clinical decision-mak- 264
265 ing [23]. These findings contribute to the evolving landscape 265
266 of transfusion medicine, promoting safer and more effective 266
267 blood product utilization. 267

Conflicts of interest

The authors declare no conflicts of interest.

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