

Improved clinical concordance of TIR and HbA1c by accounting for personal glycation factors

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	No, Nothing to disclose
X	Yes, please specify disclosures

Company / Name	Honoraria / Expense	Consulting / Advisory Board	Funded Research	Royalties / Patent	Stock Options	Ownership / Equity Position	Employee	Other (Please specify)
Abbott						Yes	Yes	

BACKGROUND

ADA Standards of Care¹ recognizes A1C can represent a large range of average glucose levels

Table 6.1—Estimated average glucose (eAG)				95% CI Range of Average Glucose (mg/dL)	95% CI Range of Average Glucose (mmol/L)	
A1C (%)	mg/dL*		mmol/L			
5	97	(76–120)	5.4	(4.2–6.7)	44	2.5
6	126	(100–152)	7.0	(5.5–8.5)	52	3.0
7	154	(123–185)	8.6	(6.8–10.3)	62	3.5
8	183	(147–217)	10.2	(8.1–12.1)	70	4.0
9	212	(170–249)	11.8	(9.4–13.9)	79	4.5
10	240	(193–282)	13.4	(10.7–15.7)	89	5.0
11	269	(217–314)	14.9	(12.0–17.5)	97	5.5
12	298	(240–347)	16.5	(13.3–19.3)	107	6.0

95% Confidence Intervals

1. *Diabetes Care* 2023;46(Suppl. 1):S97–S110

BACKGROUND

ADA Standards of Care¹ describes limitations of A1C to reflect average glycemia due to variations in **red blood cell lifespan**, and reports² indicate variation in **overall glycation rate**

“...hemolytic and other anemias, glucose-6-phosphate dehydrogenase deficiency, recent blood transfusion, use of drugs that stimulate erythropoiesis, endstage kidney disease, and pregnancy... **may result in discrepancies between the A1C result and the patient’s true mean glycemia**”

A1C has 3 major drivers:

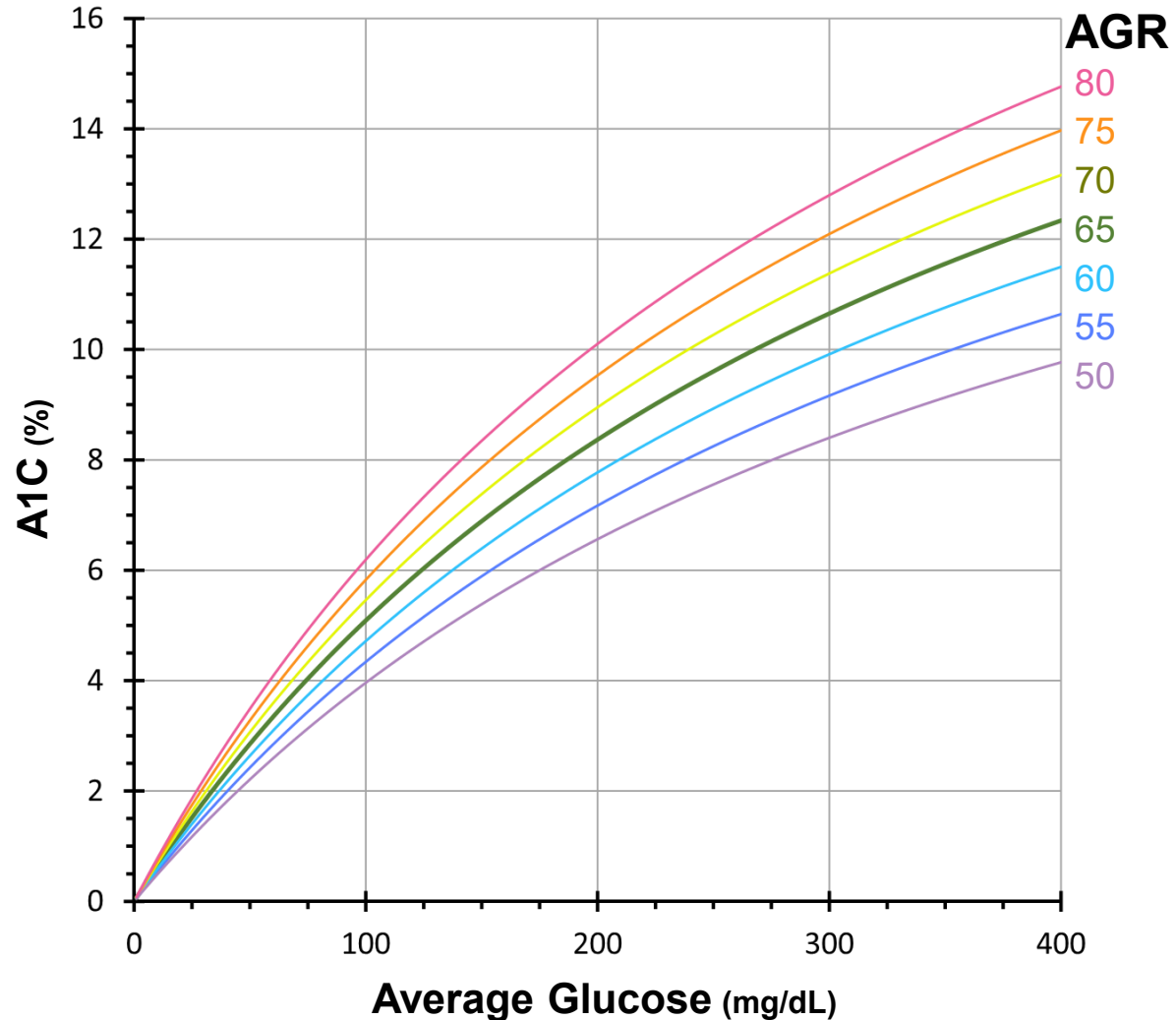
1. Average glucose
2. RBC lifespan
3. Overall Hb glycation

} Combined, these can be considered as:
“Apparent Glycation Rate”

1. *Diabetes Care* 2023;46(Suppl. 1):S97–S110
2. Ladyzynski et al. *J Trans Med.* 2014; 12:328

BACKGROUND

Using a novel model^[1-7] the AGR can be found with time-matched values of Average Glucose and A1C



$$\text{AGR} = \frac{\frac{1}{\text{Average Glucose}} + \frac{1}{K_M}}{\frac{1}{\text{A1C}/100} - 1} * 10^5$$

where:

$$K_M = 472 \text{ mg/dL} \text{ [8]}$$

- [1] Xu, et al. *J Diabetes Sci Technol*. 2021 Mar;15(2):294-302
- [2] Xu, et al. *Diabetes Technol Ther*. 2021 Jun;23(6):452-459
- [3] Xu, et al. *Diab Vasc Dis Res*. 2021 May-Jun;18(3)
- [4] Xu, et al. *elife*. 2021 Sep 13;10:e69456
- [5] Xu, et al. *Diabetes Obes Metab*. 2022 Sep;24(9):1779-1787
- [6] Xu, et al. *Diabetes Obes Metab*. 2022 Dec;24(12):2383-2390
- [7] Hirota, et al. *Diabetes Obes Metab*. 2023 Jan;25(1):319-322
- [8] Nishimura, et al. *J Biol Chem*. 1993;268(12):8514-8520.

BACKGROUND

Our novel model^[1-7] proposes a **Personalized A1C (pA1C)** adjusted to reflect average glucose levels in standard RBCs to match clinical trial findings

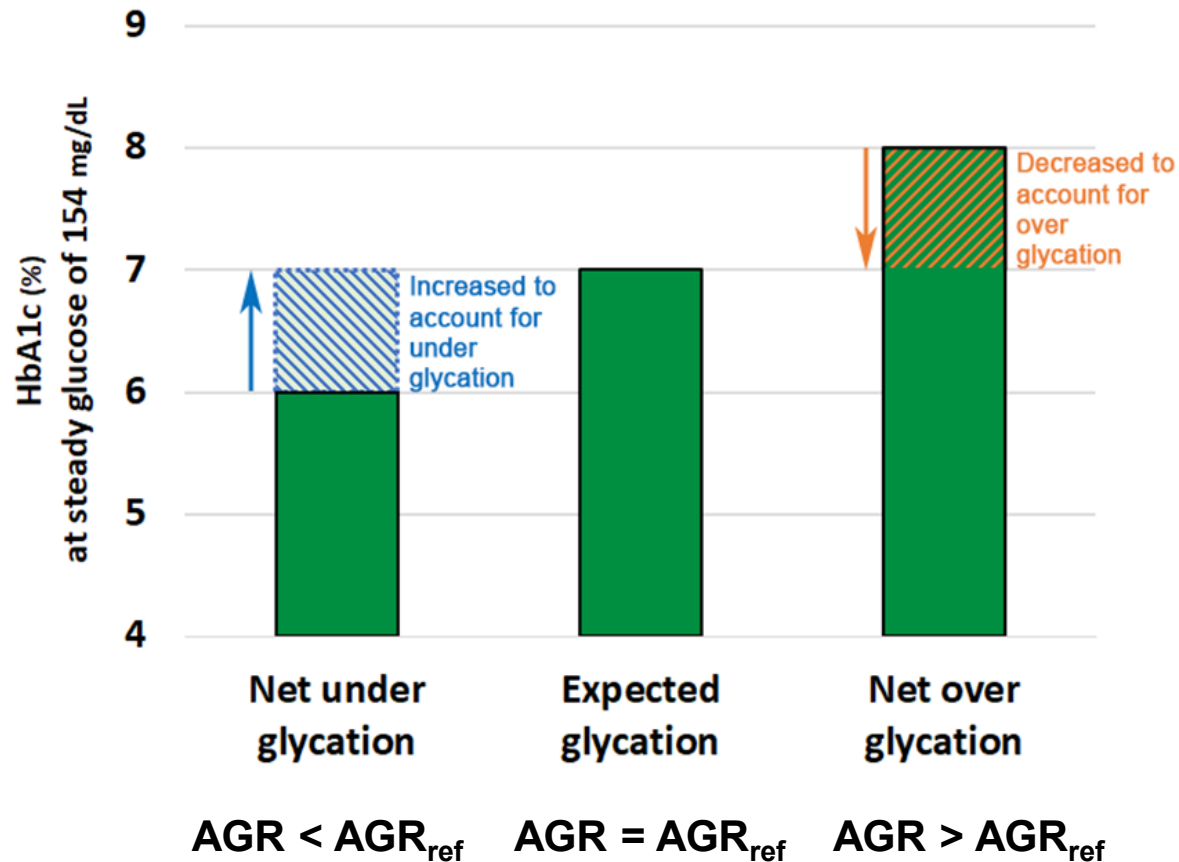
$$\text{Personalized A1C} = \text{pA1C} = \frac{100}{1 + \frac{\text{AGR}}{\text{AGR}_{\text{ref}}} \left(\frac{100}{\text{A1C}} - 1 \right)}$$

where:

$$\text{AGR}_{\text{ref}} = 65.1 \text{ dL/mg}$$

Standard RBC lifespan = 105 days^[8]

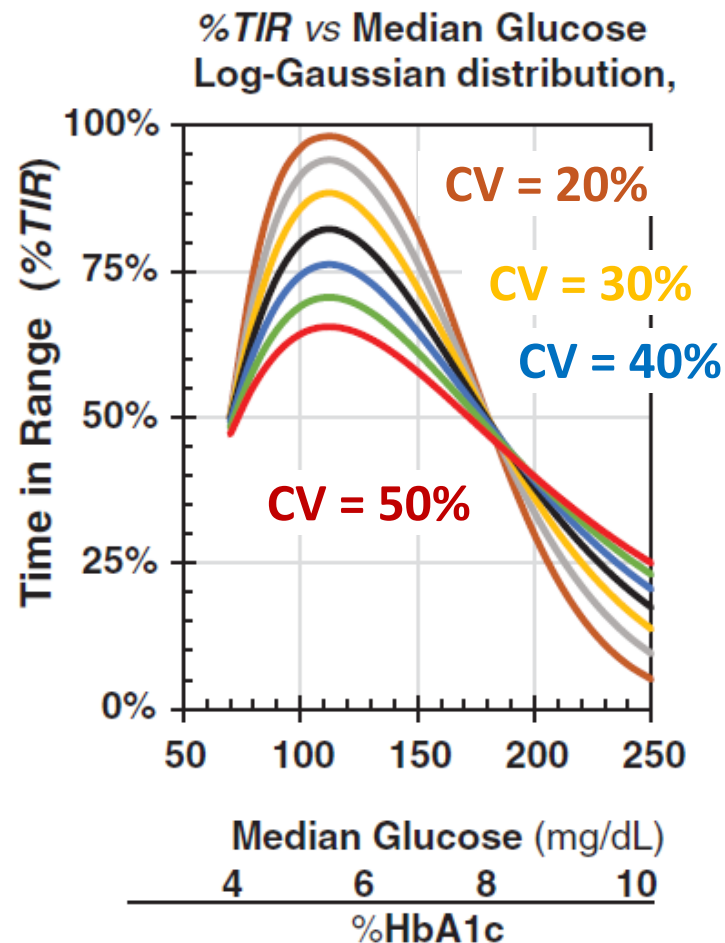
Standard RBC glycation rate constant = 6.2×10^{-6} dL/mg/day^[9]



- [1] Xu, et al. *J Diabetes Sci Technol*. 2021 Mar;15(2):294-302
- [2] Xu, et al. *Diabetes Technol Ther*. 2021 Jun;23(6):452-459
- [3] Xu, et al. *Diab Vasc Dis Res*. 2021 May-Jun;18(3)
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- [7] Hirota, et al. *Diabetes Obes Metab*. 2023 Jan;25(1):319-322
- [8] Khera, et al. *Am J Hematol*. 2015; 90:50-55
- [9] Ladyzynski et al. *J Trans Med*. 2014; 12:328

AIMS

- Evaluate improved A1C accuracy at reflecting in-range glucose exposure by adjusting for a personal RBC factor
- Investigate the effects of glycaemic variability on A1C accuracy.



- Previous work¹ has described the theoretical relationship between TIR and A1C given an underlying shape of the glucose distribution
- The TIR-A1C relationship should vary with glucose variability (CV)

METHODS

CGM and A1C data

- 3 months of CGM preceding A1C values
 - (n=1,636 periods, from 758 people in 7 clinical trials¹)
- All individuals had 2 to 5 repeated A1C-CGM periods to determine their AGR
- Calculated metrics:
 - TIR 70-180 mg/dL (90-day period)
 - Glucose CV (90-day period)
 - Personalized A1C

Statistical analysis

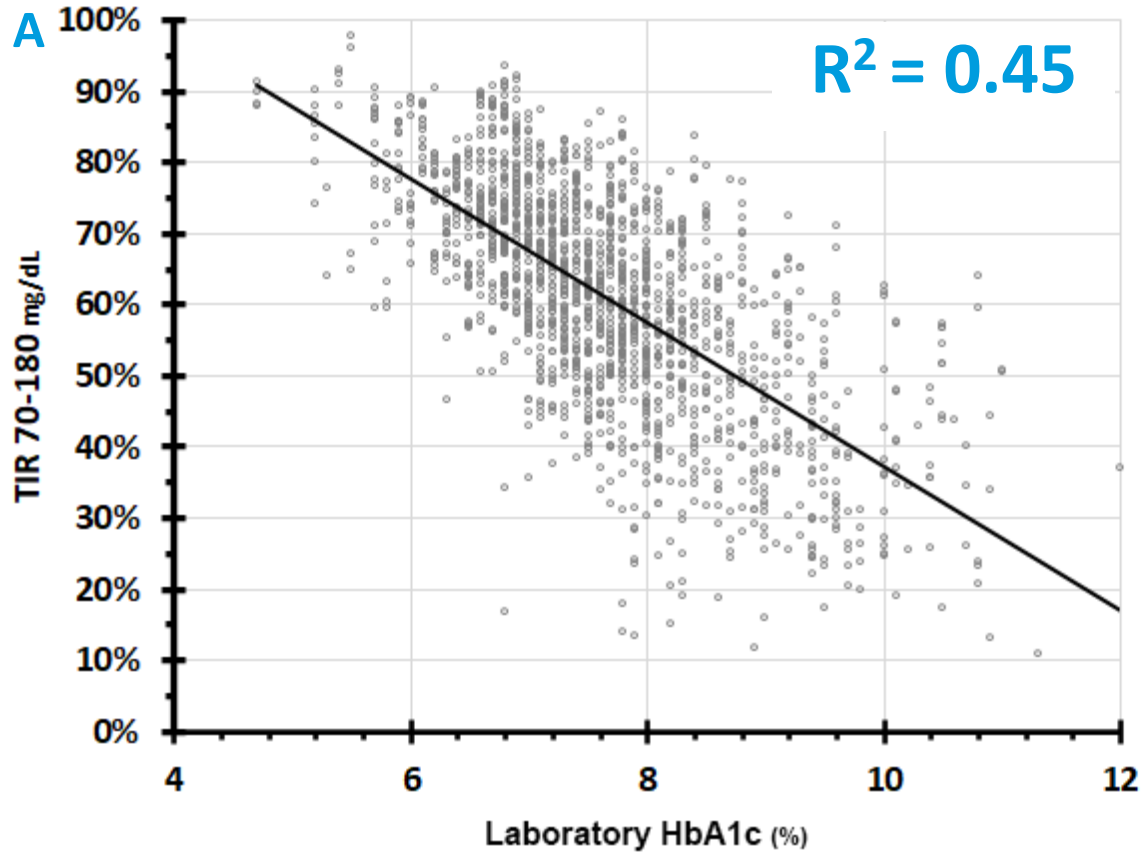
- Linear regression of TIR and A1C values
 - Overall
 - Subgroup by quartiles of glucose CV

1. Datasets available at Jaeb Center for Health Research (<https://www.jaeb.org/resources/>)

RESULTS 1 OF 2

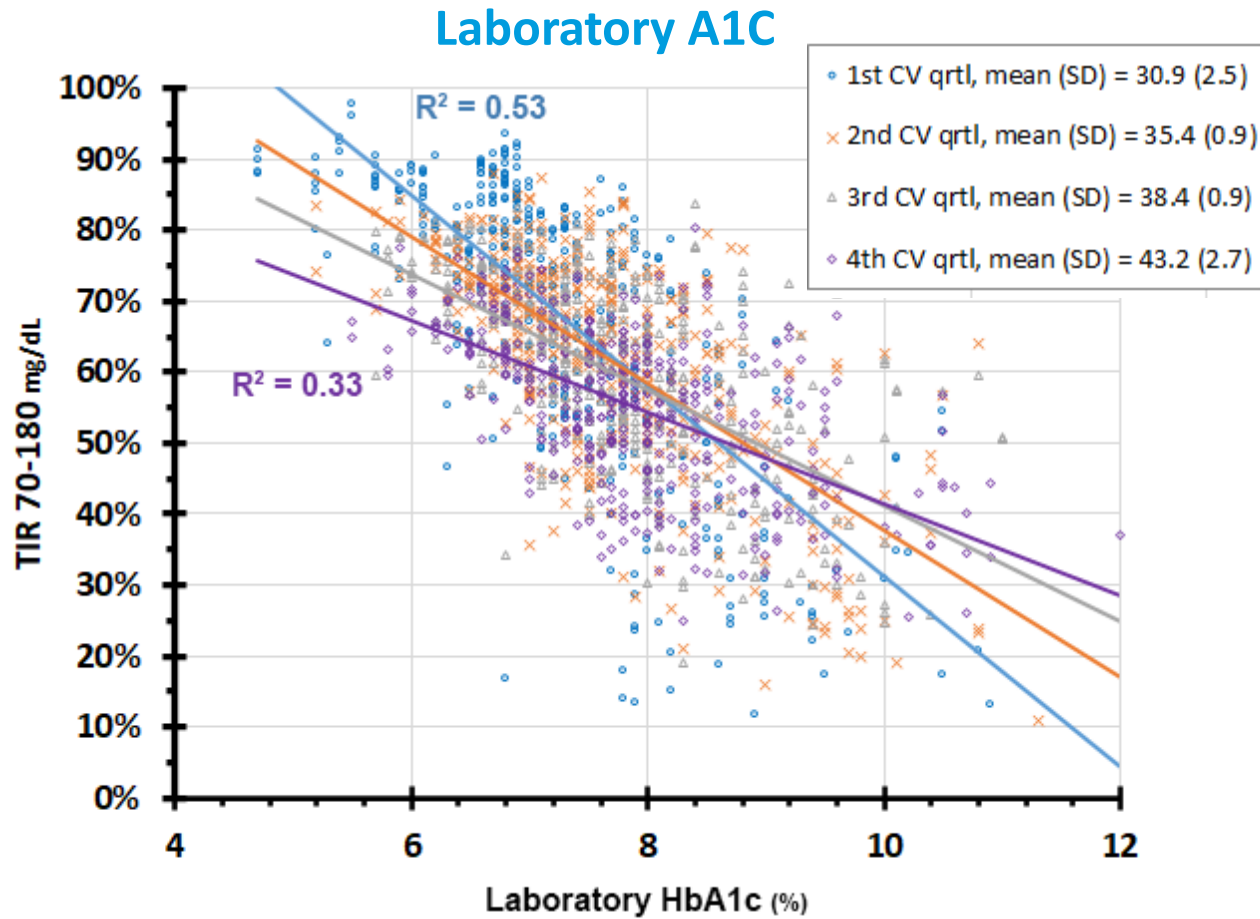
Substantially better agreement between TIR and Personalized A1C

Laboratory A1C



RESULTS 2 OF 2

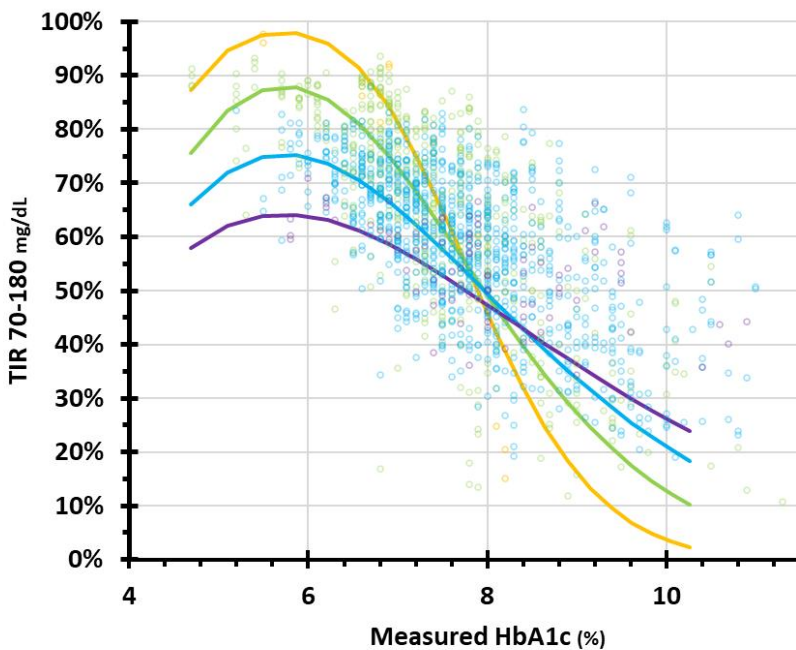
Agreement between TIR and Personalized A1C maintained at high CV



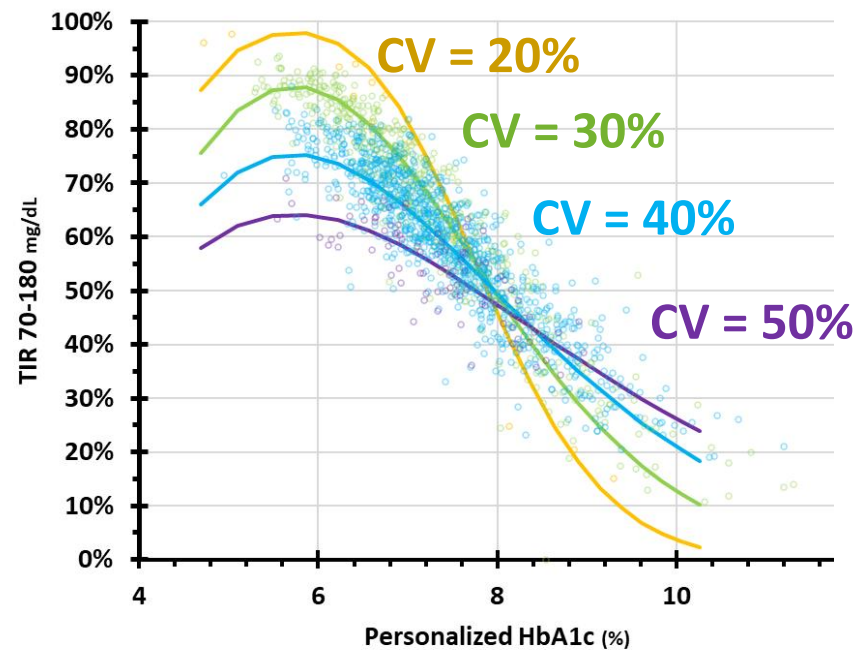
DISCUSSION

Agreement with gamma-shaped underlying glucose distribution¹

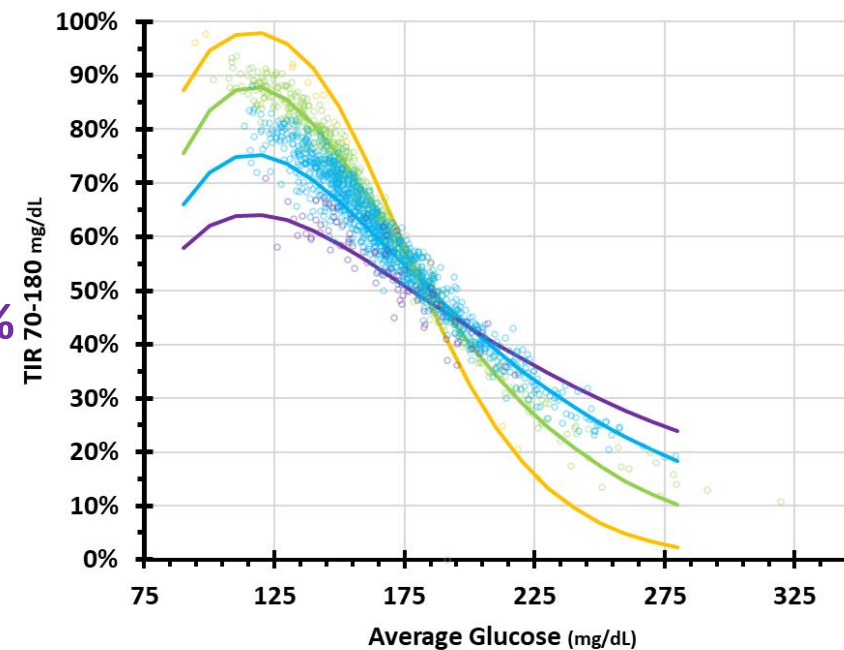
Laboratory A1C



Personalized A1C



Average Glucose



CV < 25%	0.9%
25% < CV ≤ 35%	34%
35% < CV ≤ 45%	60%
45% < CV	5%

1. Xu, Y et al. Glucose Variability Alters the Relationship between Average Glucose and Time In Range: Implications for Clinical Practice. *ATTD*, 2023, E-Poster 320

CONCLUSIONS

Accounting for a personal glycation factor improves the accuracy of A1C at reflecting glucose exposure even in the presence of high GV

- Adjustment to standard RBC factor improves agreement between TIR at Personalized A1C
- Personalization of A1C supports consistent clinical decision-making when reviewing both A1C and CGM measures



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