

Roy J¹, Morfini M², Ignas D¹, Carcao M¹, Barnes C³, Zhao X Y¹, Tanoshima R¹, Ito S¹, Blanchette V¹ and Bjorkman S⁴.

1. Hospital for Sick Children, Toronto, Canada; 2. University Hospital of Florence, Florence, Italy; 3. Royal Children's Hospital, Melbourne, Australia; 4. Uppsala University, Uppsala, Sweden

INTRODUCTION

- Pharmacokinetic (PK) studies provide better estimation of drug activity than patient characteristics alone¹
- PK of recombinant FVIII (rFVIII) traditionally use the:

ISTH 11-point protocol²: after a 72h washout administer 25-50 U/kg of rFVIII 11 samples taken at: 0, 15, 30, 60 mins AND 3, 6, 9, 24, 28, 32, 48 h
For Children do 5 point sampling instead at: 0, 1, 10, 24, 48 h
Non-compartmental model used.

- Population PK model has been proposed instead¹
 - no washout required
 - fewer samples taken
 - Bayesian forecasting
 - Excellent correlation with ISTH method
- Point-of-care software now available to do population PK

METHODS

- Data from previously published cohort of 20 boys with Haemophilia A³ was used as basis for study. That study evaluated PK of Kogenate-FS using the ISTH 11-point protocol after administering 50 IU/kg rFVIII

- From this a Population PK model was built by S. Bjorkman
 - $t_{1/2}$ and CI: correlated with vWF:Ag ($p < 0.01$) and not with patient age, weight, or mutation type

- For this study PK parameters ($t_{1/2}$, CI) were re-calculated from study described above using:

- All 11 data points and non-compartmental model
 - Population PK model with Bayesian analysis:
 - 11, 5 or 2 (24 & 28h or 28 & 32h) data points
- All these approaches to PK analysis were compared

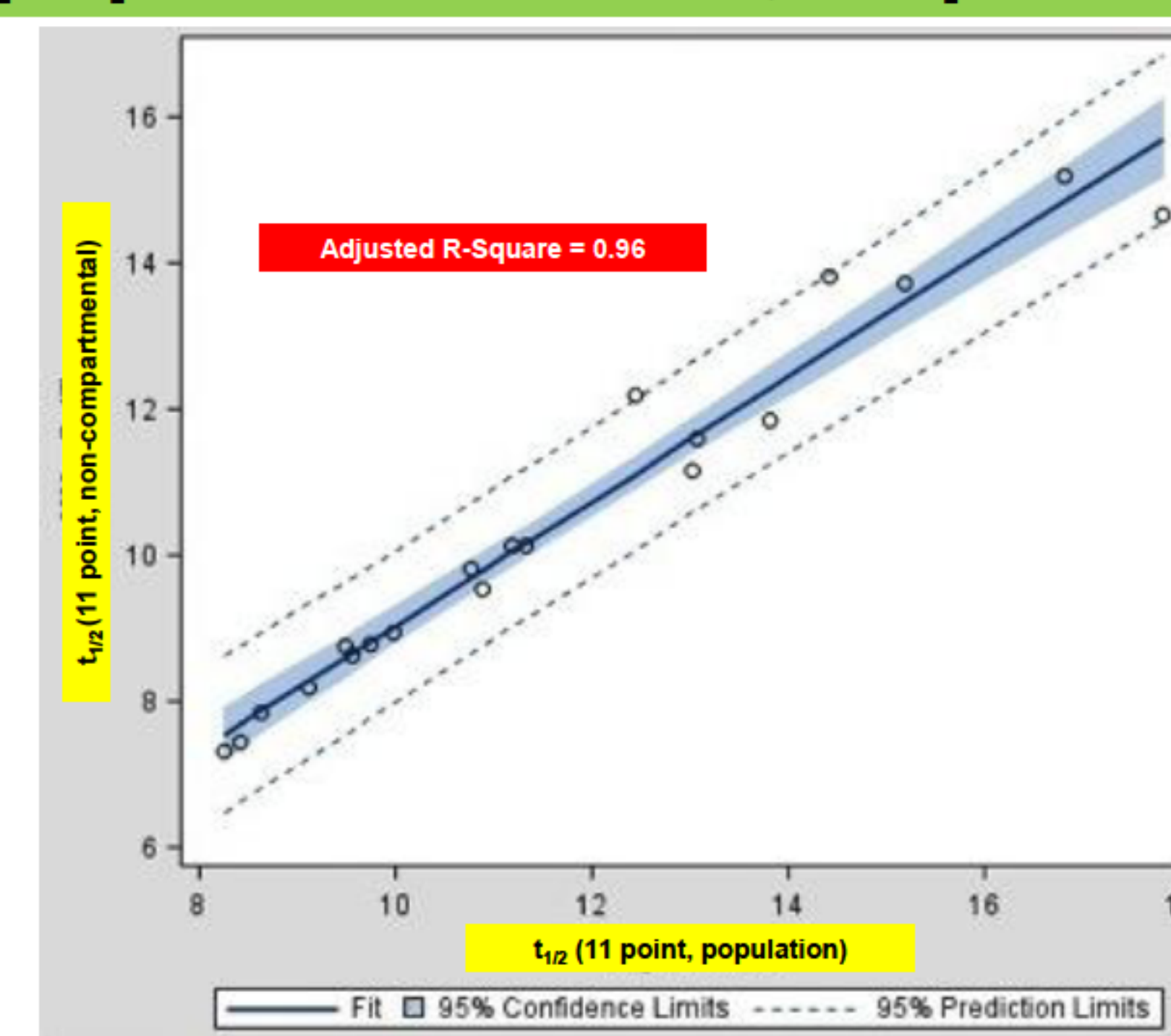
RESULTS

Variable	Interclass correlation coefficient	95% CI
Traditional vs Population PK Model (using all 11 sampling points)		
Clearance (CI)	0.97	0.92 – 0.99
Half-life ($t_{1/2}$)	0.97	0.93 – 0.99
Comparing number of sampling points (using Population PK model)		
5 vs 11, CI	0.999	0.997 – 0.999
5 vs 11, $t_{1/2}$	0.99	0.98 – 0.997
2 (24/28) vs 11, CI	0.95	0.87 – 0.98
2 (24/28) vs 11, $t_{1/2}$	0.81	0.58 – 0.92
2 (28/32) vs 11, CI	0.92	0.81 – 0.97
2 (28/32) vs 11, $t_{1/2}$	0.84	0.62 – 0.93

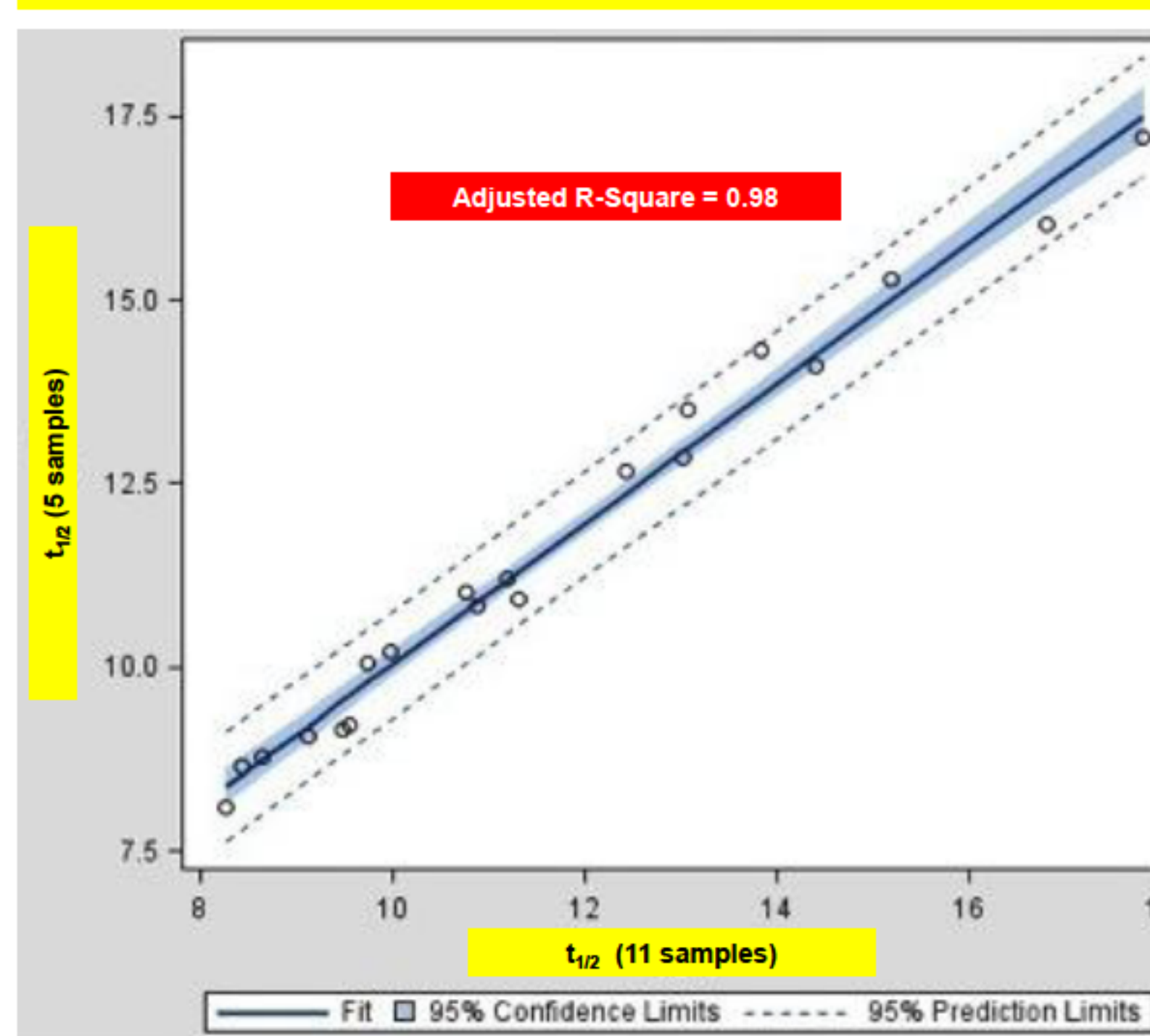
Interclass coefficient
0.7-0.8 = strong agreement
>0.8 = almost perfect agreement

Dataset from Barnes et al³:
median age 12.8y (4.4-18.1)
FVIII:C <1% n=15
FVIII:C 1-6% n=5

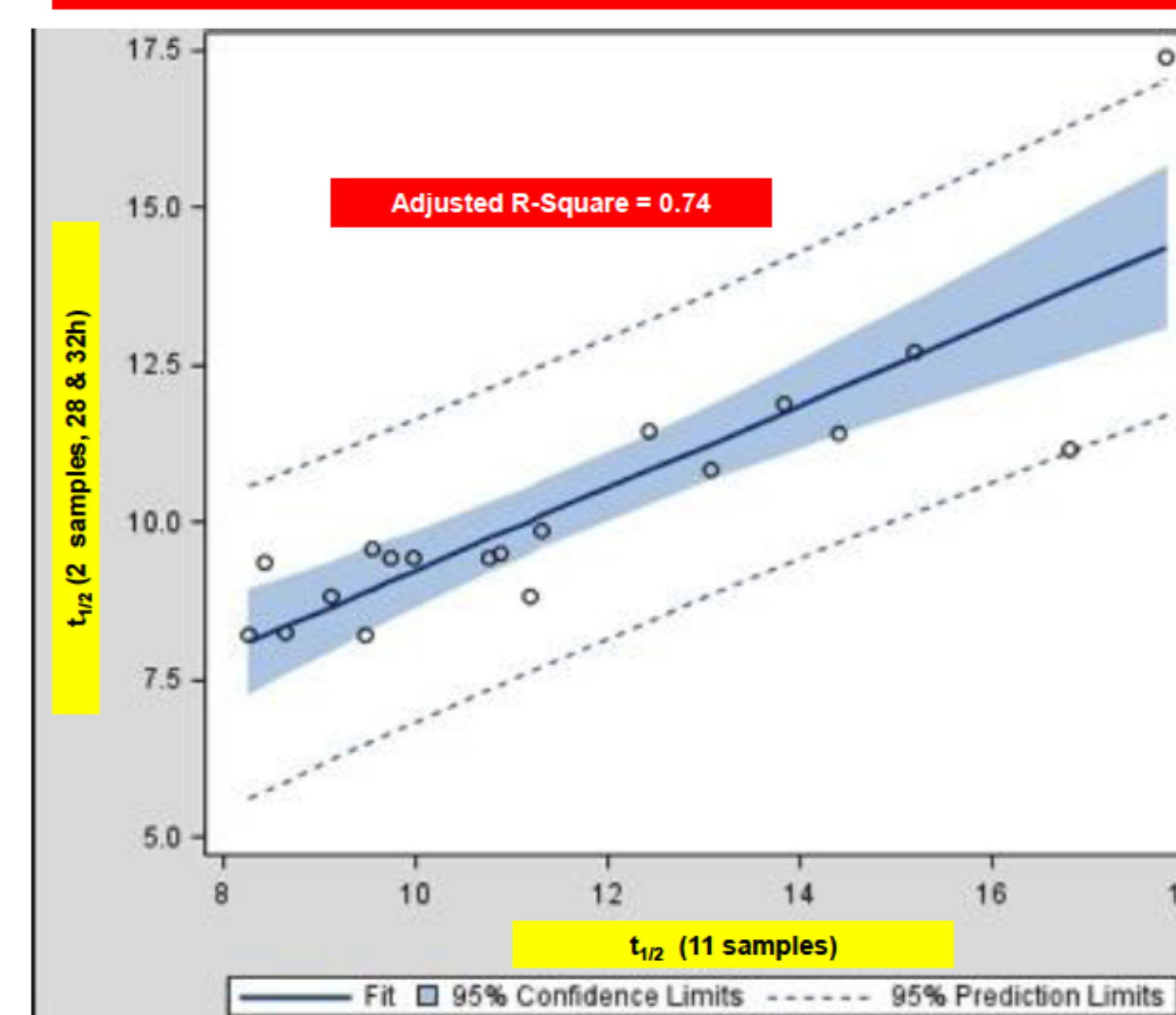
Non-compartmental vs population model, 11 points



11 vs 5 sampling points



11 vs 2 sampling points



CONCLUSIONS

- Bayesian analysis of Population PK model:
 - yields clinically similar data to traditional PK 11-point ISTH model
 - is more practical
 - no washout is required
 - fewer samples needed
 - requires prospective validation
 - of model
 - of point-of-care software, including dosing estimates

REFERENCES

1. BJÖRKMAN S. Haemophilia 2010; 119: 612–8; 2. Lee M, et al. ISTH Website 21 March, 2001; 3. Barnes C et al. Haemophilia; 2006; 12: 40–9.