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## Supplementary information

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# Thrombin generation test based on a 96-channel pipettor for evaluation of FXIa procoagulant activity in pharmaceuticals

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## **Supplementary Information**

**Title:** Thrombin generation test based on a 96 channel pipettor for evaluation of FXIa procoagulant activity in biopharmaceuticals

**Authors:** Leonid A. Parunov\*, Maria E. Shea\*, Yideng Liang\*, Stepan S. Surov, Maitreyi Chattopadhyay, Timothy K. Lee, Dorothy E. Scott, and Mikhail V. Ovanesov

\*Equal contribution

**Affiliation:** Center for Biologics Evaluation and Research, U.S. Food and Drug Administration, Silver Spring, Maryland, United States of America

**Table S1. Description of worksheets generated by running AutoTGT**

| Worksheets after AutoTGT | Description                                                                             |
|--------------------------|-----------------------------------------------------------------------------------------|
| AvrThrombin              | TG curves averaged across duplicates with standard deviations.                          |
| DatAMC                   | Fluorescence data after smoothing and drop correction and with AMC calibration applied. |
| DatAUC                   | Fluorescence data after smoothing and drop correction and with AMC calibration applied  |
| DatRFU                   | Fluorescence data after smoothing and drop correction                                   |
| DatThrombin              | TG curves, separate for each                                                            |
| DatVelocity              | Velocity taken from TG curves                                                           |
| RawData                  | Is not used in this version of script                                                   |
| ResultsTG                | Thrombin parameters averaged across duplicate wells                                     |
| ResultsTGall             | Thrombin parameters separate for each well                                              |
| USERcalibrDT             | Is not used in this version of script                                                   |
| WellDataFG               | Is not used in this version of script                                                   |
| WellDataTG               | Enter fluorescence data here for thrombin generation                                    |
| WKSHTResults             | Is not used in this version of script                                                   |
| Temp*                    | Temporary worksheets                                                                    |

**Table S2. Description of worksheet columns in ResultsTG.** Note that the SD suffix refer to the reported values standard deviation when there are duplicate wells for a sample.

| <b>ResultsTG Column Heading</b> | <b>Description</b>                                                                               |
|---------------------------------|--------------------------------------------------------------------------------------------------|
| <b>NSample</b>                  | Sample number                                                                                    |
| <b>SampleName</b>               | Sample name                                                                                      |
| <b>totalTG.1 (TG.1'SD)</b>      | AUC (area under the curve) up to first time cutoff (default 20 minutes)                          |
| <b>totalTG.2 (TG.2'SD)</b>      | AUC up to second time cutoff (default 40 minutes)                                                |
| <b>totalTG.3 (TG.3'SD)</b>      | AUC up to third time cutoff (default 60 minutes)                                                 |
| <b>totalTG.4 (TG.4'SD)</b>      | AUC up to fourth time cutoff (default 90 minutes).                                               |
| <b>maxTG (maxTG.SD)</b>         | Maximum thrombin generation, used to calculate potency TPH (thrombin peak height).               |
| <b>TimemxTG (TimemxTGSD)</b>    | Time of maximum thrombin generation, (time to thrombin peak).                                    |
| <b>maxTGrate (maxTGrateSD)</b>  | Maximum rate of thrombin generation, (velocity).                                                 |
| <b>timeMaxTGrate (tiMaTGSD)</b> | Time of max TG rate, (time to maximum velocity)                                                  |
| <b>Time.10nM (Time.10nMSD)</b>  | Time to 17% of thrombin generation curve (default), referred to as lag time, LagTime or LT later |

**Table S3. Description of worksheet columns in ResultsTGall.** Note that .avr and .SD suffixes refer to the averaged values and the average's standard deviation.

| <b>ResultsTGall Column Heading</b>           | <b>Description</b>                                                                                                  |
|----------------------------------------------|---------------------------------------------------------------------------------------------------------------------|
| <b>SampleN</b>                               | Sample number                                                                                                       |
| <b>SampleName</b>                            | Sample name                                                                                                         |
| <b>AUC.1 (AUC.1.avr, AUC.1.SD)</b>           | AUC (area under the curve) up to first time cutoff (default 20 minutes)                                             |
| <b>AUC.2 (AUC.2.avr, AUC.2.SD)</b>           | AUC up to second time cutoff (default 40 minutes)                                                                   |
| <b>AUC.3 (AUC.3.avr, AUC.3.SD)</b>           | AUC up to third time cutoff (default 60 minutes)                                                                    |
| <b>AUC.4 (AUC.4.avr, AUC.4.SD)</b>           | AUC up to fourth time cutoff (default 90 minutes). This value is used as the AUC value for the activity calculation |
| <b>TpeakHeight (TGpeakH.avr, TGpkMax.SD)</b> | Maximum thrombin generation, used to calculate activity using TPH (thrombin peak height)                            |
| <b>TtoPeak (TtoPeak.avr, TtoPeak.SD)</b>     | Time of maximum thrombin generation, used to calculate activity using TTP (time to thrombin peak)                   |
| <b>VelocityMax (VelocMx.avr, VelocMx.SD)</b> | Maximum rate of thrombin generation, used to calculate activity using VEL (velocity)                                |
| <b>TtoVMax (TtoVmax.avr, TtoVmax.SD)</b>     | Time of max TG rate, used to calculate activity using TVL (time to maximum velocity)                                |
| <b>LagTime (LagTime.avr, LagTime.SD)</b>     | Time to 17% of thrombin generation curve (default) used to calculate activity using LT (lag time)                   |

**Table S4. Description of worksheets generated by running aPotency.**

| <b>Worksheets after running aPotency</b> | <b>Description</b>                                                                                                                |
|------------------------------------------|-----------------------------------------------------------------------------------------------------------------------------------|
| <b>PotencyALL</b>                        | Activities of each sample calculated using different parameters ( PH, TTP, LT, AUC, VEL, and TVL ) are combined in one worksheet. |
| <b>PotencyAUC</b>                        | Activity calculations using area under the curve (AUC at 90 minutes, AUC.4)                                                       |
| <b>PotencyLT</b>                         | Activity calculations using lag time                                                                                              |
| <b>PotencyTPH</b>                        | Activity calculations using thrombin peak height                                                                                  |
| <b>PotencyTTP</b>                        | Activity calculations using time to peak                                                                                          |
| <b>PotencyTVL</b>                        | Activity calculations using time to maximum velocity                                                                              |
| <b>PotencyVEL</b>                        | Activity calculations using maximum velocity                                                                                      |
| <b>SerialDilAUC</b>                      | Area under the curve values for plotting                                                                                          |
| <b>SerialDilLT</b>                       | Lag time values for plotting                                                                                                      |
| <b>SerialDilTPH</b>                      | Thrombin peak height values for plotting                                                                                          |
| <b>SerialDilTTP</b>                      | Time to peak values for plotting                                                                                                  |
| <b>SerialDilTVL</b>                      | Time to maximum velocity values for plotting                                                                                      |
| <b>SerialDilVEL</b>                      | Maximum velocity values for plotting                                                                                              |

**Table S5. Description of graphs generated by running aPotency.**

| <b>Graphs after running aPotency</b>              | <b>Description</b>                                                                                                                                                                                                    |
|---------------------------------------------------|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| <b>_PotencyALL</b>                                | Activities of each sample calculated from TPH, TTP, LT, AUC, VEL, and TVL parameters plotted as a bar graph.                                                                                                          |
| <b>_SerialDilALL</b>                              | Parameter values of each sample plotted against the sample's dilution fold. Points are masked if excluded from activity analysis.                                                                                     |
| <b>_SerialDilAUC</b>                              | Area under the curve values of each sample plotted against the sample's dilution fold.                                                                                                                                |
| <b>_SerialDilLT</b>                               | Lag time values of each sample plotted against the sample's dilution fold.                                                                                                                                            |
| <b>_SerialDilTPH</b>                              | Thrombin peak height values of each sample plotted against the sample's dilution fold.                                                                                                                                |
| <b>_SerialDilTTP</b>                              | Time to peak values of each sample plotted against the sample's dilution fold.                                                                                                                                        |
| <b>_SerialDilTVL</b>                              | Time to maximum velocity values of each sample plotted against the sample's dilution fold.                                                                                                                            |
| <b>_SerialDilVEL</b>                              | Maximum velocity values of each sample plotted against the sample's dilution fold.                                                                                                                                    |
| <b>_StdCurvesAll</b>                              | Parameter values of each Standard and the fitted standard curve plotted against the Standard's dilution fold. Masking points on this graph and refreshing the standard curves will exclude these points from fitting. |
| <b>_TGcurves[<i>your Standard Lot number</i>]</b> | Thrombin generation curves for the standard are plotted.                                                                                                                                                              |
| <b>_TGcurvesALL</b>                               | Thrombin generation curves for the standard and the samples are plotted side-by-side.                                                                                                                                 |
| <b>_TGcurvesSmpl1</b>                             | Thrombin generation curves for the samples are plotted together.                                                                                                                                                      |

**Table S6. Description of worksheets columns in PotencyAUC.**

These column descriptions are the same for the PotencyLT, PotencyVEL, PotencyTTP, PotencyTPH, and PotencyTVL worksheets

| PotencyAUC (TPH,TTP, etc.)<br>Columns | Description                                                                                                                                                                                             |
|---------------------------------------|---------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| Concntrtn                             | Concentration of standard                                                                                                                                                                               |
| StdCurve (StdCurveSD)                 | Parameter value at given concentration of standard                                                                                                                                                      |
| ConcCorr                              | Log10 of concentration of standard                                                                                                                                                                      |
| StdCurveCorr                          | Value of standard at Conccorr. Depending on settings, background may be subtracted, and log transformation applied                                                                                      |
| SDCorr                                | Standard deviation of StdCurveCorr. Depending on settings, background may be equal to StdCurveSD or corrected when background is subtracted (calculated as standard deviation of two random variables). |
| StdCurveLog                           | Log transformed StdCurveCorr                                                                                                                                                                            |
| Residual                              | Residual from fit (difference between fit and observed value)                                                                                                                                           |
| RelativeResi                          | Relative residual (ratio of residual to observed value)                                                                                                                                                 |
| ConcFit                               | Concentration of standard for fitted curve. Depending on settings may be expressed in linear or logarithmic values.                                                                                     |
| StdCurveFit                           | Value of fitted standard curve at ConcFit                                                                                                                                                               |
| FitParameter                          | Fit parameter names                                                                                                                                                                                     |
| FitParValue                           | Fit parameter values                                                                                                                                                                                    |
| YStd                                  | Concentration of standard for all points on standard curve. Depending on settings may be expressed in linear or logarithmic values.                                                                     |
| XStdSignal                            | Value of fitted standard curve at given YStd                                                                                                                                                            |
| YStdLogC                              | Log10 of concentration of standard for all points on standard curve                                                                                                                                     |

|              |                                                                                  |
|--------------|----------------------------------------------------------------------------------|
| YStdConc     | Concentration of standard for all points on standard curve                       |
| XStdSignalR  | Reverse transformation from logarithmic to linear scale for XStdSignal           |
| WellN        | Number of well                                                                   |
| WellName     | Name of well                                                                     |
| WellValue    | Parameter value for given well                                                   |
| WellPotency  | Calculated activity (or potency) (unadjusted) for given well                     |
| WellDilution | Dilution fold for given well                                                     |
| WPotByDil    | Activity multiplied (WellPotency) by dilution fold (WellDilution) for given well |
| SampleN      | Number of sample                                                                 |
| SampleName   | Name of sample                                                                   |
| NReplicate   | Number of replicates for each sample                                             |
| SampleAvr    | Average value for the parameter of sample                                        |
| SampleSD     | Standard deviation of the parameter of sample                                    |
| SPotencyAvr  | Averaged activity for each sample                                                |
| SPotencySD   | Standard deviation of the activity for each sample                               |
| LotDilN      | Number of sample                                                                 |
| LotNumber    | Name of lot (pooled samples of different dilutions)                              |
| Dilutions    | Dilution factor for each sample                                                  |
| PotByDil     | Averaged activity (SPotencyAvr) multiplied by dilution fold (Dilutions)          |
| LotName      | Name of each lot                                                                 |
| POTENCY      | Activity averaged for all dilutions in a lot                                     |
| POTENCYSD    | Standard deviation of averaged activity                                          |
| CV           | Coefficient of variation (standard deviation/average activity)                   |
| N            | Number of values (dilutions) averaged to find the activity                       |

**Table S7. Description of worksheet columns in PotencyALL.** Note that the SD suffix refer to the reported activity's standard deviation.

| PotencyALL Column Heading | Description                                        |
|---------------------------|----------------------------------------------------|
| <b>LotName</b>            | Name of Sample Lot                                 |
| <b>TPH (TPH.SD)</b>       | Activity calculated using thrombin peak height     |
| <b>TTP (TTP.SD)</b>       | Activity calculated using time to peak             |
| <b>LT (LT.SD)</b>         | Activity calculated using lag time                 |
| <b>AUC (AUC.SD)</b>       | Activity calculated using area under the curve     |
| <b>VEL (VEL.SD)</b>       | Activity calculated using maximum velocity         |
| <b>TVL (TVL.SD)</b>       | Activity calculated using time to maximum velocity |
| <b>Avr (Avr.SD)</b>       | Average activity across all parameters             |