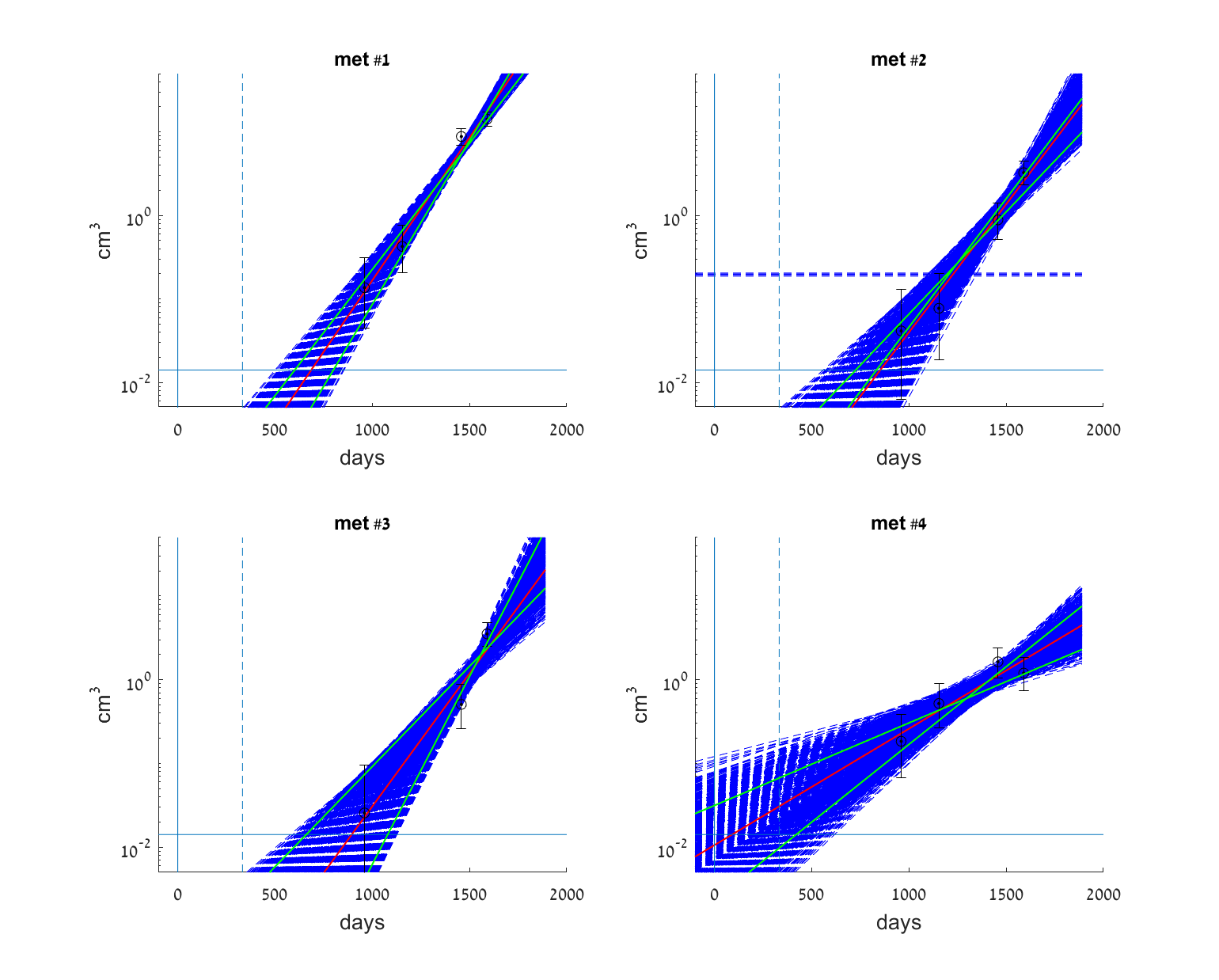
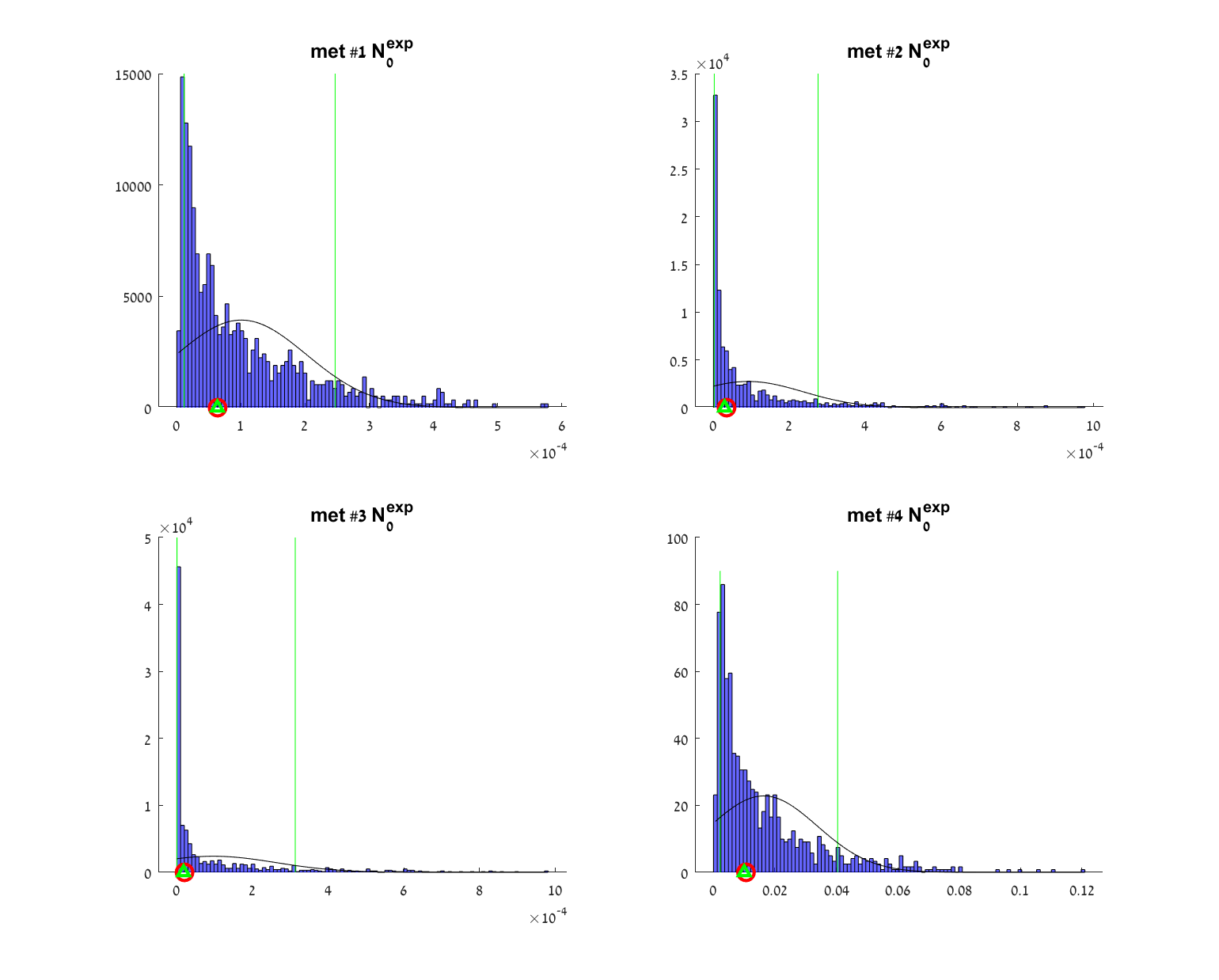
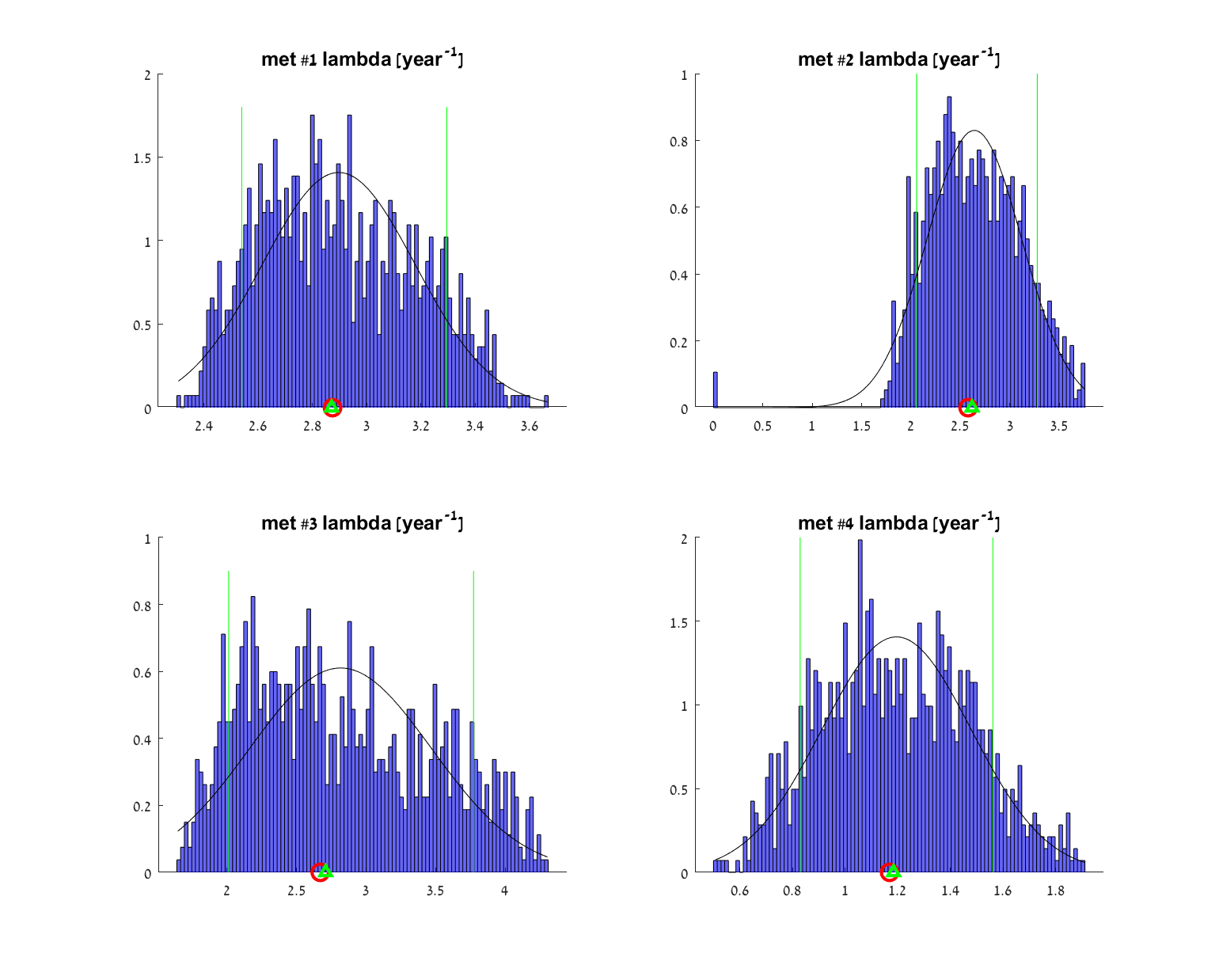
***Sensitivity Analysis Figures***



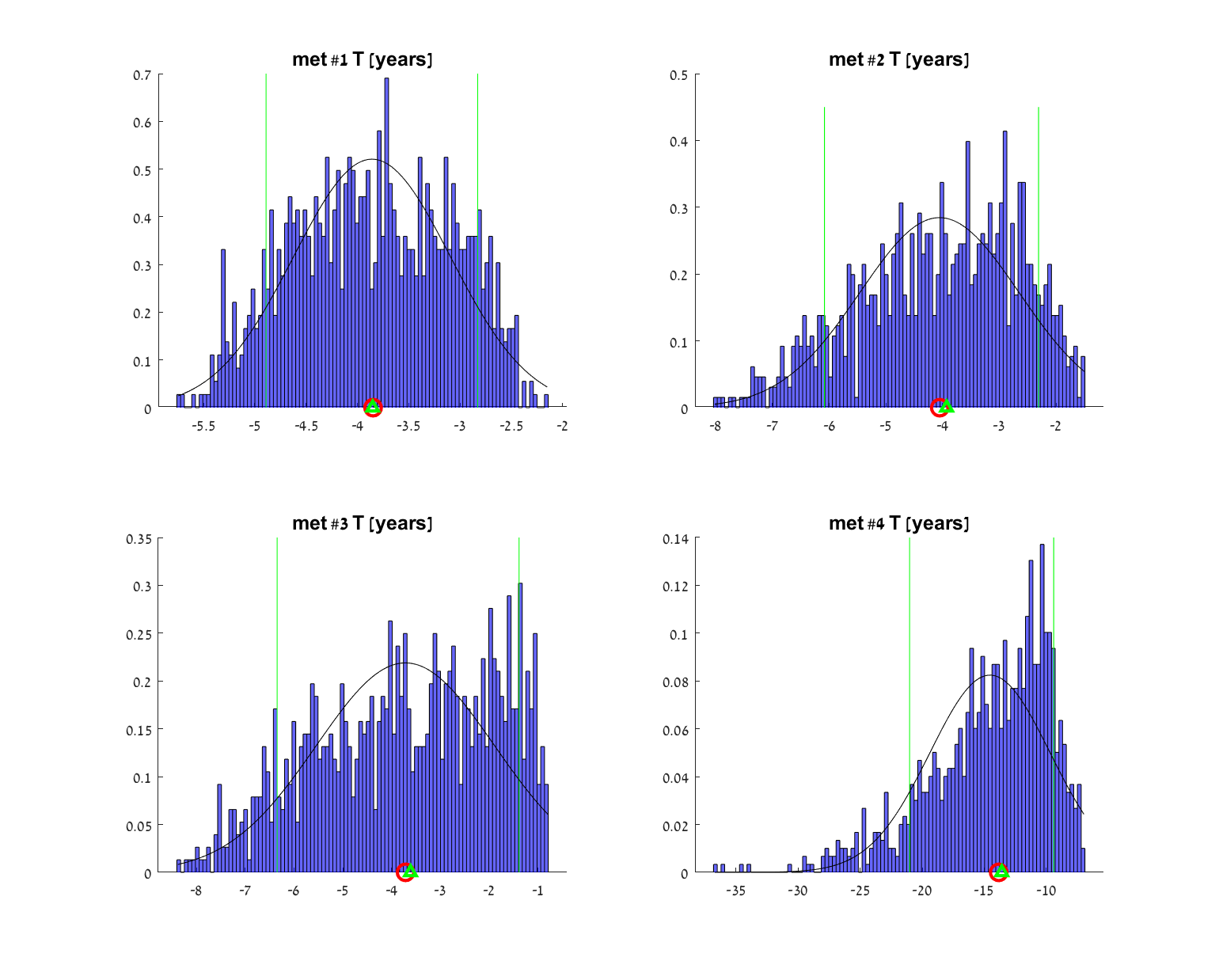
***Figure S1: Exponential model - Sensitivity analysis of the fit. Blue dashed lines: exponential models, fitted to a 1000 random different measurement samples within the measured data error bars. Black circles: Clinical data measurements with measurement error bars. Red curve: The model fitted to the reported measurement data. Green curves: interdecile range (fitted models with 10% and 90% of the sorted values).***

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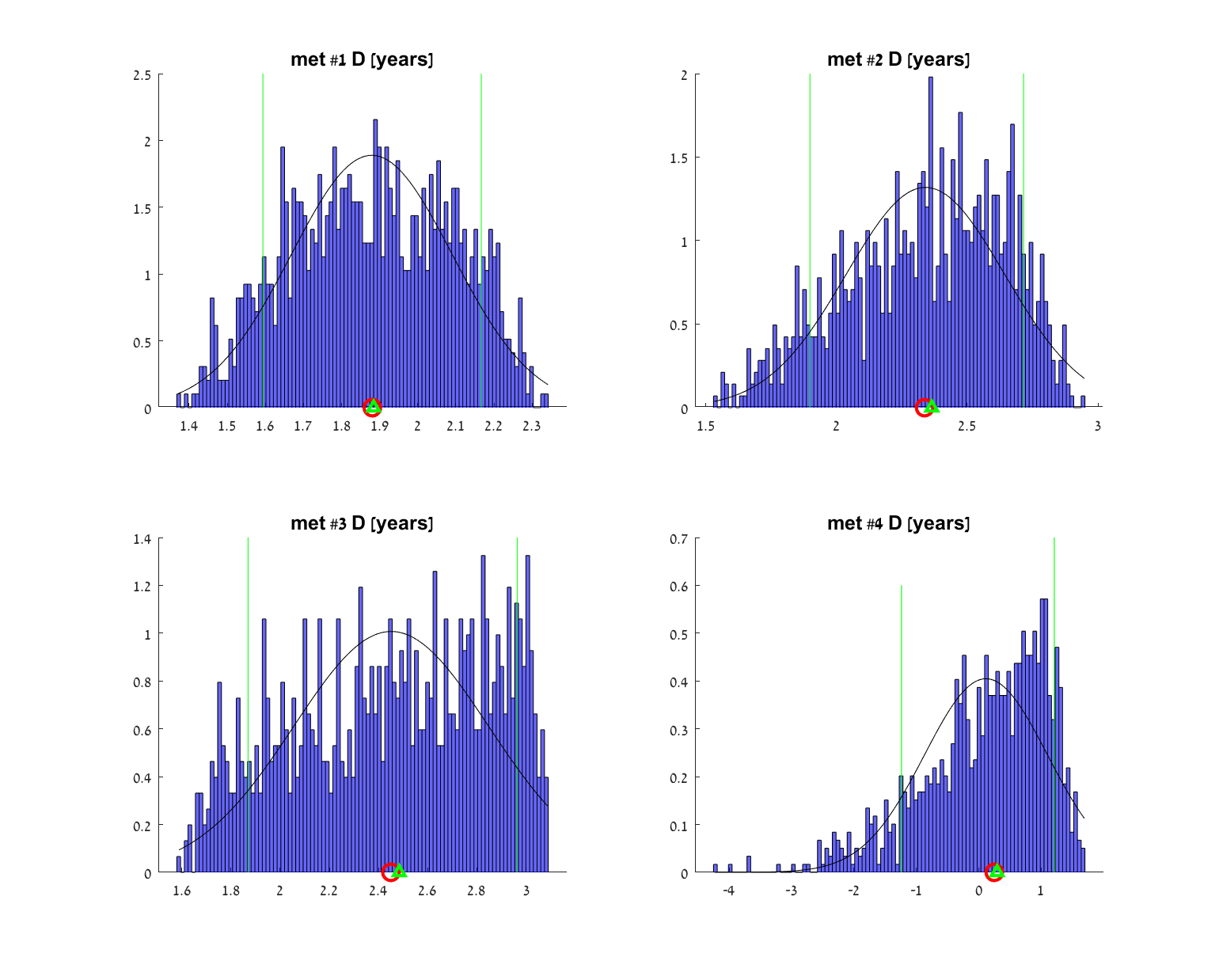
***Figure S2: Exponential model - Sensitivity analysis of the fit. Blue bars: A histogram of parameter values, fitted to a 1000 random measurement samples within the measured data error bars. Black curve: A PDF plot assuming normal distribution of the values. Green triangle: Median value. Green vertical lines: interdecile range (10% and 90% of the sorted values). Red circle: The value fitted to the reported measured data.***

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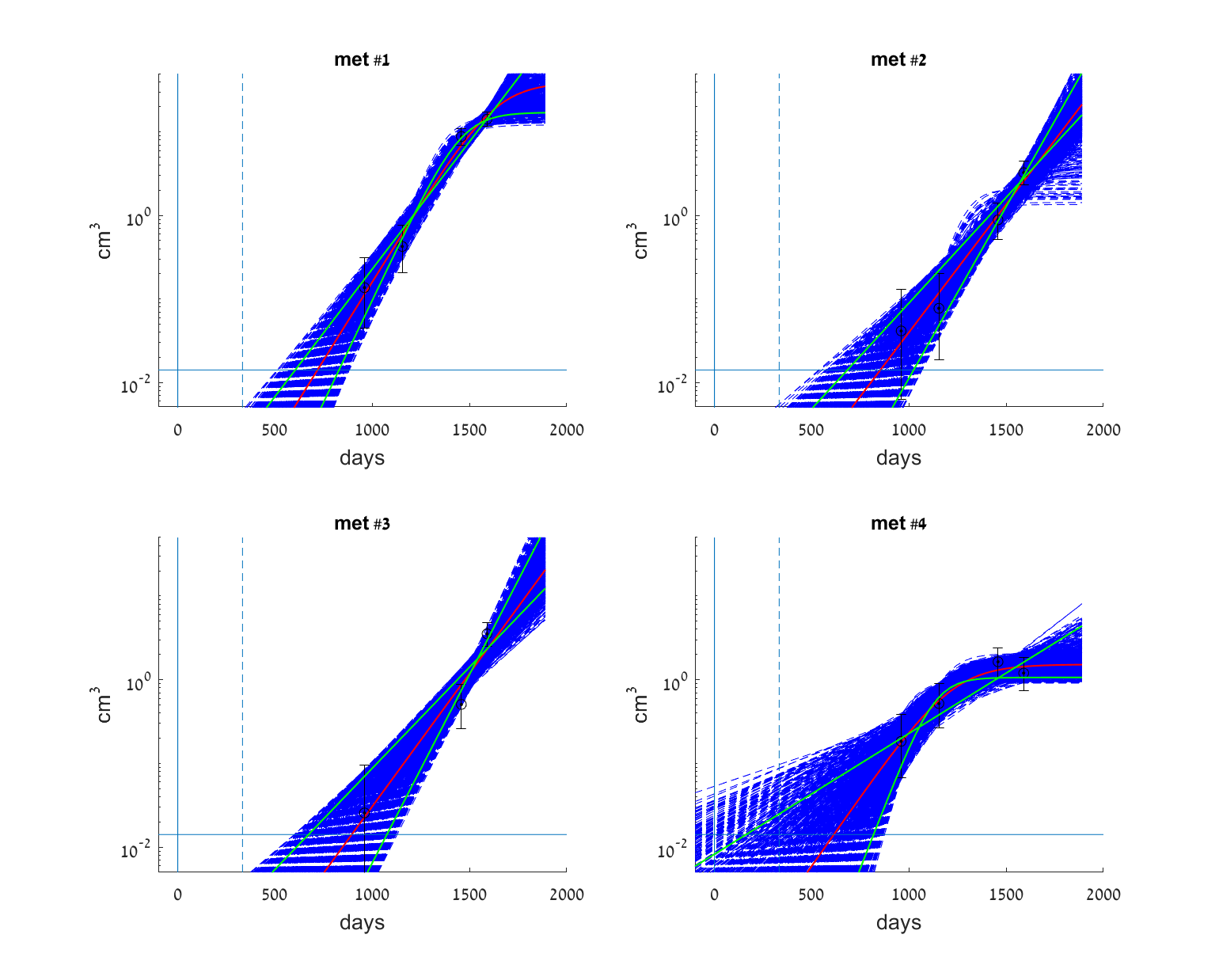
***Figure S3: Exponential model - Sensitivity analysis of the fit. Blue bars: A histogram of parameter values, fitted to a 1000 random measurement samples within the measured data error bars. Black curve: A PDF plot assuming normal distribution of the values. Green triangle: Median value. Green vertical lines: interdecile range (10% and 90% of the sorted values). Red circle: The value fitted to the reported measured data.***

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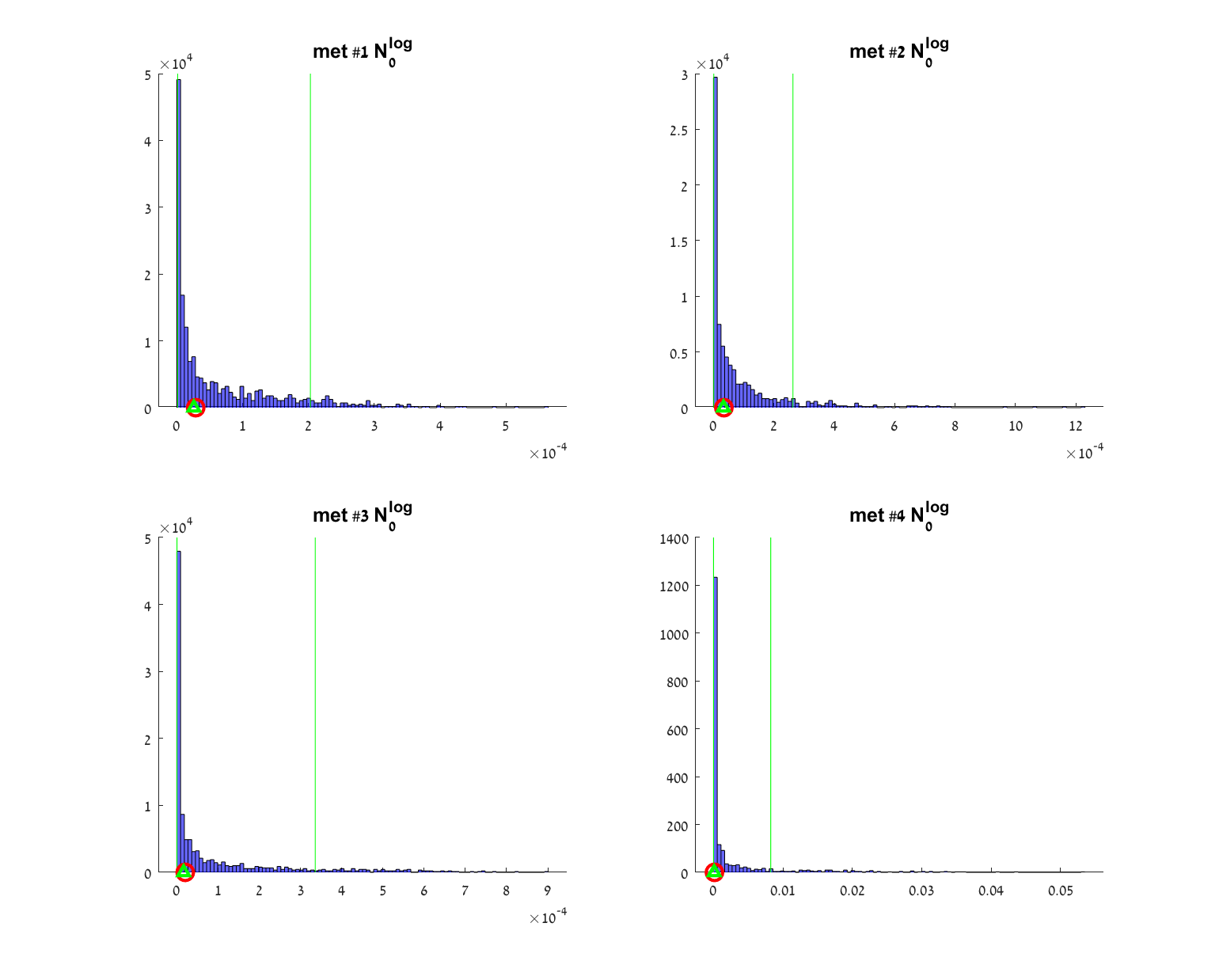
***Figure S4: Exponential model - Sensitivity analysis of the fit. Blue bars: A histogram of metastasis onset time () values, calculated from values of model parameters that were fitted to a 1000 random measurement samples within the measured data error bars. Black curve: A PDF plot assuming normal distribution of the values. Green triangle: Median value. Green vertical lines: interdecile range (10% and 90% of the sorted values). Red circle: The value resulting from fit to the reported measured data.***

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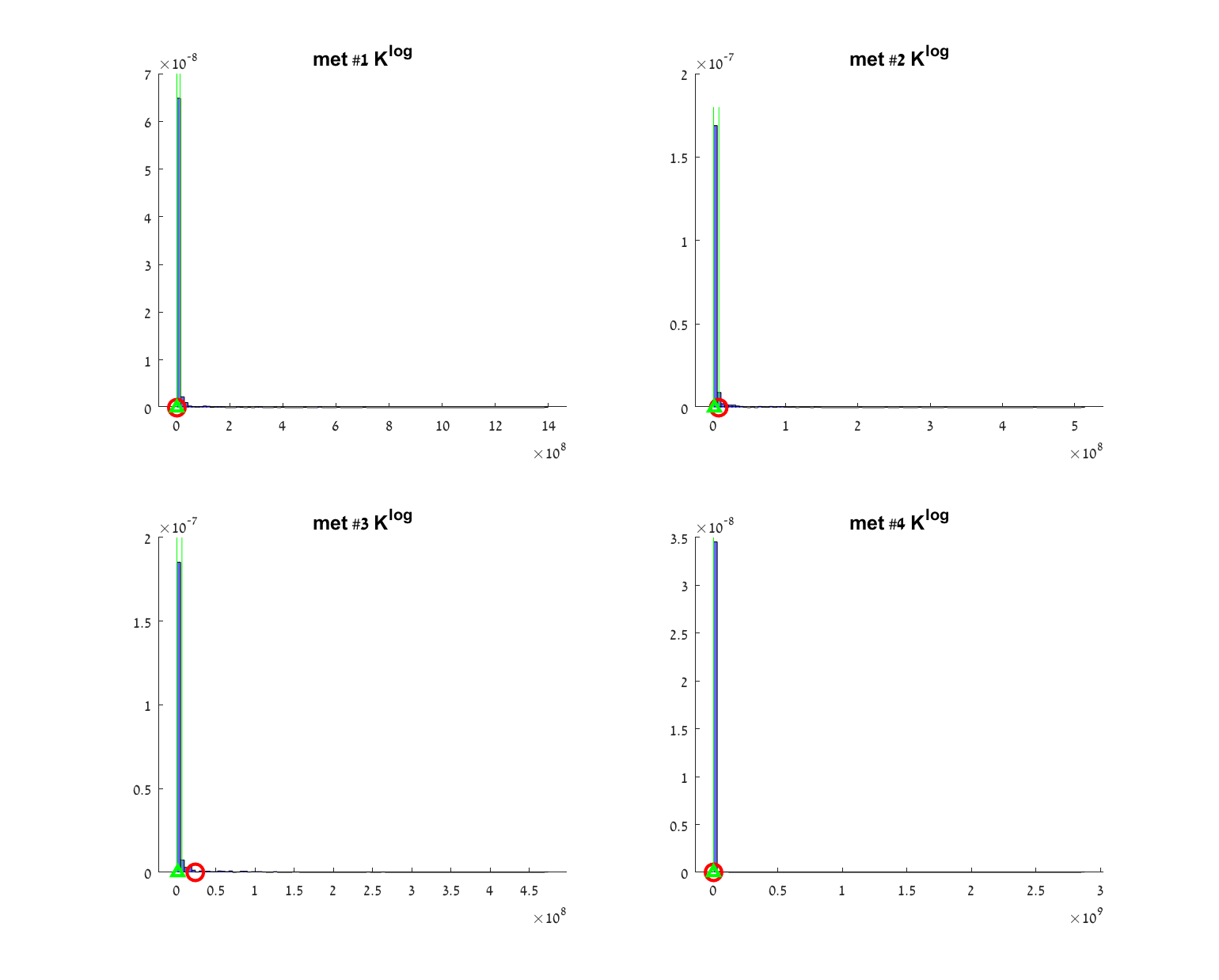
***Figure S5: Exponential model - Sensitivity analysis of the fit. Blue bars: A histogram of earliest possible detection time () values, calculated from values of model parameters that were fitted to a 1000 random measurement samples within the measured data error bars. Black curve: A PDF plot assuming normal distribution of the values. Green triangle: Median value. Green vertical lines: interdecile range (10% and 90% of the sorted values). Red circle: The value resulting from fit to the reported measured data.***

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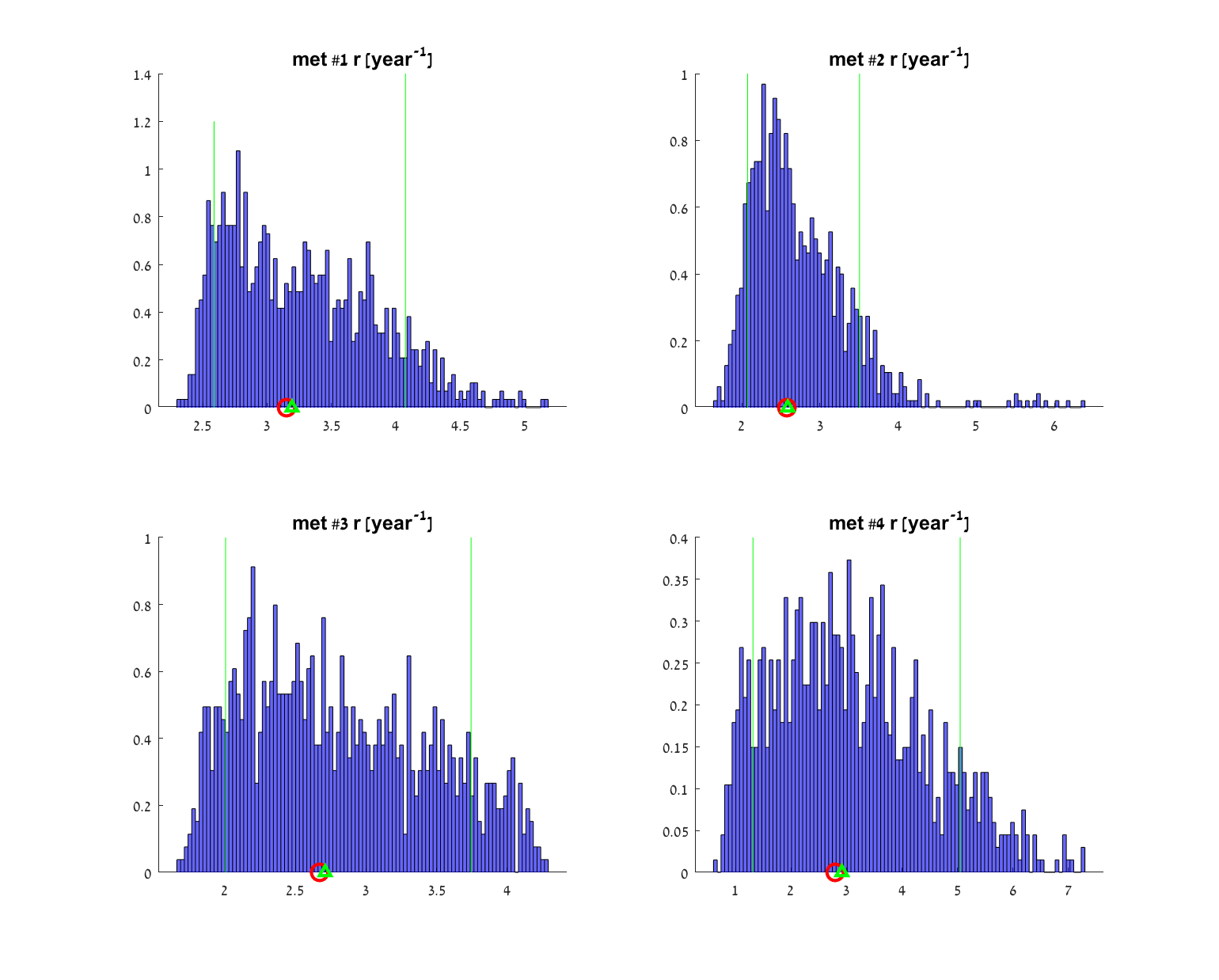
***Figure S6: Logistic model - Sensitivity analysis of the fit. Blue dashed lines: logistic models, fitted to a 1000 random different measurement samples within the measured data error bars. Black circles: Clinical data measurements with measurement error bars. Red curve: The model fitted to the reported measurement data. Green curves: interdecile range (fitted models with 10% and 90% of the sorted values).***



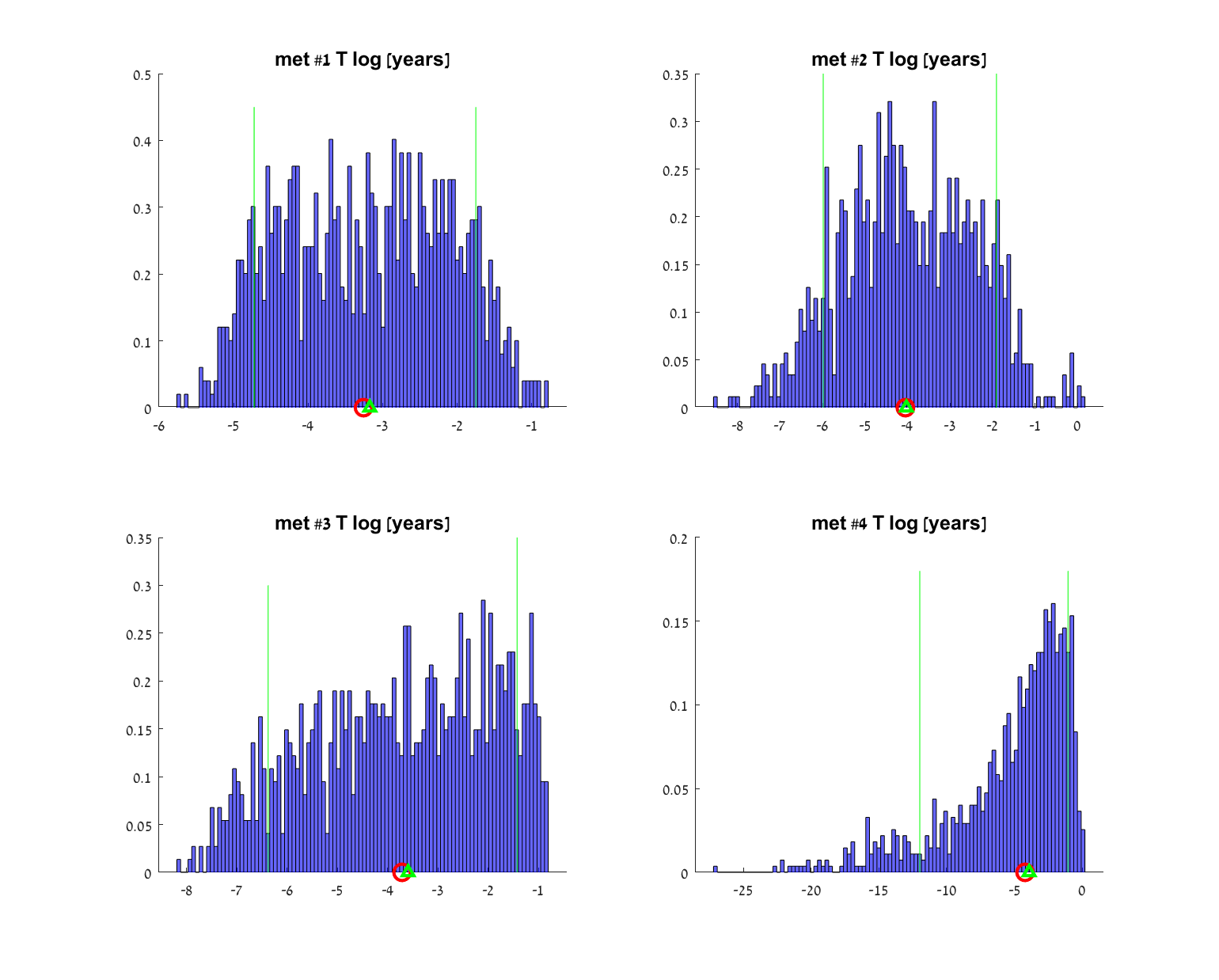
***Figure S7: Logistic model - Sensitivity analysis of the fit. Blue bars: A histogram of parameter values, fitted to a 1000 random measurement samples within the measured data error bars. Green triangle: Median value. Green vertical lines: interdecile range (10% and 90% of the sorted values). Red circle: The value fitted to the reported measured data.***

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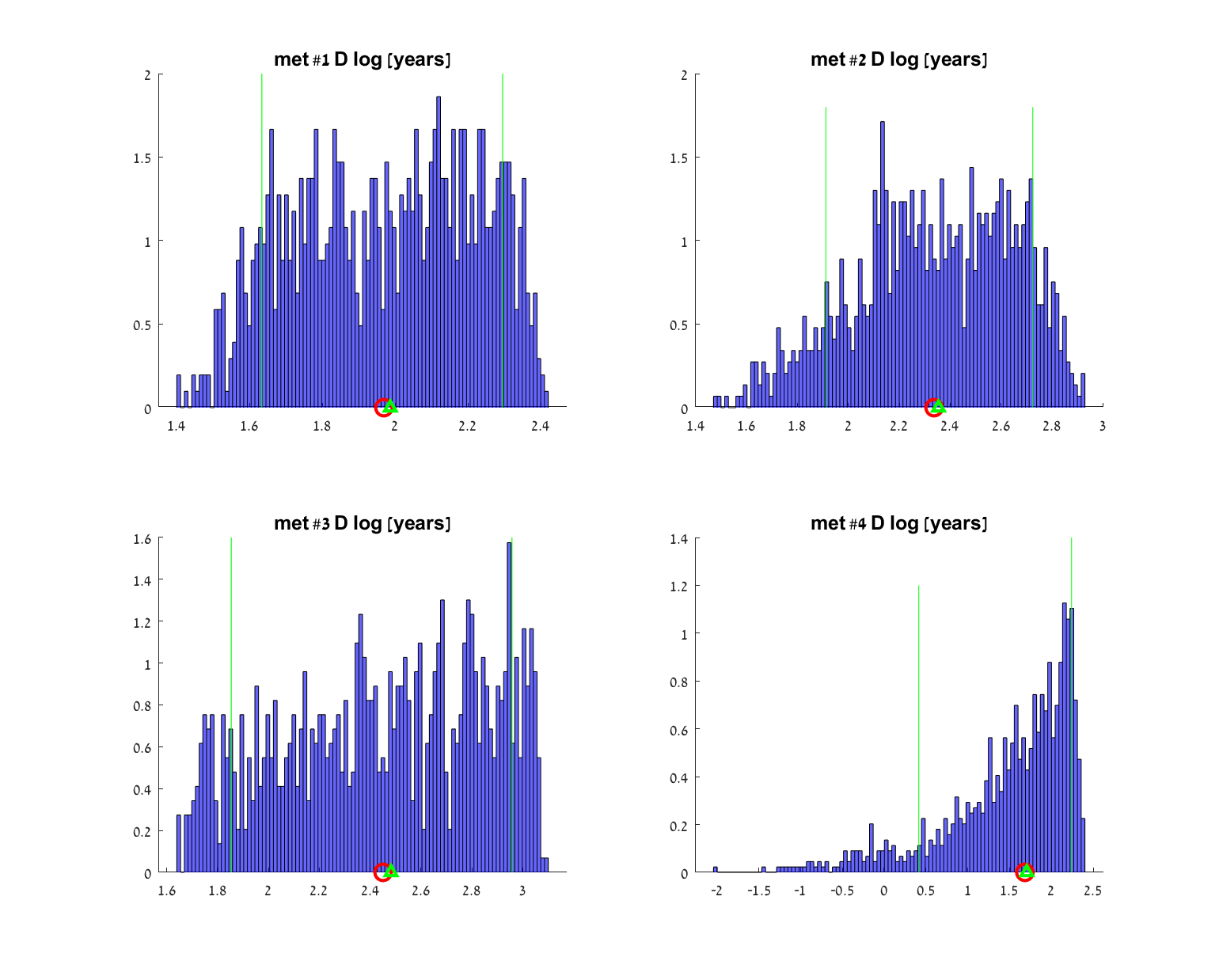
***Figure S8: Logistic model - Sensitivity analysis of the fit. Blue bars: A histogram of parameter values, fitted to a 1000 random measurement samples within the measured data error bars. Green triangle: Median value. Green vertical lines: interdecile range (10% and 90% of the sorted values). Red circle: The value fitted to the reported measured data.***

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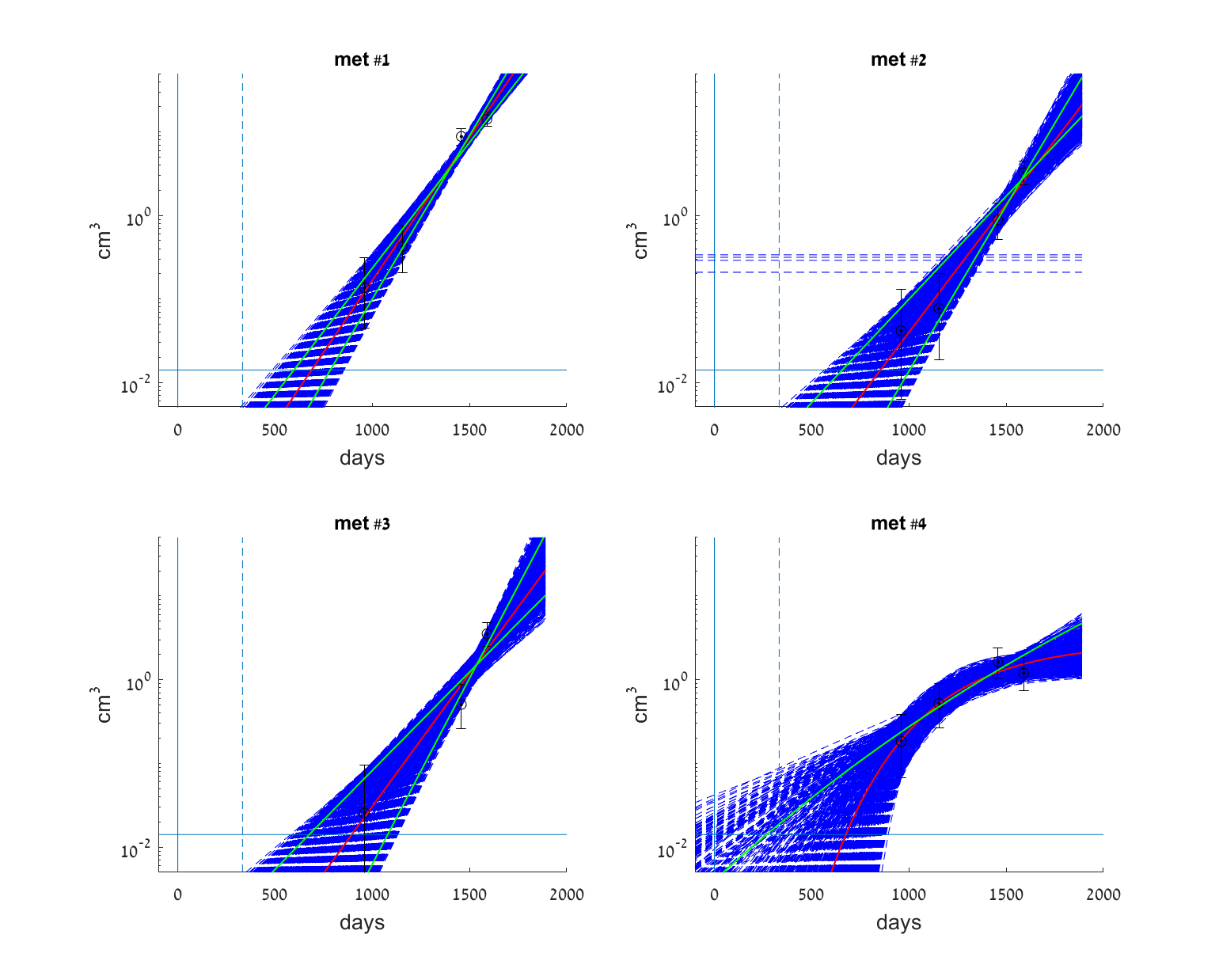
***Figure S9: Logistic model - Sensitivity analysis of the fit. Blue bars: A histogram of parameter values, fitted to a 1000 random measurement samples within the measured data error bars. Green triangle: Median value. Green vertical lines: interdecile range (10% and 90% of the sorted values). Red circle: The value fitted to the reported measured data.***

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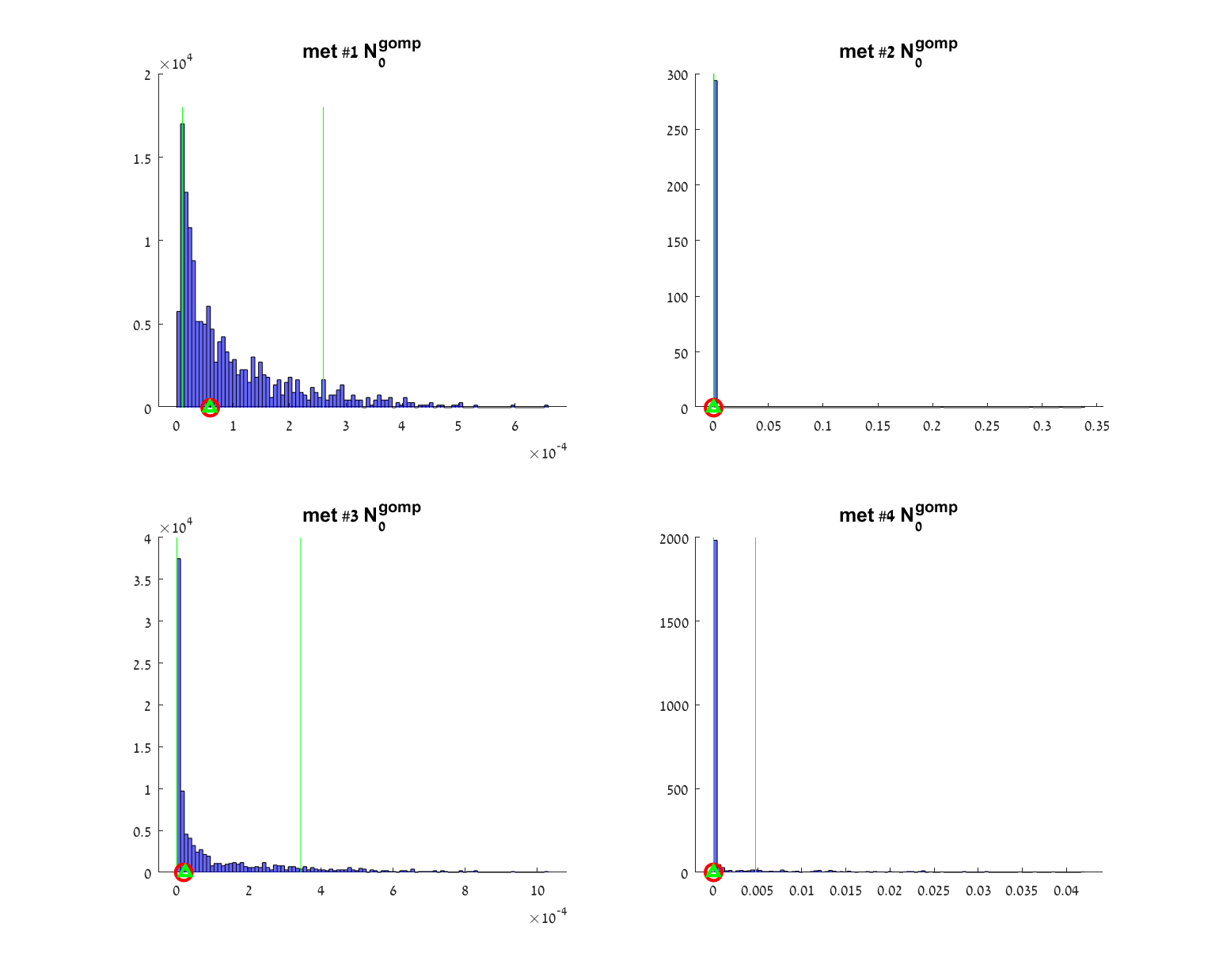
***Figure S10: Logistic model - Sensitivity analysis of the fit. Blue bars: A histogram of metastasis onset time () values, calculated from values of model parameters that were fitted to a 1000 random measurement samples within the measured data error bars. Green triangle: Median value. Green vertical lines: interdecile range (10% and 90% of the sorted values). Red circle: The value resulting from fit to the reported measured data.***

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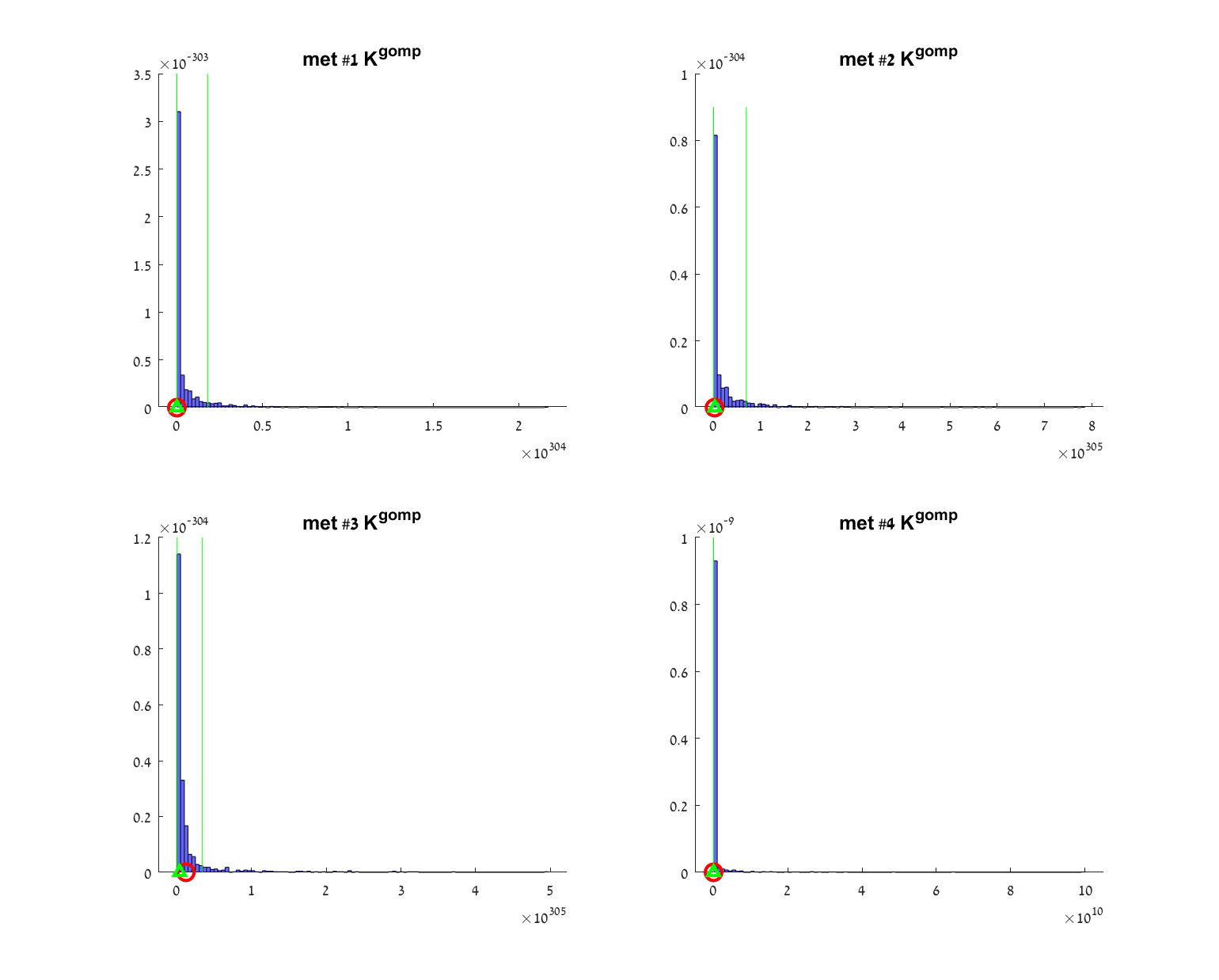
***Figure S11: Logistic model - Sensitivity analysis of the fit. Blue bars: A histogram of earliest possible detection time () values, calculated from values of model parameters that were fitted to a 1000 random measurement samples within the measured data error bars. Green triangle: Median value. Green vertical lines: interdecile range (10% and 90% of the sorted values). Red circle: The value resulting from fit to the reported measured data.***



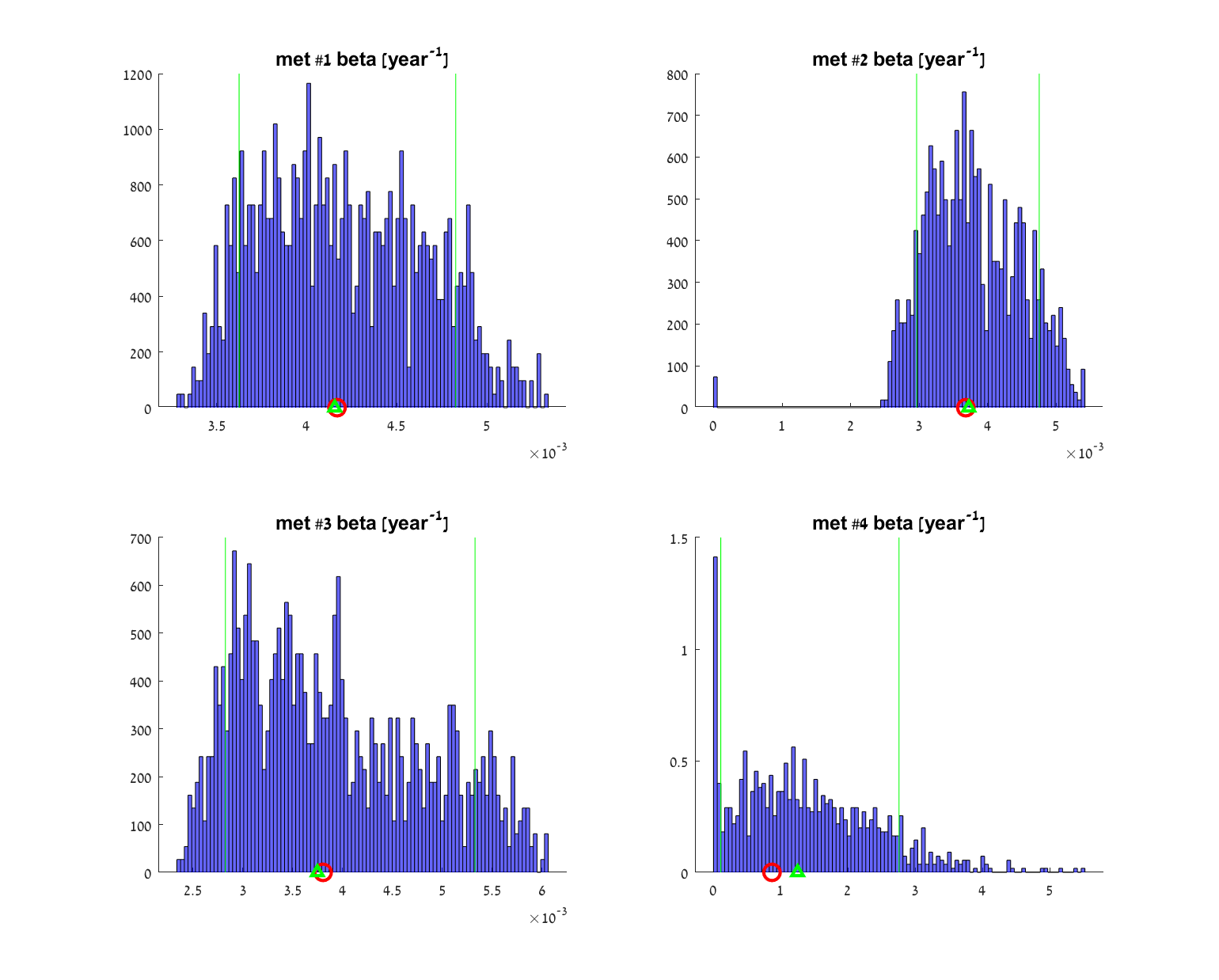
***Figure S12: Gompertz model – Sensitivity analysis of the fit. Blue dashed lines: Gompertz models, fitted to a 1000 random different measurement samples within the measured data error bars. Black circles: Clinical data measurements with measurement error bars. Red curve: The model fitted to the reported measurement data. Green curves: interdecile range (fitted models with 10% and 90% of the sorted values).***



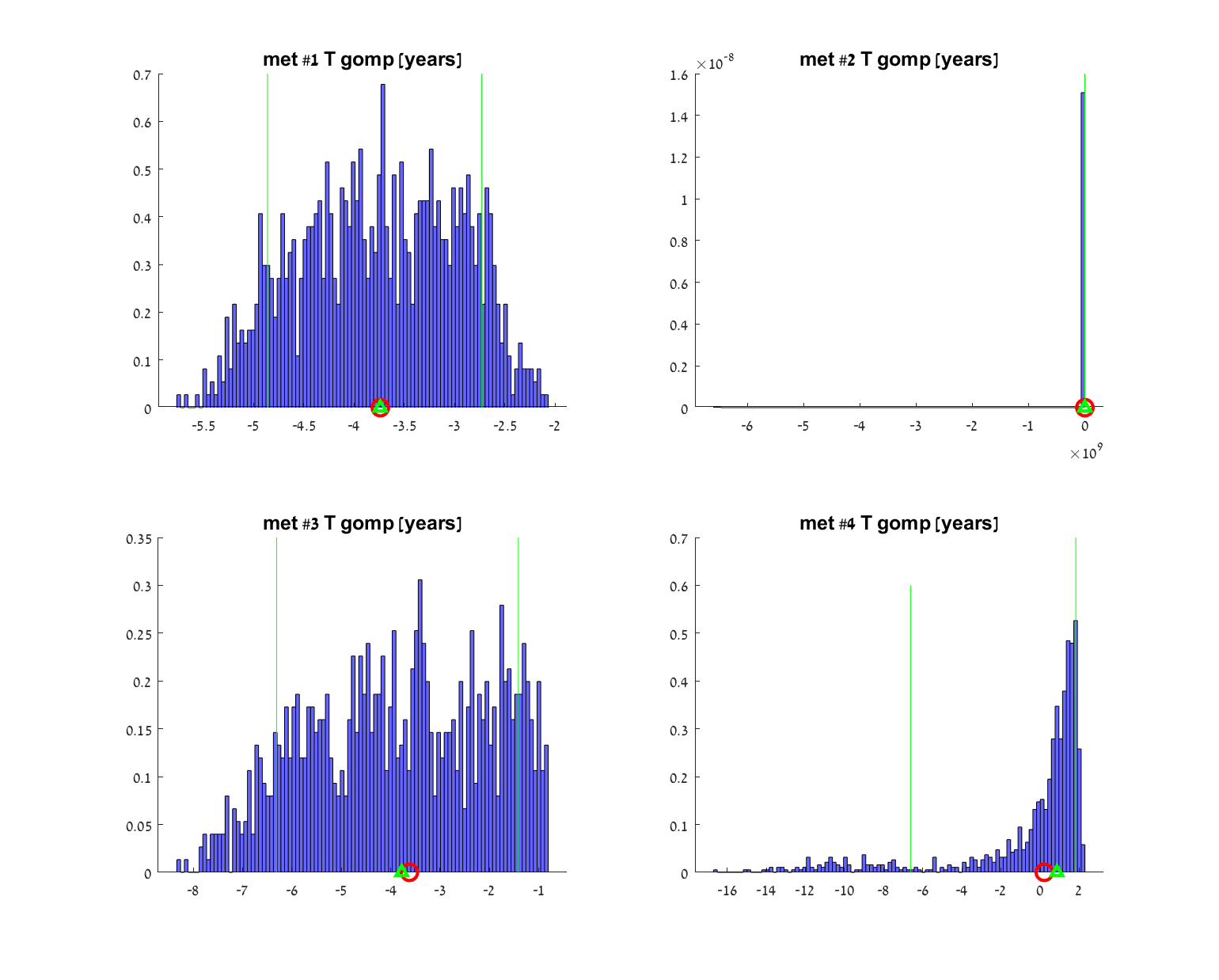
***Figure S13: Gompertz model - Sensitivity analysis of the fit. Blue bars: A histogram of parameter values, fitted to a 1000 random measurement samples within the measured data error bars. Green triangle: Median value. Green vertical lines: interdecile range (10% and 90% of the sorted values). Red circle: The value fitted to the reported measured data.***



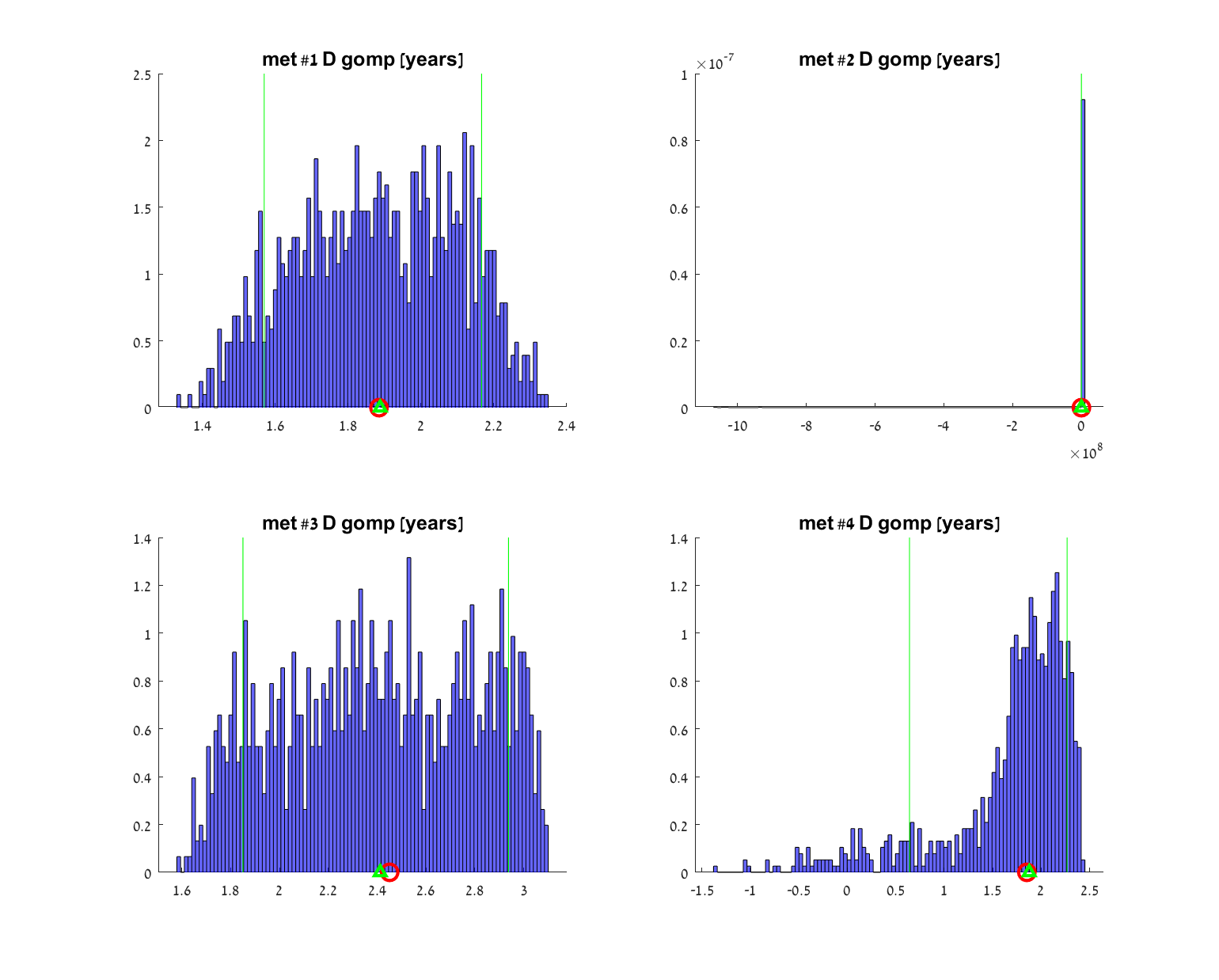
***Figure S14: Gompertz model - Sensitivity analysis of the fit. Blue bars: A histogram of parameter values, fitted to a 1000 random measurement samples within the measured data error bars. Green triangle: Median value. Green vertical lines: interdecile range (10% and 90% of the sorted values). Red circle: The value fitted to the reported measured data.***



***Figure S15: Gompertz model - Sensitivity analysis of the fit. Blue bars: A histogram of parameter values, fitted to a 1000 random measurement samples within the measured data error bars. Green triangle: Median value. Green vertical lines: interdecile range (10% and 90% of the sorted values). Red circle: The value fitted to the reported measured data.***



***Figure S16: Gompertz model - Sensitivity analysis of the fit. Blue bars: A histogram of metastasis onset time () values, calculated from values of model parameters that were fitted to a 1000 random measurement samples within the measured data error bars. Green triangle: Median value. Green vertical lines: interdecile range (10% and 90% of the sorted values). Red circle: The value resulting from fit to the reported measured data.***



***Figure S17: Gompertz model - Sensitivity analysis of the fit. Blue bars: A histogram of earliest possible detection time () values, calculated from values of model parameters that were fitted to a 1000 random measurement samples within the measured data error bars. Green triangle: Median value. Green vertical lines: interdecile range (10% and 90% of the sorted values). Red circle: The value resulting from fit to the reported measured data.***