Open Systems Pharmacology Suite - 10
Folder Comparison

gibpk

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Chapter 1

Folder Comparison Results

Overall Comparison Result: Invalid
Number of Compared Files: 146

1.1 Comparison Results

Overall Comparison Result
Invalid

Old Folder
D:\Outputs_9.1

New Folder
D:\Outputs_10.0

Using exclusions
|Interstitial|P-gp|
|Intracellular|P-gp|
|Plasma|OATP1B1|
|Interstitial|OATP1B1|
|Intracellular|OATP1B1|
|Plasma|TRANS|
|Interstitial|TRANS|
|Intracellular|TRANS|

1.1.1 Invalid Simulations (16/146)

Simulation: DDI_MultipleCombinations-05_MM_Mechanismbased_Mechanismbased
Result of the validation: Invalid
Absolute Tolerance: 1.00E-10
Relative Tolerance: 1.00E-5

Output Path: Organism|Liver|Periportal|Interstitial|Erythromycin|Concentration in container
Deviation for 'Organism|Liver|Periportal|Interstitial|Erythromycin|Concentration in container' is 45.33% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.45

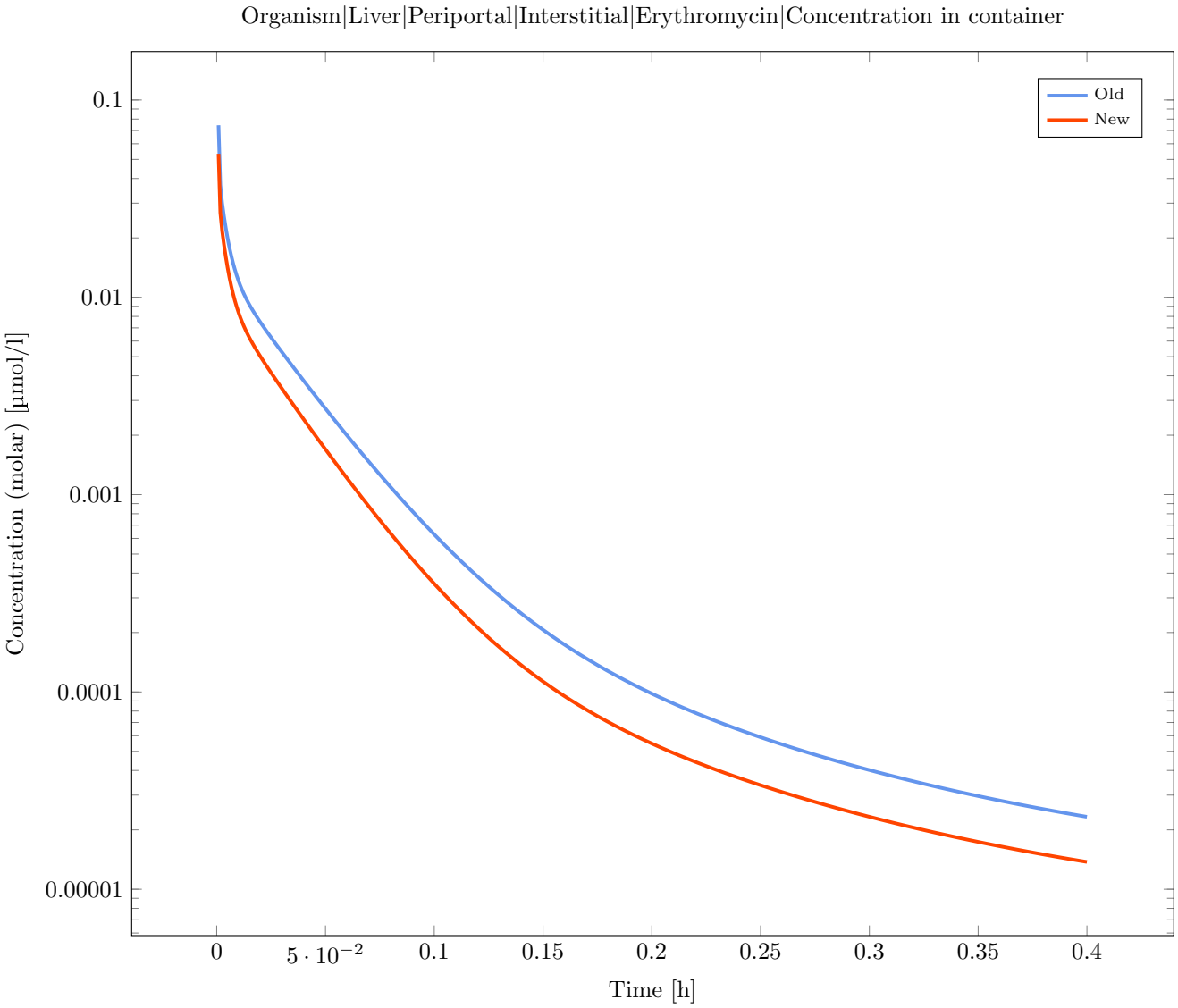
Organism|Liver|Periportal|Interstitial|Erythromycin|Concentration in container
Old
New
Concentration (molar) [µmol/l]
0.1
0.01
0.001
0.0001
0.00001
0
$5 \cdot 10^{-2}$
0.1
0.15
0.2
0.25
0.3
0.35
0.4
Time [h]

Figure 1.1

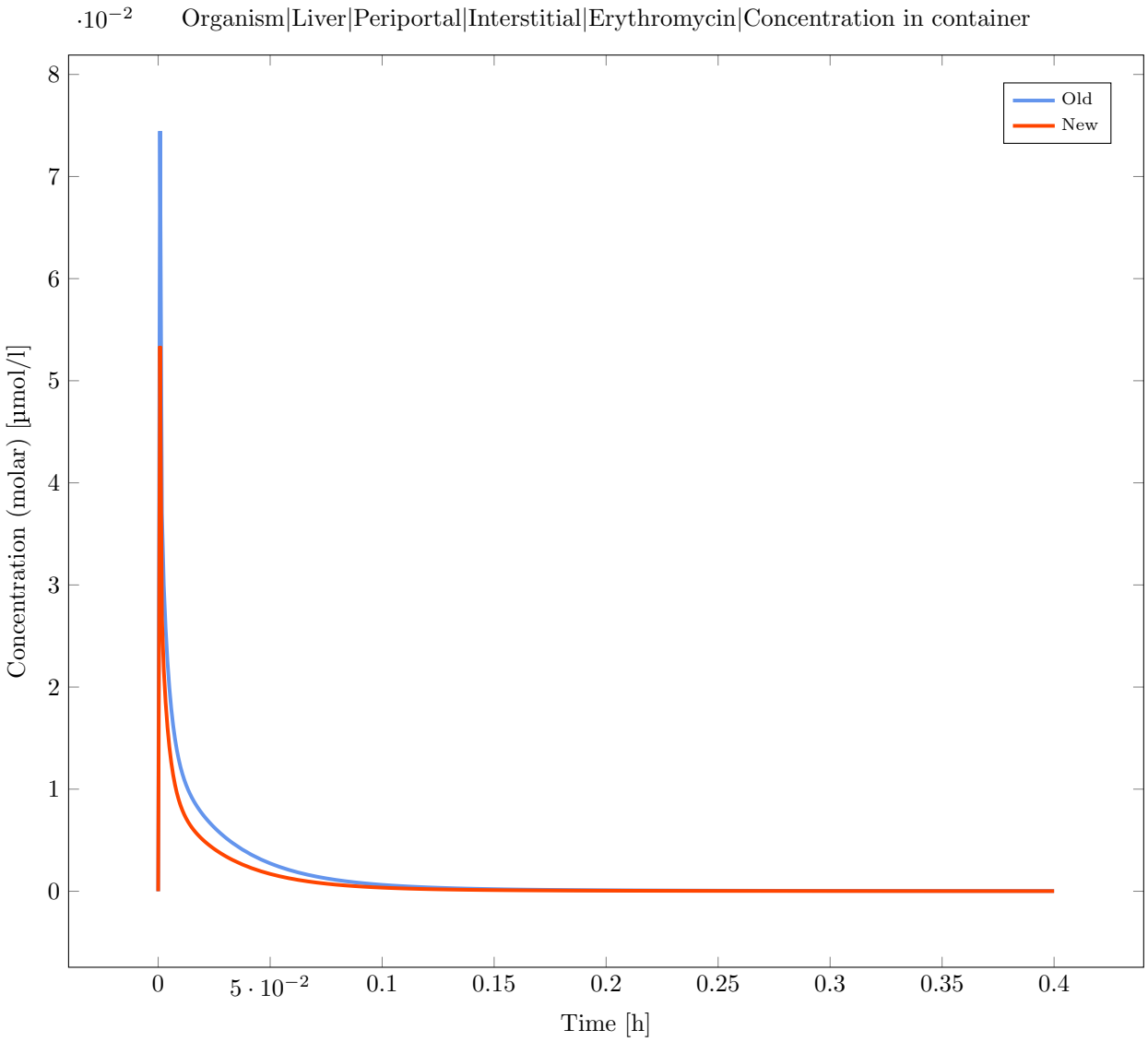

Figure 1.2

Simulation: DDI_MultipleCombinations-06_MM_Induction_Induction
Result of the validation: Invalid
Absolute Tolerance: 1.00E-10
Relative Tolerance: 1.00E-5

Output Path: Organism|Lumen|ColonDescendens|Rifampicin|Fraction of oral drug mass absorbed into mucosa segment
Deviation for 'Organism|Lumen|ColonDescendens|Rifampicin|Fraction of oral drug mass absorbed into mucosa segment' is 1122.72% and is greater than the allowed max. tolerance of 3.00%
Deviation: 11.23

Organism|Lumen|ColonDescendens|Rifampicin|Fraction of oral drug mass absorbed into mucosa segment

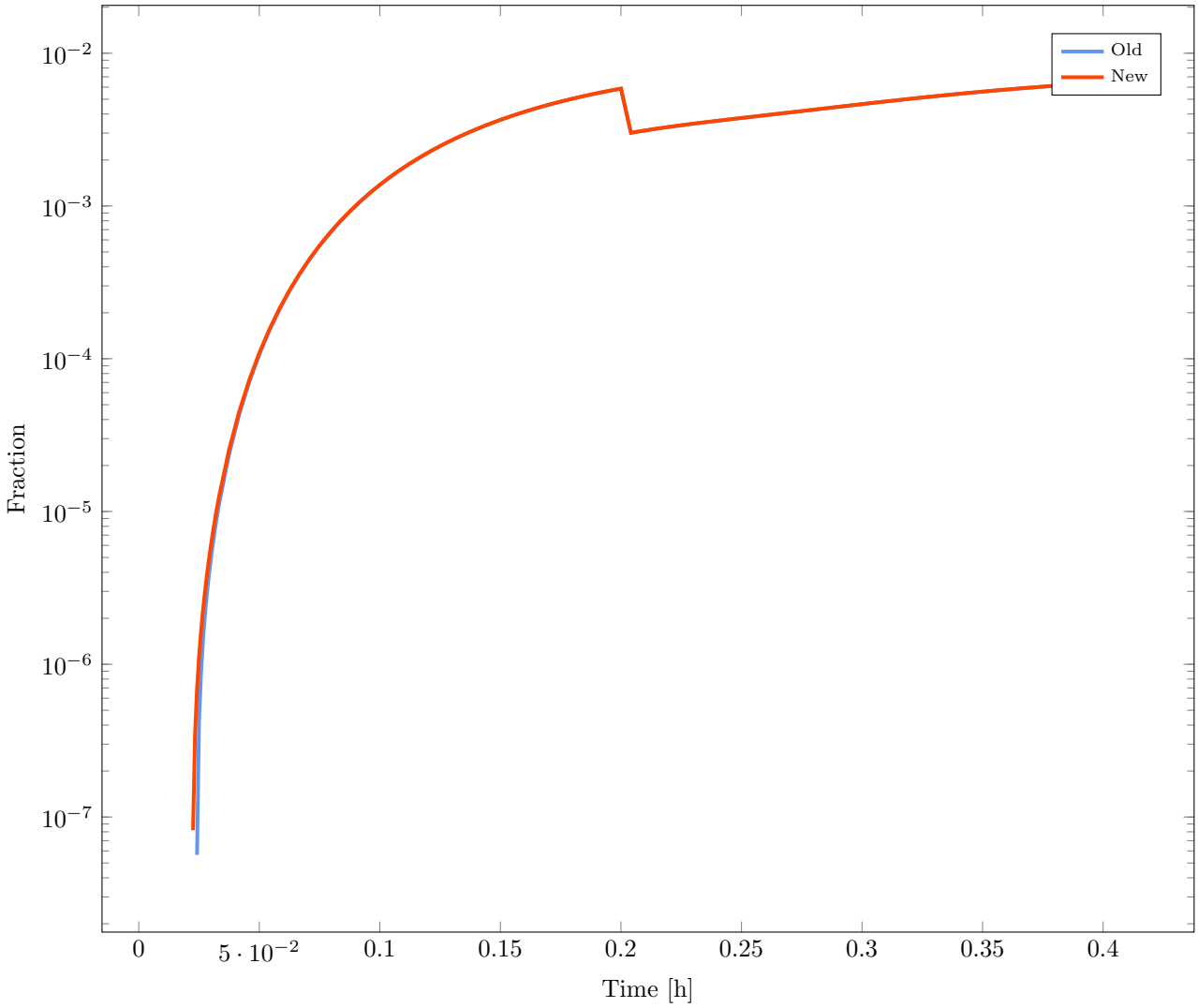

Figure 1.3

Organism|Lumen|ColonDescendens|Rifampicin|Fraction of oral drug mass absorbed into mucosa segment

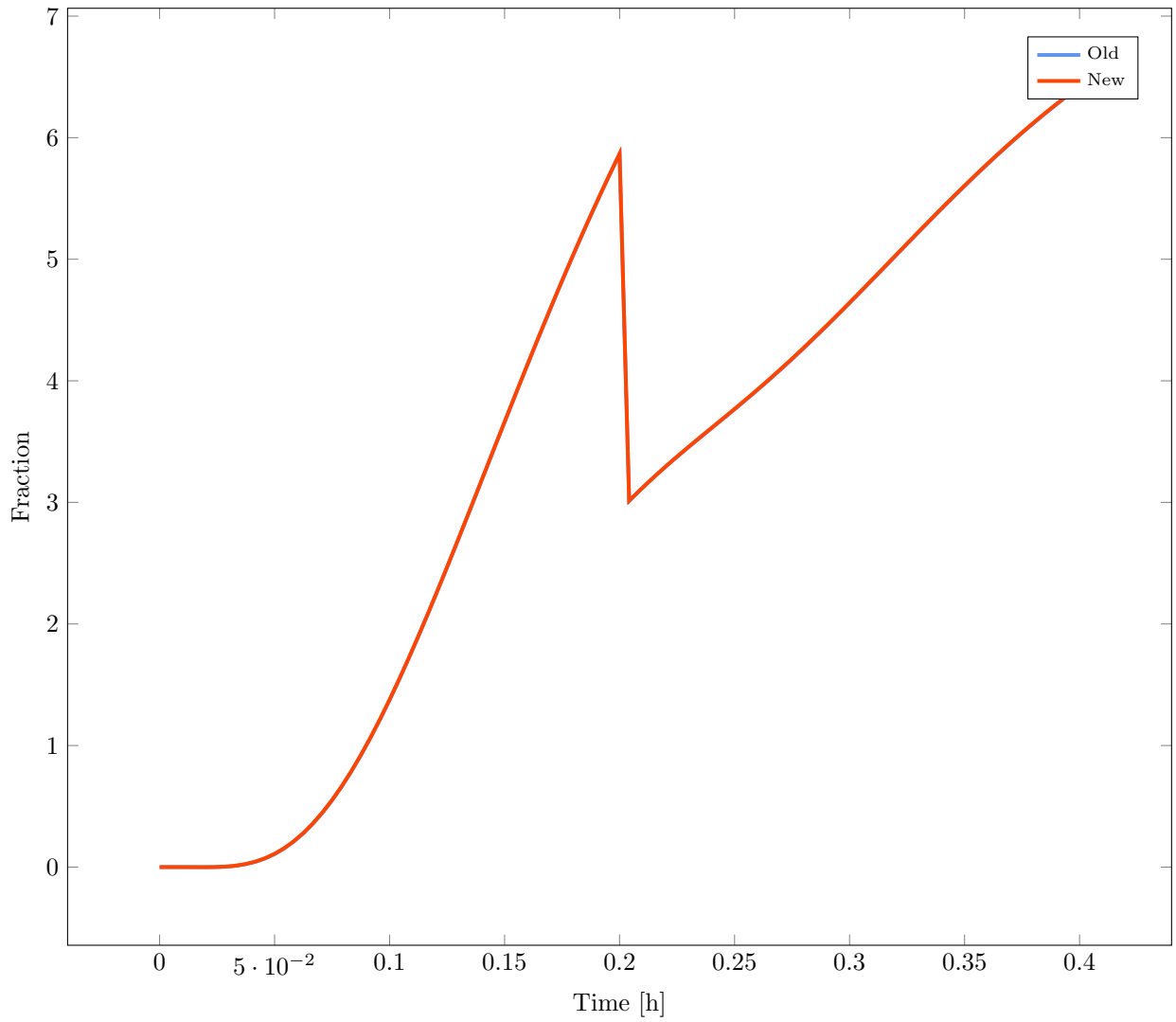

Figure 1.4

Simulation: DDI_MultipleCombinations-07_MM_Competitive_Competitive_Mechanismbased_Mechanismbas
Result of the validation: Invalid
Absolute Tolerance: 1.00E-10
Relative Tolerance: 1.00E-5

Output Path: Organism|Liver|Periportal|Interstitial|Erythromycin|Concentration in container
Deviation for 'Organism|Liver|Periportal|Interstitial|Erythromycin|Concentration in container' is 56.47% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.56

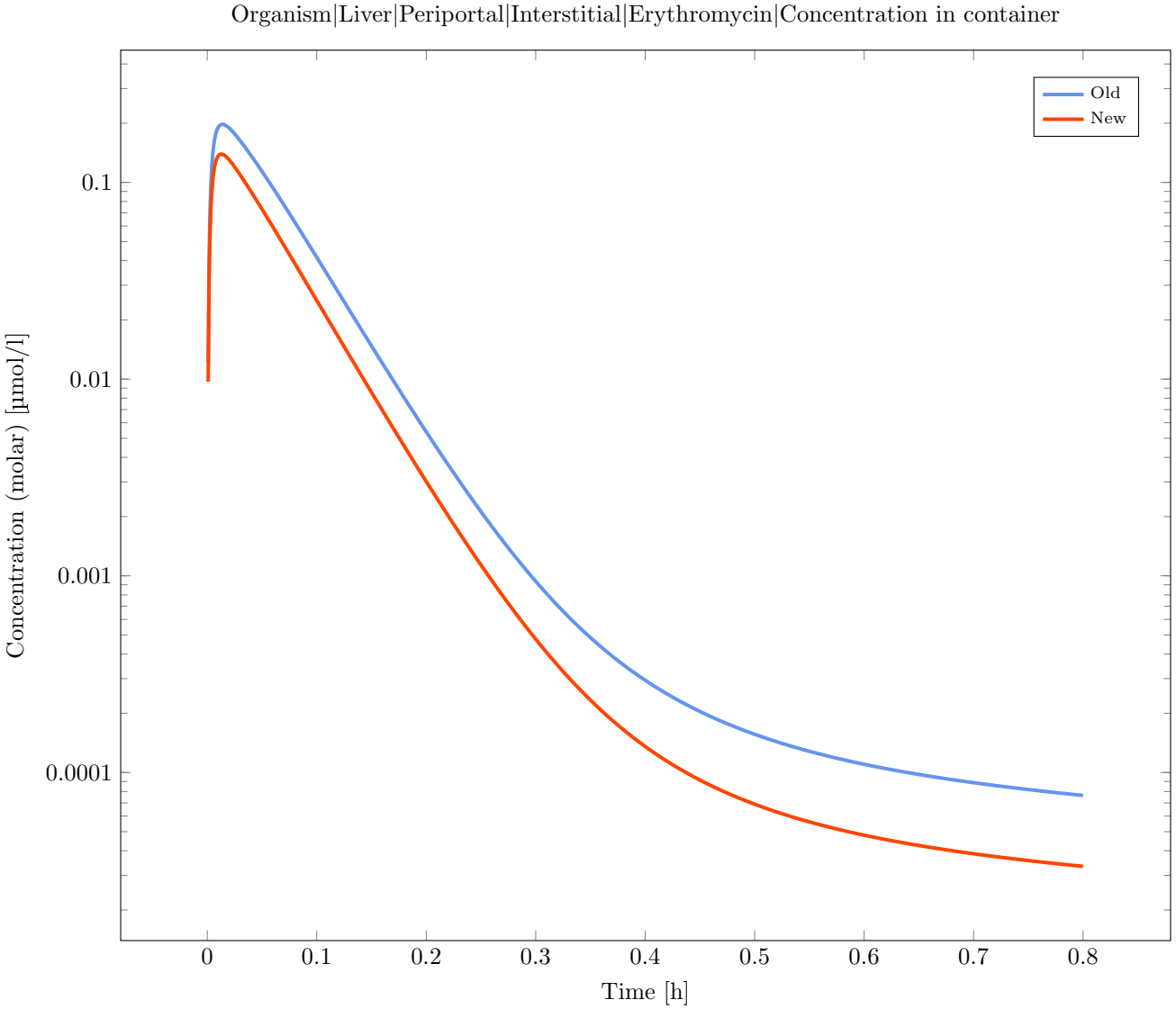
Organism|Liver|Periportal|Interstitial|Erythromycin|Concentration in container
Old
New
Concentration (molar) [µmol/l]
0.1
0.01
0.001
0.0001
0
0.1
0.2
0.3
0.4
0.5
0.6
0.7
0.8
Time [h]

Figure 1.5

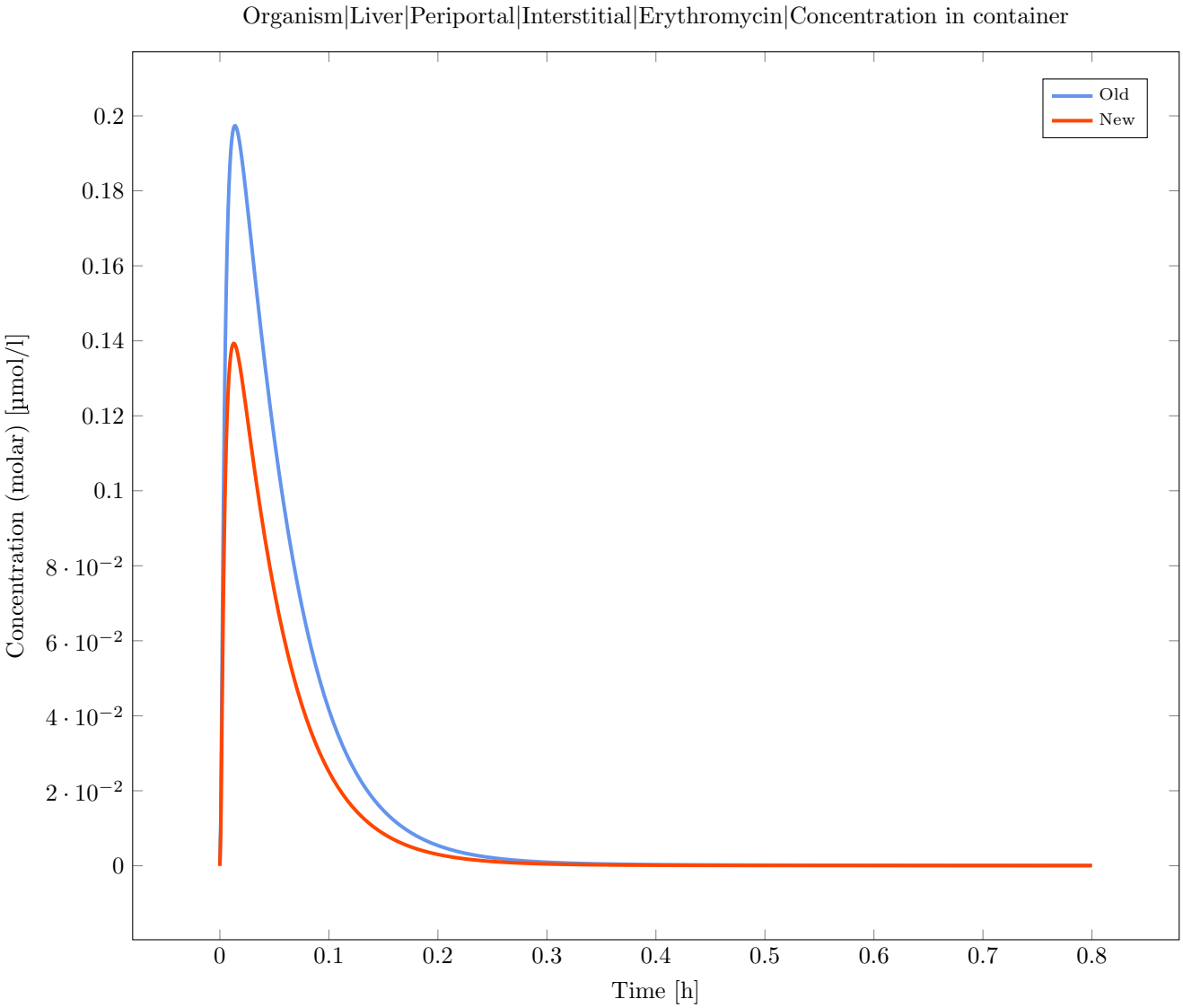

Figure 1.6

Simulation: DDI_MultipleCombinations-08_MM_Uncompetitive_Uncompetitive_Mechanismbased_-Mechanismbased
Result of the validation: Invalid
Absolute Tolerance: 1.00E-10
Relative Tolerance: 1.00E-5

Output Path: Organism|Liver|Periportal|Interstitial|MechanismBased_4|Concentration in container
Deviation for 'Organism|Liver|Periportal|Interstitial|MechanismBased_4|Concentration in container' is 49.46% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.49

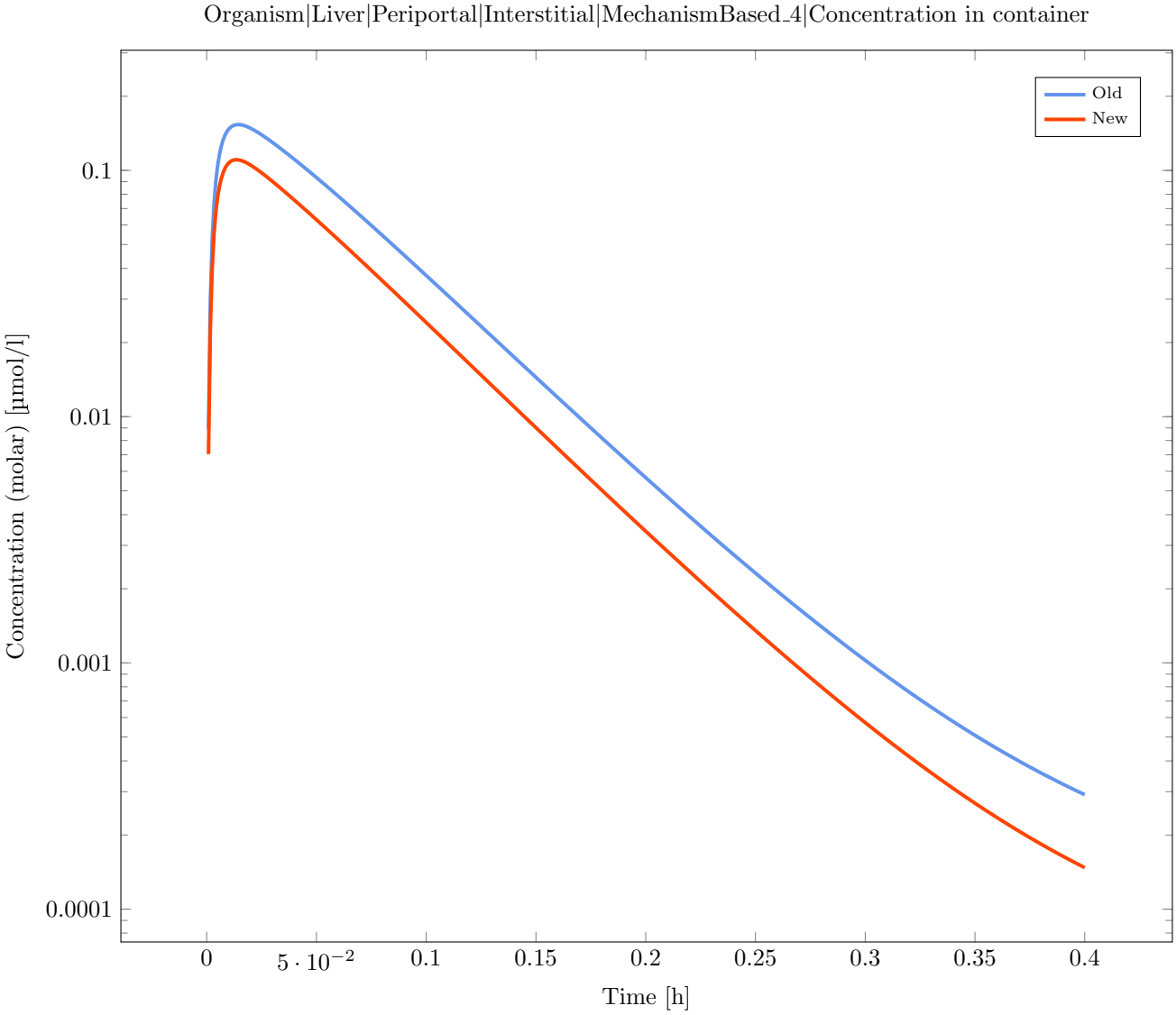
Organism|Liver|Periportal|Interstitial|MechanismBased_4|Concentration in container
Old
New
Concentration (molar) [µmol/l]
0.1
0.01
0.001
0.0001
0
5 · 10−2
0.1
0.15
0.2
0.25
0.3
0.35
0.4
Time [h]

Figure 1.7

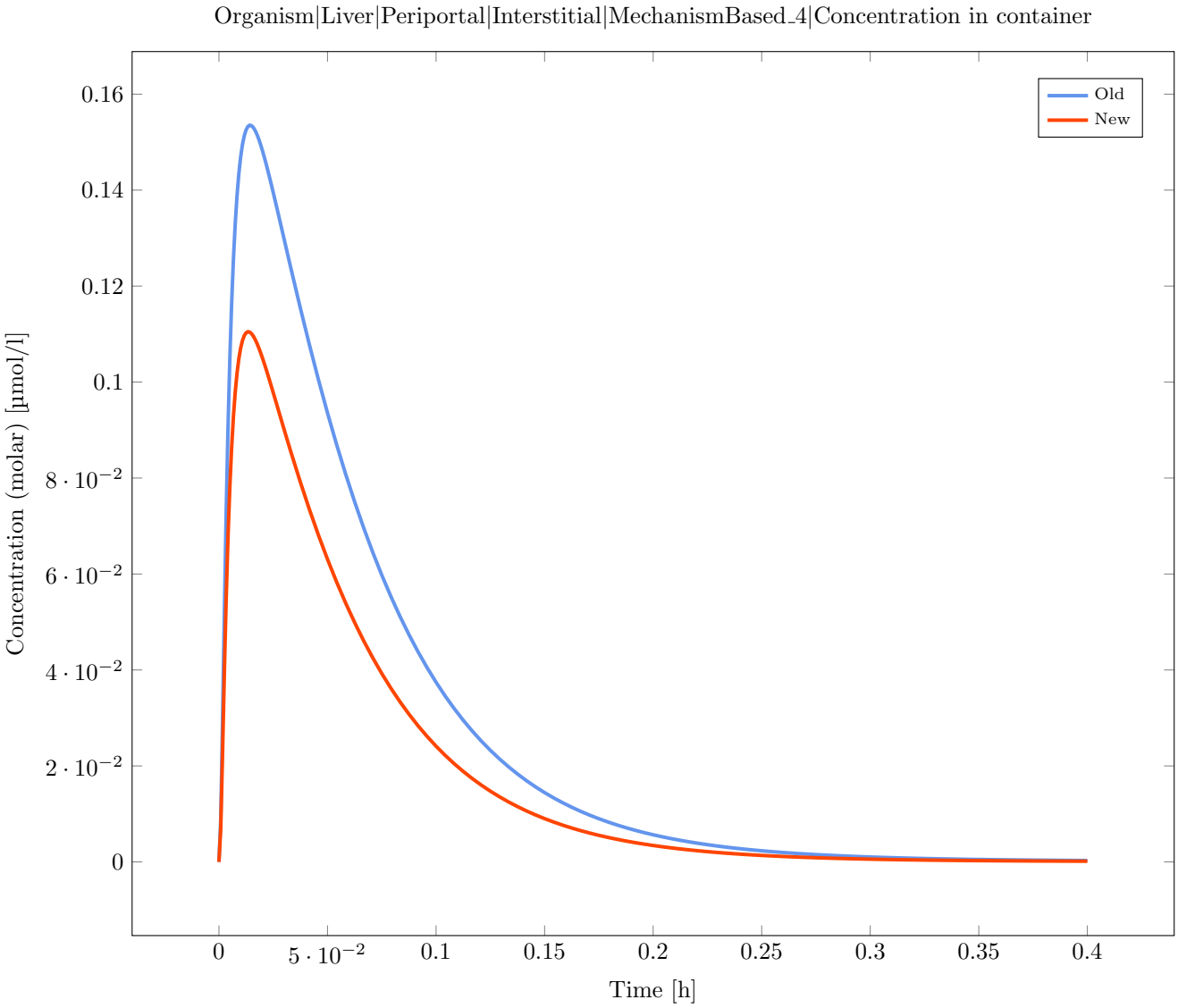

Figure 1.8

Simulation: DDI_MultipleCombinations-09_MM_Noncompetitive_Noncompetitive_Mechanismbased_-Mechanismbased
Result of the validation: Invalid
Absolute Tolerance: 1.00E-10
Relative Tolerance: 1.00E-5

Output Path: Organism|Liver|Periportal|Interstitial|Erythromycin|Concentration in container
Deviation for 'Organism|Liver|Periportal|Interstitial|Erythromycin|Concentration in container' is 39.95% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.40

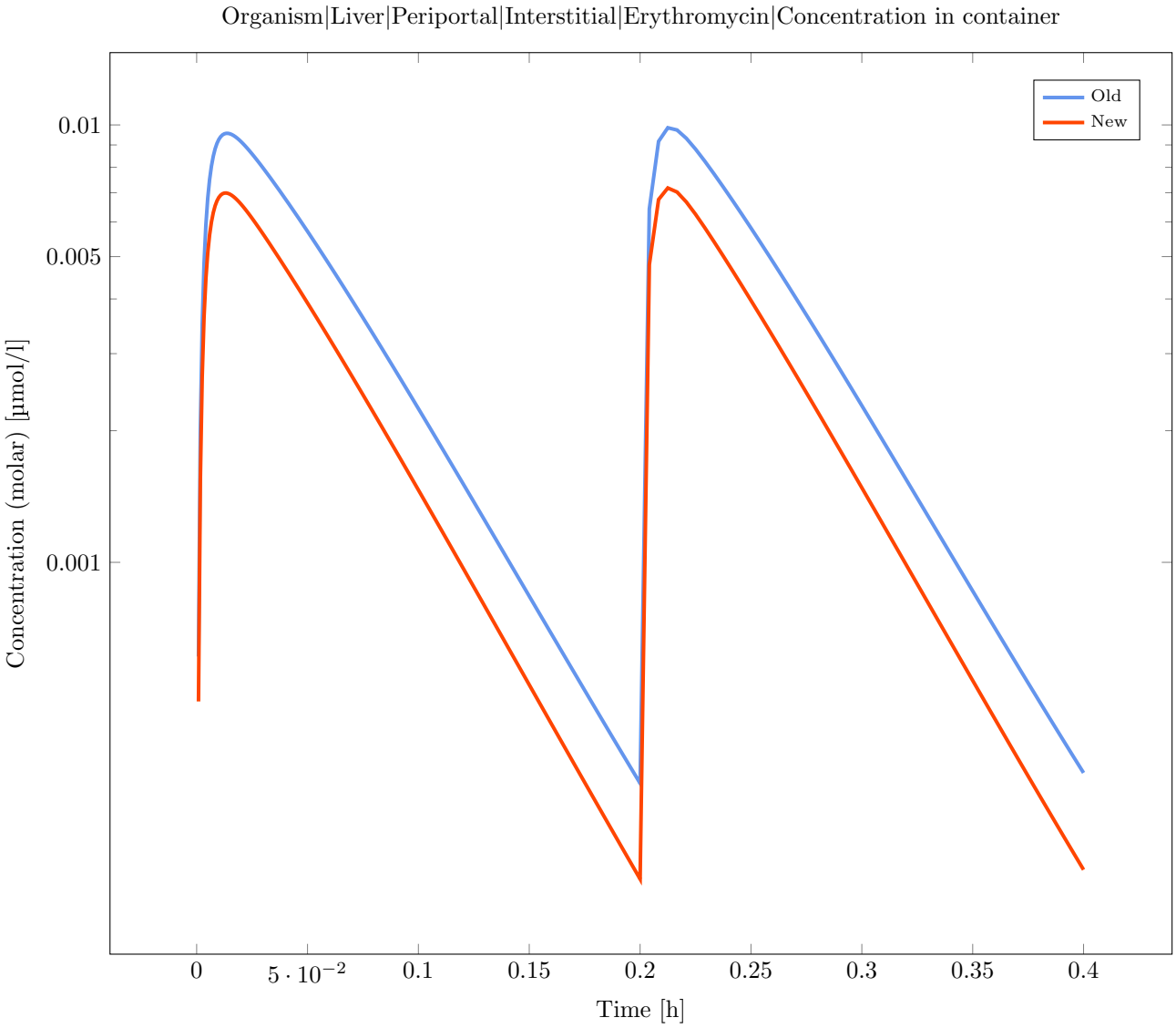
Organism|Liver|Periportal|Interstitial|Erythromycin|Concentration in container
Old
New
0.01
0.005
0.001
Concentration (molar) [µmol/l]
0
$5 \cdot 10^{-2}$
0.1
0.15
0.2
0.25
0.3
0.35
0.4
Time [h]

Figure 1.9

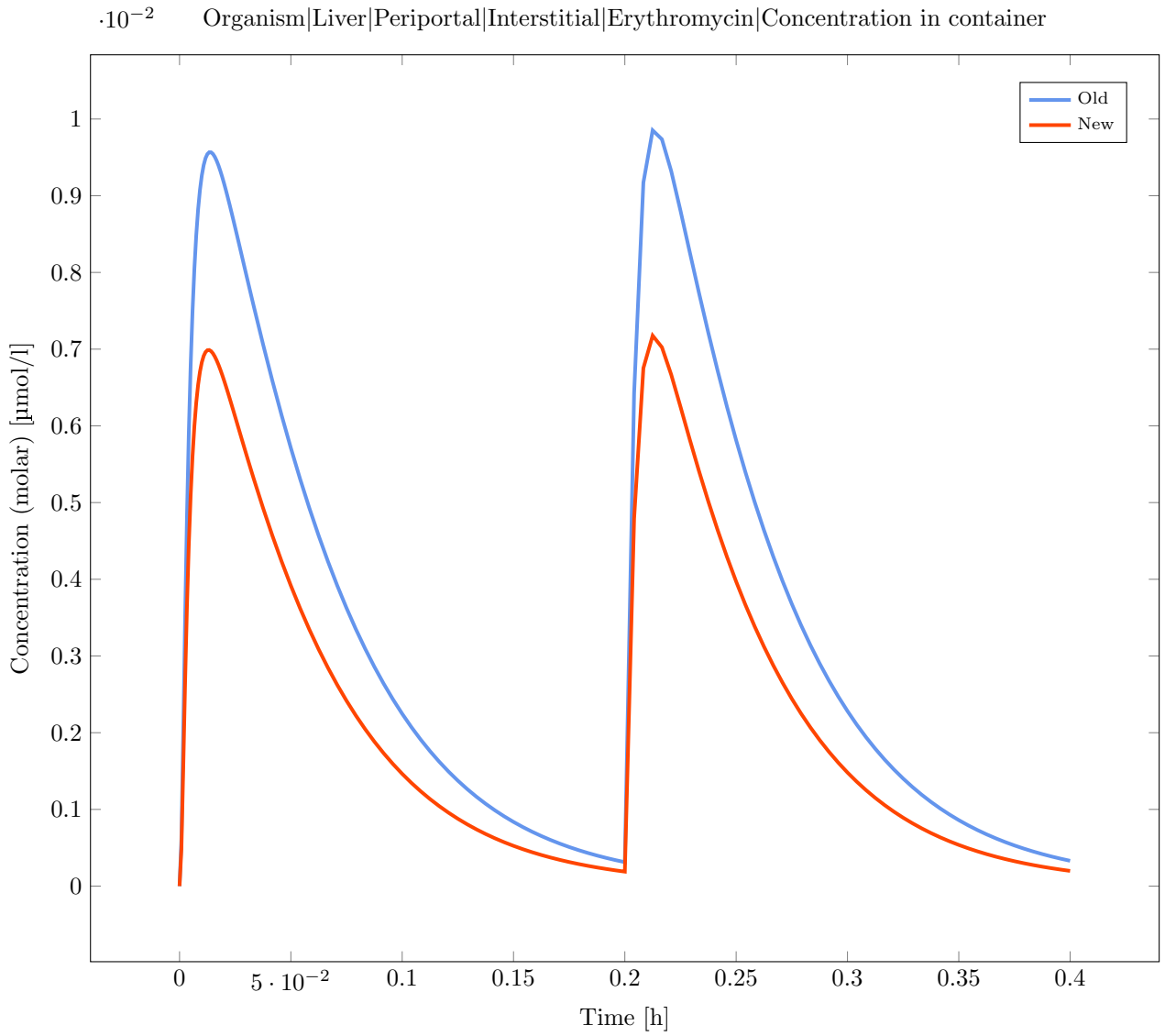

Figure 1.10

Simulation: DDI_MultipleCombinations-10_MM_Mixed_Mixed_Mechanismbased_Mechanismbased
Result of the validation: Invalid
Absolute Tolerance: 1.00E-10
Relative Tolerance: 1.00E-5

Output Path: Organism|Liver|Periportal|MechanismBased_3|Interstitial Unbound
Deviation for 'Organism|Liver|Periportal|MechanismBased_3|Interstitial Unbound' is 48.79% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.49

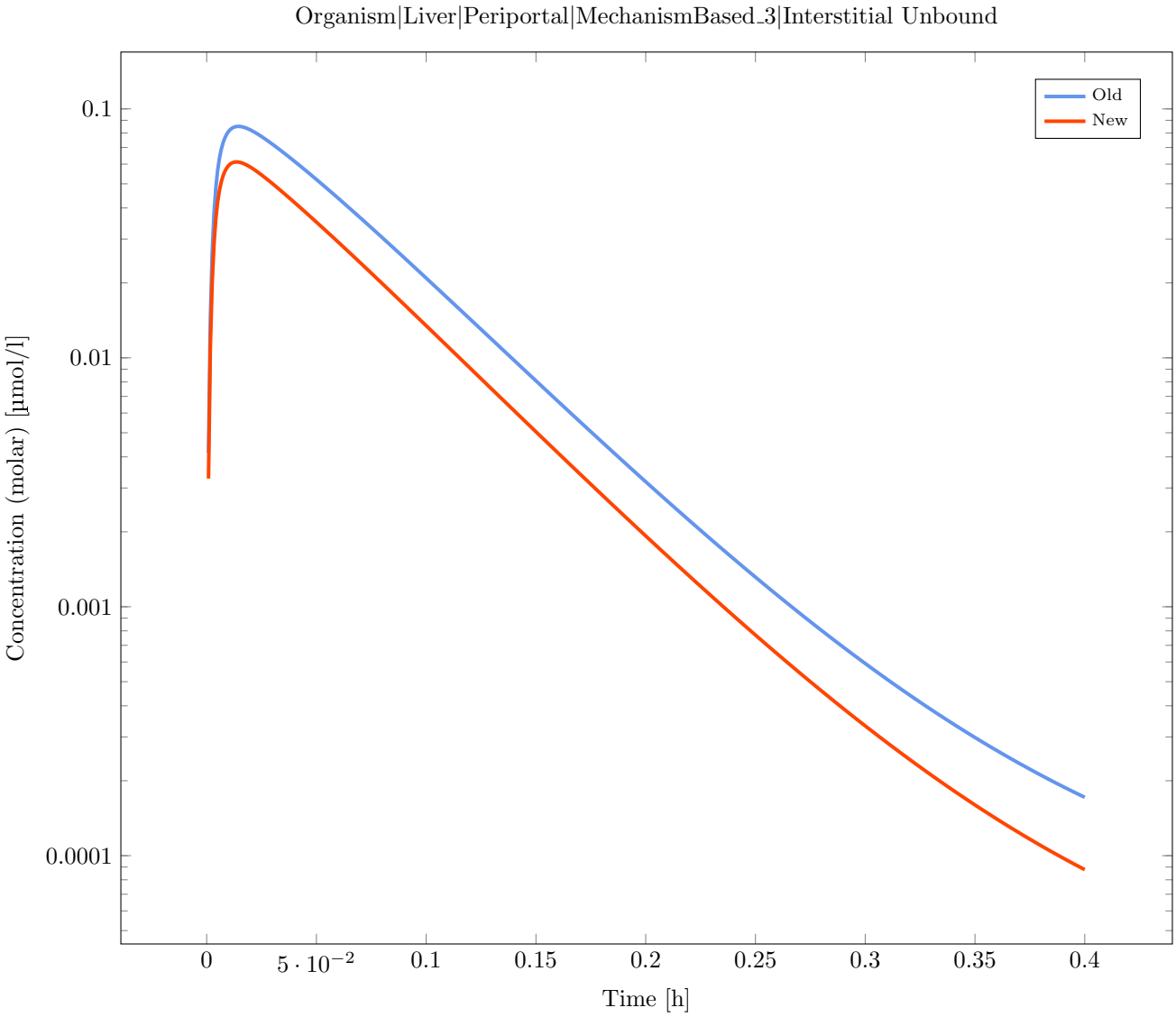
Organism|Liver|Periportal|MechanismBased_3|Interstitial Unbound
Old
New
Concentration (molar) [µmol/l]
0.1
0.01
0.001
0.0001
0
5 · 10−2
0.1
0.15
0.2
0.25
0.3
0.35
0.4
Time [h]

Figure 1.11

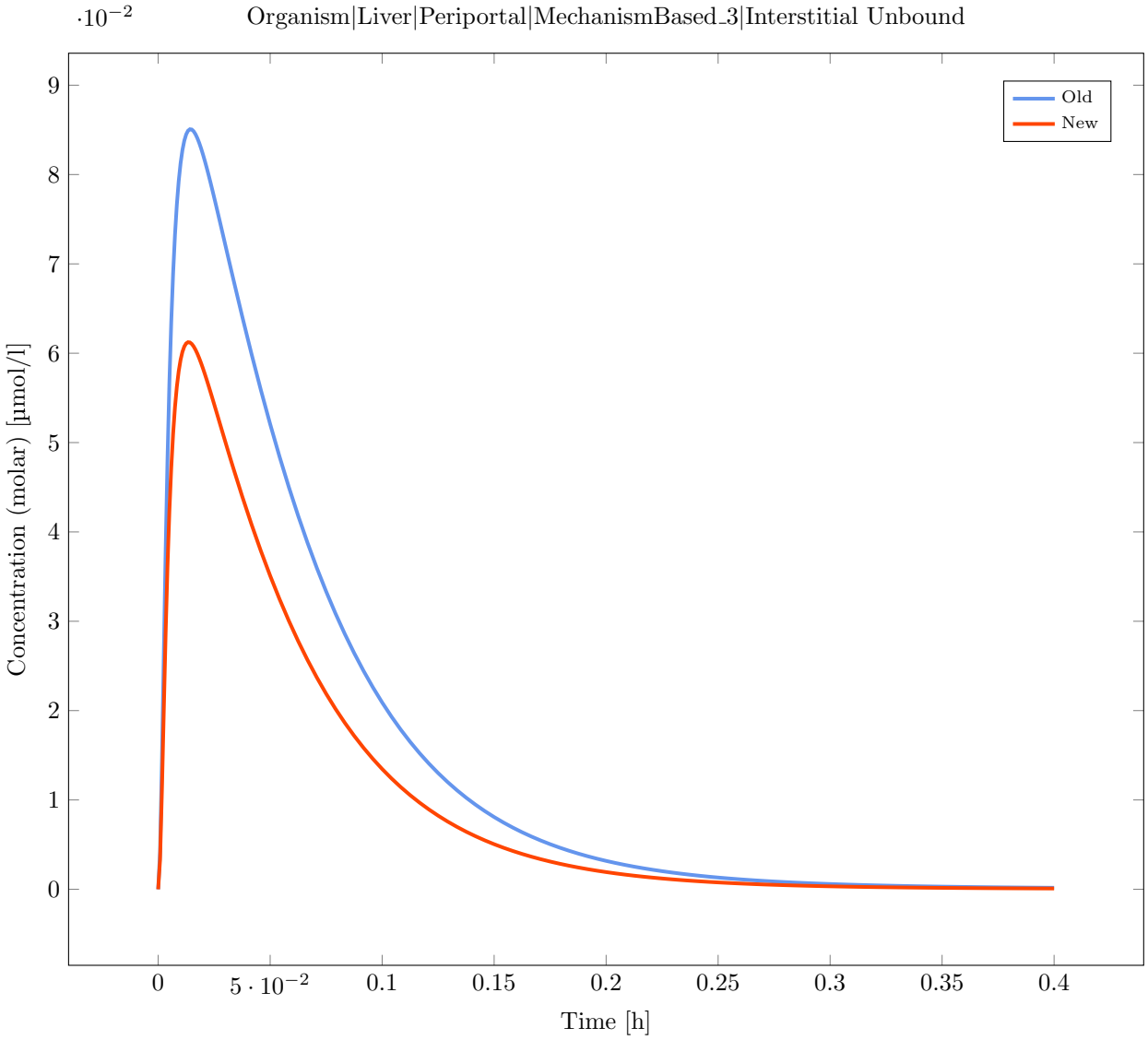

Figure 1.12

Simulation: DDI_MultipleCombinations-11_MM_Mechanismbased_Mechanismbased_Induction_-Induction
Result of the validation: Invalid
Absolute Tolerance: 1.00E-10
Relative Tolerance: 1.00E-5

Output Path: Organism|Lumen|ColonSigmoid|Rifampicin|Fraction of oral drug mass absorbed into mucosa segment
Deviation for 'Organism|Lumen|ColonSigmoid|Rifampicin|Fraction of oral drug mass absorbed into mucosa segment' is 1897.69% and is greater than the allowed max. tolerance of 3.00%
Deviation: 18.98

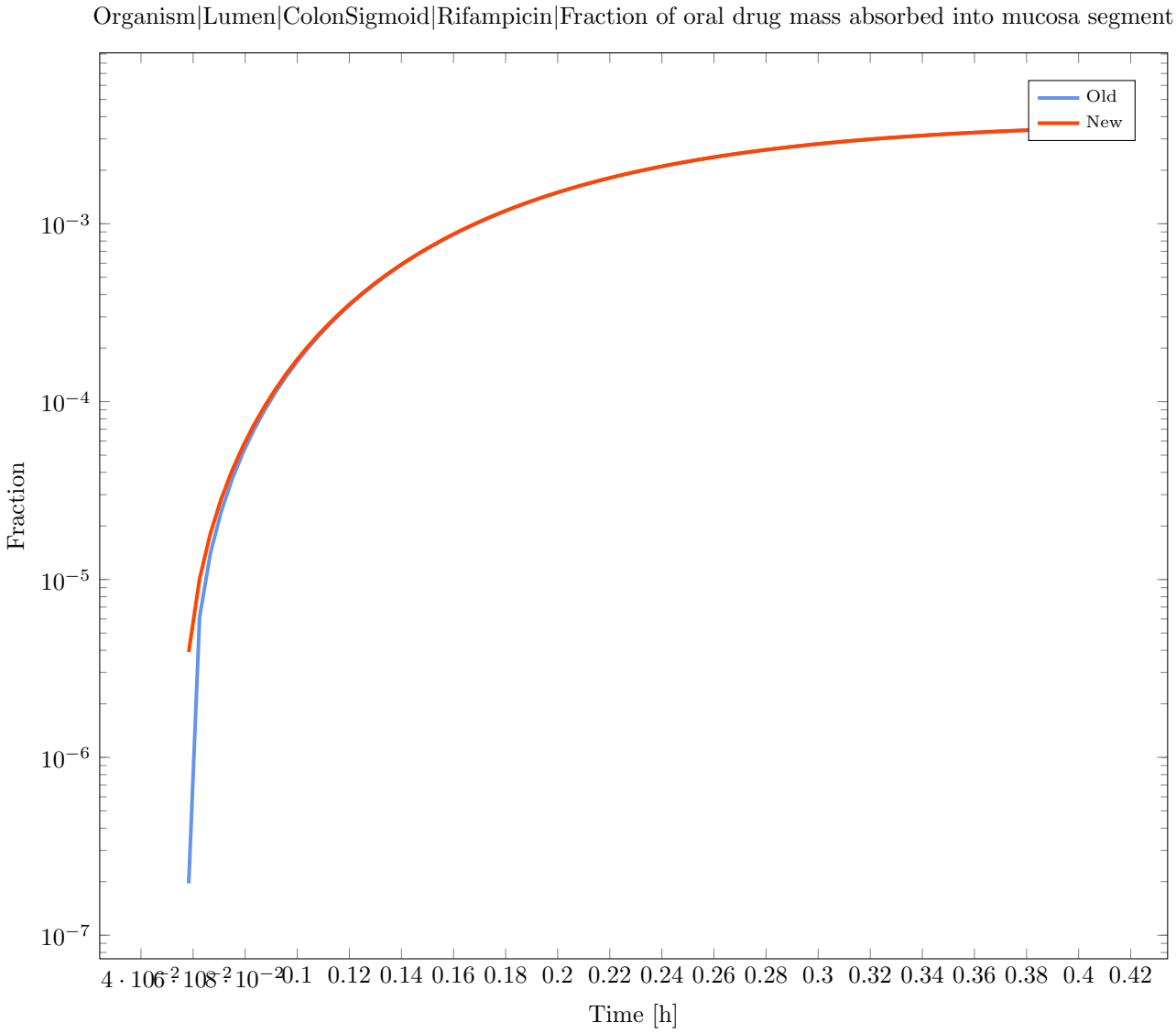
Organism|Lumen|ColonSigmoid|Rifampicin|Fraction of oral drug mass absorbed into mucosa segment
Old
New
Fraction
10^{-3}
10^{-4}
10^{-5}
10^{-6}
10^{-7}
$4 \cdot 10^{-2}$
$6 \cdot 10^{-2}$
$8 \cdot 10^{-2}$
10^{-1}
0.1
0.12
0.14
0.16
0.18
0.2
0.22
0.24
0.26
0.28
0.3
0.32
0.34
0.36
0.38
0.4
0.42
Time [h]

Figure 1.13

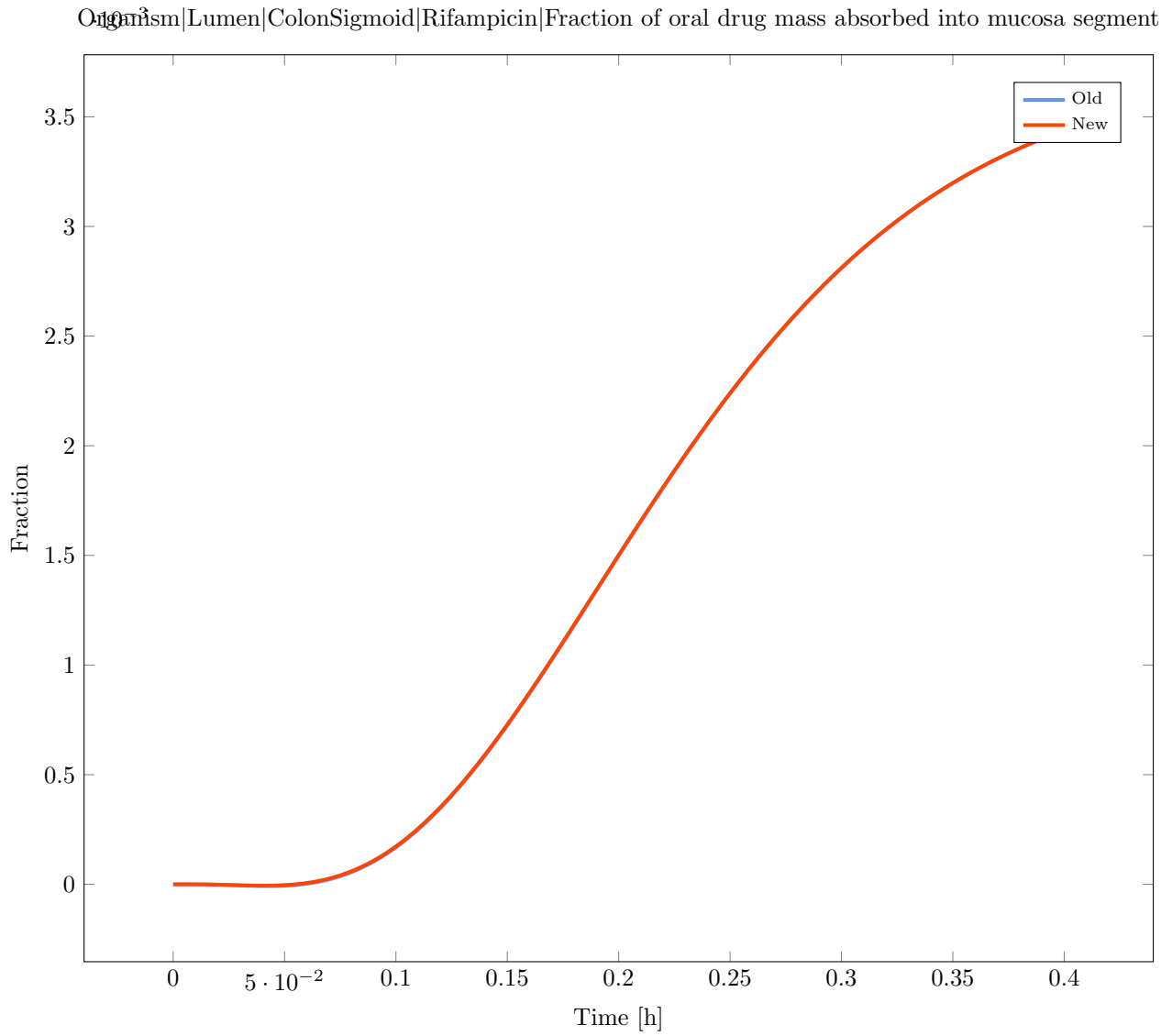

Figure 1.14

Simulation: DDI_MultipleCombinations-12_MM_All_DDI_Types
Result of the validation: Invalid
Absolute Tolerance: 1.00E-10
Relative Tolerance: 1.00E-5

Output Path: Organism|Liver|Periportal|Interstitial|MechanismBased_3|Concentration in container
Deviation for 'Organism|Liver|Periportal|Interstitial|MechanismBased_3|Concentration in container' is 48.14% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.48

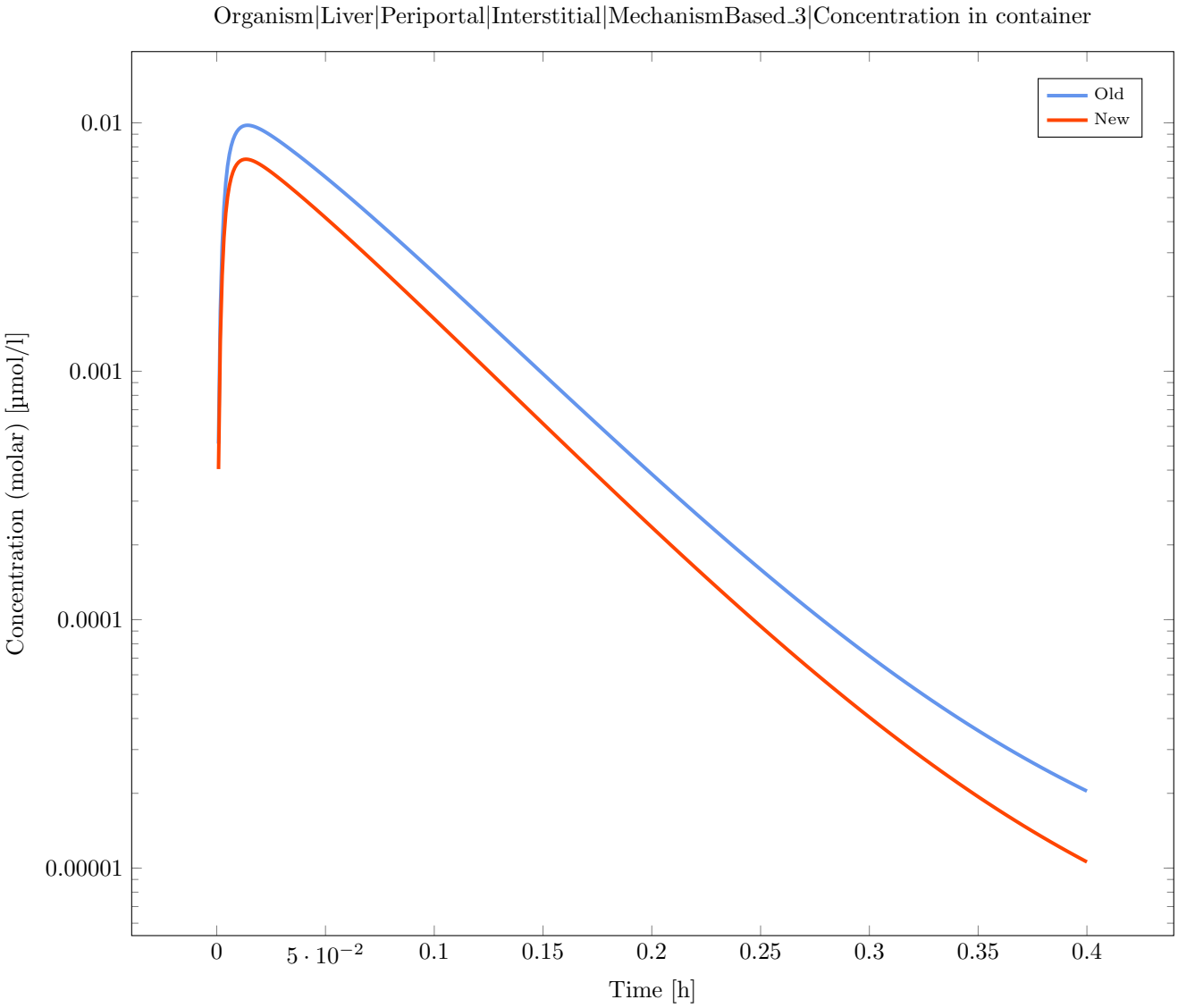
Organism|Liver|Periportal|Interstitial|MechanismBased_3|Concentration in container
Old
New
Concentration (molar) [µmol/l]
0.01
0.001
0.0001
0.00001
0
5 · 10−2
0.1
0.15
0.2
0.25
0.3
0.35
0.4
Time [h]

Figure 1.15

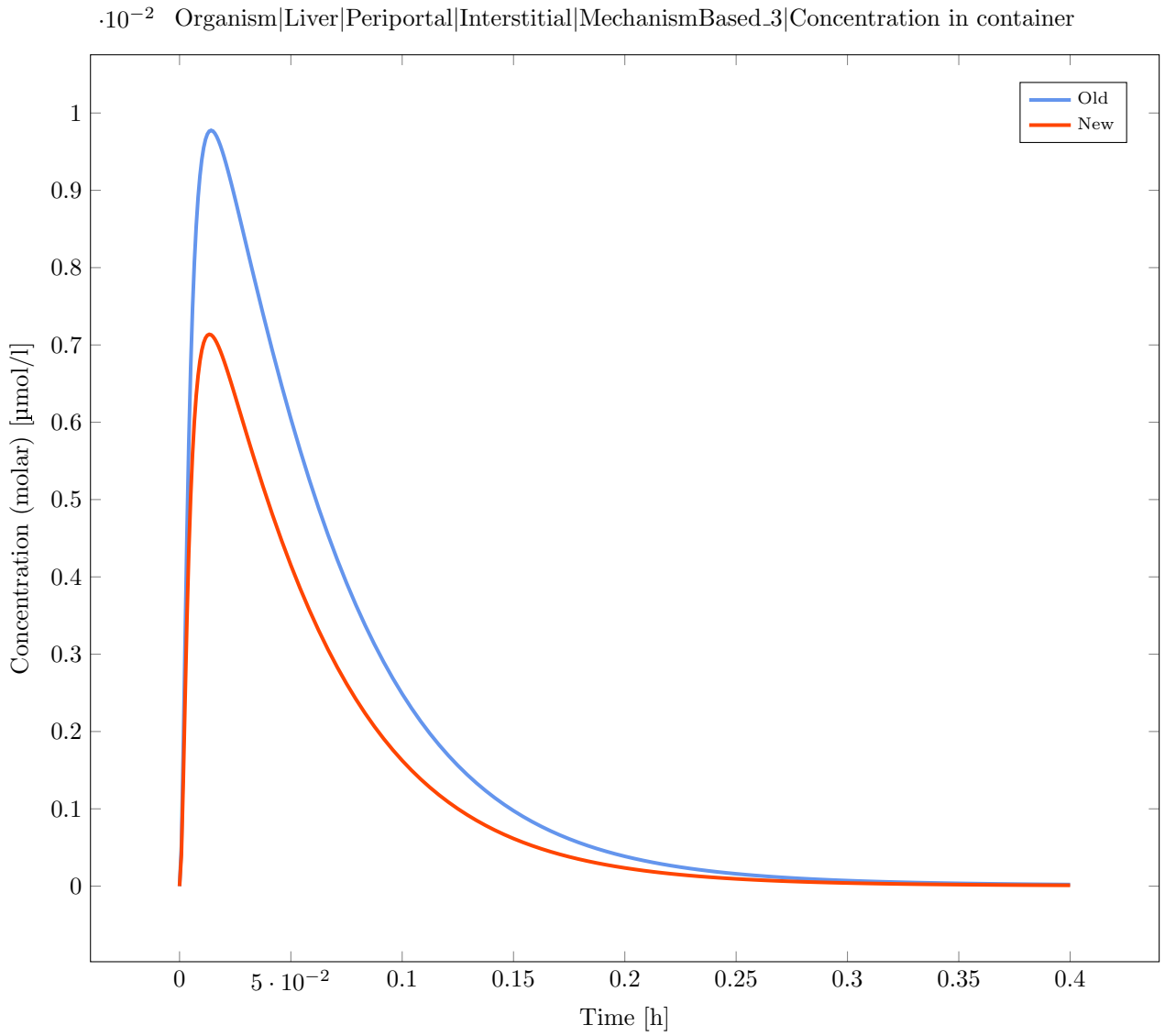

Figure 1.16

Simulation: DDI_MultipleCombinations-25_1st_Mechanismbased_Mechanismbased
Result of the validation: Invalid
Absolute Tolerance: 1.00E-10
Relative Tolerance: 1.00E-5

Output Path: Organism|Liver|Periportal|Erythromycin|Interstitial Unbound
Deviation for 'Organism|Liver|Periportal|Erythromycin|Interstitial Unbound' is 45.33% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.45

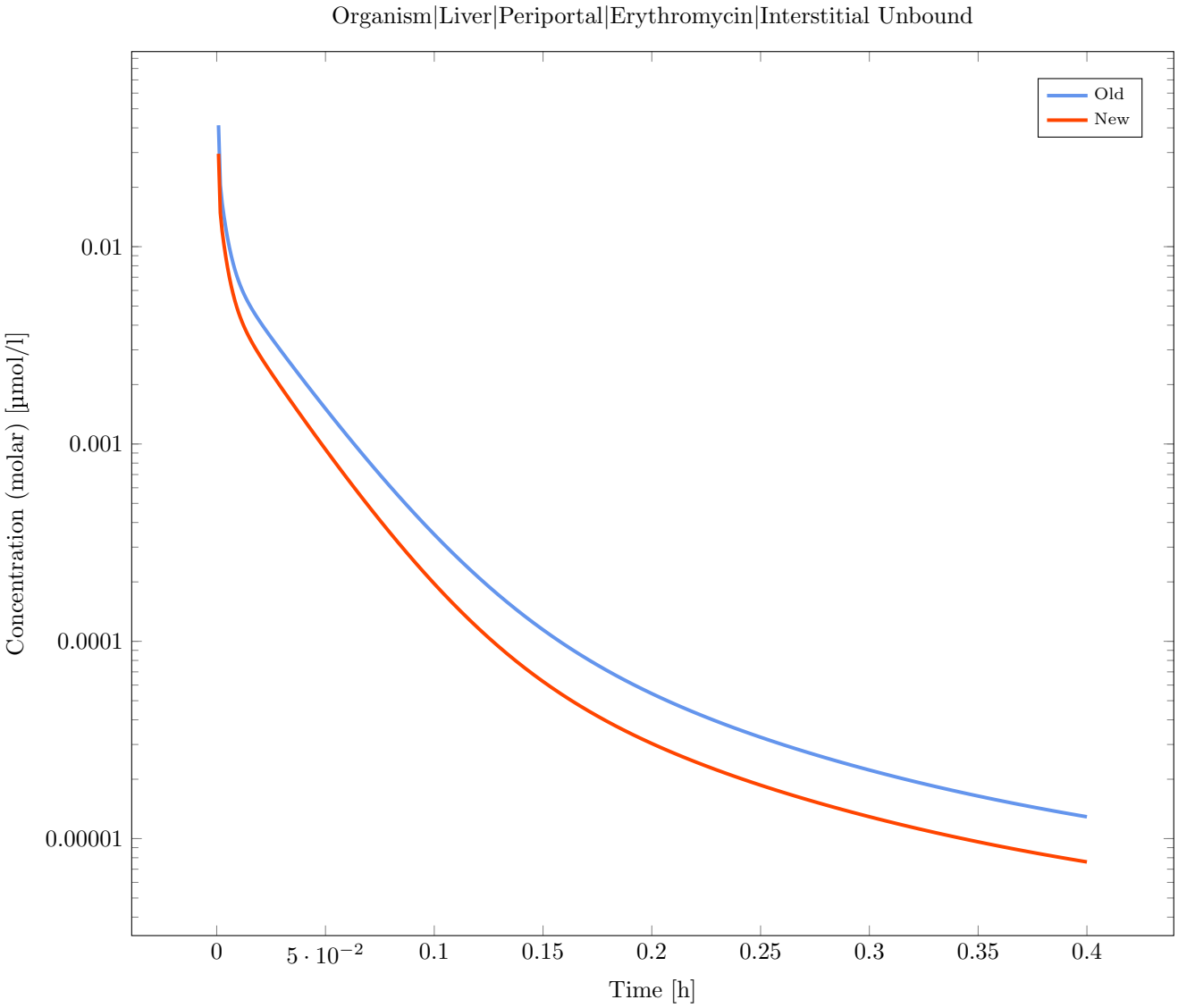
Organism|Liver|Periportal|Erythromycin|Interstitial Unbound
Old
New
Concentration (molar) [µmol/l]
0.01
0.001
0.0001
0.00001
0
5 · 10−2
0.1
0.15
0.2
0.25
0.3
0.35
0.4
Time [h]

Figure 1.17

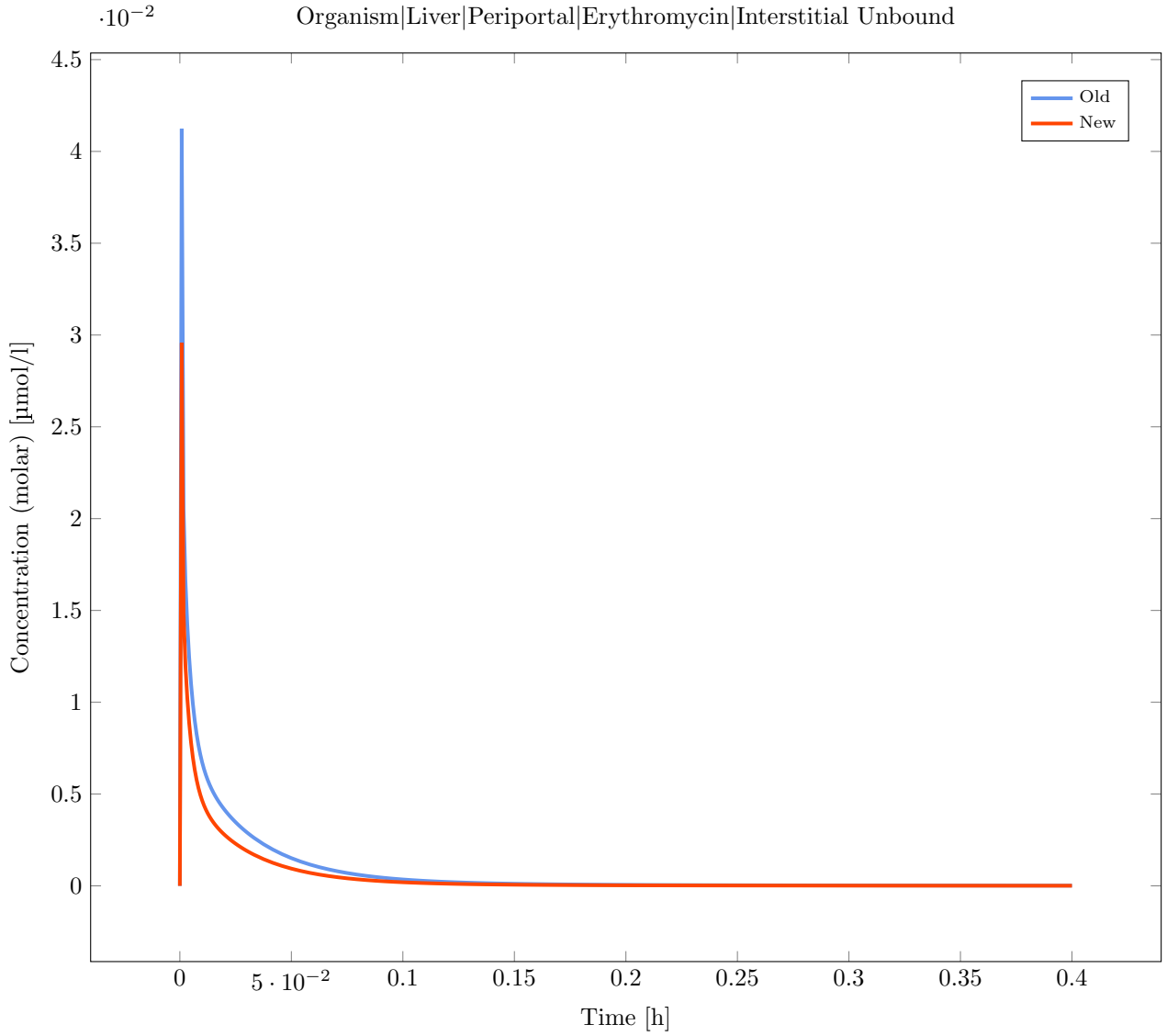

Figure 1.18

Simulation: DDI_MultipleCombinations-26_1st_Induction_Induction
Result of the validation: Invalid
Absolute Tolerance: 1.00E-10
Relative Tolerance: 1.00E-5

Output Path: Organism|Lumen|ColonDescendens|Rifampicin|Fraction of oral drug mass absorbed into mucosa segment
Deviation for 'Organism|Lumen|ColonDescendens|Rifampicin|Fraction of oral drug mass absorbed into mucosa segment' is 1122.69% and is greater than the allowed max. tolerance of 3.00%
Deviation: 11.23

Organism|Lumen|ColonDescendens|Rifampicin|Fraction of oral drug mass absorbed into mucosa segment

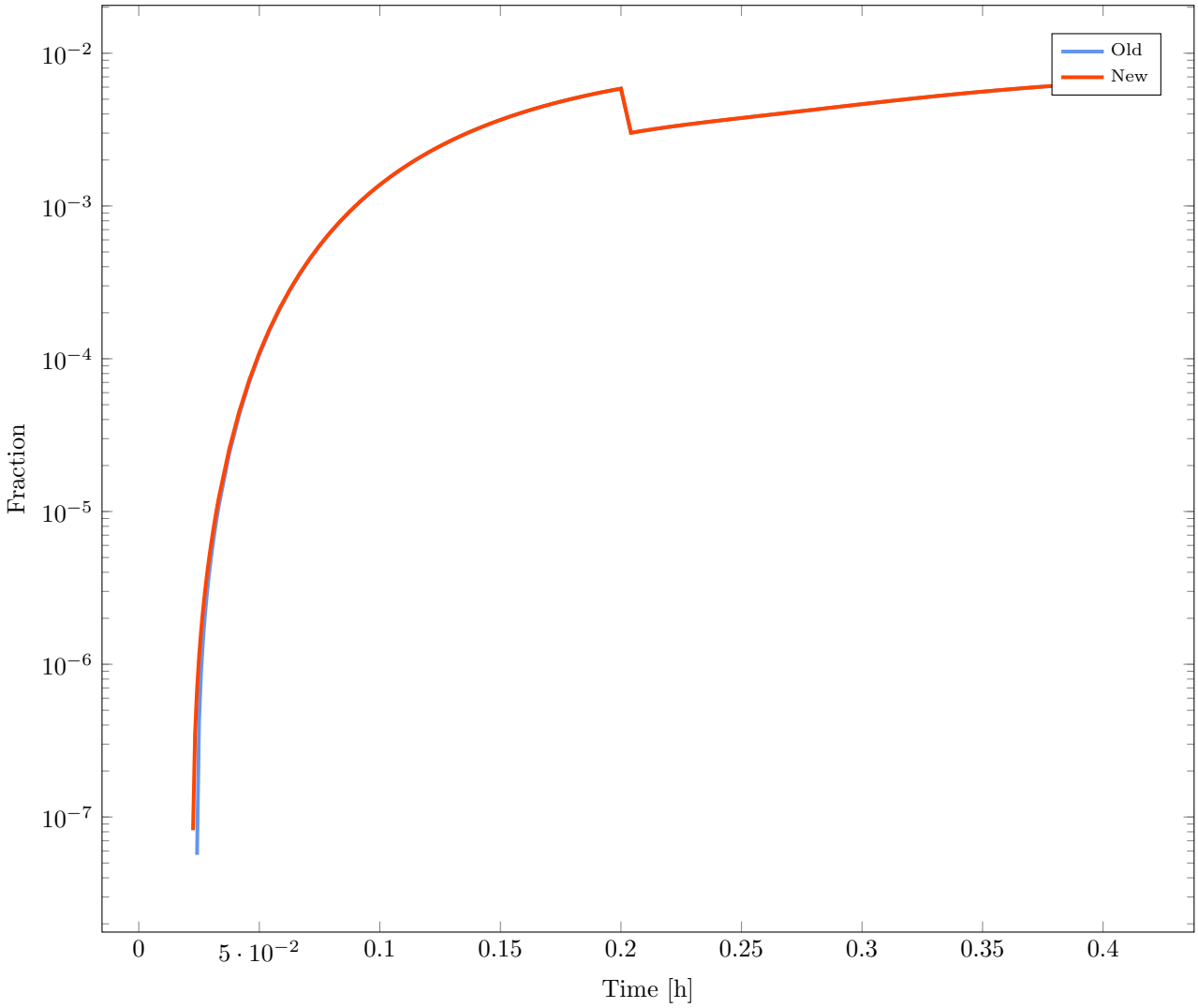

Figure 1.19

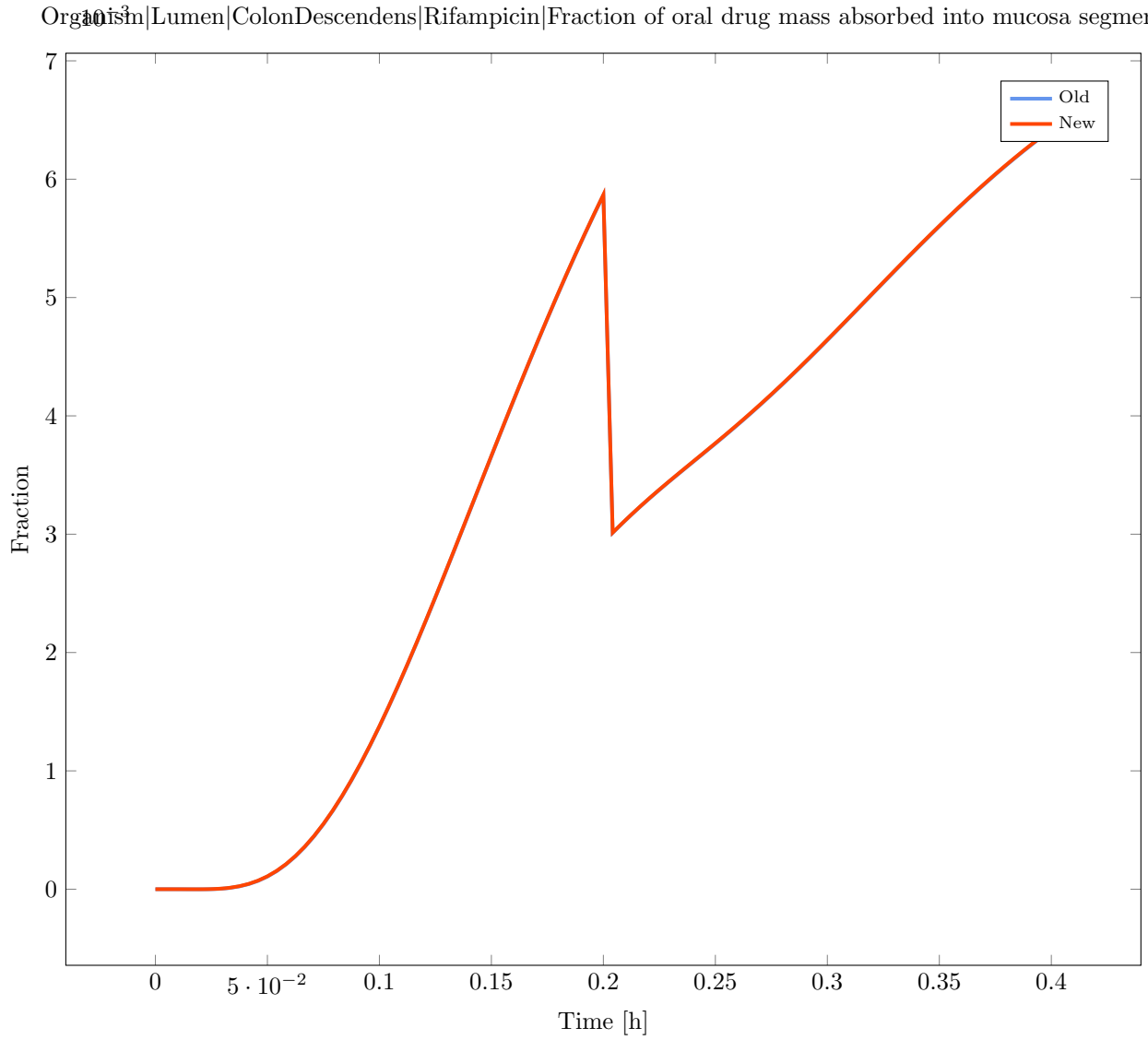

Figure 1.20

Simulation: DDI_MultipleCombinations-27_1st_Competitive_Competitive_Mechanismbased_Mechanismbased
Result of the validation: Invalid
Absolute Tolerance: 1.00E-10
Relative Tolerance: 1.00E-5

Output Path: Organism|Liver|Periportal|Interstitial|Erythromycin|Concentration in container
Deviation for 'Organism|Liver|Periportal|Interstitial|Erythromycin|Concentration in container' is 56.47% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.56

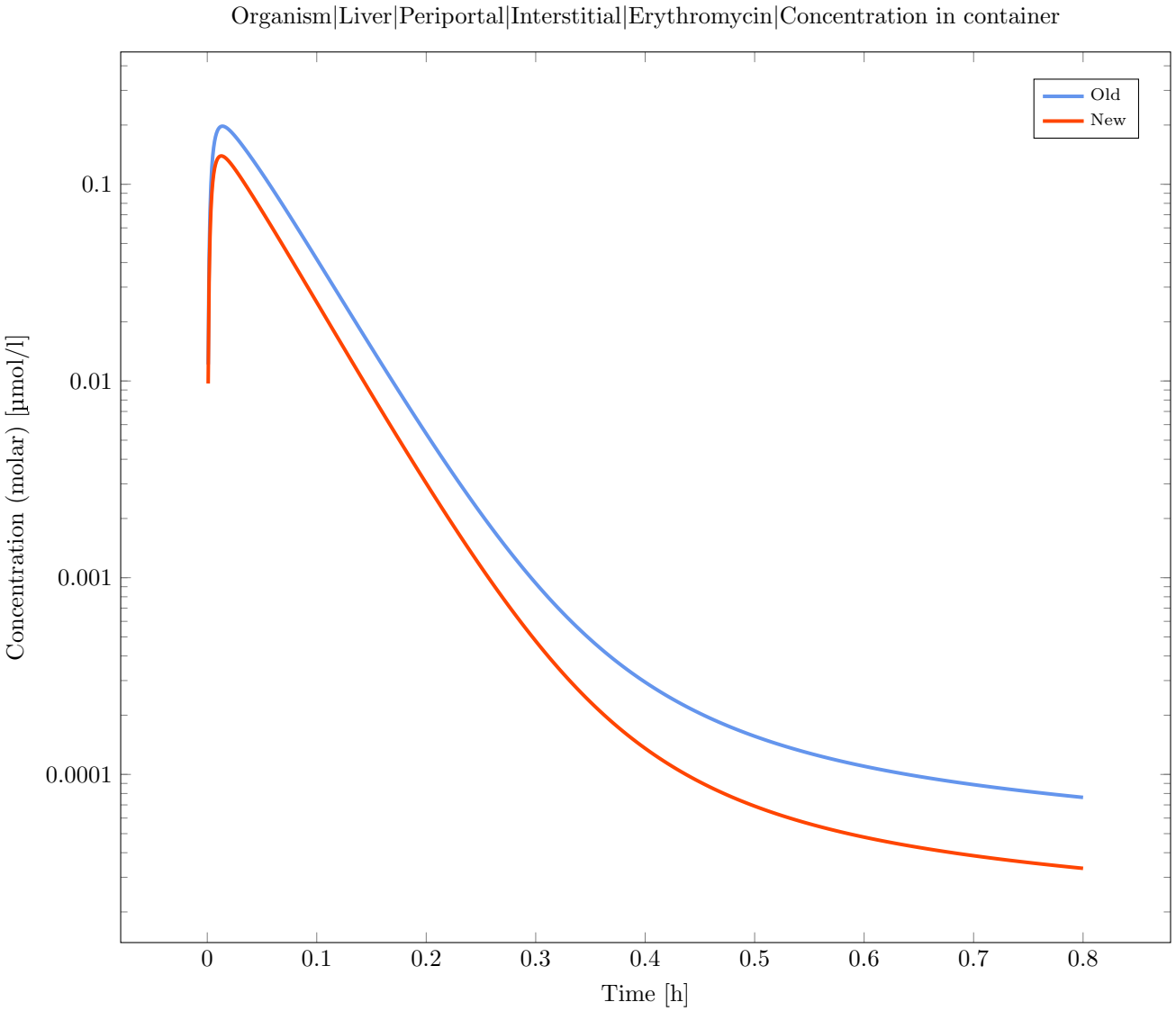
Organism|Liver|Periportal|Interstitial|Erythromycin|Concentration in container
Old
New
Concentration (molar) [µmol/l]
0.1
0.01
0.001
0.0001
0
0.1
0.2
0.3
0.4
0.5
0.6
0.7
0.8
Time [h]

Figure 1.21

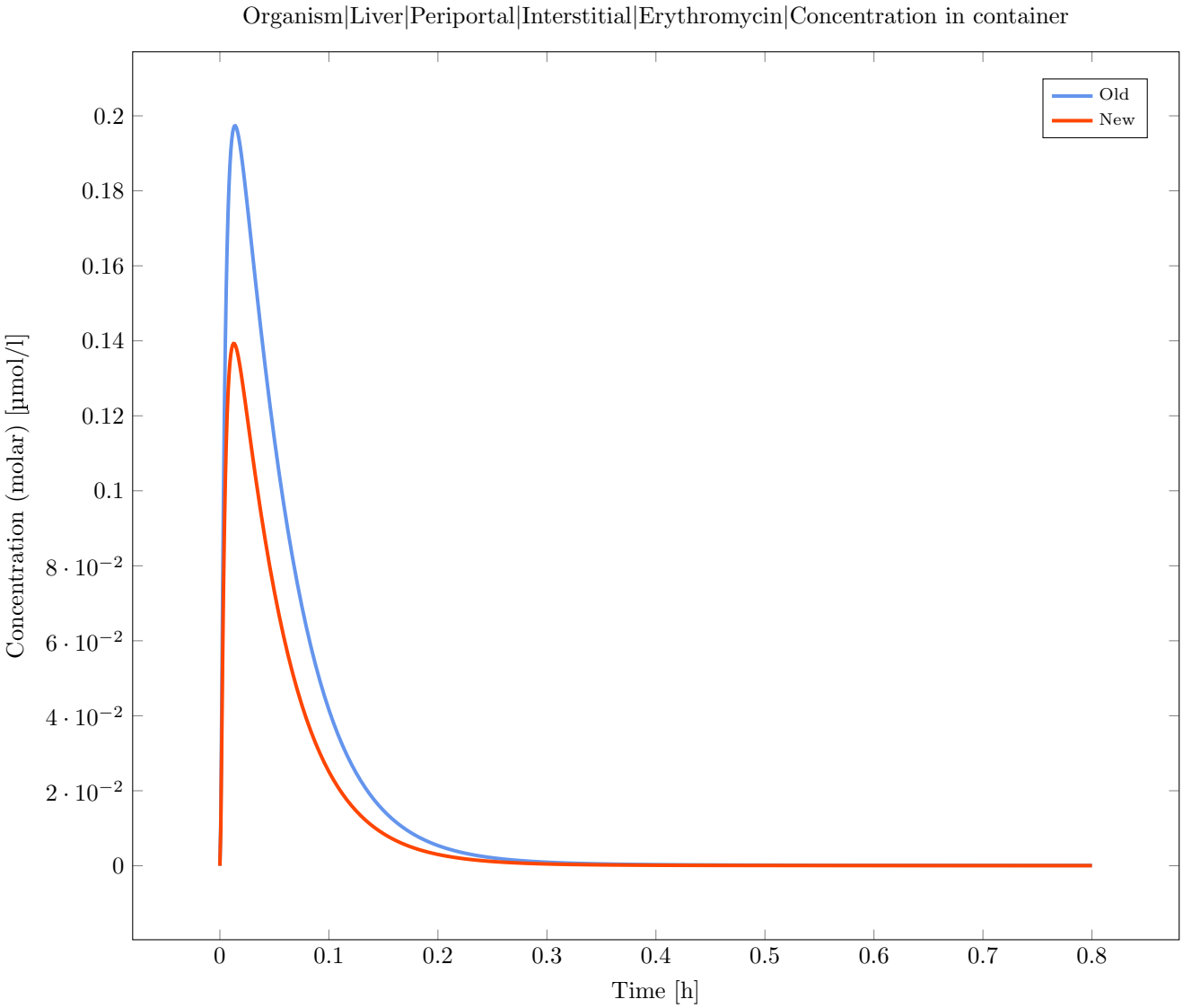

Figure 1.22

Simulation: DDI_MultipleCombinations-28_1st_Uncompetitive_Uncompetitive_Mechanismbased_-Mechanismbased
Result of the validation: Invalid
Absolute Tolerance: 1.00E-10
Relative Tolerance: 1.00E-5

Output Path: Organism|Liver|Periportal|Interstitial|MechanismBased_4|Concentration in container
Deviation for 'Organism|Liver|Periportal|Interstitial|MechanismBased_4|Concentration in container' is 49.46% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.49

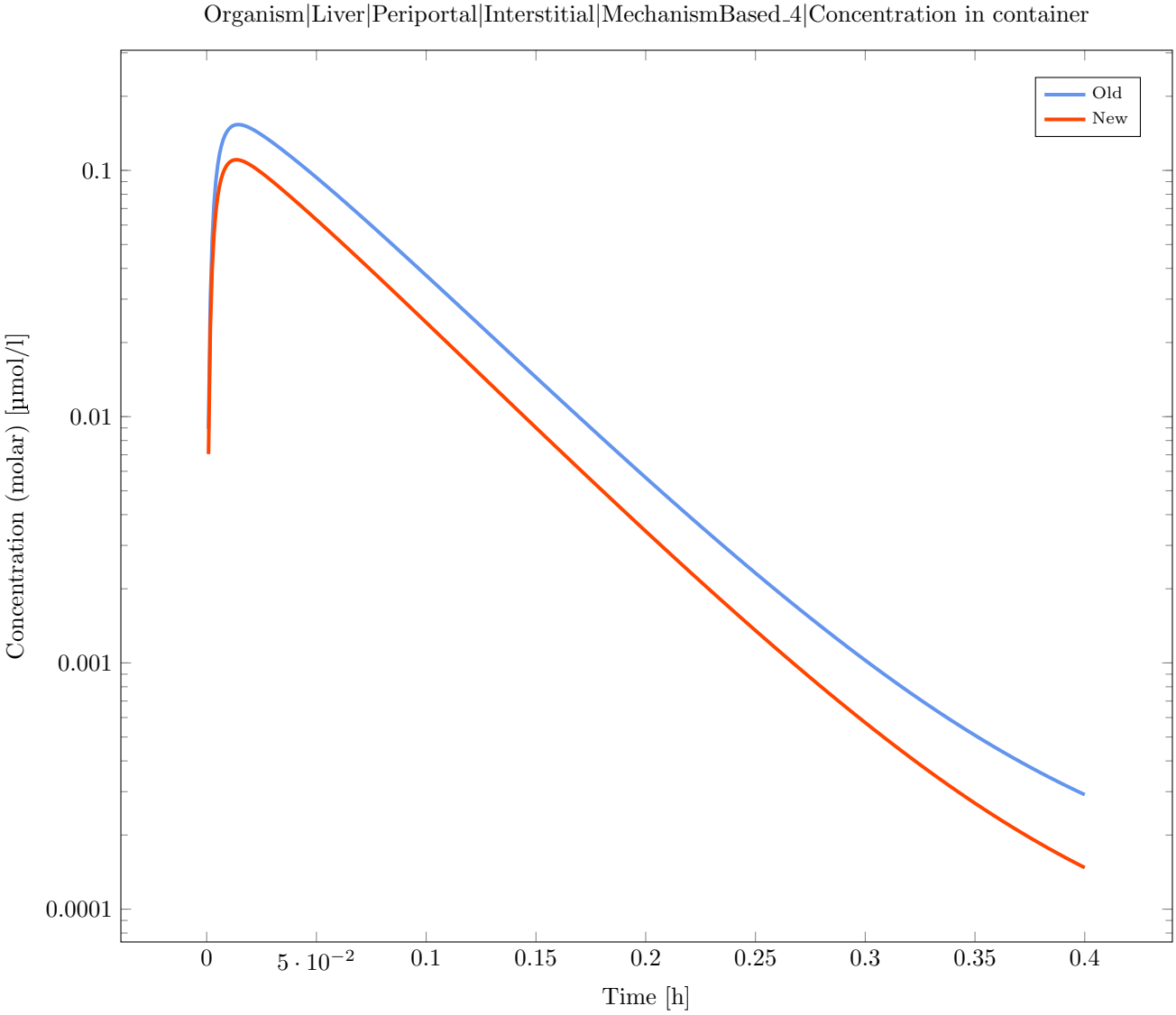
Organism|Liver|Periportal|Interstitial|MechanismBased_4|Concentration in container
Old
New
Concentration (molar) [µmol/l]
0.1
0.01
0.001
0.0001
0
5 · 10−2
0.1
0.15
0.2
0.25
0.3
0.35
0.4
Time [h]

Figure 1.23

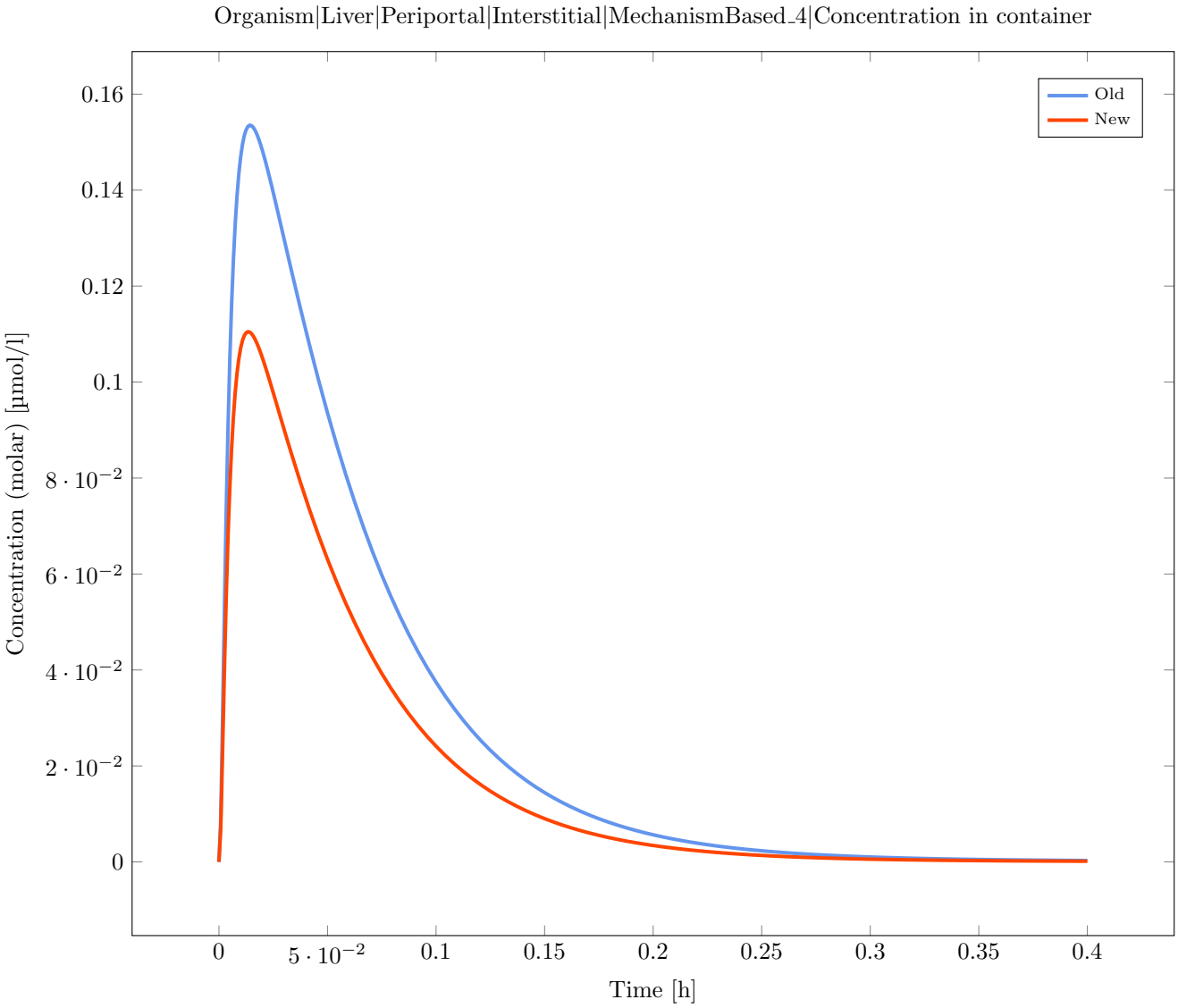

Figure 1.24

Simulation: DDI_MultipleCombinations-29_1st_Noncompetitive_Noncompetitive_Mechanismbased_-Mechanismbased
Result of the validation: Invalid
Absolute Tolerance: 1.00E-10
Relative Tolerance: 1.00E-5

Output Path: Organism|Liver|Periportal|Interstitial|Erythromycin|Concentration in container
Deviation for 'Organism|Liver|Periportal|Interstitial|Erythromycin|Concentration in container' is 39.95% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.40

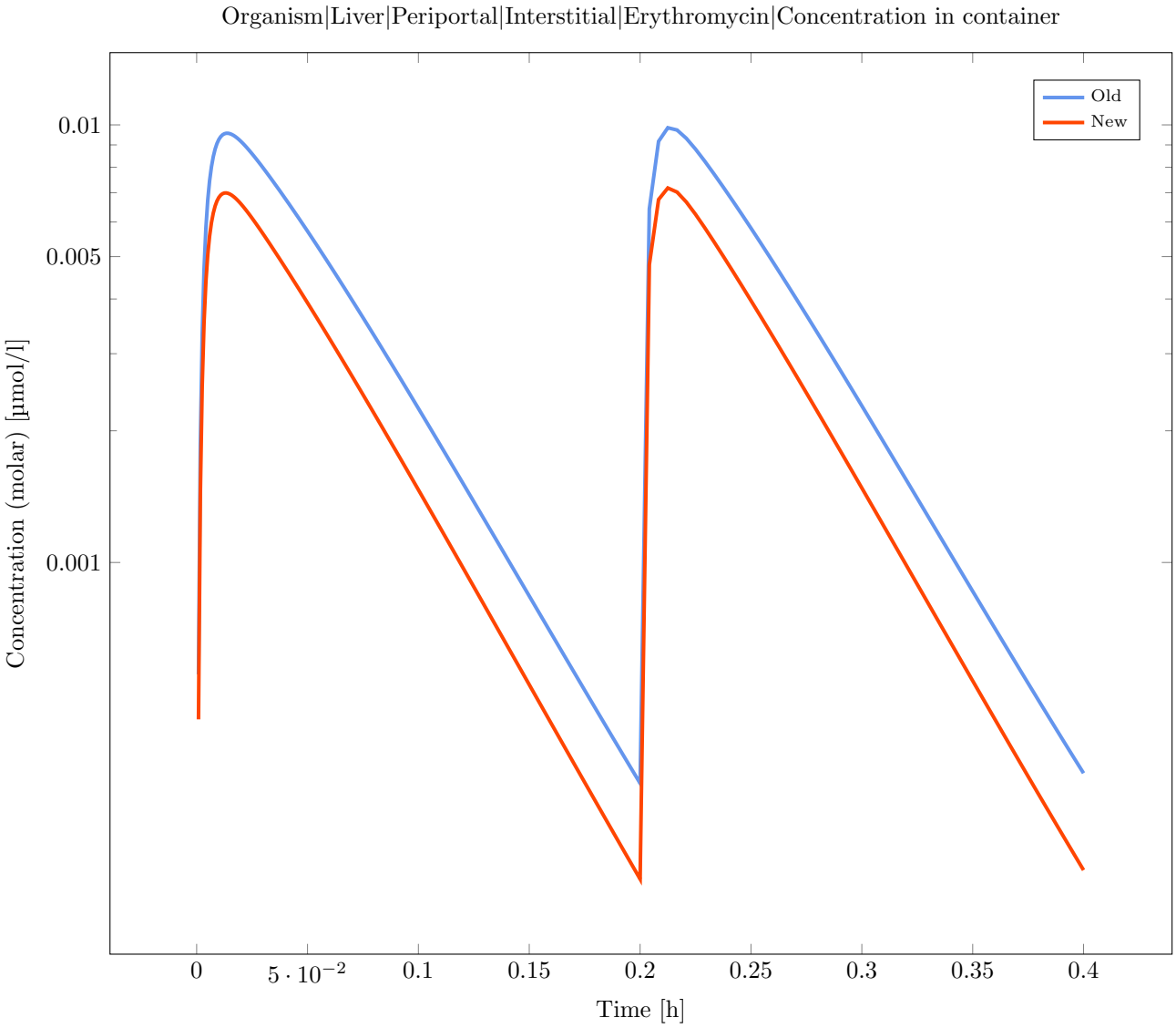
Organism|Liver|Periportal|Interstitial|Erythromycin|Concentration in container
Old
New
0.01
0.005
0.001
Concentration (molar) [µmol/l]
0
5 · 10−2
0.1
0.15
0.2
0.25
0.3
0.35
0.4
Time [h]

Figure 1.25

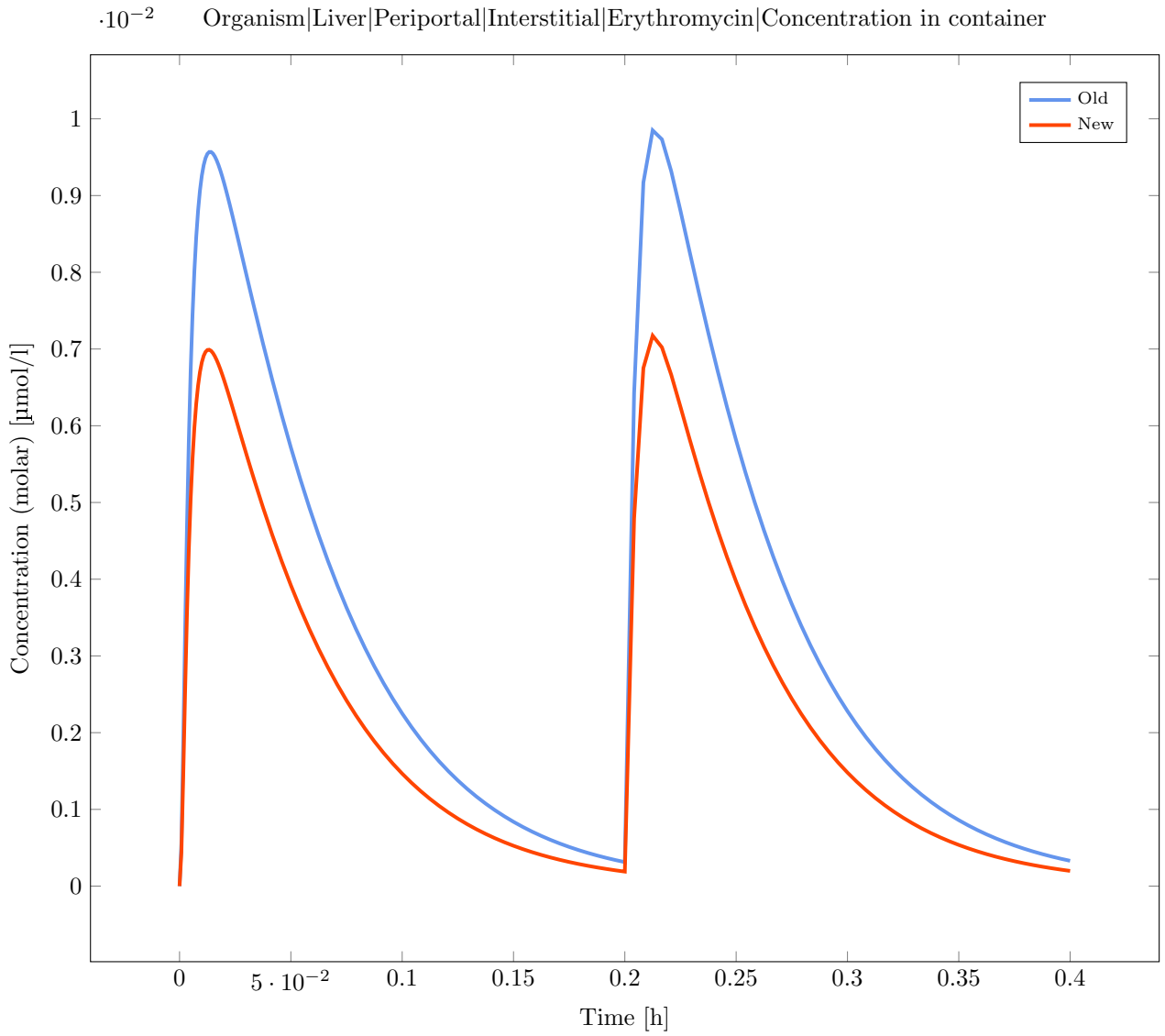

Figure 1.26

Simulation: DDI_MultipleCombinations-30_1st_Mixed_Mixed_Mechanismbased_Mechanismbased
Result of the validation: Invalid
Absolute Tolerance: 1.00E-10
Relative Tolerance: 1.00E-5

Output Path: Organism|Liver|Periportal|Interstitial|MechanismBased_3|Concentration in container
Deviation for 'Organism|Liver|Periportal|Interstitial|MechanismBased_3|Concentration in container' is 48.80% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.49

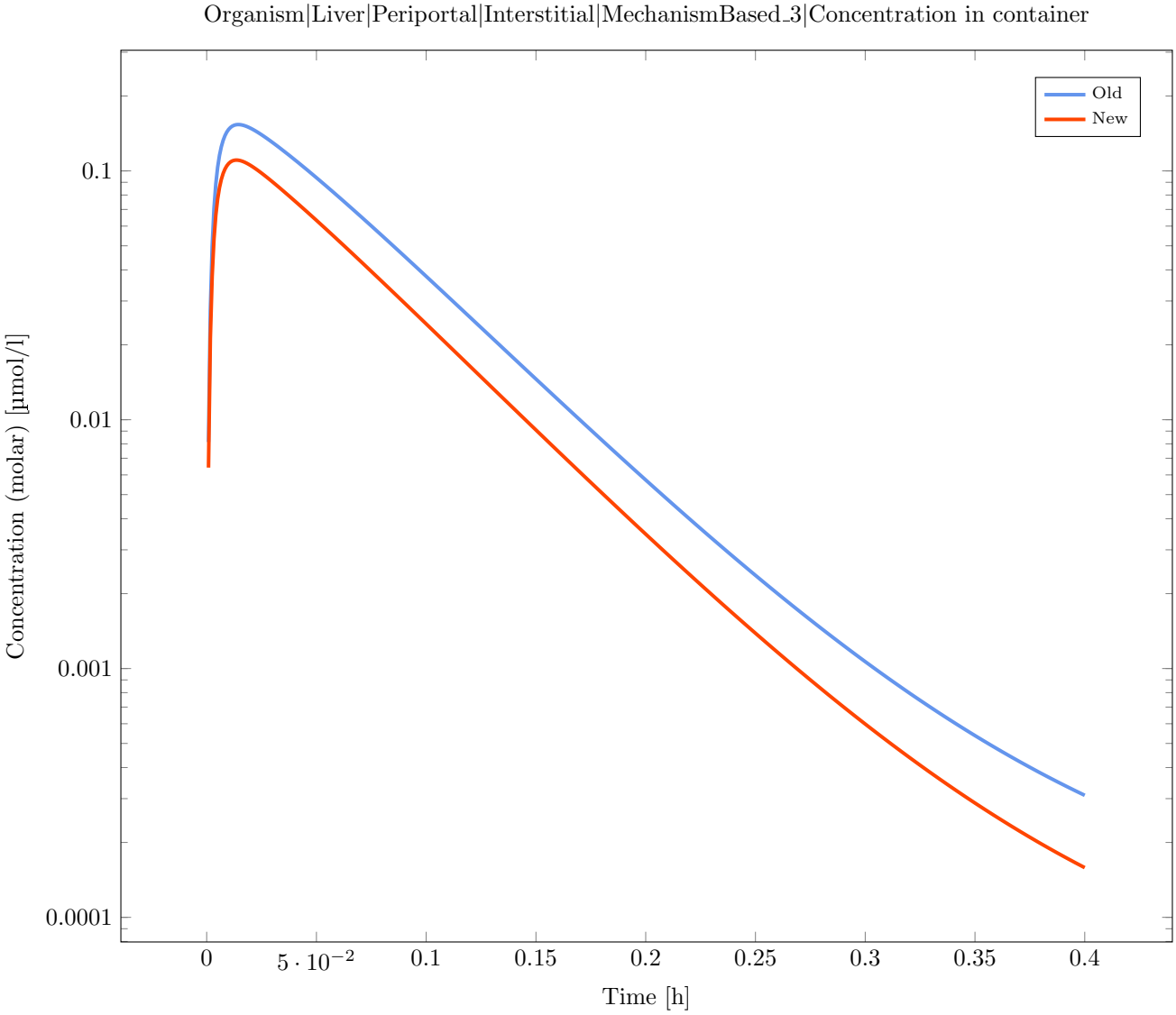
Organism|Liver|Periportal|Interstitial|MechanismBased_3|Concentration in container
Old
New
Concentration (molar) [µmol/l]
0.1
0.01
0.001
0.0001
0
5 · 10−2
0.1
0.15
0.2
0.25
0.3
0.35
0.4
Time [h]

Figure 1.27

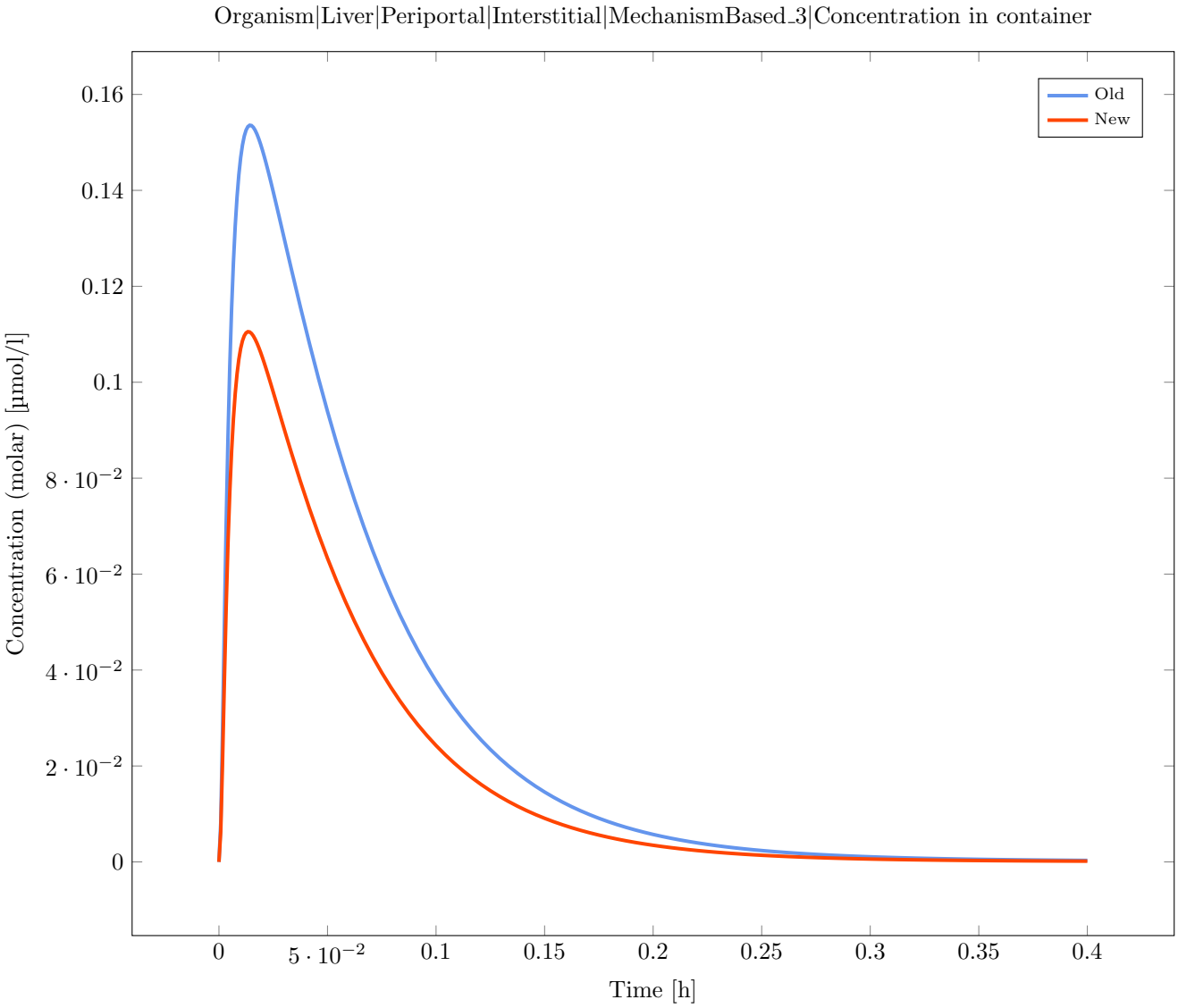

Figure 1.28

Simulation: DDI_MultipleCombinations-31_1st_Mechanismbased_Mechanismbased_Induction_Induction
Result of the validation: Invalid
Absolute Tolerance: 1.00E-10
Relative Tolerance: 1.00E-5

Output Path: Organism|Lumen|ColonDescendens|Rifampicin|Fraction of oral drug mass absorbed into mucosa segment
Deviation for 'Organism|Lumen|ColonDescendens|Rifampicin|Fraction of oral drug mass absorbed into mucosa segment' is 825.21% and is greater than the allowed max. tolerance of 3.00%
Deviation: 8.25

Organism|Lumen|ColonDescendens|Rifampicin|Fraction of oral drug mass absorbed into mucosa segment

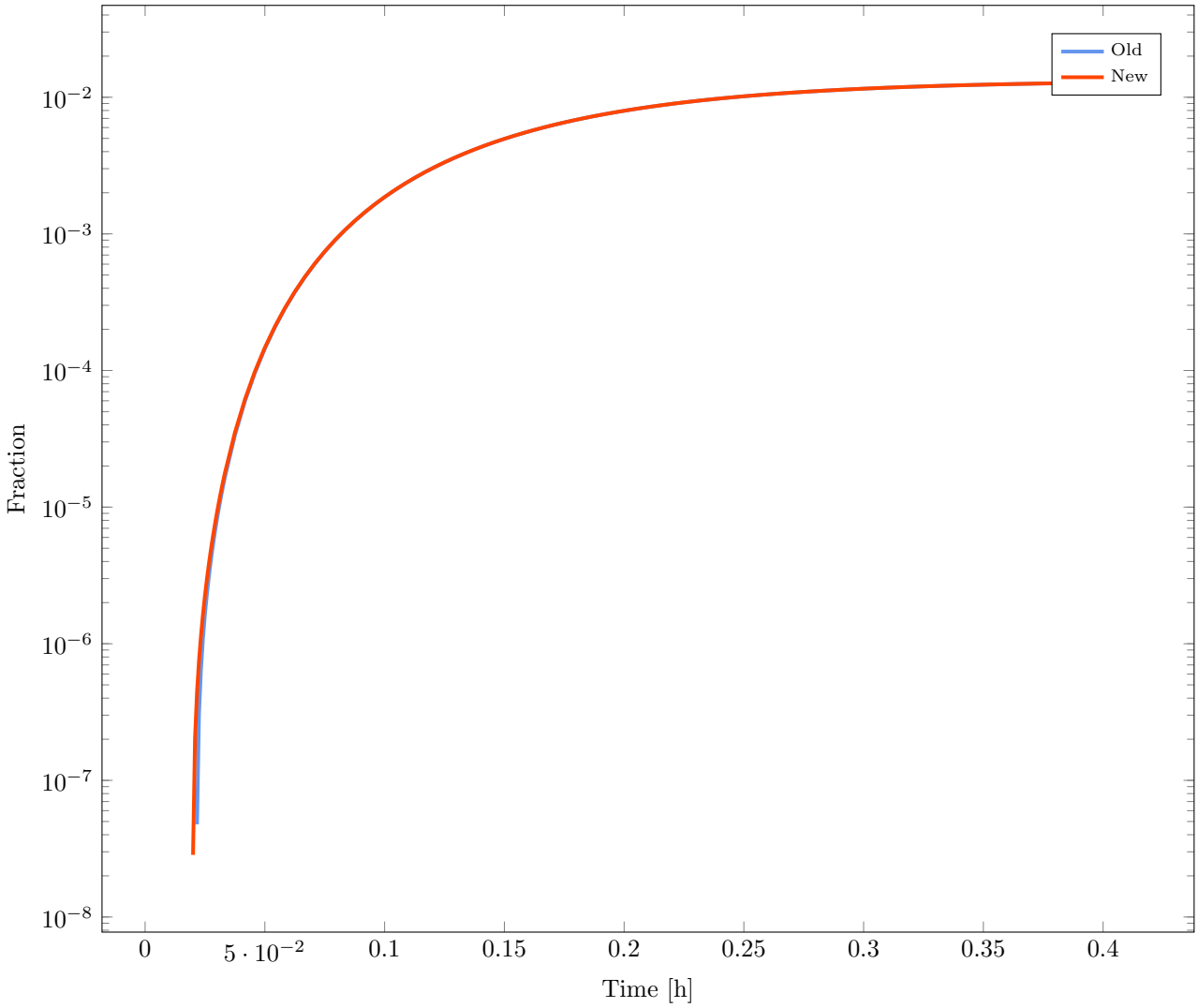

Figure 1.29

Organism|Lumen|ColonDescendens|Rifampicin|Fraction of oral drug mass absorbed into mucosa segment

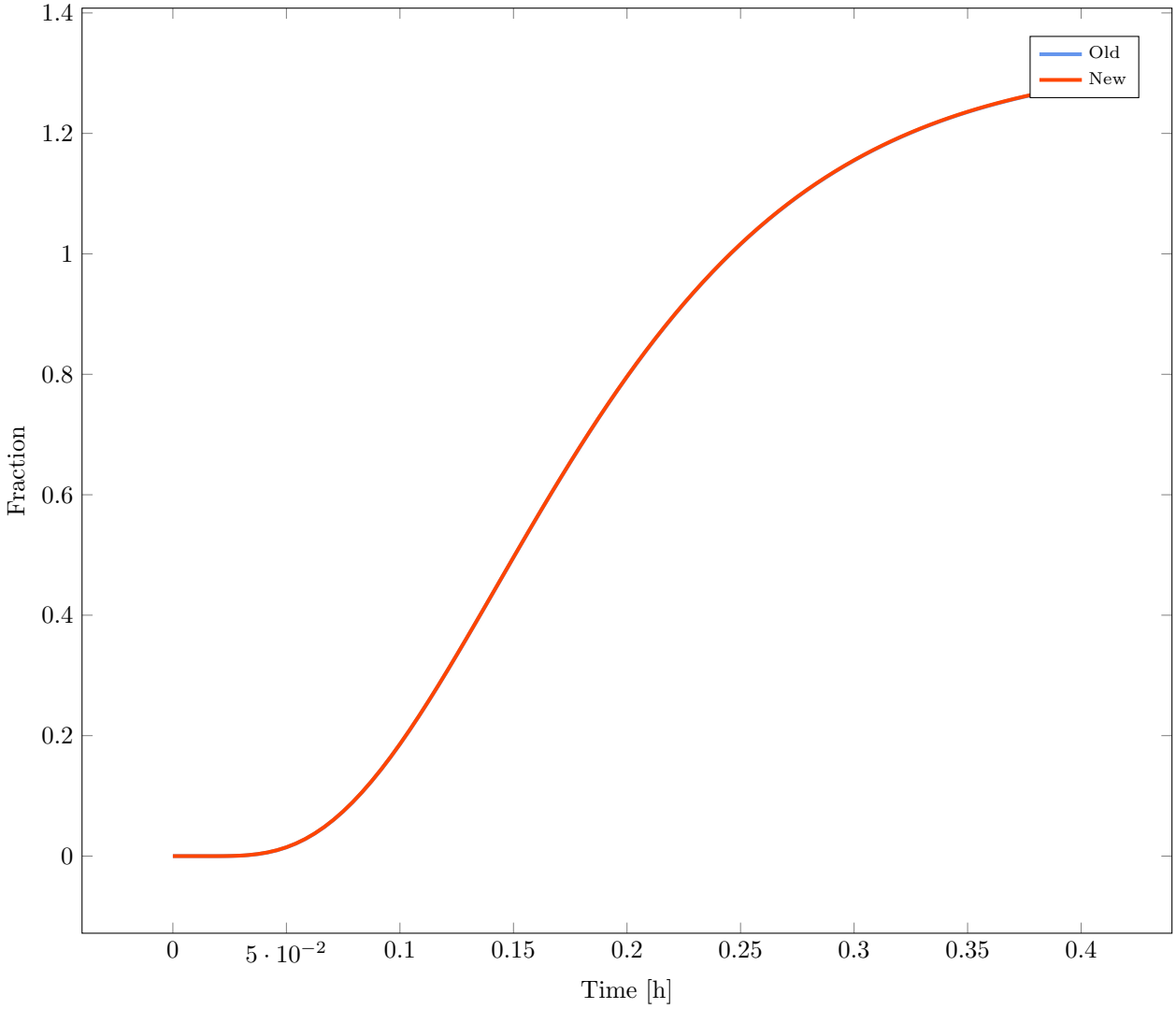

Figure 1.30

Simulation: DDI_MultipleCombinations-32_1st_All_DDI_Types
Result of the validation: Invalid
Absolute Tolerance: 1.00E-10
Relative Tolerance: 1.00E-5

Output Path: Organism|Liver|Periportal|Interstitial|MechanismBased_3|Concentration in container
Deviation for 'Organism|Liver|Periportal|Interstitial|MechanismBased_3|Concentration in container' is 48.14% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.48

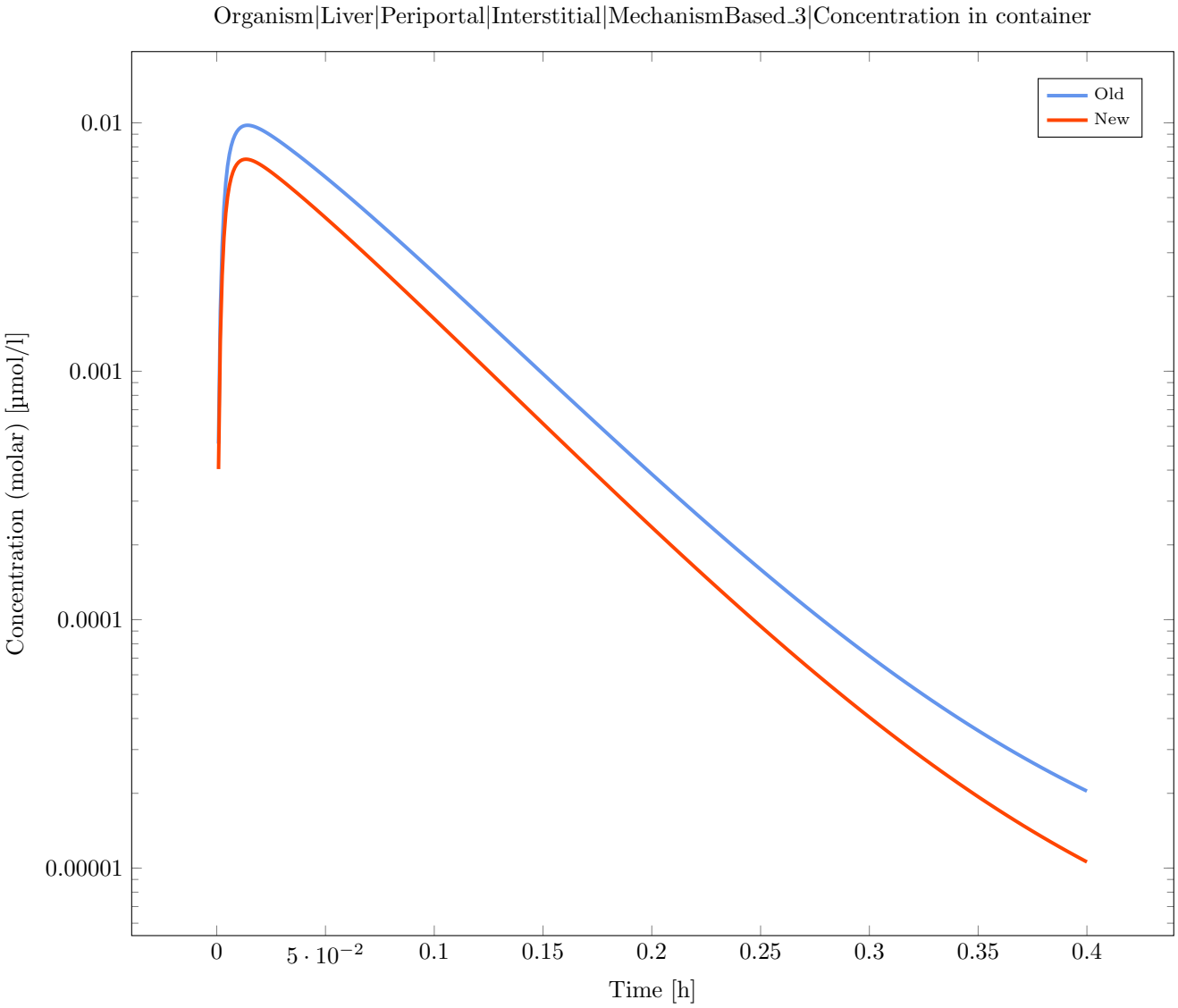
Organism|Liver|Periportal|Interstitial|MechanismBased_3|Concentration in container
Old
New
Concentration (molar) [µmol/l]
0.01
0.001
0.0001
0.00001
0
5 · 10−2
0.1
0.15
0.2
0.25
0.3
0.35
0.4
Time [h]

Figure 1.31

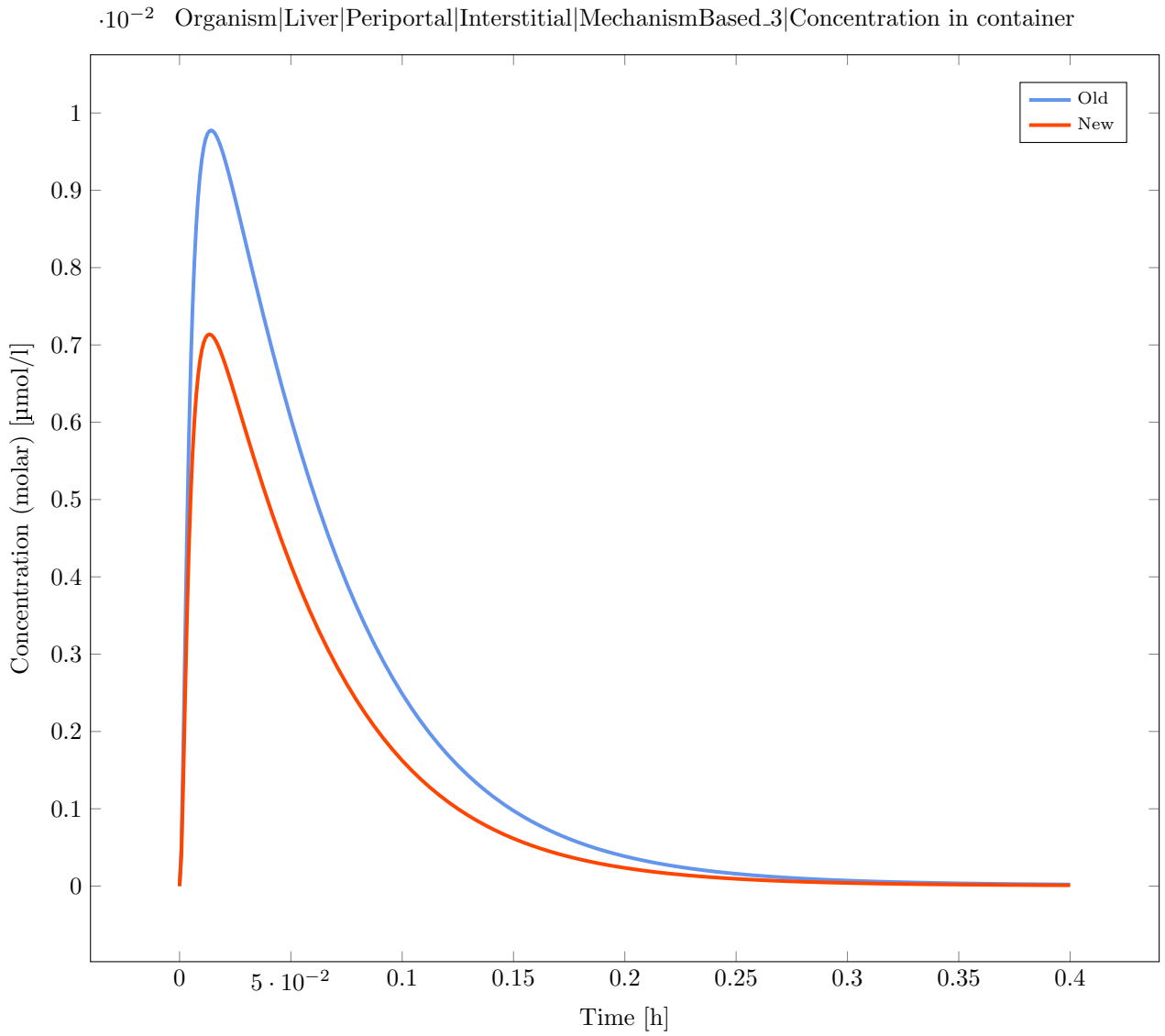

Figure 1.32

1.1.2 Valid Simulations (130/146)

Simulation: Beagle_SingleORAL_Dissolved-Beagle_SingleORAL_Dissolved
Result of the validation: Valid

Simulation: Beagle_SingleORAL_Dissolved-Beagle_SingleORAL_Dissolved_MW_200_fu_0.2_LogP_-5
Result of the validation: Valid

Simulation: Beagle_SingleORAL_Dissolved-Beagle_SingleORAL_Dissolved_MW_800_fu_0.6_LogP_--5
Result of the validation: Valid

Simulation: DDI_MultipleCombinations-01_MM_Competitive_Competitive
Result of the validation: Valid

Simulation: DDI_MultipleCombinations-02_MM_Uncompetitive_Uncompetitive
Result of the validation: Valid

Simulation: DDI_MultipleCombinations-03_MM_Noncompetitive_Noncompetitive
Result of the validation: Valid

Simulation: DDI_MultipleCombinations-04_MM_Mixed_Mixed
Result of the validation: Valid

Simulation: DDI_MultipleCombinations-21_1st_Competitive_Competitive
Result of the validation: Valid

Simulation: DDI_MultipleCombinations-23_1st_Noncompetitive_Noncompetitive
Result of the validation: Valid

Simulation: DDI_MultipleCombinations-24_1st_Mixed_Mixed
Result of the validation: Valid

Simulation: Dog_MultiORAL_12_12_Dissolved-Dog_MultiORAL_12_12_Dissolved
Result of the validation: Valid

Simulation: Dog_MultiORAL_24_Dissolved-Dog_MultiORAL_24_Dissolved
Result of the validation: Valid

Simulation: European_SingleORAL_Age_0_CYP3A4-European_SingleORAL_Age_0_CYP3A4
Result of the validation: Valid

Simulation: European_SingleORAL_Age_0_GFR-European_SingleORAL_Age_0_GFR
Result of the validation: Valid

Simulation: European_SingleORAL_Age_1_CYP3A4-European_SingleORAL_Age_1_CYP3A4
Result of the validation: Valid

Simulation: European_SingleORAL_Age_1_GFR-European_SingleORAL_Age_1_GFR
Result of the validation: Valid

Simulation: Human_CompetitiveInhibition-Human_CompetitiveInhibition
Result of the validation: Valid

Simulation: Human_ICRP_AGP-01_ICRP_0y_Male
Result of the validation: Valid

Simulation: Human_ICRP_AGP-02_ICRP_0.05y_Female
Result of the validation: Valid

Simulation: Human_ICRP_AGP-03_ICRP_0.18y_Male
Result of the validation: Valid

Simulation: Human_ICRP_AGP-04_ICRP_1y_Female
Result of the validation: Valid

Simulation: Human_ICRP_AGP-05_ICRP_12y_Male
Result of the validation: Valid

Simulation: Human_ICRP_AGP-06_ICRP_30y_Female
Result of the validation: Valid

Simulation: Human_ICRP_AGP-07_ICRP_100y_Male
Result of the validation: Valid

Simulation: Human_IrreversibleInhibition-Human_IrreversibleInhibition
Result of the validation: Valid

Simulation: Human_MixedInhibition-Human_MixedInhibition
Result of the validation: Valid

Simulation: Human_MultiIV_6_6_12-Human_MultiIV_6_6_12
Result of the validation: Valid

Simulation: Human_MultiORAL_6_12_12_Dissolved-Human_MultiORAL_6_12_12_Dissolved
Result of the validation: Valid

Simulation: Human_MultiORAL_6_12_12_Dissolved-Human_MultiORAL_6_12_12_Dissolved_absorption_-sink_conditions
Result of the validation: Valid

Simulation: Human_MultiORAL_6_12_12_Dissolved-Human_MultiORAL_6_12_12_Dissolved_EHC_-continuous_fraction_0.5
Result of the validation: Valid

Simulation: Human_MultiORAL_6_12_12_Dissolved-Human_MultiORAL_6_12_12_Dissolved_EHC_-continuous_fraction_1
Result of the validation: Valid

Simulation: Human_MultiORAL_6_12_12_Dissolved-Human_MultiORAL_6_12_12_Dissolved_pKa-dependent penalty factor
Result of the validation: Valid

Simulation: Human_MultiORAL_6_12_12_Dissolved-Human_MultiORAL_6_12_12_Dissolved_solubility
Result of the validation: Valid

Simulation: Human_NonCompetitiveInhibition-Human_NonCompetitiveInhibition
Result of the validation: Valid

Simulation: Human_Oral_BiDaily_TableFormulation-S1_suspension
Result of the validation: Valid

Simulation: Human_Oral_BiDaily_TableFormulation-S2_NoSuspension
Result of the validation: Valid

Simulation: Human_pH_SolubilityTable-S1_Table
Result of the validation: Valid

Simulation: Human_pH_SolubilityTable-S2_Measurement
Result of the validation: Valid

Simulation: Human_pH_SolubilityTable-S3_Table_SolubilityChanged
Result of the validation: Valid

Simulation: Human_pH_SolubilityTable-S4_Table_SolubilityTableChanged
Result of the validation: Valid

Simulation: Human_SingleIV_Configuration-Human_SingleIV_Configuration
Result of the validation: Valid

Simulation: Human_SingleIV-Human_SingleIV
Result of the validation: Valid

Simulation: Human_SingleIV-Human_SingleIV_MW_200_fu_0.2_LogP_5
Result of the validation: Valid

Simulation: Human_SingleIV-Human_SingleIV_MW_800_fu_0.6_LogP_-5
Result of the validation: Valid

Simulation: Human_SingleORAL_Dissolved_PlasmaClearance-Human_SingleORAL_Dissolved_PlasmaClearan
Result of the validation: Valid

Simulation: Human_SingleORAL_Dissolved_PlasmaClearance-Human_SingleORAL_Dissolved_PlasmaClearan MW_200_fu_0.2_LogP_5
Result of the validation: Valid

Simulation: Human_SingleORAL_Dissolved_PlasmaClearance-Human_SingleORAL_Dissolved_PlasmaClearan MW_800_fu_0.6_LogP_-5
Result of the validation: Valid

Simulation: Human_SingleORAL_Dissolved-Human_SingleORAL_Dissolved
Result of the validation: Valid

Simulation: Human_SingleORAL_Dissolved-Human_SingleORAL_Dissolved_MW_200_fu_0.2_LogP_-5
Result of the validation: Valid

Simulation: Human_SingleORAL_Dissolved-Human_SingleORAL_Dissolved_MW_800_fu_0.6_LogP_--5
Result of the validation: Valid

Simulation: Human_SingleORAL_Lint80_AsSuspention-Human_SingleORAL_Lint80_AsSuspention
Result of the validation: Valid

Simulation: Human_SingleORAL_Lint80-Human_SingleORAL_Lint80
Result of the validation: Valid

Simulation: Human_SingleORAL_MonoParticles_AsSuspention-Human_SingleORAL_MonoParticles_-AsSuspention
Result of the validation: Valid

Simulation: Human_SingleORAL_PolyParticlesLogNormal_AsSuspention-Human_SingleORAL_-PolyParticlesLogNormal_AsSuspention
Result of the validation: Valid

Simulation: Human_SingleORAL_PolyParticlesNormal_AsSuspention-Human_SingleORAL_PolyParticlesNo AsSuspention
Result of the validation: Valid

Simulation: Human_SingleORAL_PolyParticlesNormal_AsSuspention-Human_SingleORAL_PolyParticlesNo
AsSuspention_dissolved_radius
Result of the validation: Valid

Simulation: Human_SingleORAL_PolyParticlesNormal_AsSuspention-Human_SingleORAL_PolyParticlesNo
AsSuspention_treat_precipated_drug_as_soluble
Result of the validation: Valid

Simulation: Human_SingleORAL_Weibull_AsSuspention-Human_SingleORAL_Weibull_AsSuspention
Result of the validation: Valid

Simulation: Human_SingleORAL_Weibull_AsSuspention-Human_SingleORAL_Weibull_AsSuspention_-MW_200_fu_0.2_LogP_5
Result of the validation: Valid

Simulation: Human_SingleORAL_Weibull_AsSuspention-Human_SingleORAL_Weibull_AsSuspention_-MW_800_fu_0.6_LogP_-5
Result of the validation: Valid

Simulation: Human_SingleORAL_Weibull-Human_SingleORAL_Weibull
Result of the validation: Valid

Simulation: Human_SingleORAL_Weibull-Human_SingleORAL_Weibull_MW_200_fu_0.2_LogP_5
Result of the validation: Valid

Simulation: Human_SingleORAL_Weibull-Human_SingleORAL_Weibull_MW_800_fu_0.6_LogP_-5
Result of the validation: Valid

Simulation: Human_UncompetitiveInhibition-Human_UncompetitiveInhibition
Result of the validation: Valid

Simulation: Minipig_SingleORAL_Dissolved-Minipig_SingleORAL_Dissolved
Result of the validation: Valid

Simulation: Minipig_SingleORAL_Dissolved-Minipig_SingleORAL_Dissolved_MW_200_fu_0.2_LogP_-5
Result of the validation: Valid

Simulation: Minipig_SingleORAL_Dissolved-Minipig_SingleORAL_Dissolved_MW_800_fu_0.6_LogP_--5
Result of the validation: Valid

Simulation: Monkey_SingleORAL_Dissolved-Monkey_SingleORAL_Dissolved
Result of the validation: Valid

Simulation: Monkey_SingleORAL_Dissolved-Monkey_SingleORAL_Dissolved_MW_200_fu_0.2_LogP_-5
Result of the validation: Valid

Simulation: Monkey_SingleORAL_Dissolved-Monkey_SingleORAL_Dissolved_MW_800_fu_0.6_LogP_--5
Result of the validation: Valid

Simulation: Mouse_SingleORAL_Dissolved-Mouse_SingleORAL_Dissolved
Result of the validation: Valid

Simulation: Mouse_SingleORAL_Dissolved-Mouse_SingleORAL_Dissolved_MW_200_fu_0.2_LogP_-5
Result of the validation: Valid

Simulation: Mouse_SingleORAL_Dissolved-Mouse_SingleORAL_Dissolved_MW_800_fu_0.6_LogP_--5
Result of the validation: Valid

Simulation: Preterm_SingleIV_Age_0_GA_32_CYP3A4-Preterm_SingleIV_Age_0_GA_32_CYP3A4
Result of the validation: Valid

Simulation: Preterm_SingleIV_Age_0_GA_32_GFR-Preterm_SingleIV_Age_0_GA_32_GFR
Result of the validation: Valid

Simulation: Preterm_SingleIV_Age_15_GA_32_CYP3A4-Preterm_SingleIV_Age_15_GA_32_CYP3A4
Result of the validation: Valid

Simulation: Preterm_SingleIV_Age_15_GA_32_GFR-Preterm_SingleIV_Age_15_GA_32_GFR
Result of the validation: Valid

Simulation: Rabbit_SingleORAL_Dissolved-Rabbit_SingleORAL_Dissolved
Result of the validation: Valid

Simulation: Rabbit_SingleORAL_Dissolved-Rabbit_SingleORAL_Dissolved_MW_200_fu_0.2_LogP_-5
Result of the validation: Valid

Simulation: Rabbit_SingleORAL_Dissolved-Rabbit_SingleORAL_Dissolved_MW_800_fu_0.6_LogP_--5
Result of the validation: Valid

Simulation: Rat_MultiORAL_6_6_12_Dissolved-Rat_MultiORAL_6_6_12_Dissolved
Result of the validation: Valid

Simulation: Rat_MultiORAL_6_6_6_6_Dissolved-Rat_MultiORAL_6_6_6_6_Dissolved
Result of the validation: Valid

Simulation: Rat_MultiORAL_8_8_8_Dissolved-Rat_MultiORAL_8_8_8_Dissolved
Result of the validation: Valid

Simulation: SingleIV_2Pores_Human-SingleIV_2Pores_Human
Result of the validation: Valid

Simulation: SingleIV_2Pores_Human-SingleIV_2Pores_Human_SimulationC
Result of the validation: Valid

Simulation: SingleIV_2Pores_Human-SingleIV_2Pores_Human_SimulationD
Result of the validation: Valid

Simulation: SingleIV_2Pores_Human-SingleIV_2Pores_Human_SimulationF
Result of the validation: Valid

Simulation: SingleIV_2Pores_Monkey-SingleIV_2Pores_Monkey
Result of the validation: Valid

Simulation: SingleIV_2Pores_Monkey-SingleIV_2Pores_Monkey_SimulationG
Result of the validation: Valid

Simulation: SingleIV_2Pores_Monkey-SingleIV_2Pores_Monkey_SimulationH
Result of the validation: Valid

Simulation: SingleIV_2Pores_Mouse-SingleIV_2Pores_Mouse
Result of the validation: Valid

Simulation: SingleIV_2Pores_Mouse-SingleIV_2Pores_Mouse_SimulationA
Result of the validation: Valid

Simulation: SingleIV_2Pores_Mouse-SingleIV_2Pores_Mouse_SimulationB
Result of the validation: Valid

Simulation: SingleIV_2Pores_Mouse-SingleIV_2Pores_Mouse_SimulationE
Result of the validation: Valid

Simulation: SingleIV_C1_4Comp_standard_standard_standard-SingleIV_C1_4Comp_standard_standard_standard
Result of the validation: Valid

Simulation: SingleIV_C2_4Comp_PT_standard_standard-SingleIV_C2_4Comp_PT_standard_standard
Result of the validation: Valid

Simulation: SingleIV_C2_4Comp_RR_standard_standard-SingleIV_C2_4Comp_RR_standard_standard
Result of the validation: Valid

Simulation: SingleIV_C2_4Comp_standard_schmitt_standard-SingleIV_C2_4Comp_standard_schmitt_standard
Result of the validation: Valid

Simulation: SingleIV_C3_4Comp_RR_schmitt_standard-SingleIV_C3_4Comp_RR_schmitt_standard
Result of the validation: Valid

Simulation: SingleIV_C3_4Comp_standard_schmittnormlized_standard-SingleIV_C3_4Comp_standard_schmittnormlized_standard
Result of the validation: Valid

Simulation: SingleIV_C4_2Pores_RR_standard_standard-SingleIV_C4_2Pores_RR_standard_standard
Result of the validation: Valid

Simulation: SingleIV_C4_4Comp_Ber_standard_standard-SingleIV_C4_4Comp_Ber_standard_standard
Result of the validation: Valid

Simulation: SingleIV_C5_2Pores_Ber_standard_standard-SingleIV_C5_2Pores_Ber_standard_standard
Result of the validation: Valid

Simulation: SingleIV_C5_2Pores_PT_standard_standard-SingleIV_C5_2Pores_PT_standard_standard
Result of the validation: Valid

Simulation: SingleIV_C5_2Pores_RR_schmitt_standard-SingleIV_C5_2Pores_RR_schmitt_standard
Result of the validation: Valid

Simulation: SingleIV_C6_2Pores_standard_standard_standard-SingleIV_C6_2Pores_standard_standard_-standard
Result of the validation: Valid

Simulation: SingleIV_C7_2Pores_standard_schmitt_standard-SingleIV_C7_2Pores_standard_schmitt_-standard
Result of the validation: Valid

Simulation: SingleIV_C7_4Comp_schmitt_standard_standard-SingleIV_C7_4Comp_schmitt_standard_-standard
Result of the validation: Valid

Simulation: SingleIV_C8_2Pores_standard_schmittnormalized_standard-SingleIV_C8_2Pores_standard_-schmittnormalized_standard
Result of the validation: Valid

Simulation: SingleIV_C9_2Pores_schmitt_standard_standard-SingleIV_C9_2Pores_schmitt_standard_-standard
Result of the validation: Valid

Simulation: SingleORAL_C10_4Comp_PT_standard_standard-SingleORAL_C10_4Comp_PT_standard_-standard
Result of the validation: Valid

Simulation: SingleORAL_C11_4Comp_schmitt_standard_standard-SingleORAL_C11_4Comp_schmitt_-standard_standard
Result of the validation: Valid

Simulation: SingleORAL_C11_4Comp_standard_standard_standard-SingleORAL_C11_4Comp_standard_-standard_standard
Result of the validation: Valid

Simulation: SingleORAL_C12_4Comp_standard_schmitt_standard-SingleORAL_C12_4Comp_standard_-schmitt_standard
Result of the validation: Valid

Simulation: SingleORAL_C13_2Pores_schmitt_standard_standard-SingleORAL_C13_2Pores_schmitt_-standard_standard
Result of the validation: Valid

Simulation: SingleORAL_C13_4Comp_standard_schmittnormalized_standard-SingleORAL_C13_-4Comp_standard_schmittnormalized_standard
Result of the validation: Valid

Simulation: SingleORAL_C14_2Pores_PT_standard_standard-SingleORAL_C14_2Pores_PT_standard_-standard
Result of the validation: Valid

Simulation: SingleORAL_C2_2Pores_standard_standard_standard-SingleORAL_C2_2Pores_standard_-standard_standard
Result of the validation: Valid

Simulation: SingleORAL_C3_2Pores_standard_schmitt_standard-SingleORAL_C3_2Pores_standard_-schmitt_standard
Result of the validation: Valid

Simulation: SingleORAL_C4_2Pores_standard_schmittnormalized_standard-SingleORAL_C4_2Pores_-standard_schmittnormalized_standard
Result of the validation: Valid

Simulation: SingleORAL_C6_4Comp_Ber_standard_standard-SingleORAL_C6_4Comp_Ber_standard_-standard
Result of the validation: Valid

Simulation: SingleORAL_C6_4Comp_RR_standard_standard-SingleORAL_C6_4Comp_RR_standard_-standard
Result of the validation: Valid

Simulation: SingleORAL_C7_2Pores_Ber_standard_standard-SingleORAL_C7_2Pores_Ber_standard_-standard
Result of the validation: Valid

Simulation: SingleORAL_C7_4Comp_RR_schmitt_standard-SingleORAL_C7_4Comp_RR_schmitt_-standard
Result of the validation: Valid

Simulation: SingleORAL_C8_2Pores_RR_standard_standard-SingleORAL_C8_2Pores_RR_standard_-standard
Result of the validation: Valid

Simulation: SingleORAL_C9_2Pores_RR_schmitt_standard-SingleORAL_C9_2Pores_RR_schmitt_-standard
Result of the validation: Valid

Simulation: Test 18.1_I1_C1_A1_Config1-Test 18.1_I1_C1_A1_Config1
Result of the validation: Valid

Simulation: Test 18.1_I2_C1_A1_Config2-Test 18.1_I2_C1_A1_Config2
Result of the validation: Valid

Simulation: Test 18.1_I2_C3_A1_Config2-Test 18.1_I2_C3_A1_Config2
Result of the validation: Valid

Simulation: Test 18.1_I3_C3_A3_Config2-Test 18.1_I3_C3_A3_Config2
Result of the validation: Valid