Open Systems Pharmacology Suite - 11
Folder Comparison

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Generated using Installation Validator® 11.0.136.

Contents

Chapter 1

Folder Comparison Results

Overall Comparison Result: Invalid
Number of Compared Files: 155

1.1 Comparison Results

Overall Comparison Result
Invalid

Old Folder
D:\Outputs_10.0

New Folder
D:\Outputs_11.0

Using exclusions
**|drug|Fraction of oral drug mass absorbed into mucosa segment
**|Receptor Occupancy-drug-BIND-Lab Complex

1.1.1 Invalid Simulations (13/155)

Simulation: DDI_MultipleCombinations-21_1st_Competitive_Competitive
Result of the validation: Invalid
Absolute Tolerance: 1.00E-8
Relative Tolerance: 1.00E-4

Output Path: Organism|SmallIntestine|Mucosa|UpperJejunum|Intracellular|C1|Concentration in container
Deviation for 'Organism|SmallIntestine|Mucosa|UpperJejunum|Intracellular|C1|Concentration in container' is 6.24% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.06

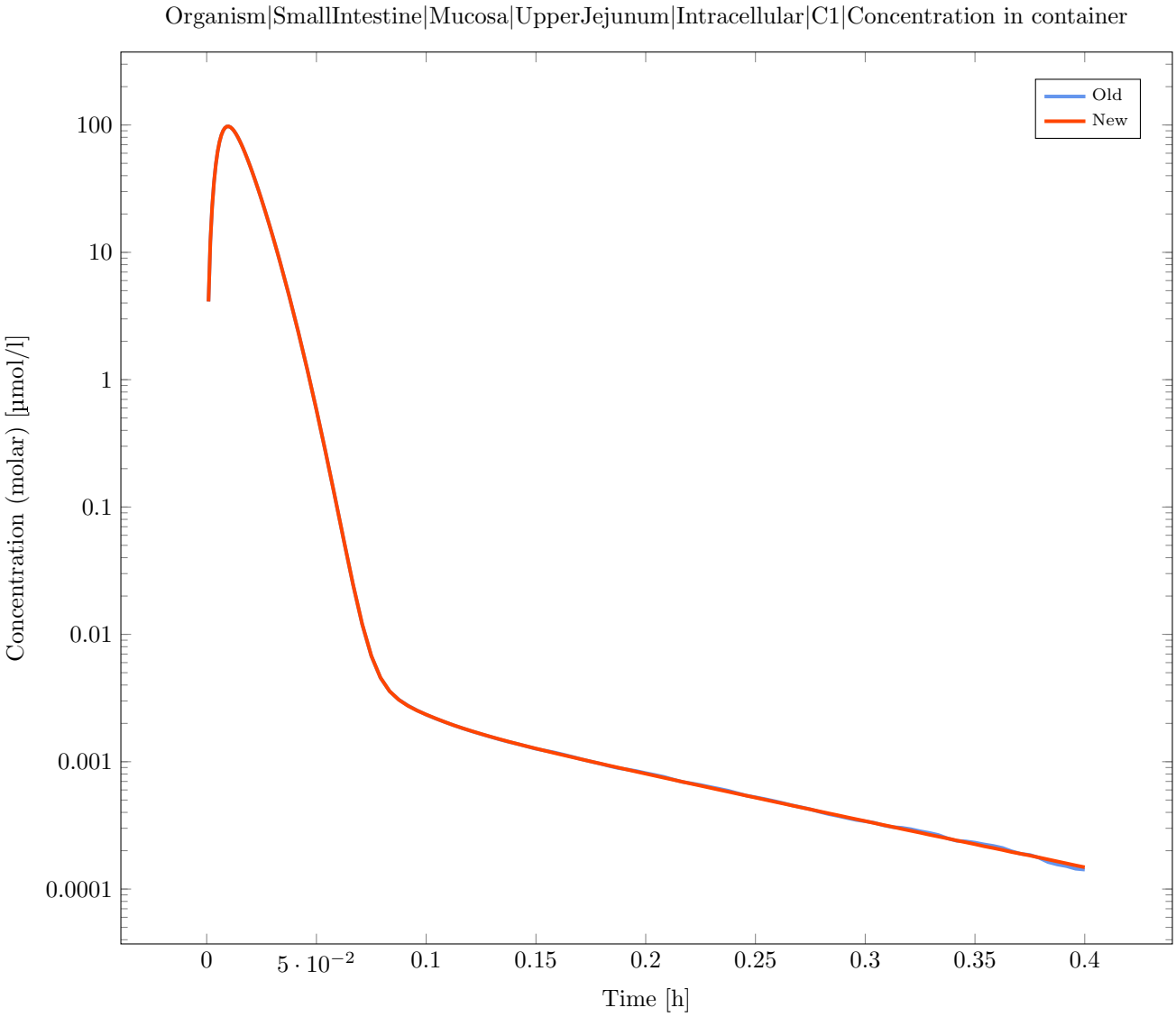
Organism|SmallIntestine|Mucosa|UpperJejunum|Intracellular|C1|Concentration in container
Old
New
Concentration (molar) [µmol/l]
100
10
1
0.1
0.01
0.001
0.0001
0
5 · 10−2
0.1
0.15
0.2
0.25
0.3
0.35
0.4
Time [h]

Figure 1.1

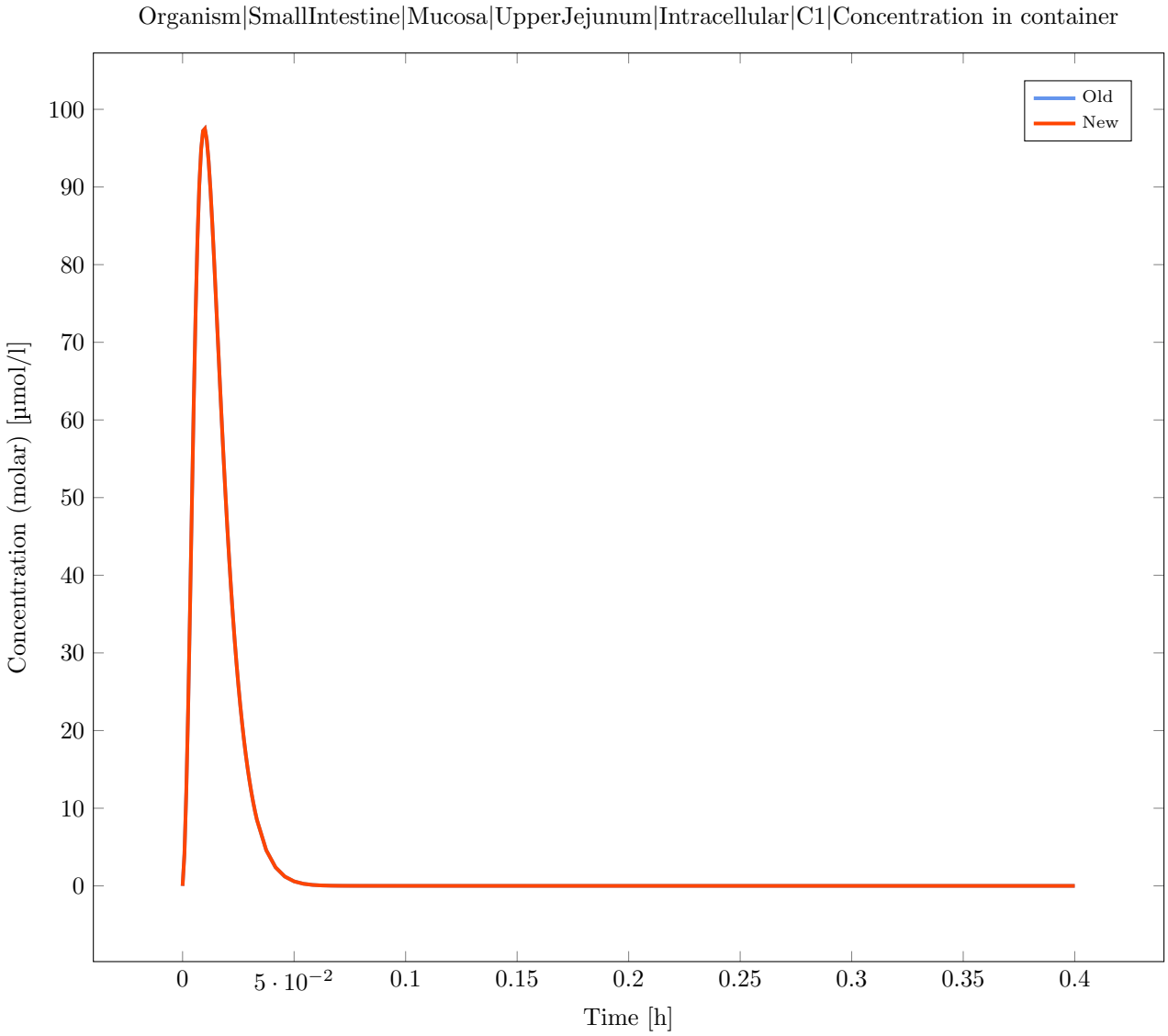

Figure 1.2

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

Output Path: Organism|Heart|Intracellular|drug|Concentration in container
Deviation for 'Organism|Heart|Intracellular|drug|Concentration in container' is 13.22% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.13

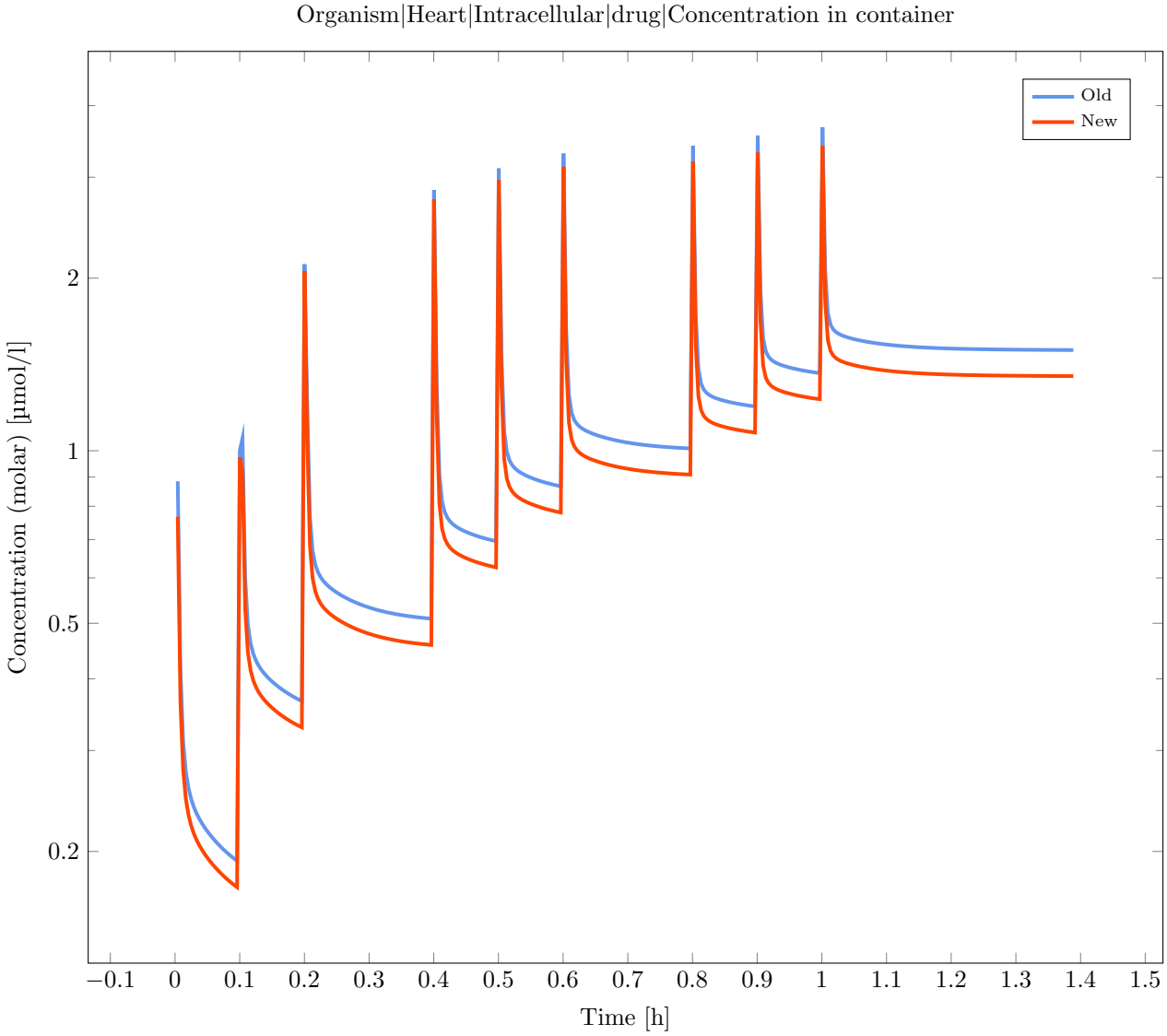
Organism|Heart|Intracellular|drug|Concentration in container
Old
New
Concentration (molar) [µmol/l]
2
1
0.5
0.2
−0.1
0
0.1
0.2
0.3
0.4
0.5
0.6
0.7
0.8
0.9
1
1.1
1.2
1.3
1.4
1.5
Time [h]

Figure 1.3

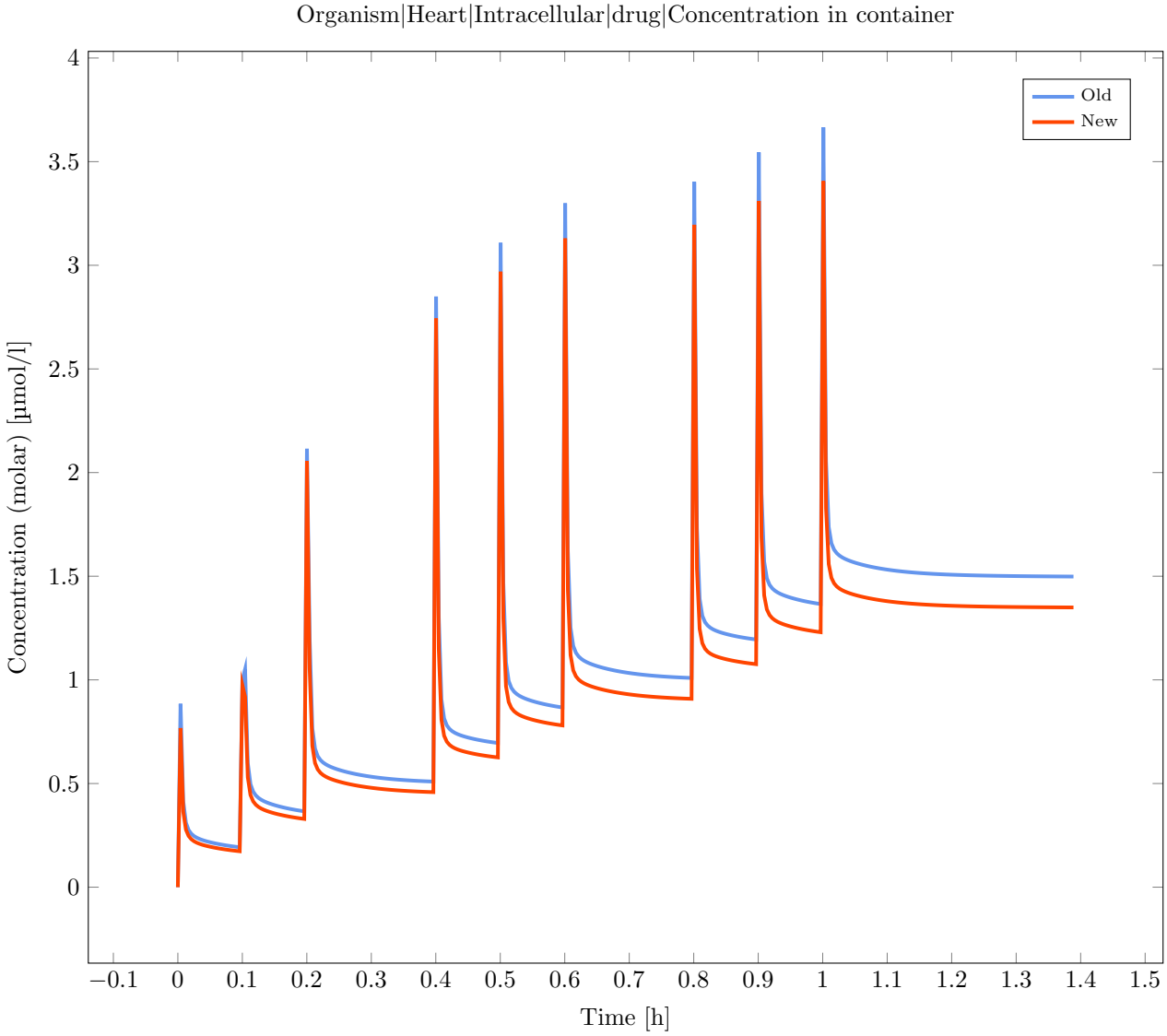

Figure 1.4

Output Path: Organism|Heart|drug|Intracellular Unbound
Deviation for 'Organism|Heart|drug|Intracellular Unbound' is 13.22% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.13

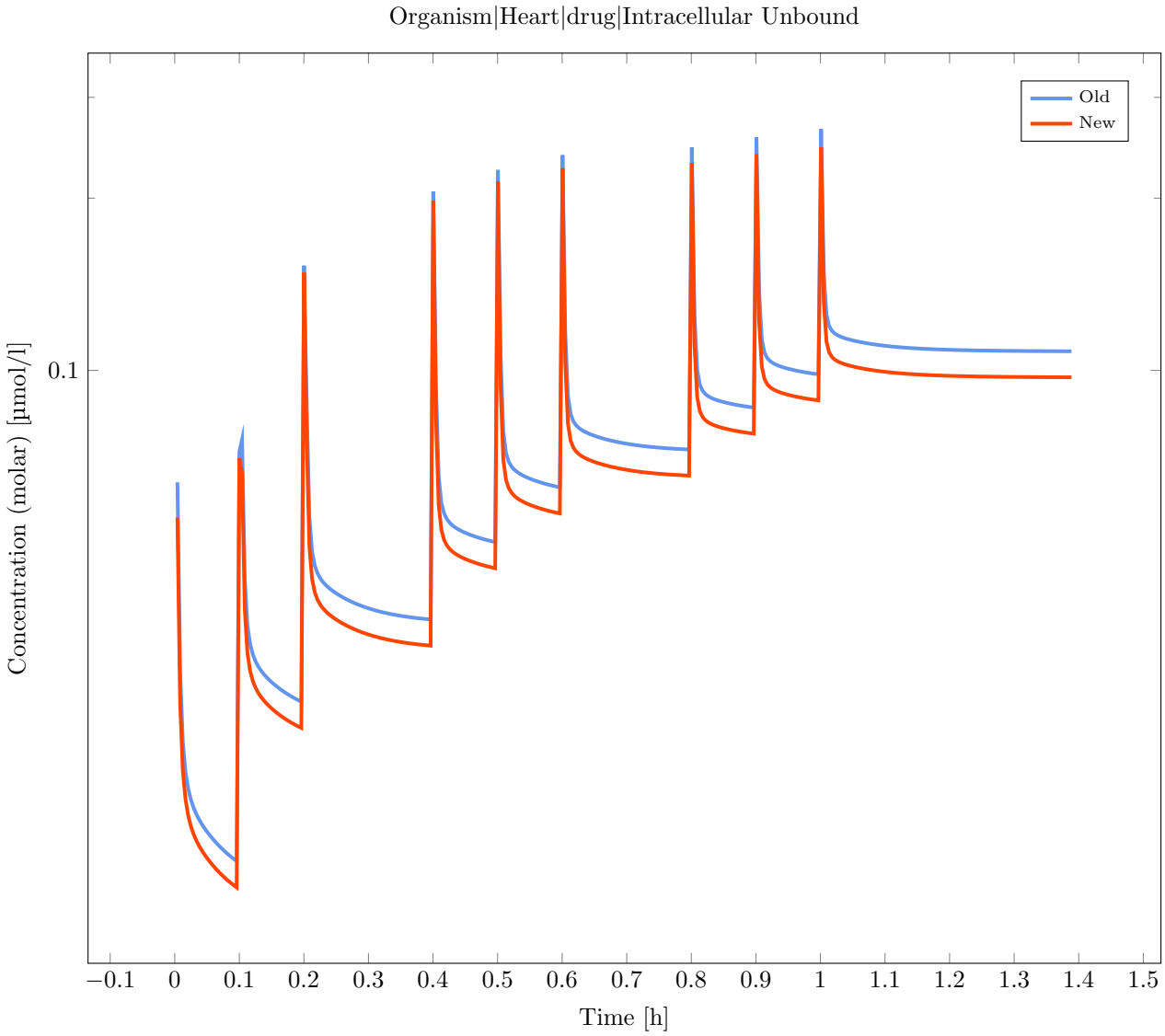
Organism|Heart|drug|Intracellular Unbound
Old
New
Concentration (molar) [µmol/l]
0.1
−0.1
0
0.1
0.2
0.3
0.4
0.5
0.6
0.7
0.8
0.9
1
1.1
1.2
1.3
1.4
1.5
Time [h]

Figure 1.5

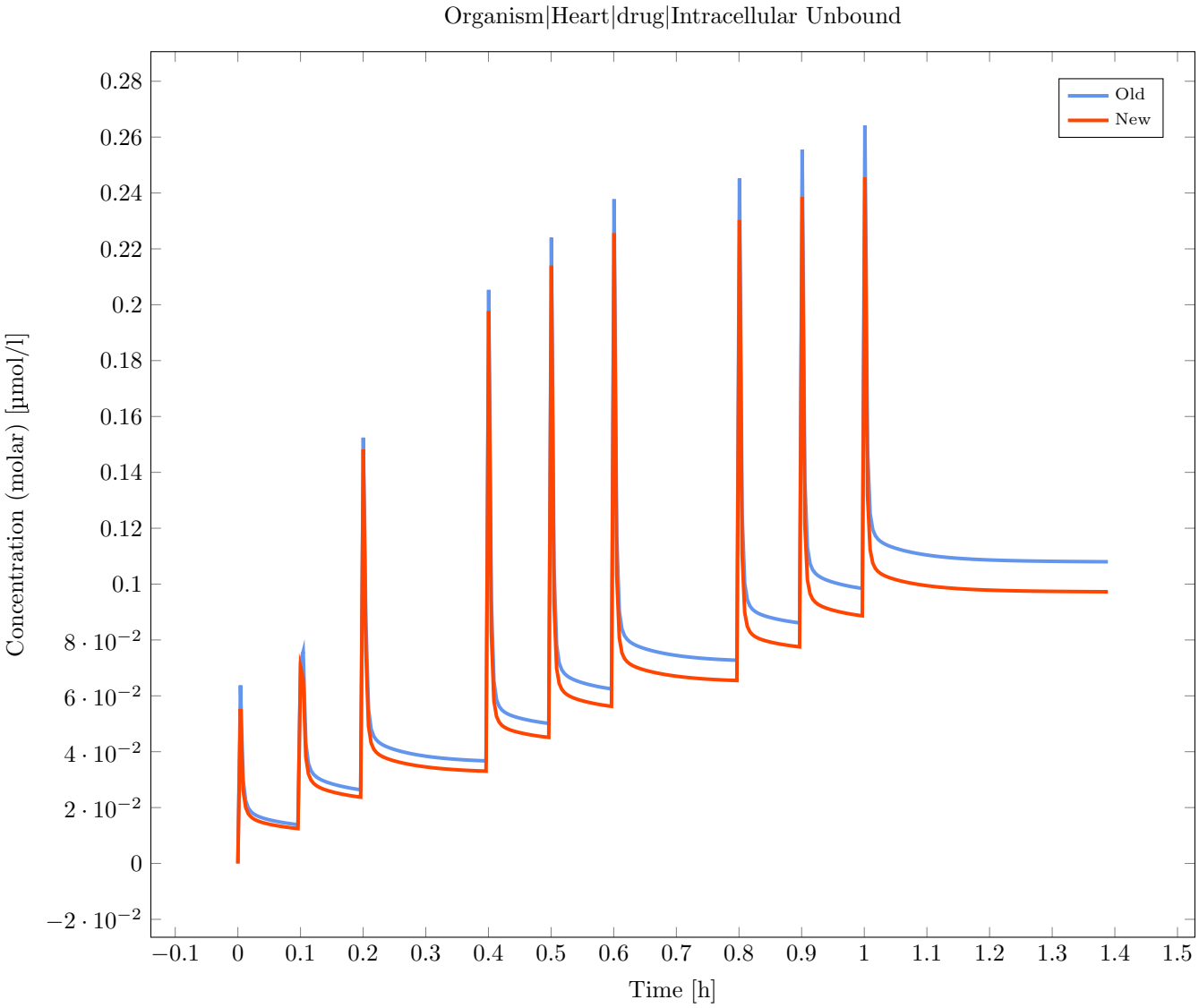

Figure 1.6

Output Path: Organism|Heart|drug|Tissue
Deviation for 'Organism|Heart|drug|Tissue' is 13.13% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.13

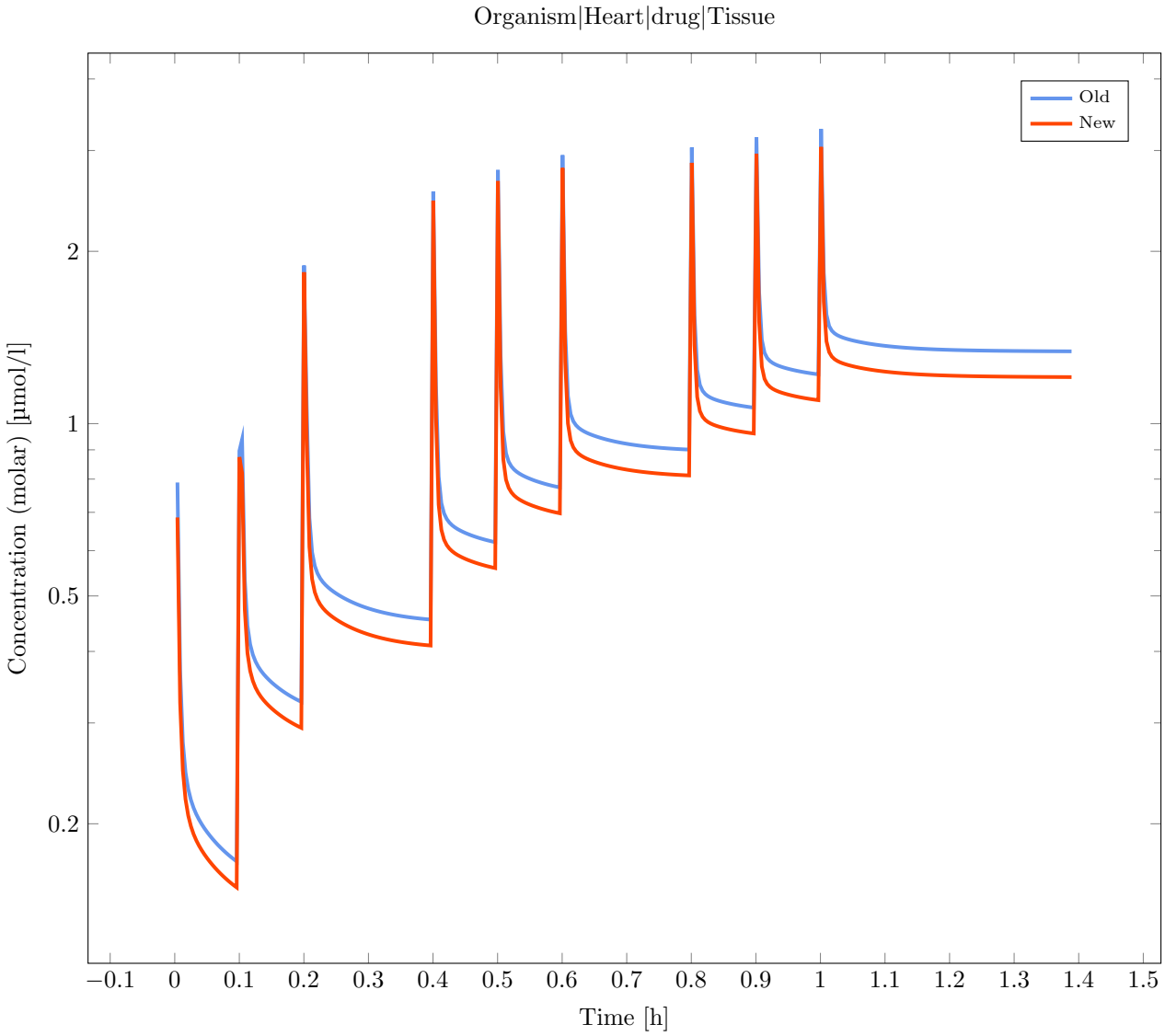
Organism|Heart|drug|Tissue
Old
New
Concentration (molar) [µmol/l]
2
1
0.5
0.2
−0.1
0
0.1
0.2
0.3
0.4
0.5
0.6
0.7
0.8
0.9
1
1.1
1.2
1.3
1.4
1.5
Time [h]

Figure 1.7

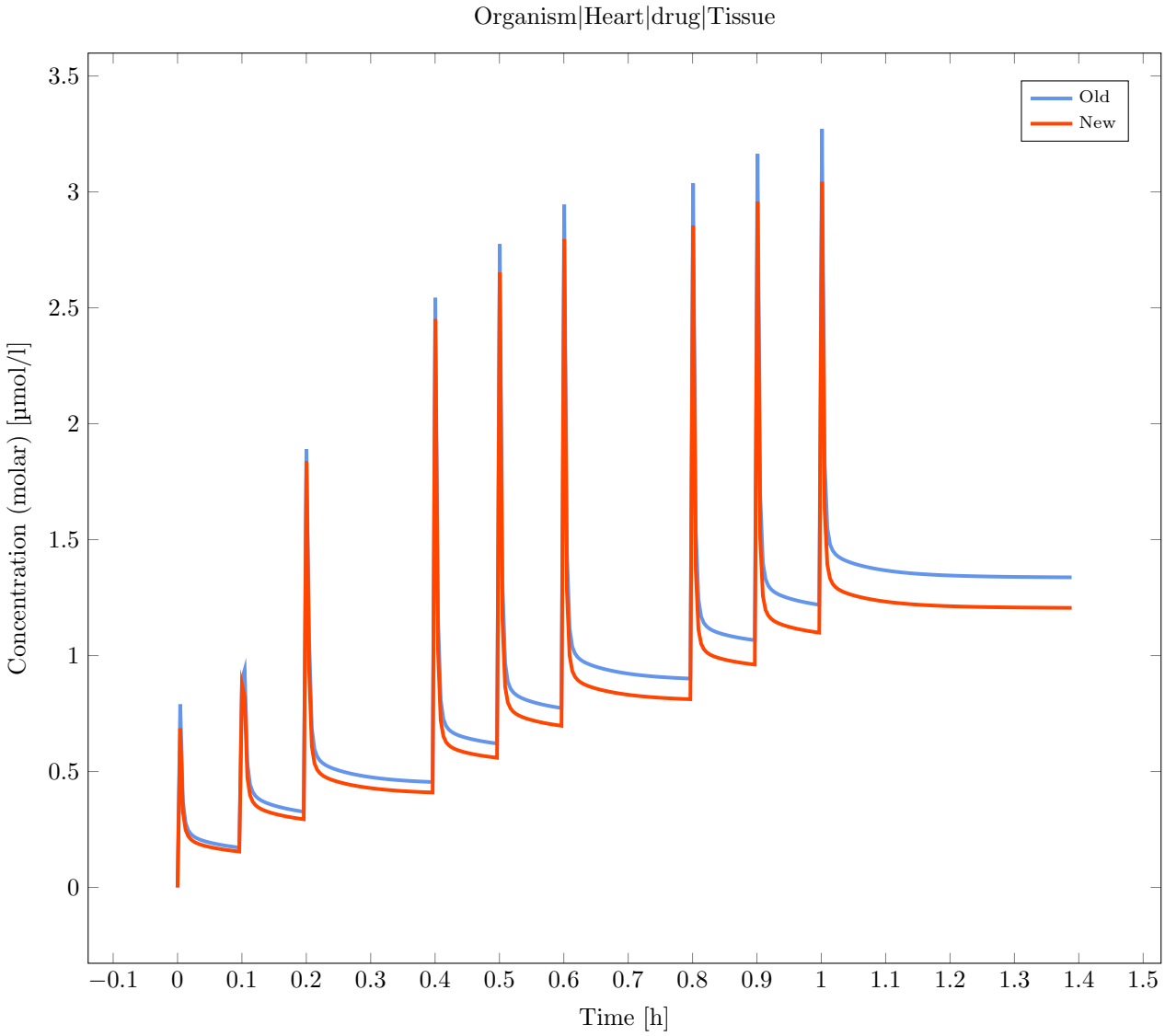

Figure 1.8

Output Path: Organism|Heart|drug|Interstitial Unbound
Deviation for 'Organism|Heart|drug|Interstitial Unbound' is 7.51% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.08

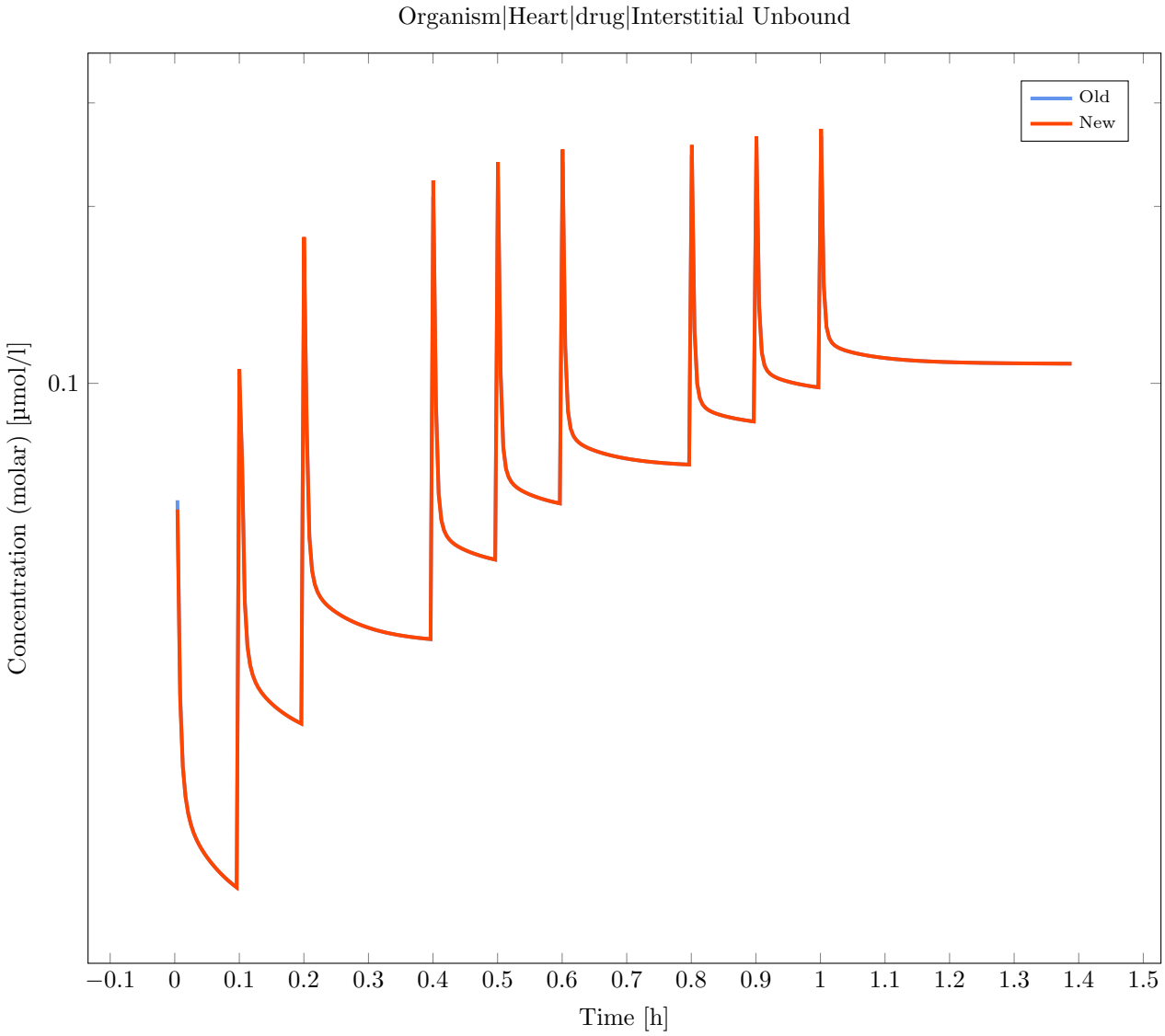
Organism|Heart|drug|Interstitial Unbound
Old
New
Concentration (molar) [µmol/l]
0.1
−0.1
0
0.1
0.2
0.3
0.4
0.5
0.6
0.7
0.8
0.9
1
1.1
1.2
1.3
1.4
1.5
Time [h]

Figure 1.9

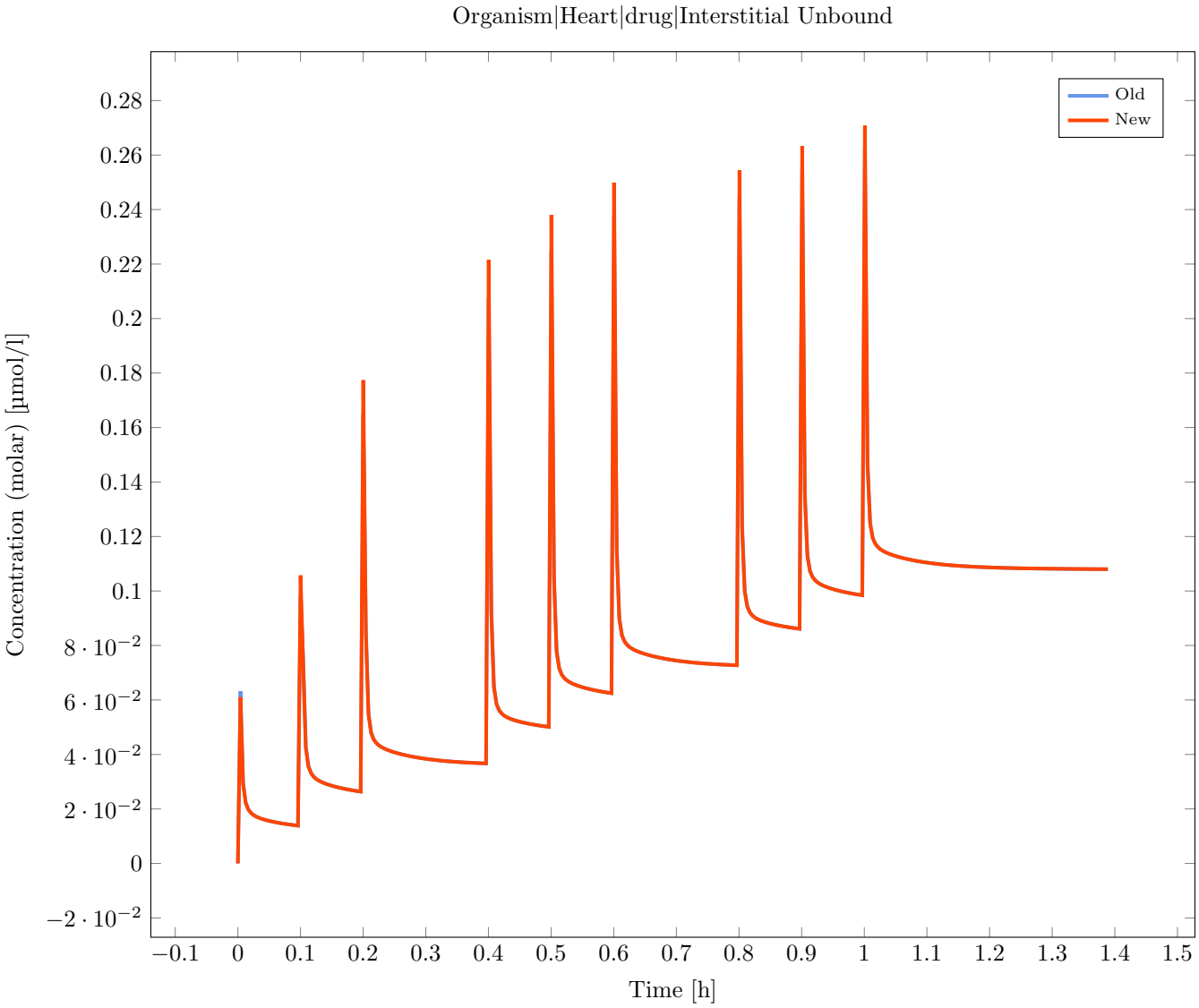

Figure 1.10

Output Path: Organism|Heart|Interstitial|drug|Concentration in container
Deviation for 'Organism|Heart|Interstitial|drug|Concentration in container' is 7.51% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.08

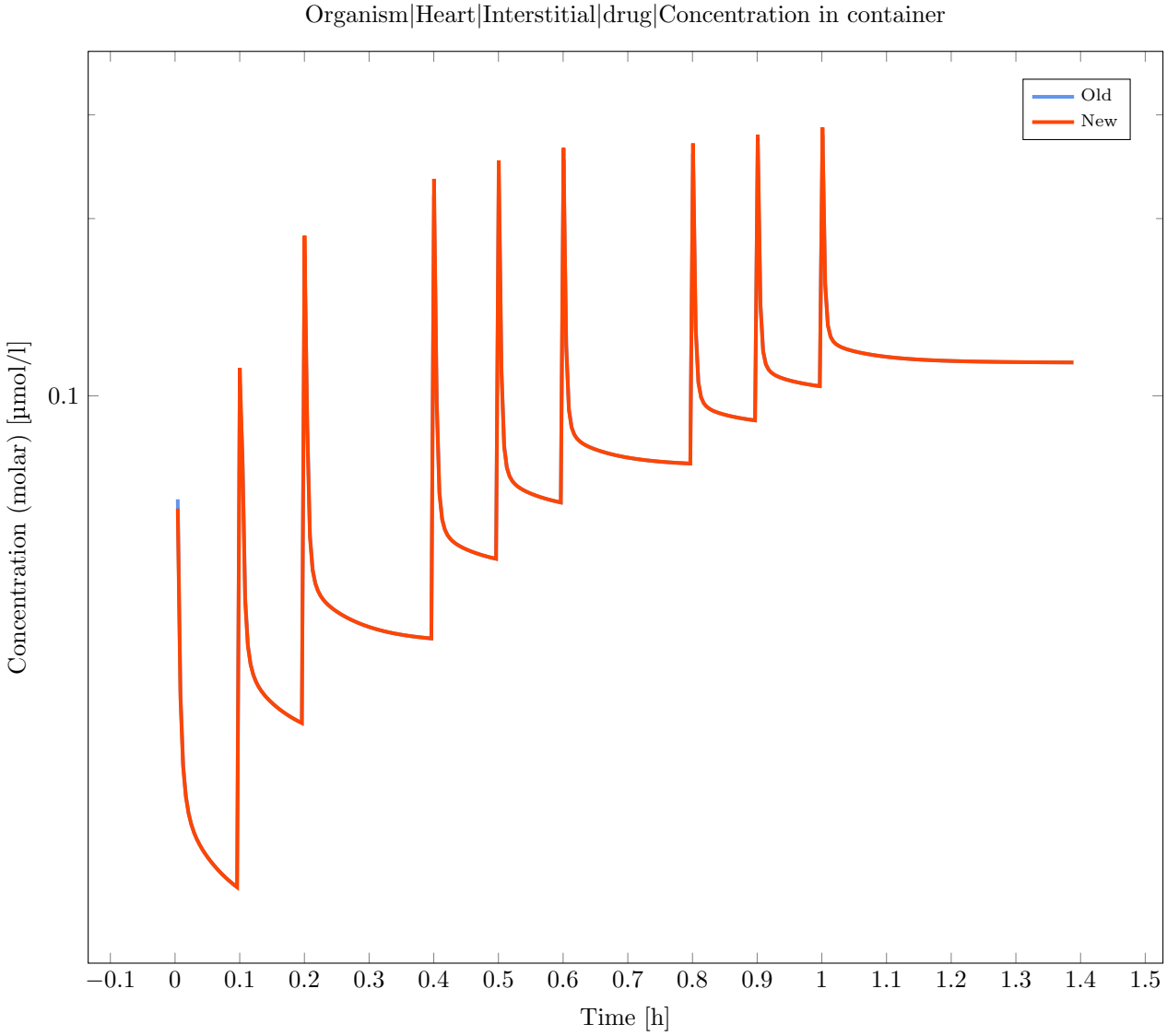
Organism|Heart|Interstitial|drug|Concentration in container
Old
New
Concentration (molar) [µmol/l]
0.1
−0.1
0
0.1
0.2
0.3
0.4
0.5
0.6
0.7
0.8
0.9
1
1.1
1.2
1.3
1.4
1.5
Time [h]

Figure 1.11

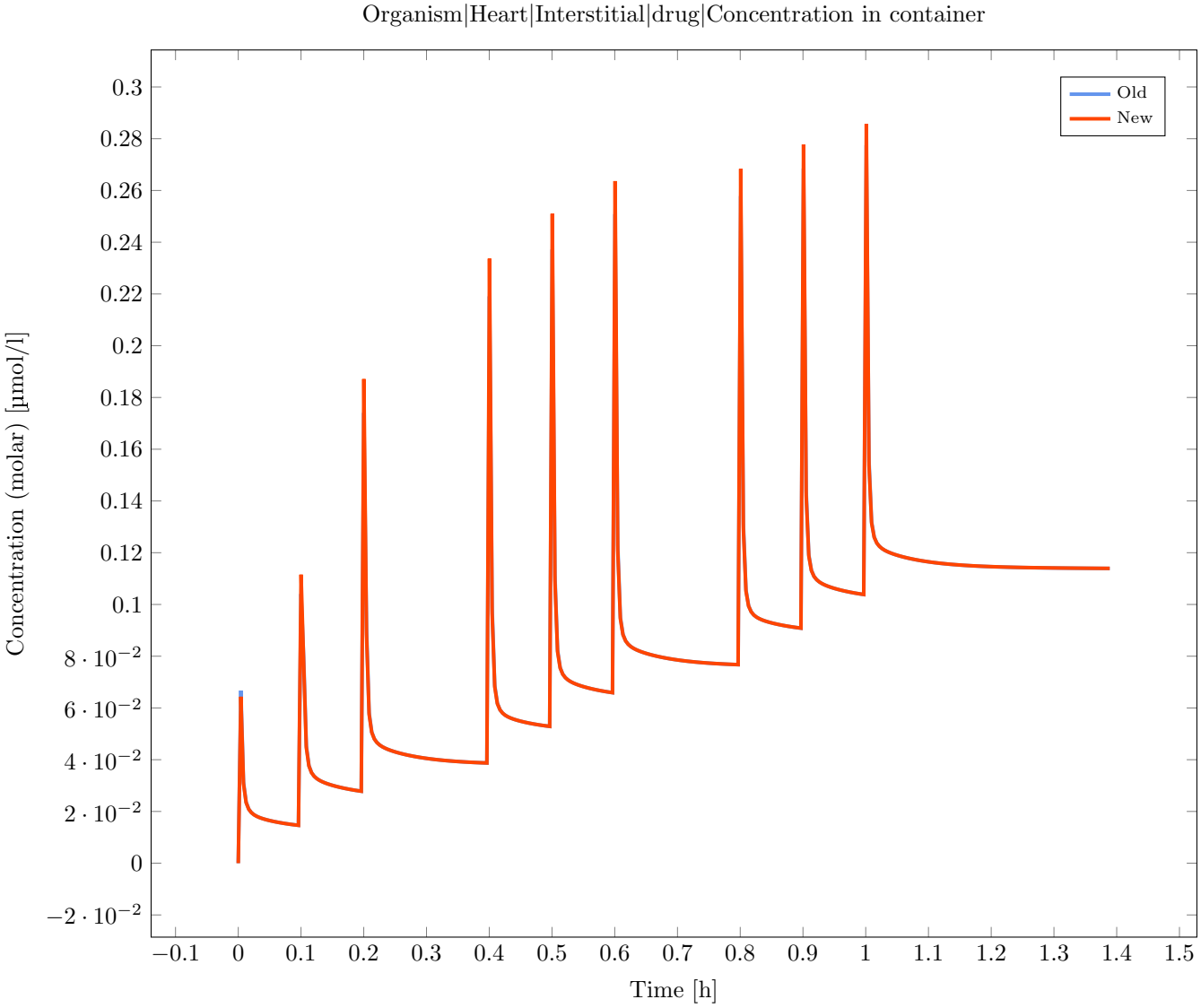

Figure 1.12

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

Output Path: Organism|Heart|drug|Intracellular Unbound
Deviation for 'Organism|Heart|drug|Intracellular Unbound' is 13.93% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.14

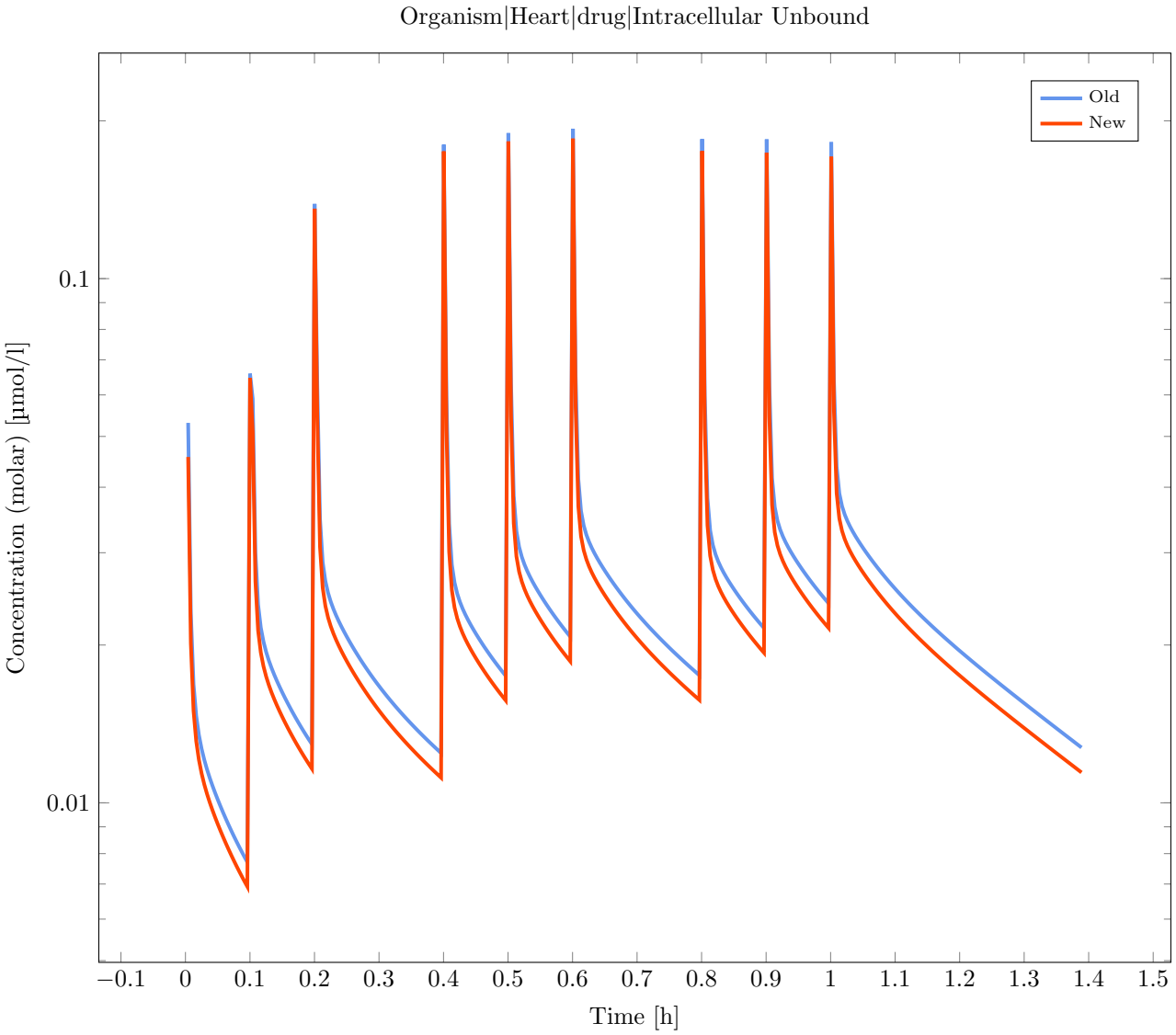
Organism|Heart|drug|Intracellular Unbound
Old
New
Concentration (molar) [µmol/l]
0.1
0.01
−0.1
0
0.1
0.2
0.3
0.4
0.5
0.6
0.7
0.8
0.9
1
1.1
1.2
1.3
1.4
1.5
Time [h]

Figure 1.13

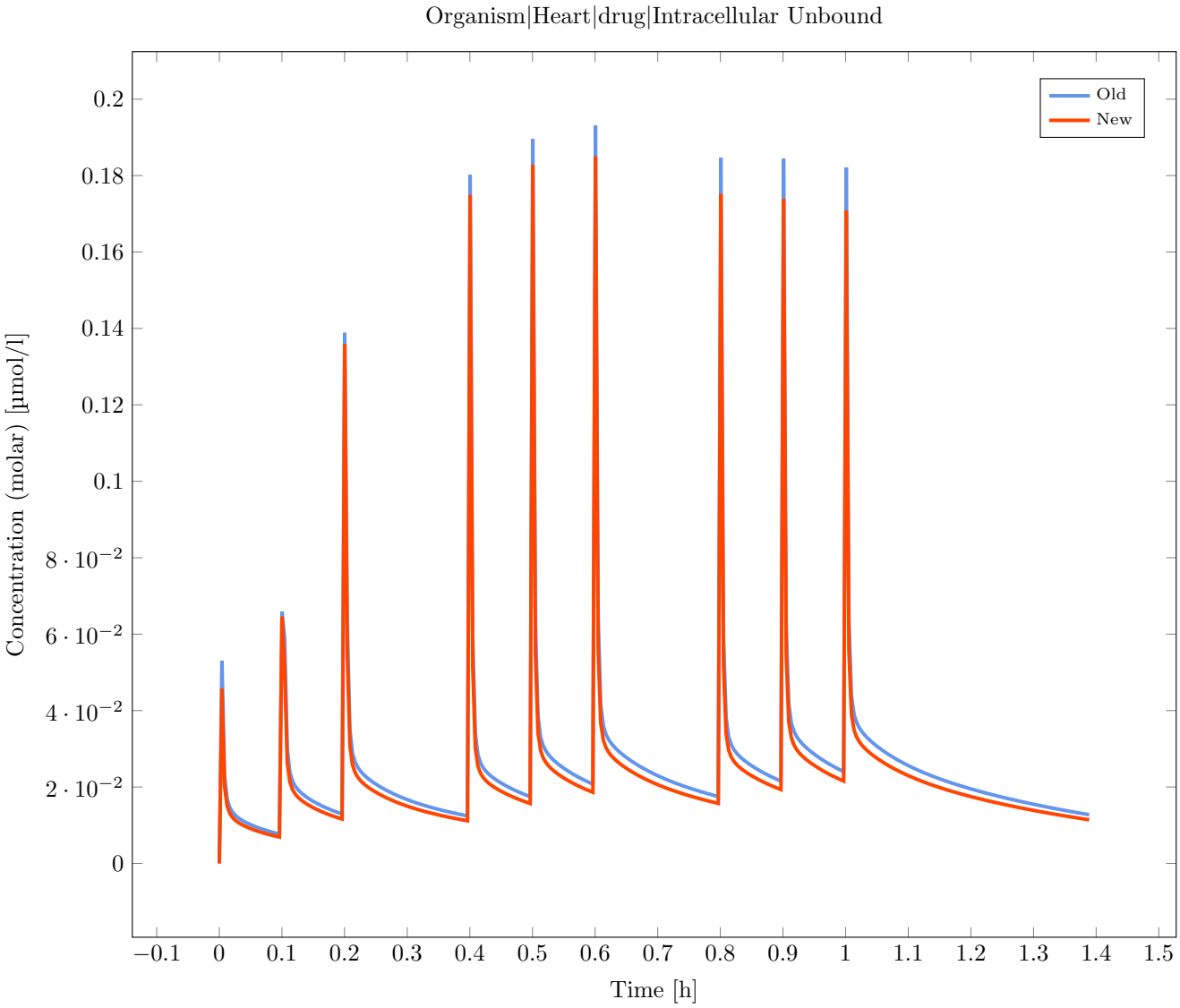

Figure 1.14

Output Path: Organism|Heart|Intracellular|drug|Concentration in container
Deviation for 'Organism|Heart|Intracellular|drug|Concentration in container' is 13.93% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.14

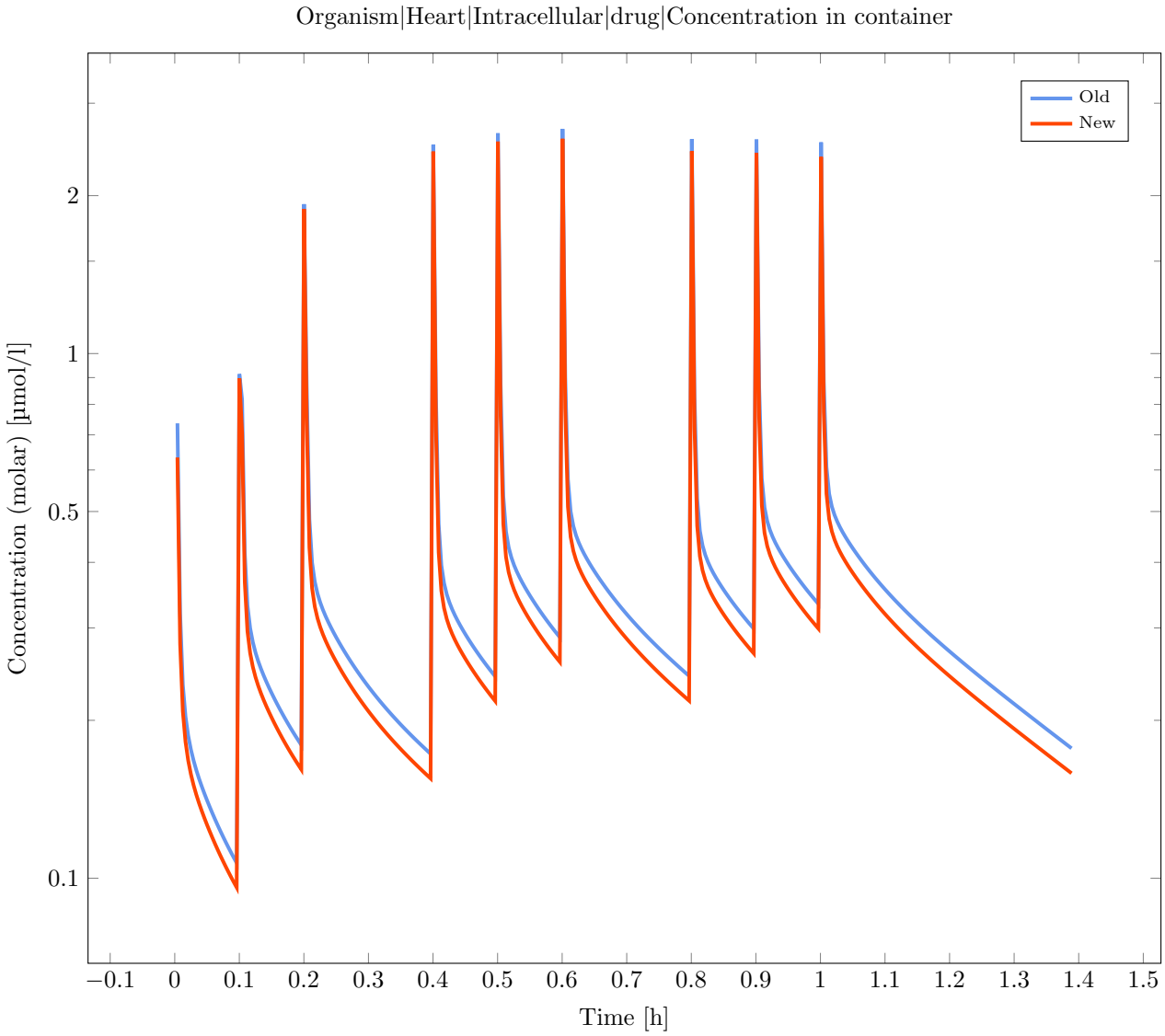
Organism|Heart|Intracellular|drug|Concentration in container
Old
New
Concentration (molar) [µmol/l]
2
1
0.5
0.1
−0.1
0
0.1
0.2
0.3
0.4
0.5
0.6
0.7
0.8
0.9
1
1.1
1.2
1.3
1.4
1.5
Time [h]

Figure 1.15

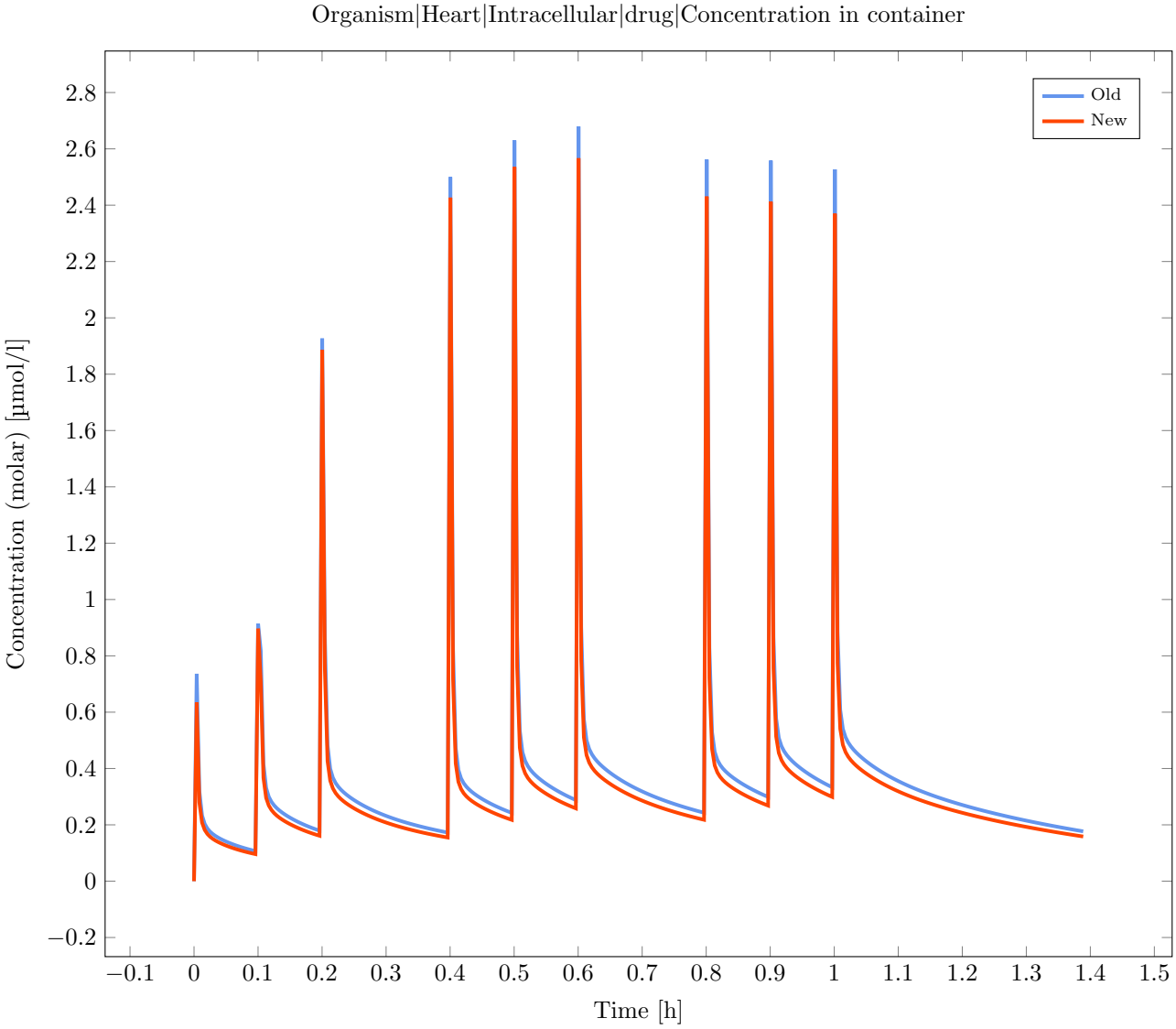

Figure 1.16

Output Path: Organism|Heart|drug|Tissue
Deviation for 'Organism|Heart|drug|Tissue' is 13.83% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.14

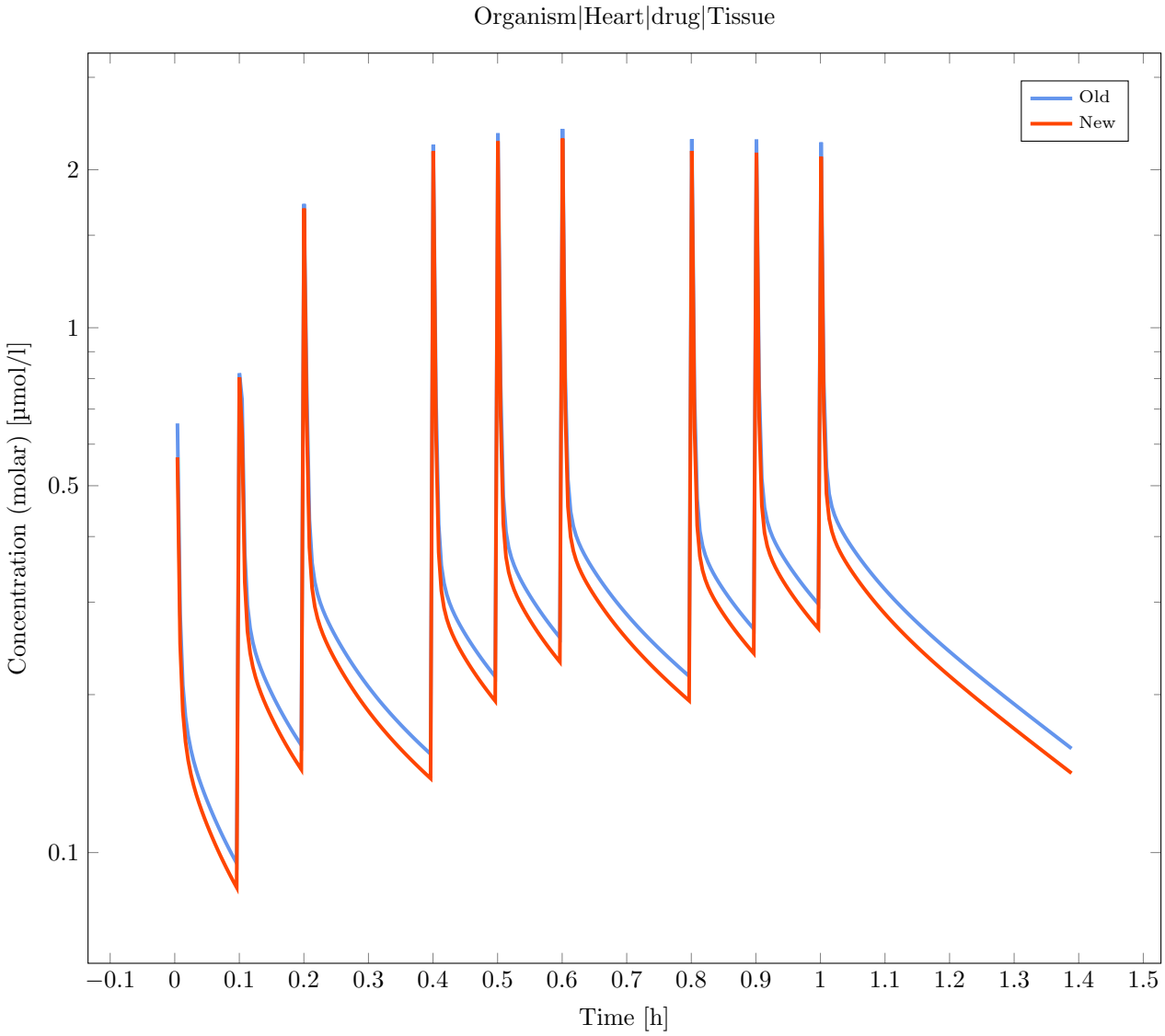
Organism|Heart|drug|Tissue
Old
New
Concentration (molar) [µmol/l]
2
1
0.5
0.1
−0.1
0
0.1
0.2
0.3
0.4
0.5
0.6
0.7
0.8
0.9
1
1.1
1.2
1.3
1.4
1.5
Time [h]

Figure 1.17

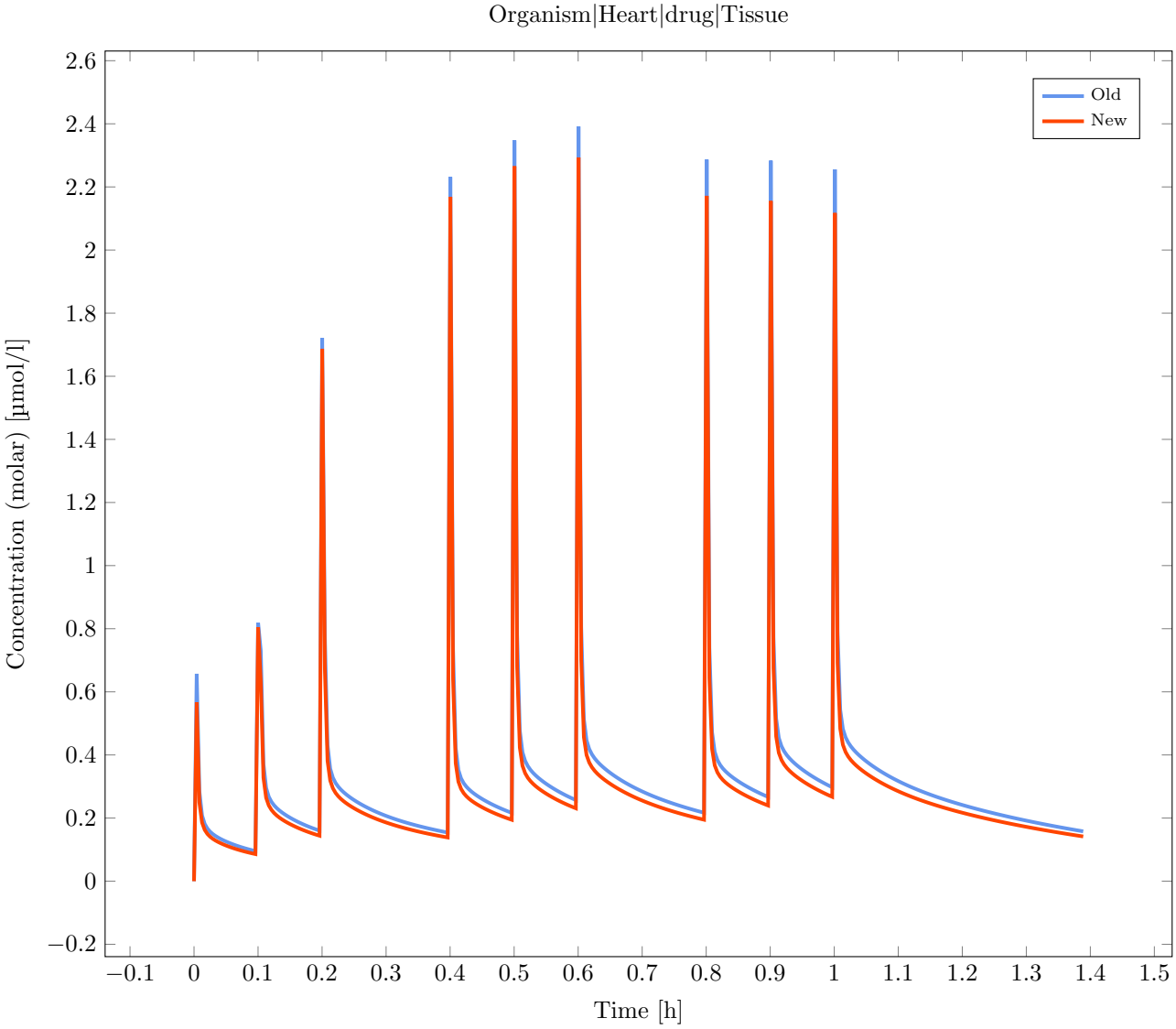

Figure 1.18

Output Path: Organism|Heart|drug|Interstitial Unbound
Deviation for 'Organism|Heart|drug|Interstitial Unbound' is 8.18% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.08

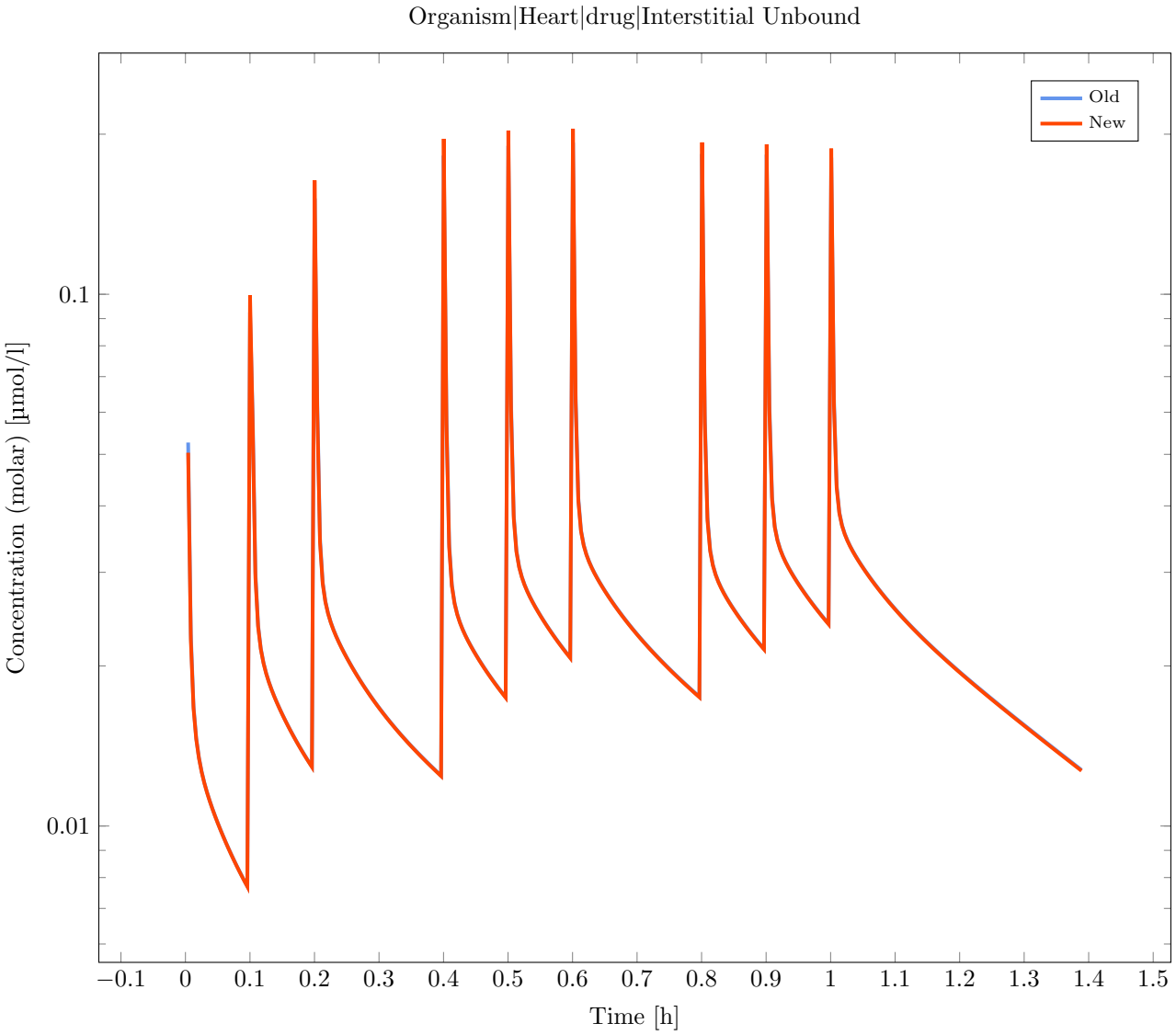
Organism|Heart|drug|Interstitial Unbound
Old
New
Concentration (molar) [µmol/l]
0.1
0.01
−0.1
0
0.1
0.2
0.3
0.4
0.5
0.6
0.7
0.8
0.9
1
1.1
1.2
1.3
1.4
1.5
Time [h]

Figure 1.19

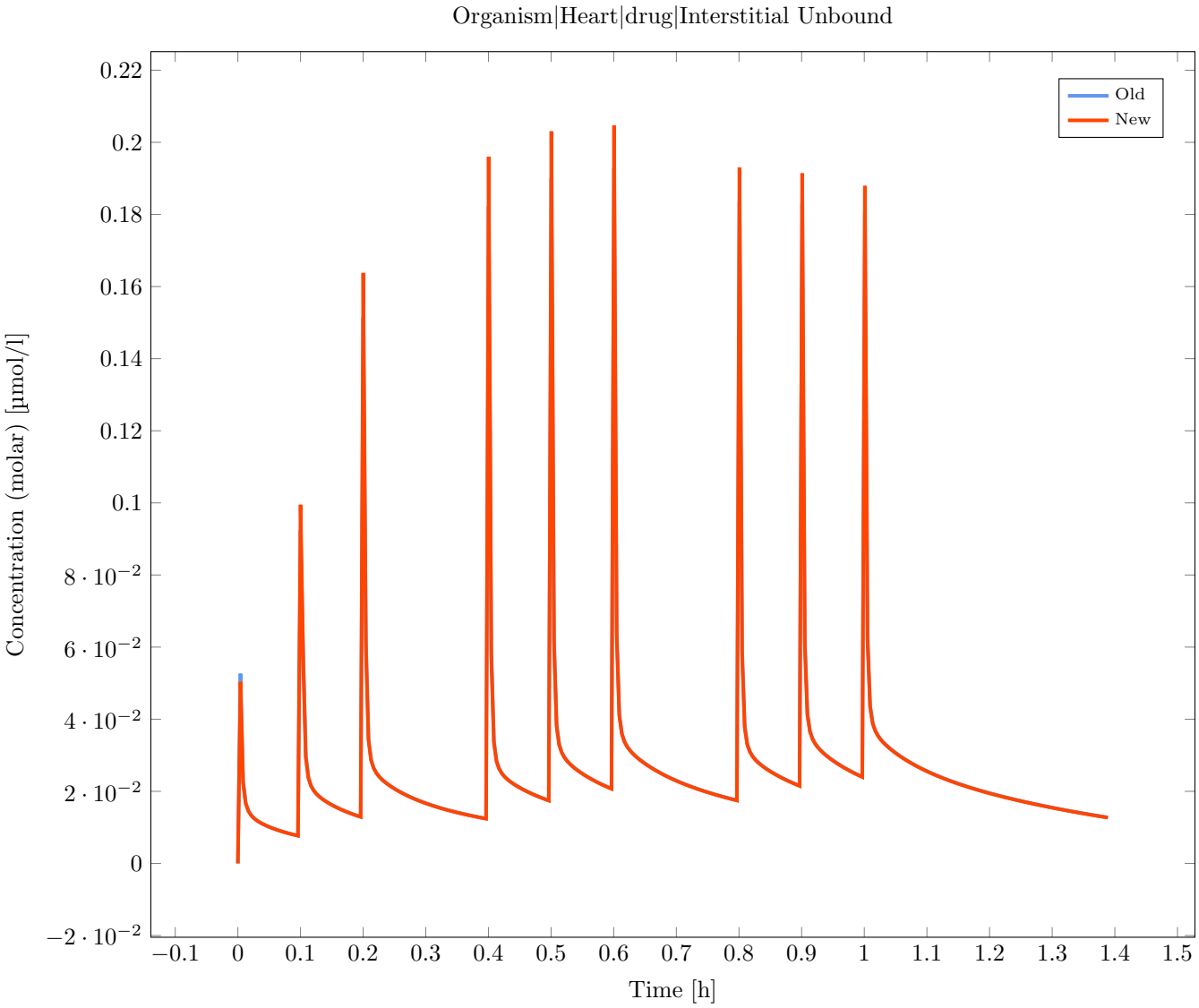

Figure 1.20

Output Path: Organism|Heart|Interstitial|drug|Concentration in container
Deviation for 'Organism|Heart|Interstitial|drug|Concentration in container' is 8.18% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.08

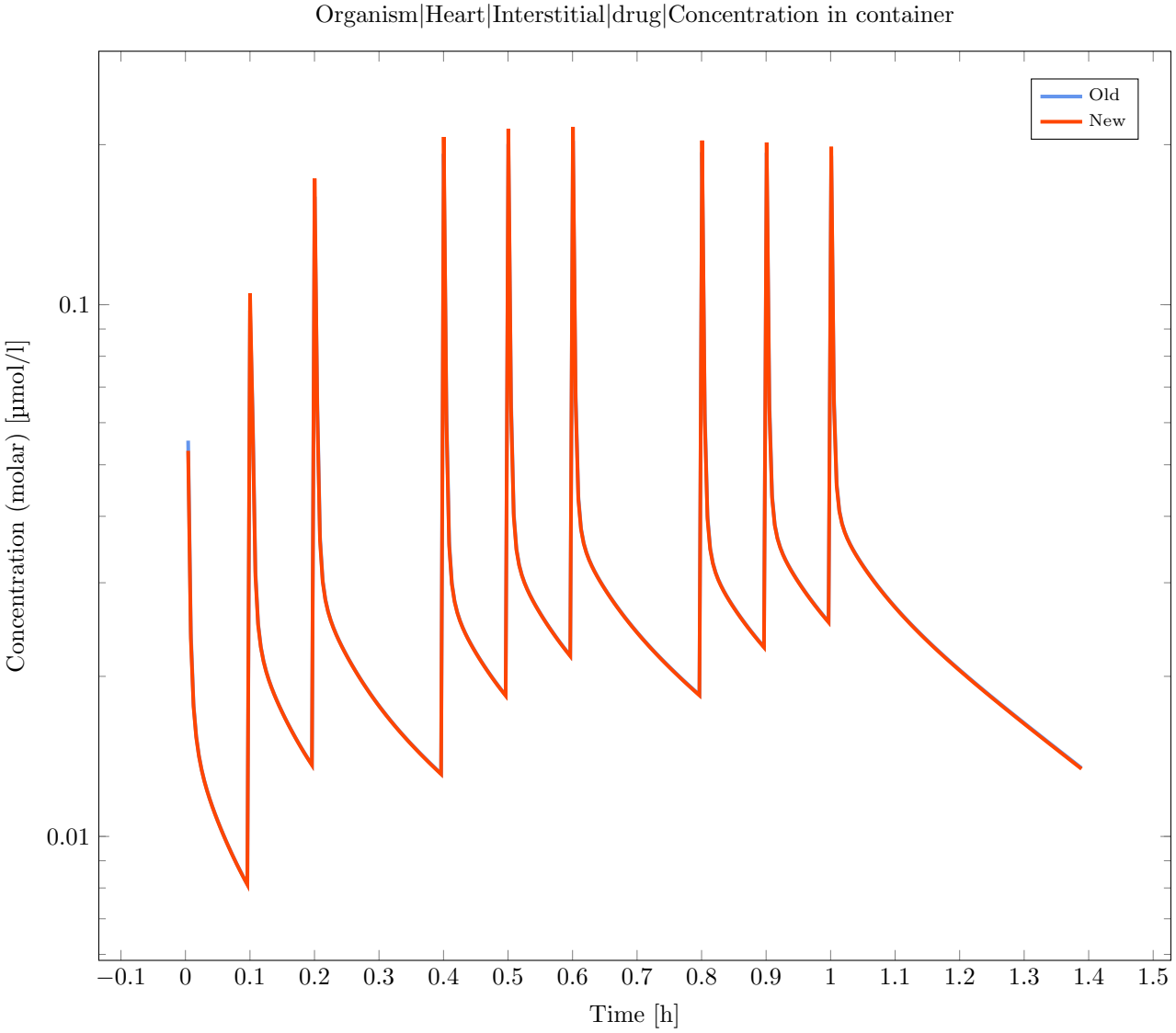
Organism|Heart|Interstitial|drug|Concentration in container
Old
New
Concentration (molar) [µmol/l]
0.1
0.01
−0.1
0
0.1
0.2
0.3
0.4
0.5
0.6
0.7
0.8
0.9
1
1.1
1.2
1.3
1.4
1.5
Time [h]

Figure 1.21

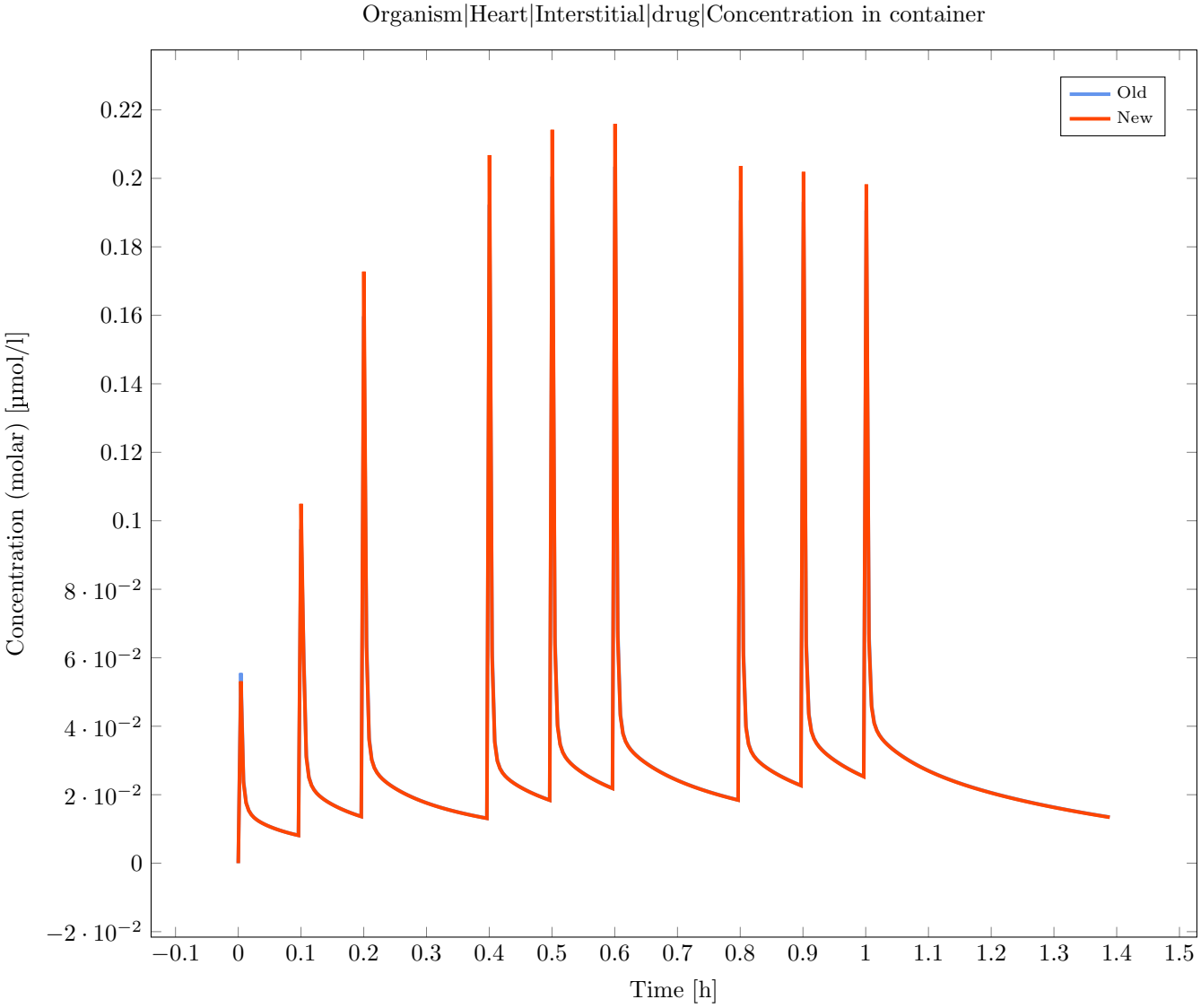

Figure 1.22

Simulation: Human_Oral_BiDaily_TableFormulation-S2_NoSuspension
Result of the validation: Invalid
Absolute Tolerance: 1.00E-10
Relative Tolerance: 1.00E-5

Output Path: Organism|Lumen|Duodenum|C1|Concentration in lumen
Deviation for 'Organism|Lumen|Duodenum|C1|Concentration in lumen' is 22.20% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.22

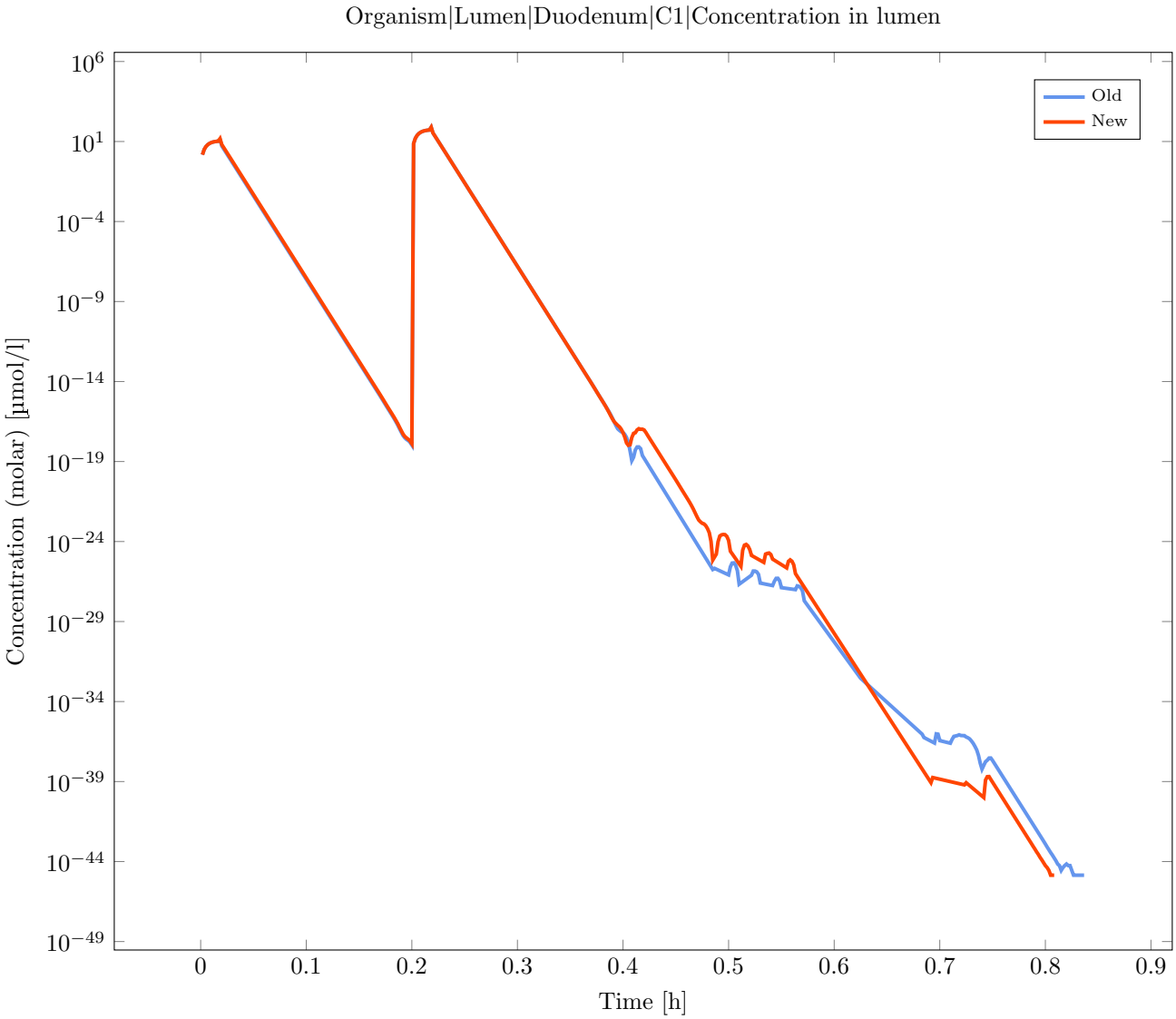
Organism|Lumen|Duodenum|C1|Concentration in lumen
Old
New
Concentration (molar) [µmol/l]
10^{6}
10^{1}
10^{-4}
10^{-9}
10^{-14}
10^{-19}
10^{-24}
10^{-29}
10^{-34}
10^{-39}
10^{-44}
10^{-49}
0
0.1
0.2
0.3
0.4
0.5
0.6
0.7
0.8
0.9
Time [h]

Figure 1.23

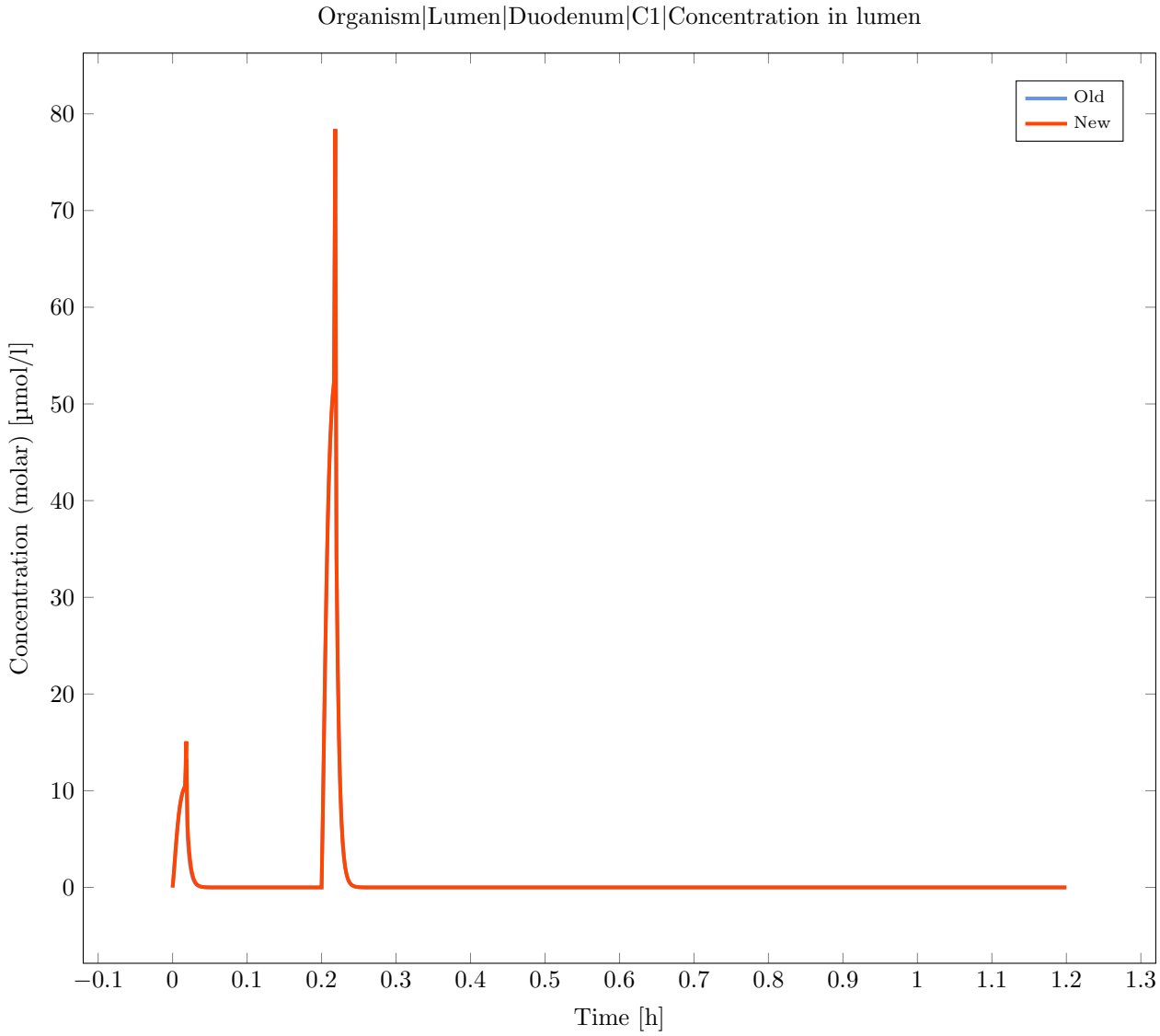

Figure 1.24

Output Path: Organism|Lumen|Stomach|C1|Concentration in lumen
Deviation for 'Organism|Lumen|Stomach|C1|Concentration in lumen' is 22.20% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.22

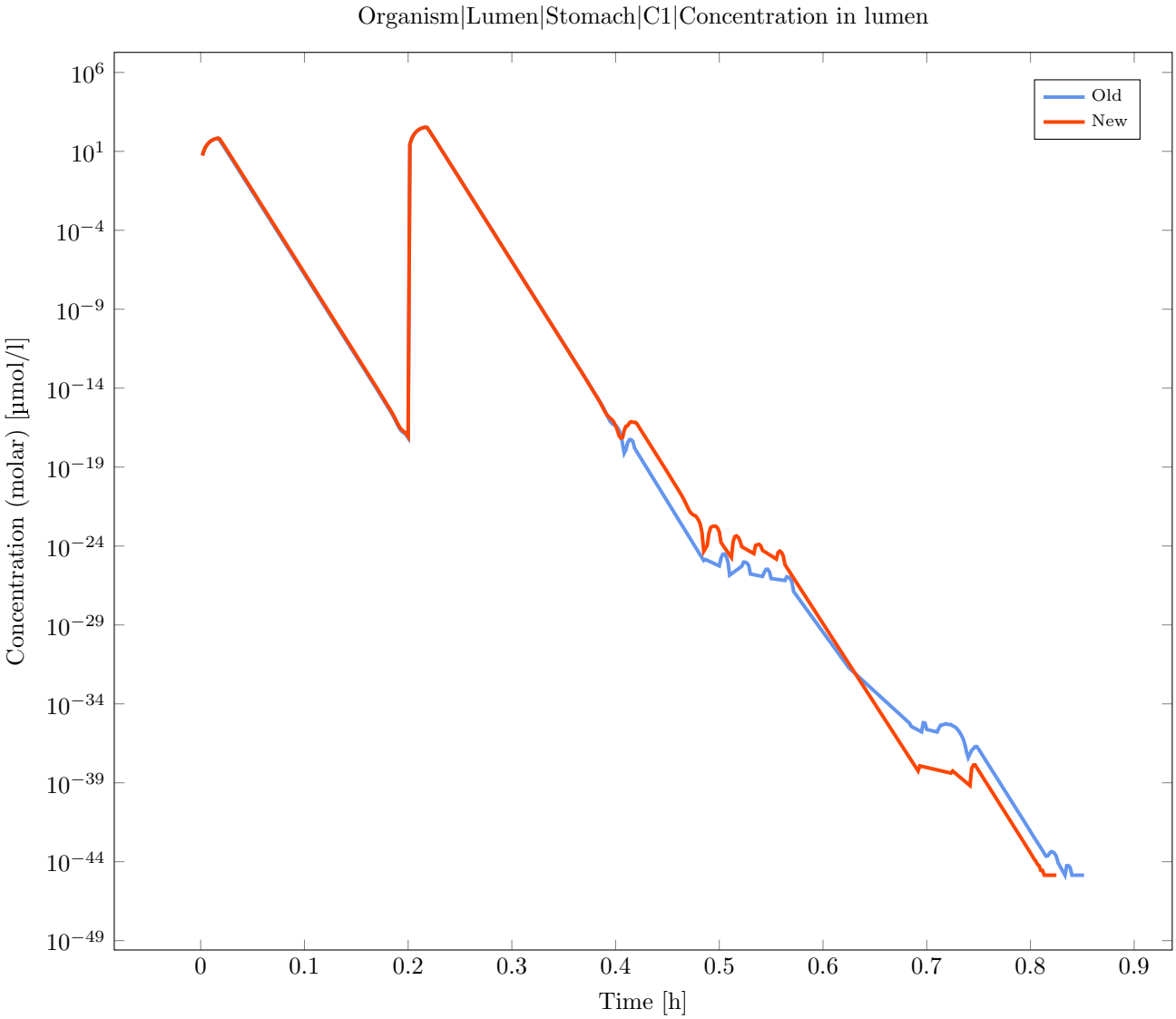
Organism|Lumen|Stomach|C1|Concentration in lumen
Old
New
Concentration (molar) [µmol/l]
10^{6}
10^{1}
10^{-4}
10^{-9}
10^{-14}
10^{-19}
10^{-24}
10^{-29}
10^{-34}
10^{-39}
10^{-44}
10^{-49}
0
0.1
0.2
0.3
0.4
0.5
0.6
0.7
0.8
0.9
Time [h]

Figure 1.25

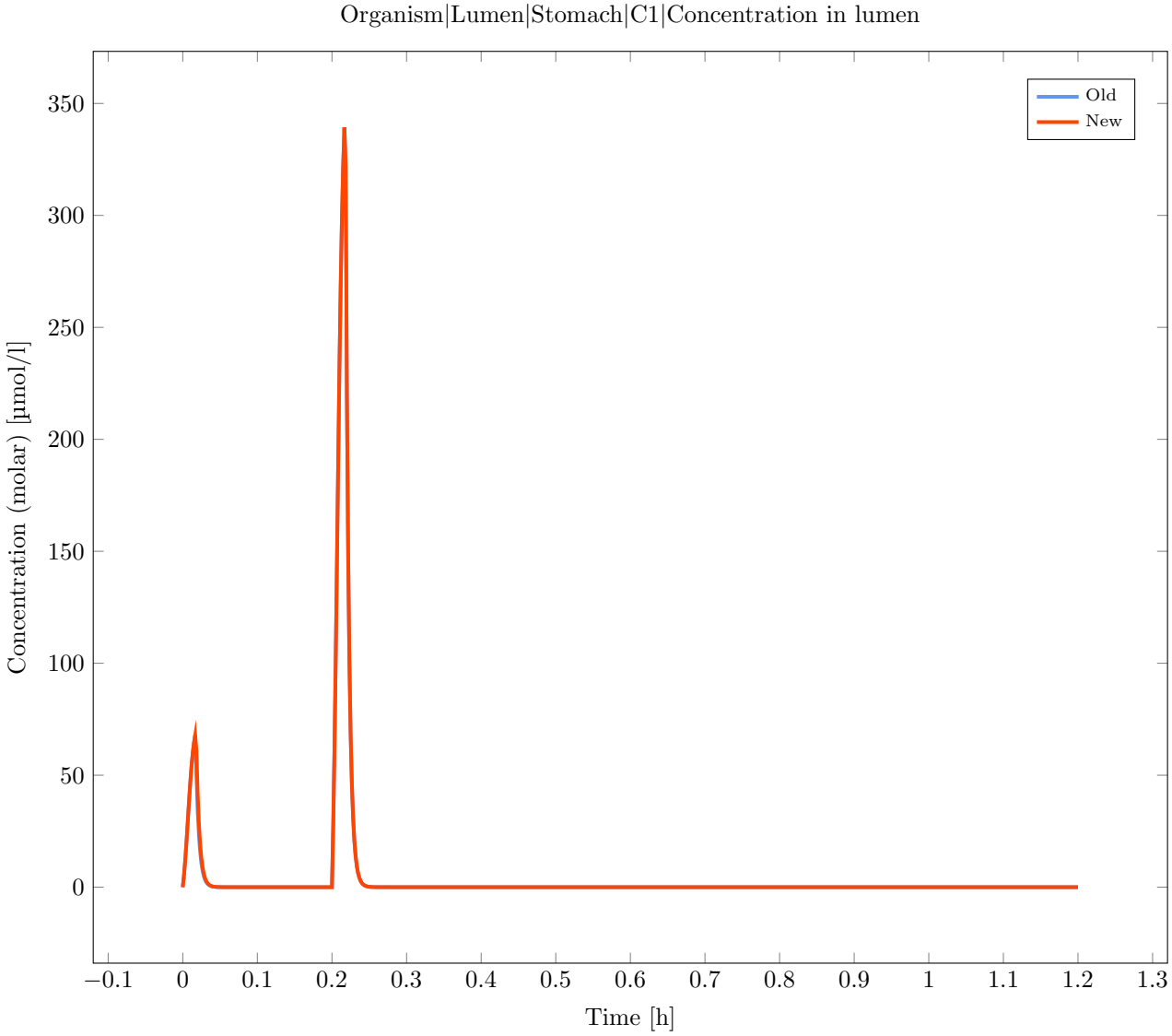

Figure 1.26

Output Path: Organism|Lumen|LowerIleum|C1|Concentration in lumen
Deviation for 'Organism|Lumen|LowerIleum|C1|Concentration in lumen' is 13.15% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.13

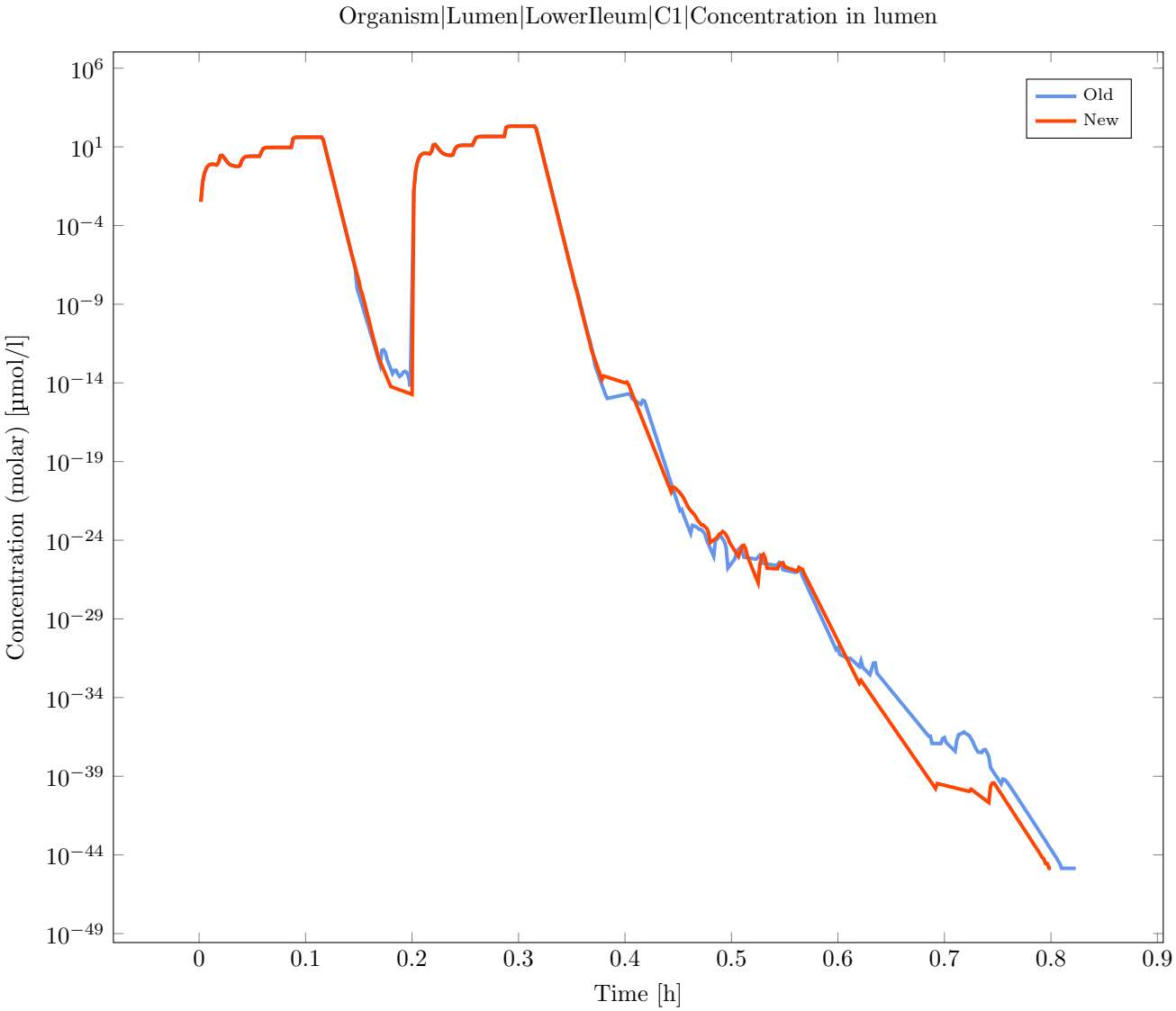
Organism|Lumen|LowerIleum|C1|Concentration in lumen
Old
New
Concentration (molar) [µmol/l]
10^{6}
10^{1}
10^{-4}
10^{-9}
10^{-14}
10^{-19}
10^{-24}
10^{-29}
10^{-34}
10^{-39}
10^{-44}
10^{-49}
0
0.1
0.2
0.3
0.4
0.5
0.6
0.7
0.8
0.9
Time [h]

Figure 1.27

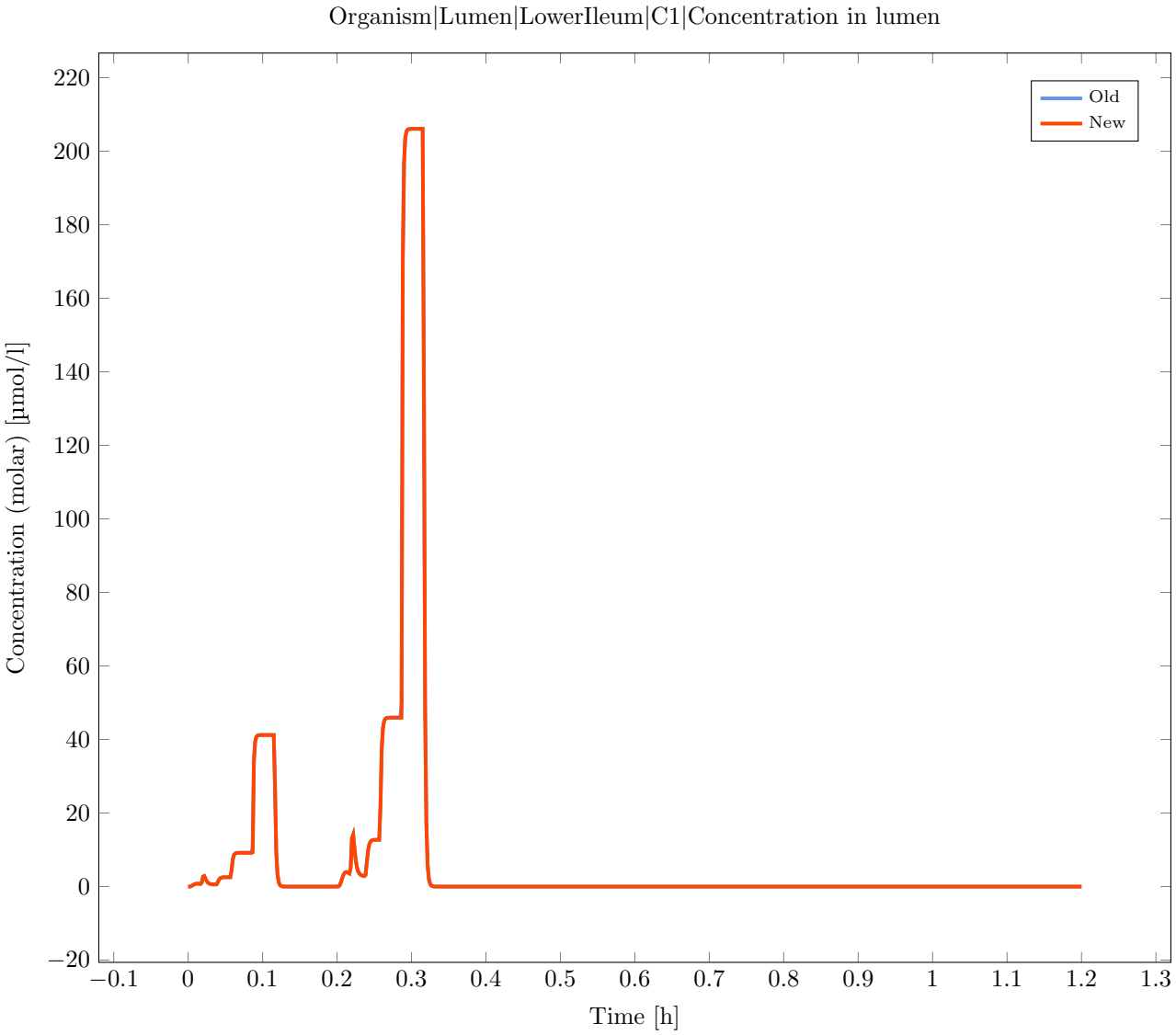

Figure 1.28

Output Path: Organism|Lumen|UpperJejunum|C1|Concentration in lumen
Deviation for 'Organism|Lumen|UpperJejunum|C1|Concentration in lumen' is 7.78% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.08

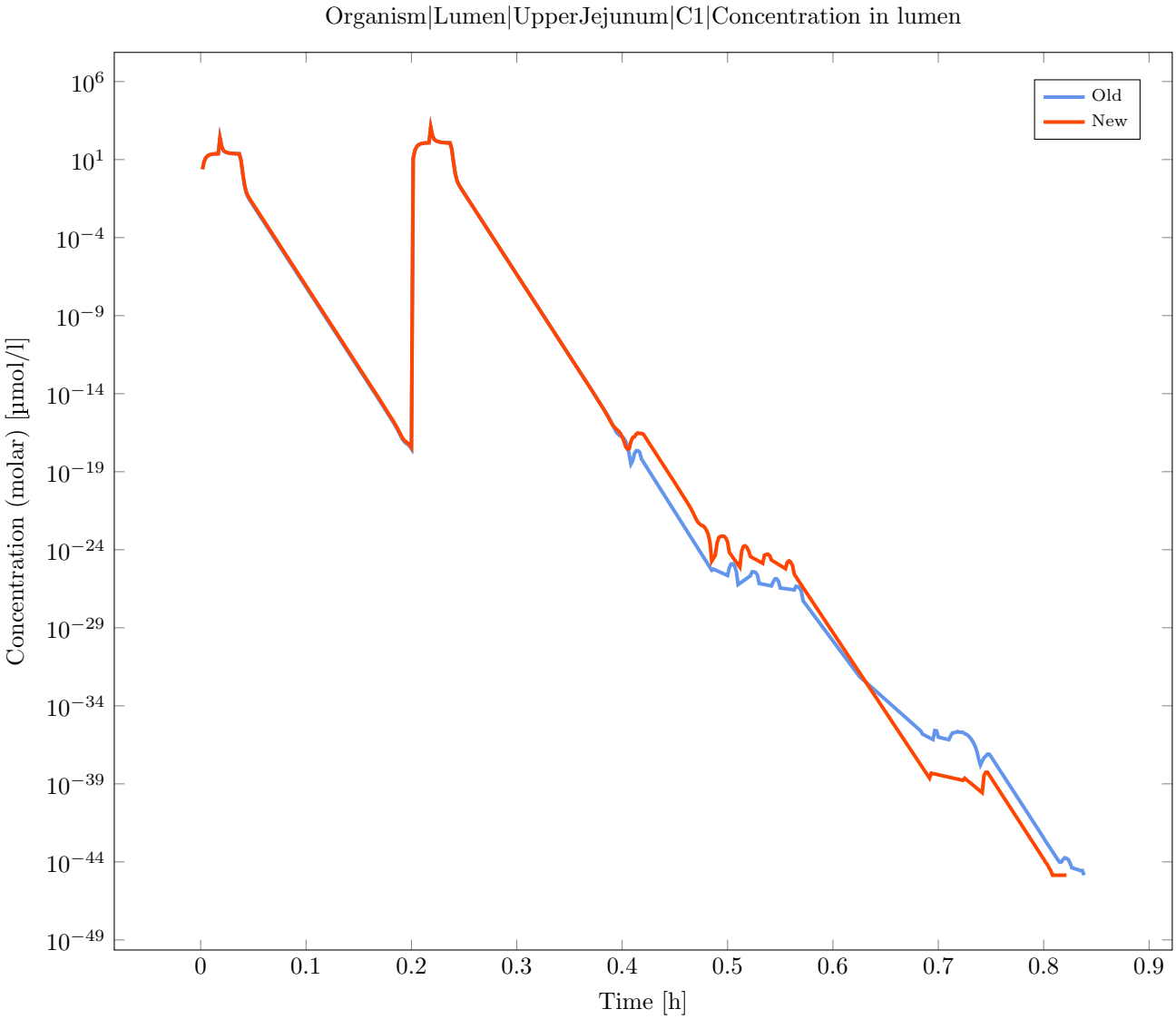
Organism|Lumen|UpperJejunum|C1|Concentration in lumen
Old
New
Concentration (molar) [µmol/l]
10^{6}
10^{1}
10^{-4}
10^{-9}
10^{-14}
10^{-19}
10^{-24}
10^{-29}
10^{-34}
10^{-39}
10^{-44}
10^{-49}
0
0.1
0.2
0.3
0.4
0.5
0.6
0.7
0.8
0.9
Time [h]

Figure 1.29

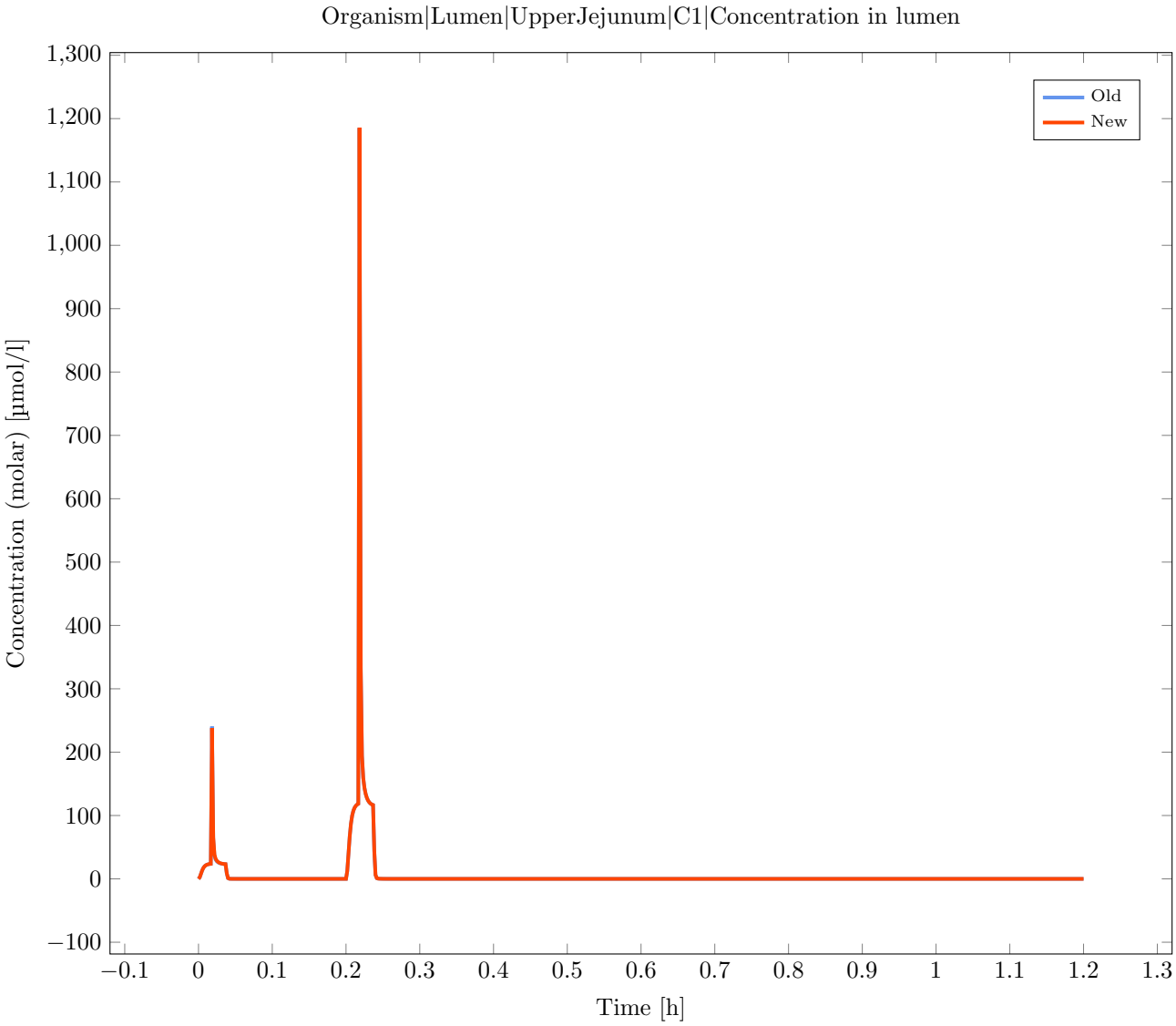

Figure 1.30

Output Path: Organism|Lumen|Rectum|C1|Concentration in lumen
Deviation for 'Organism|Lumen|Rectum|C1|Concentration in lumen' is 6.77% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.07

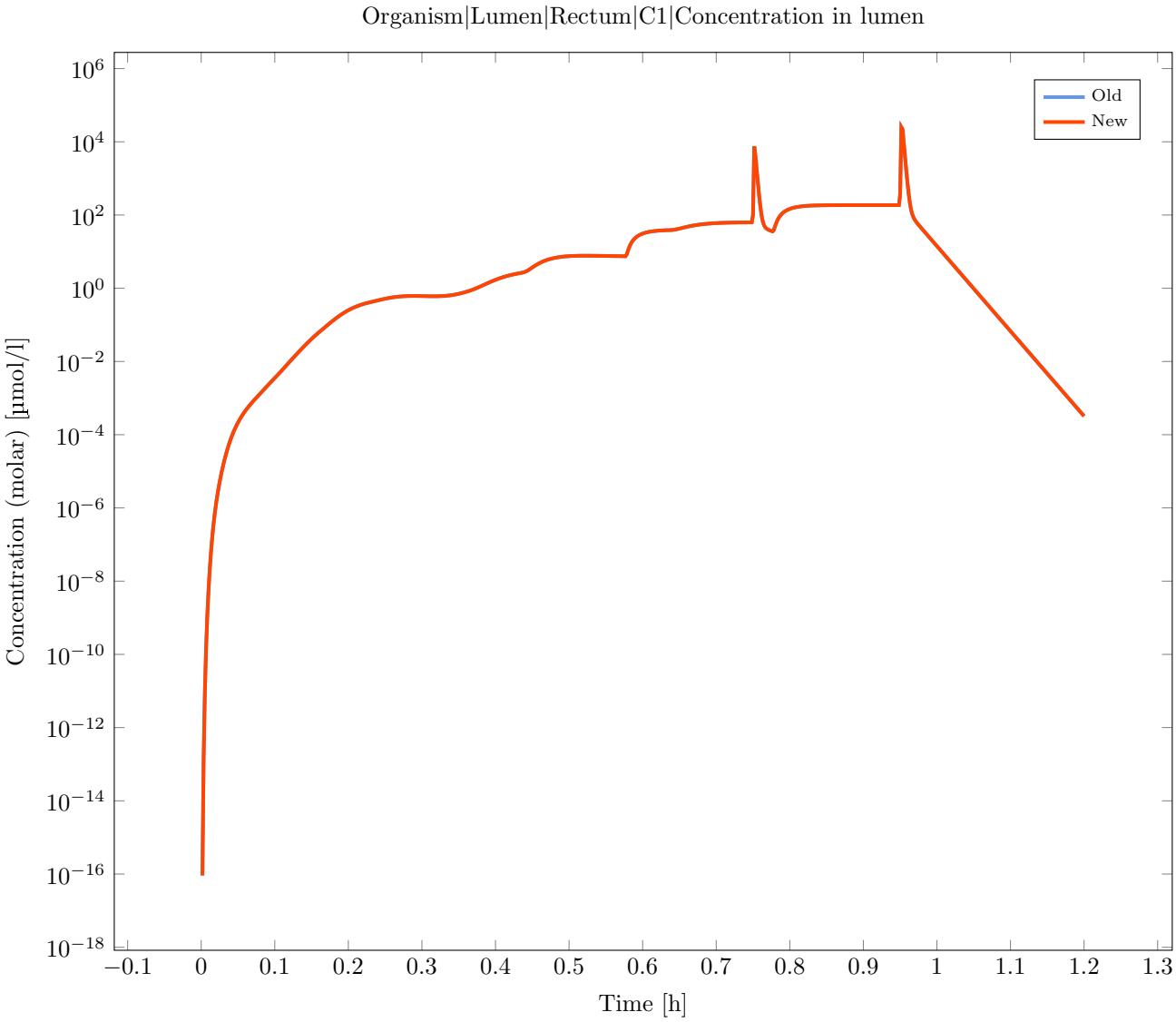
Organism|Lumen|Rectum|C1|Concentration in lumen
Old
New
Concentration (molar) [µmol/l]
10^{6}
10^{4}
10^{2}
10^{0}
10^{-2}
10^{-4}
10^{-6}
10^{-8}
10^{-10}
10^{-12}
10^{-14}
10^{-16}
10^{-18}
−0.1
0
0.1
0.2
0.3
0.4
0.5
0.6
0.7
0.8
0.9
1
1.1
1.2
1.3
Time [h]

Figure 1.31

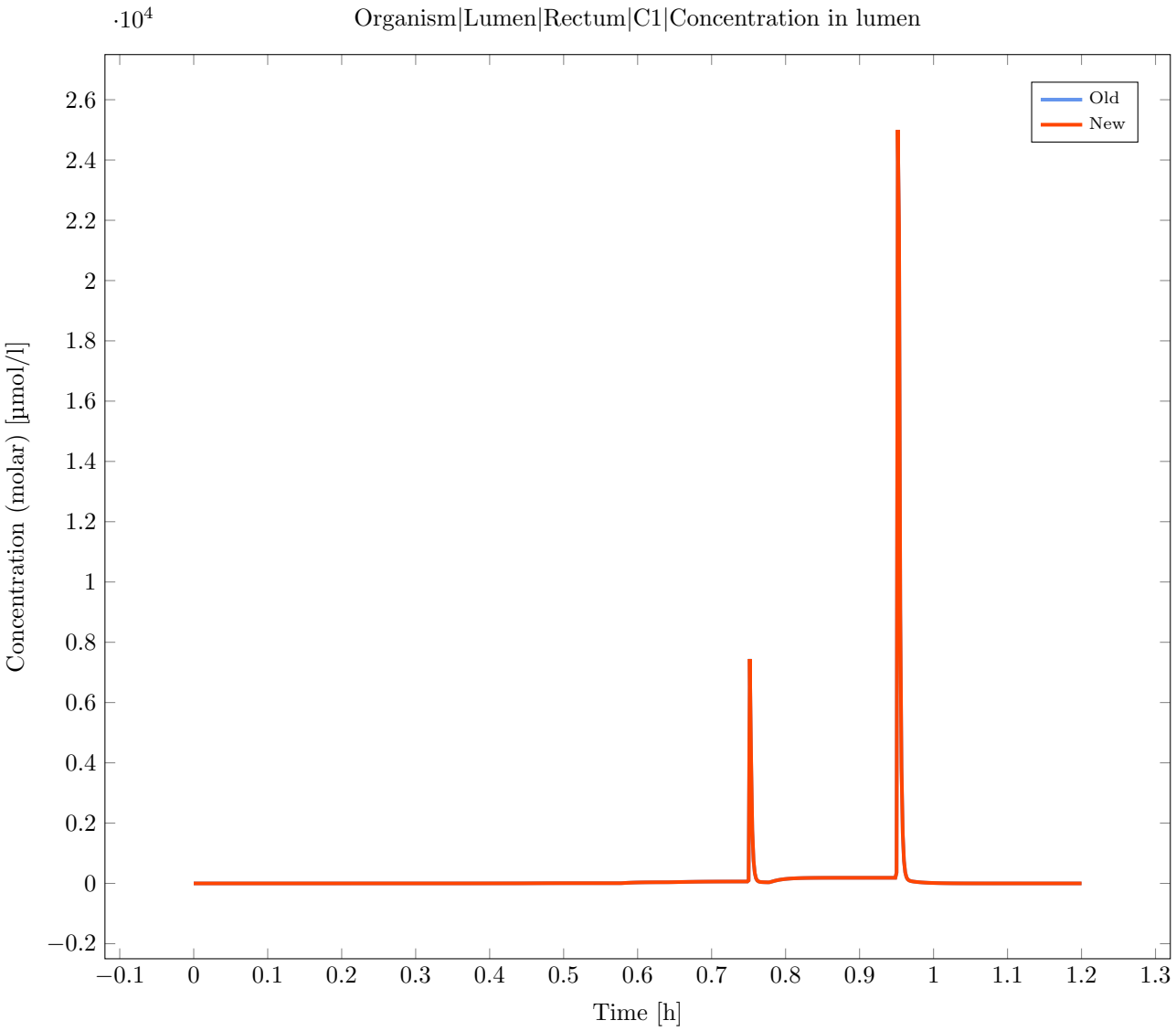

Figure 1.32

Simulation: Human_SingleORAL_MonoParticles_AsSuspention-Human_SingleORAL_MonoParticles_-AsSuspention
Result of the validation: Invalid
Absolute Tolerance: 1.00E-10
Relative Tolerance: 1.00E-5

Output Path: Organism|LargeIntestine|Mucosa|Caecum|Intracellular|drug|Concentration in container
Deviation for 'Organism|LargeIntestine|Mucosa|Caecum|Intracellular|drug|Concentration in container' is 32.90% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.33

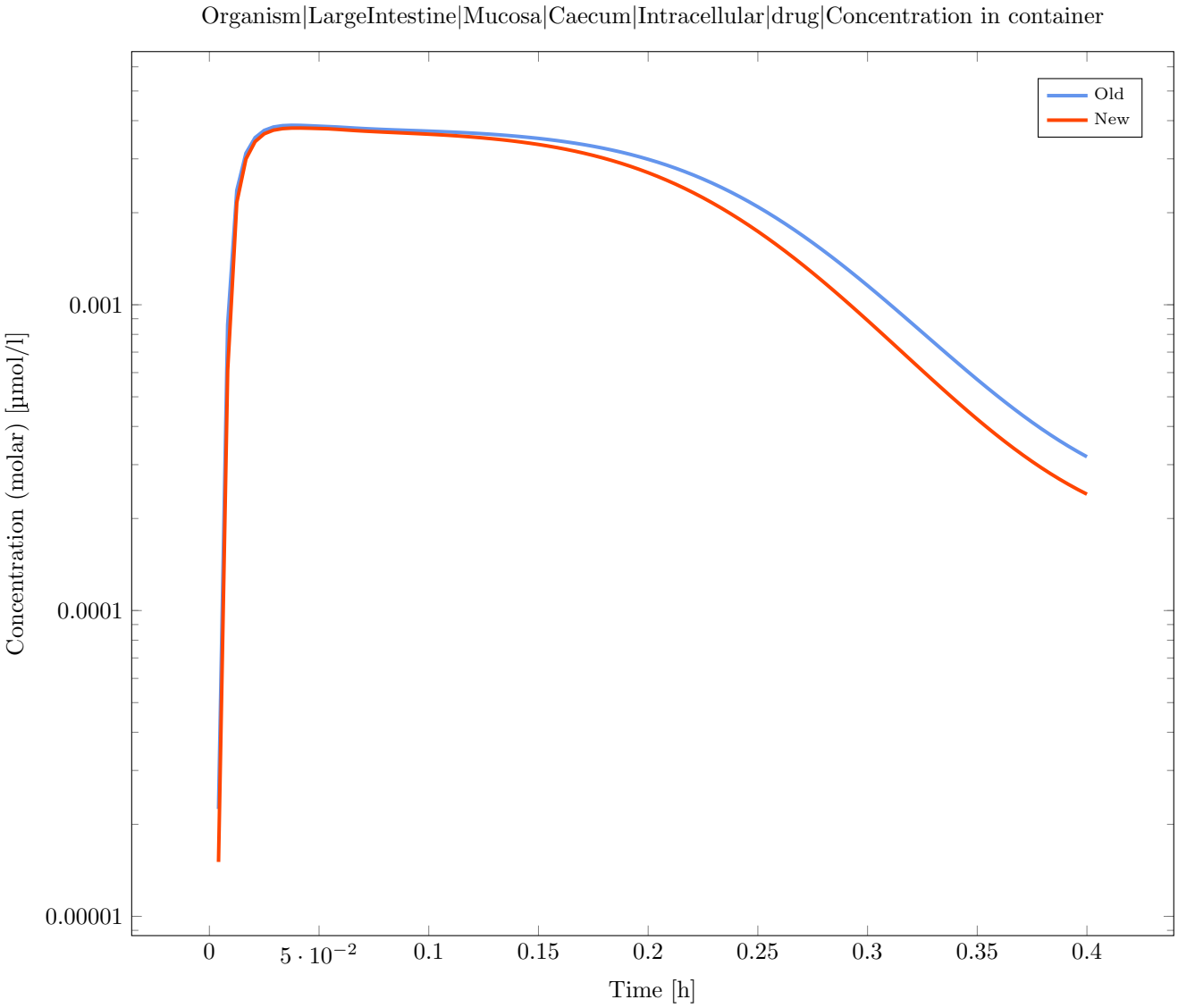
Organism|LargeIntestine|Mucosa|Caecum|Intracellular|drug|Concentration in container
Old
New
Concentration (molar) [µmol/l]
0.001
0.0001
0.00001
0
$5 \cdot 10^{-2}$
0.1
0.15
0.2
0.25
0.3
0.35
0.4
Time [h]

Figure 1.33

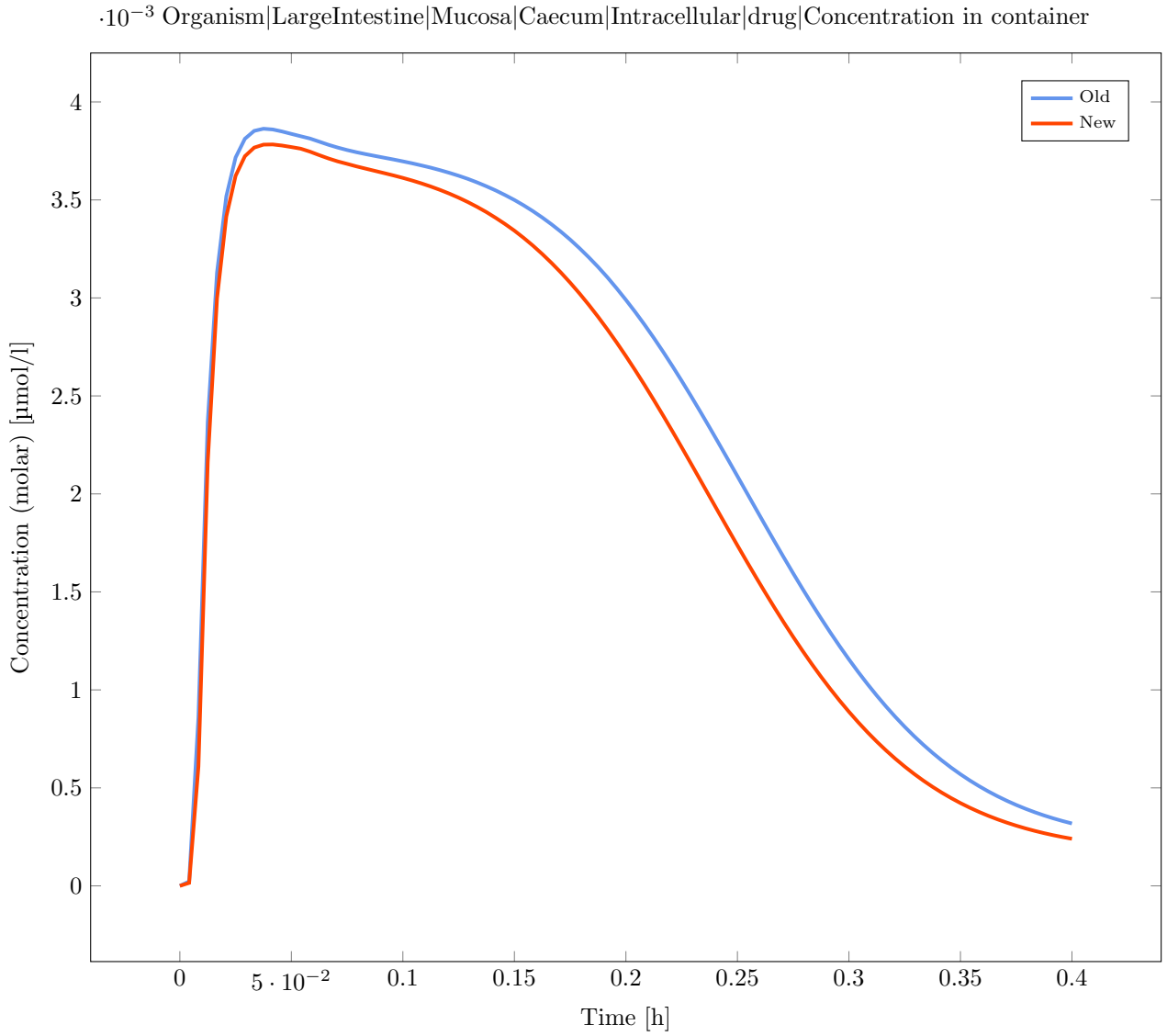

Figure 1.34

Output Path: Organism|SmallIntestine|Mucosa|LowerIleum|Intracellular|drug|Concentration in container
Deviation for 'Organism|SmallIntestine|Mucosa|LowerIleum|Intracellular|drug|Concentration in container' is 32.82% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.33

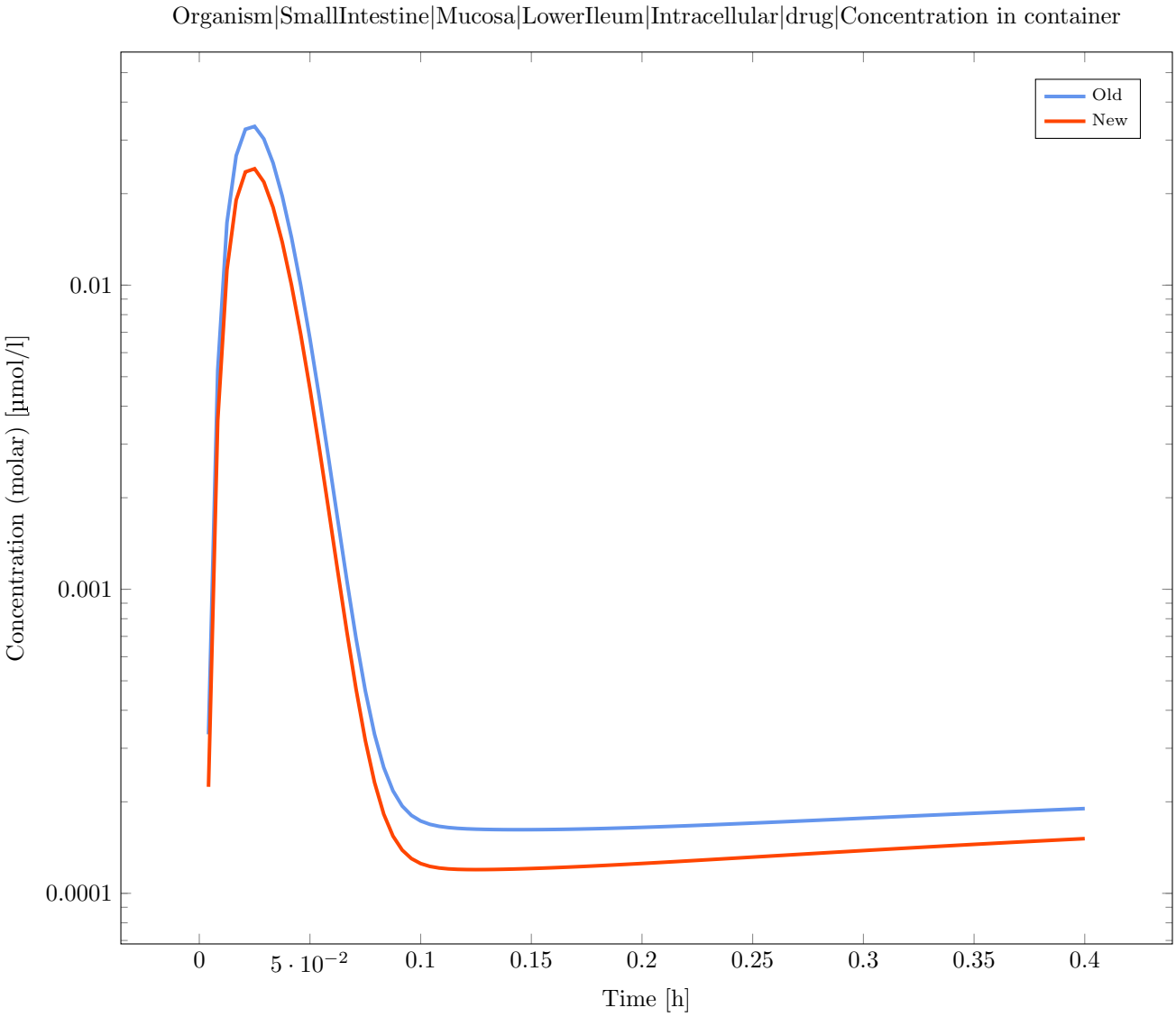
Organism|SmallIntestine|Mucosa|LowerIleum|Intracellular|drug|Concentration in container
Old
New
Concentration (molar) [µmol/l]
0.01
0.001
0.0001
0
$5 \cdot 10^{-2}$
0.1
0.15
0.2
0.25
0.3
0.35
0.4
Time [h]

Figure 1.35

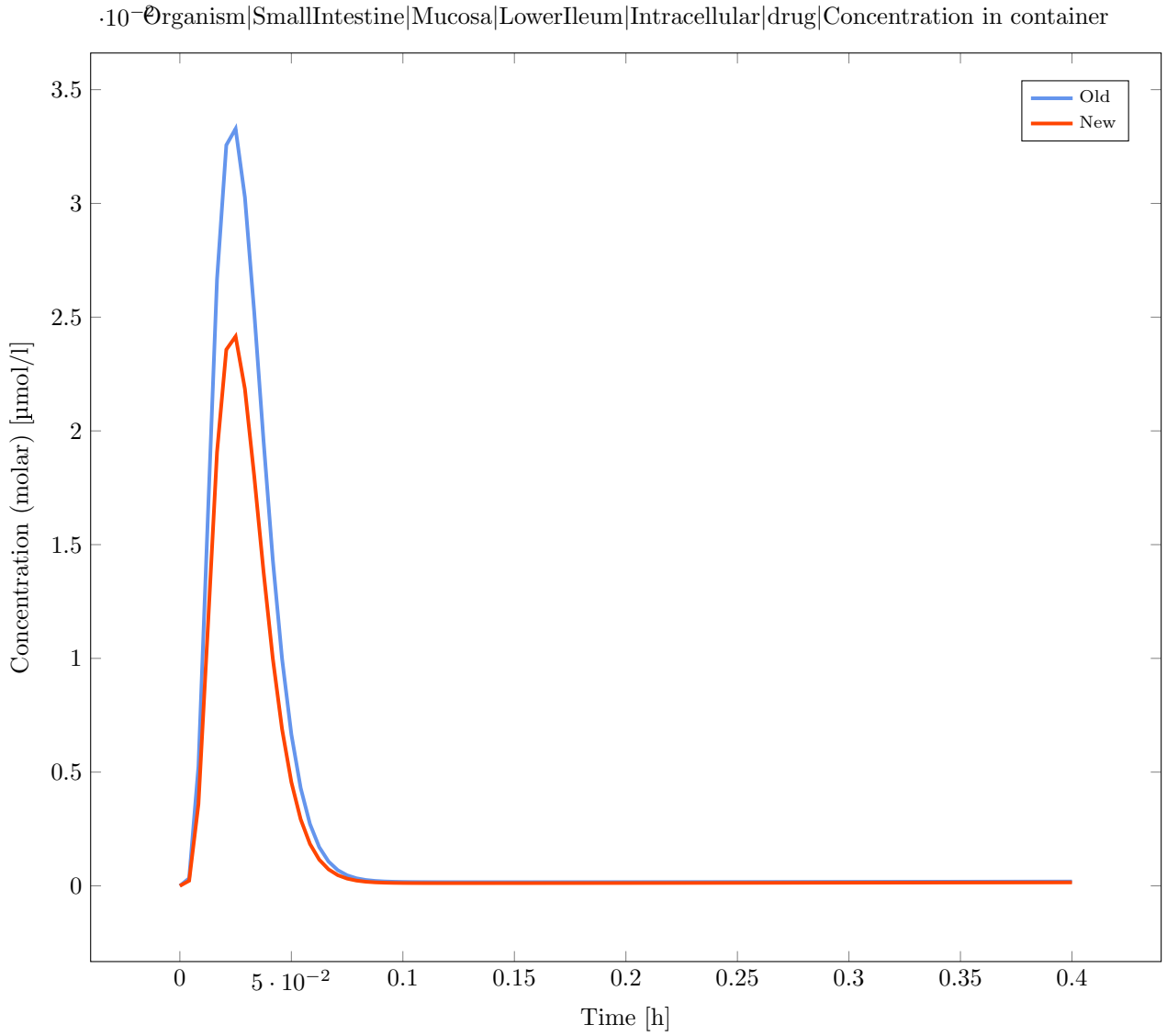

Figure 1.36

Output Path: Organism|SmallIntestine|Mucosa|LowerIleum|Interstitial|drug|Concentration in container
Deviation for 'Organism|SmallIntestine|Mucosa|LowerIleum|Interstitial|drug|Concentration in container' is 32.71% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.33

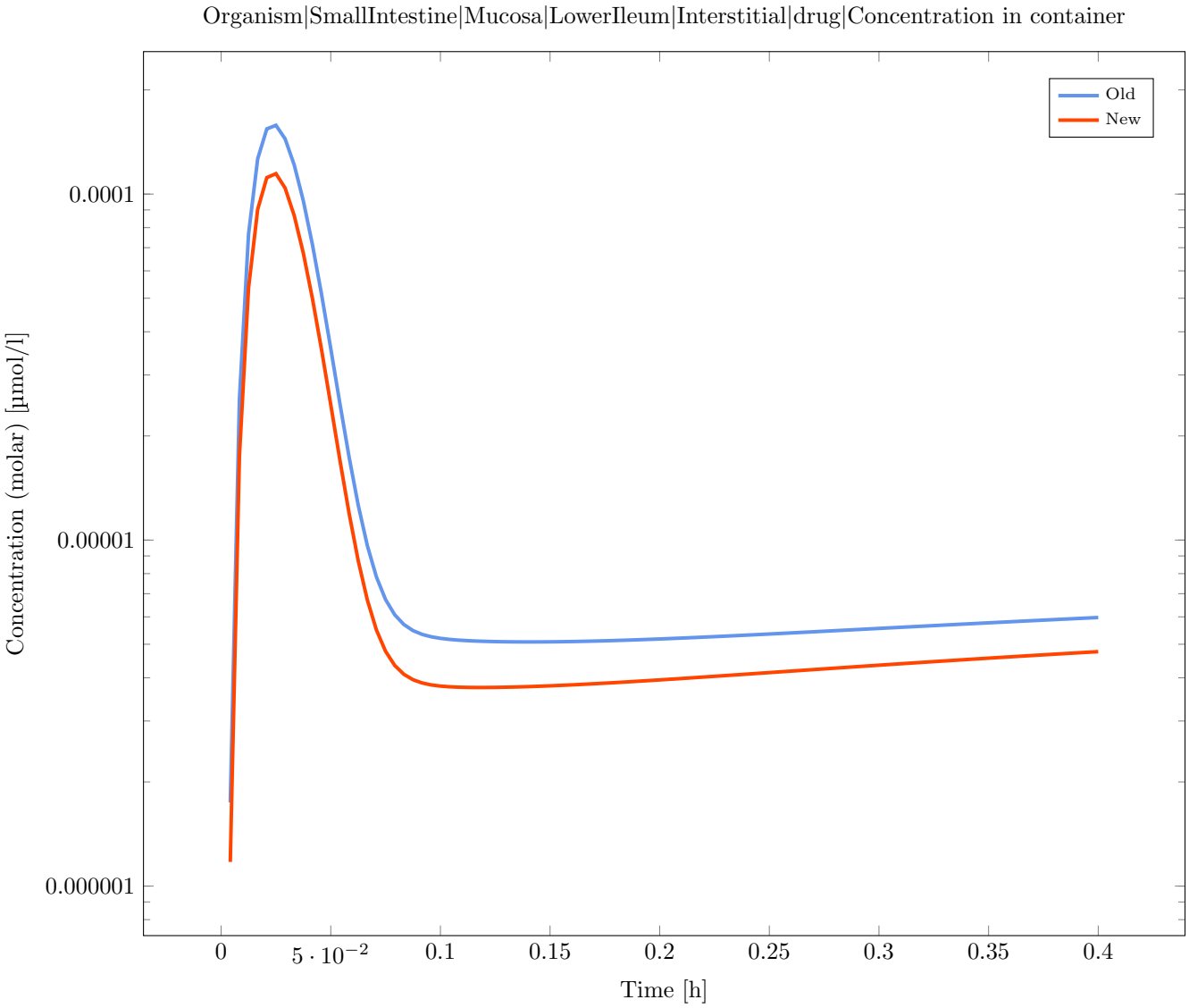
Organism|SmallIntestine|Mucosa|LowerIleum|Interstitial|drug|Concentration in container
Old
New
Concentration (molar) [µmol/l]
0.0001
0.00001
0.000001
0
5 · 10−2
0.1
0.15
0.2
0.25
0.3
0.35
0.4
Time [h]

Figure 1.37

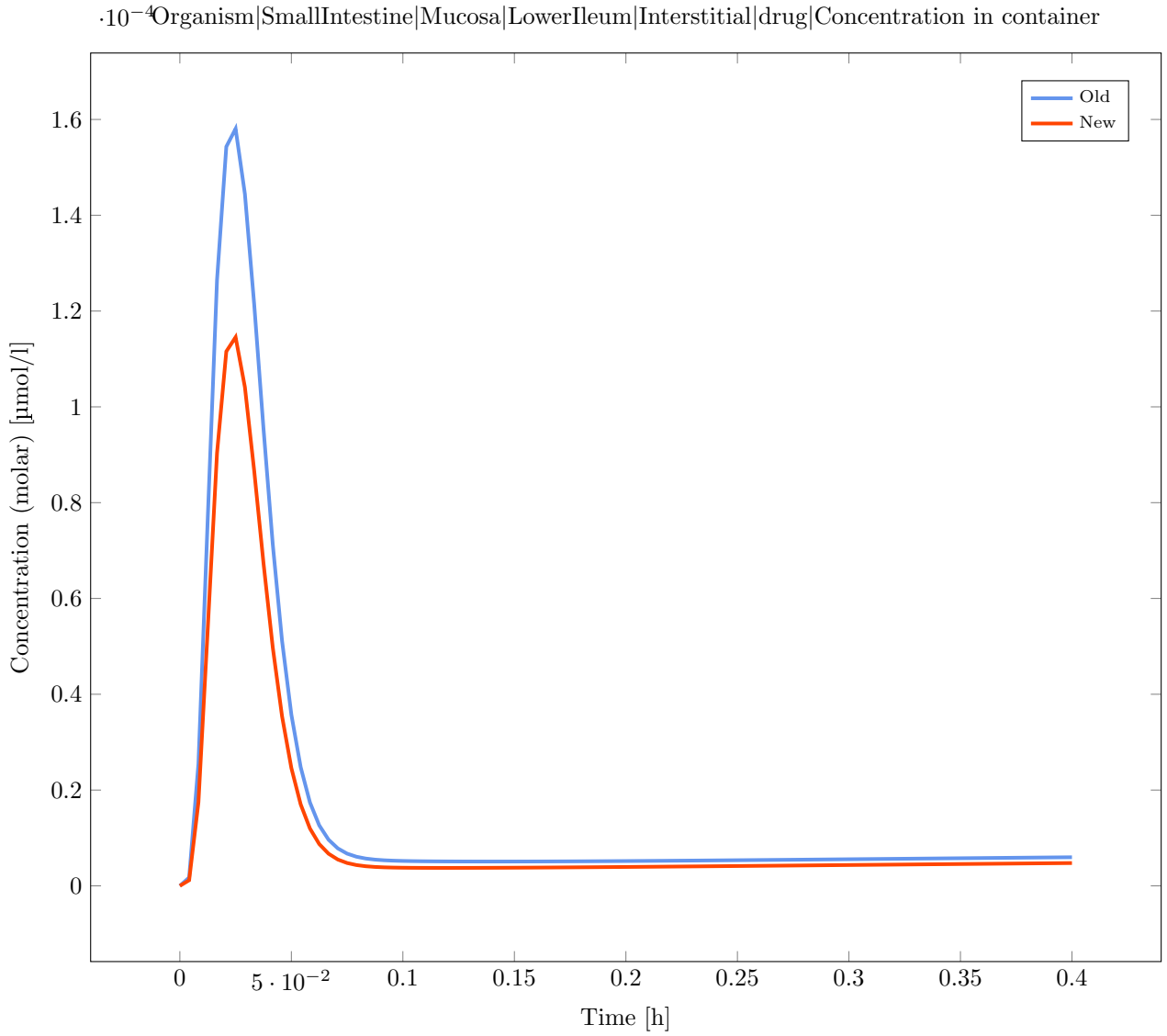

Figure 1.38

Output Path: Organism|SmallIntestine|Mucosa|LowerIleum|Plasma|drug|Concentration in container
Deviation for 'Organism|SmallIntestine|Mucosa|LowerIleum|Plasma|drug|Concentration in container' is 32.71% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.33

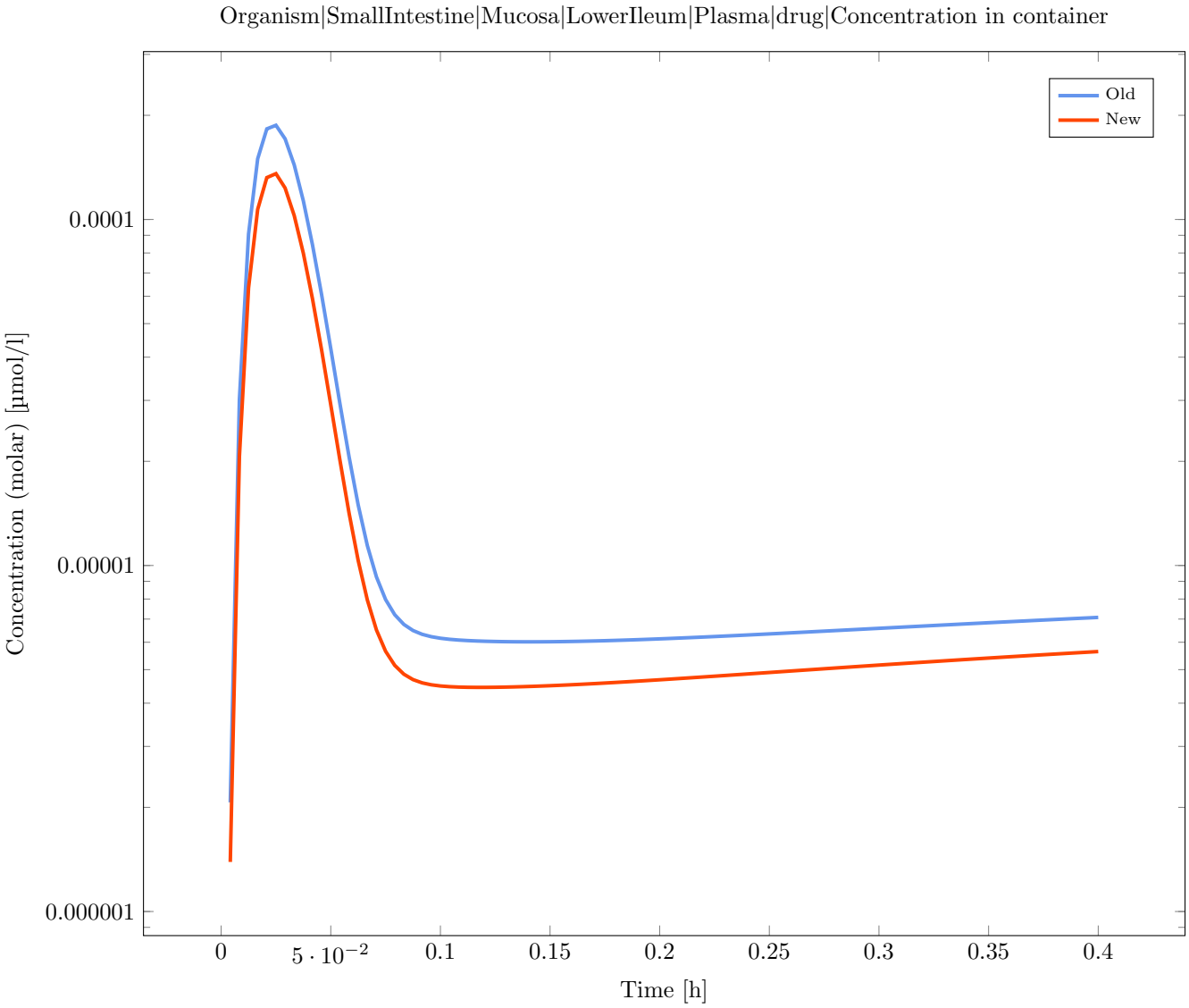
Organism|SmallIntestine|Mucosa|LowerIleum|Plasma|drug|Concentration in container
Old
New
Concentration (molar) [µmol/l]
0.0001
0.00001
0.000001
0
$5 \cdot 10^{-2}$
0.1
0.15
0.2
0.25
0.3
0.35
0.4
Time [h]

Figure 1.39

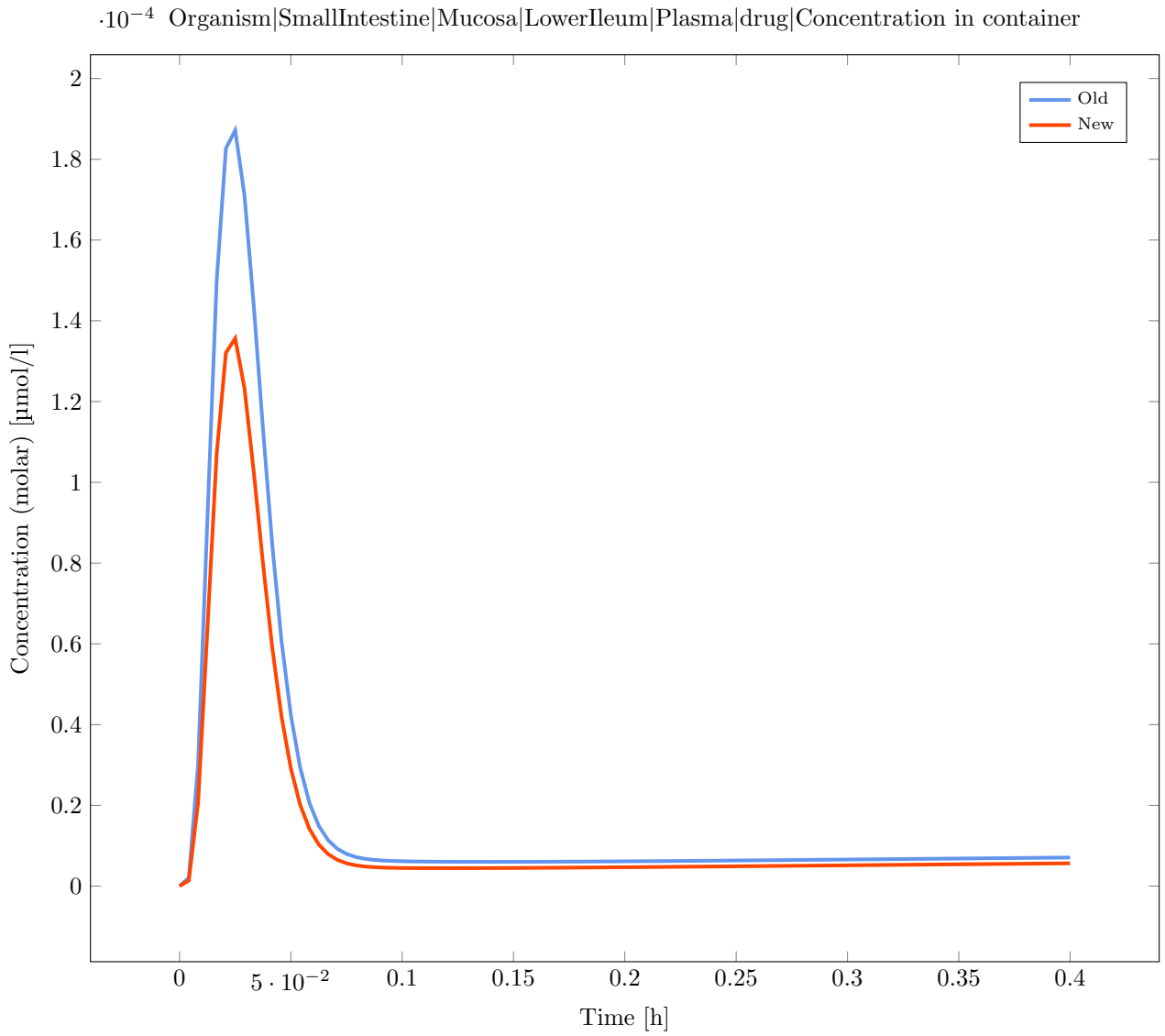

Figure 1.40

Output Path: Organism|Skin|drug|Intracellular Unbound
Deviation for 'Organism|Skin|drug|Intracellular Unbound' is 32.45% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.32

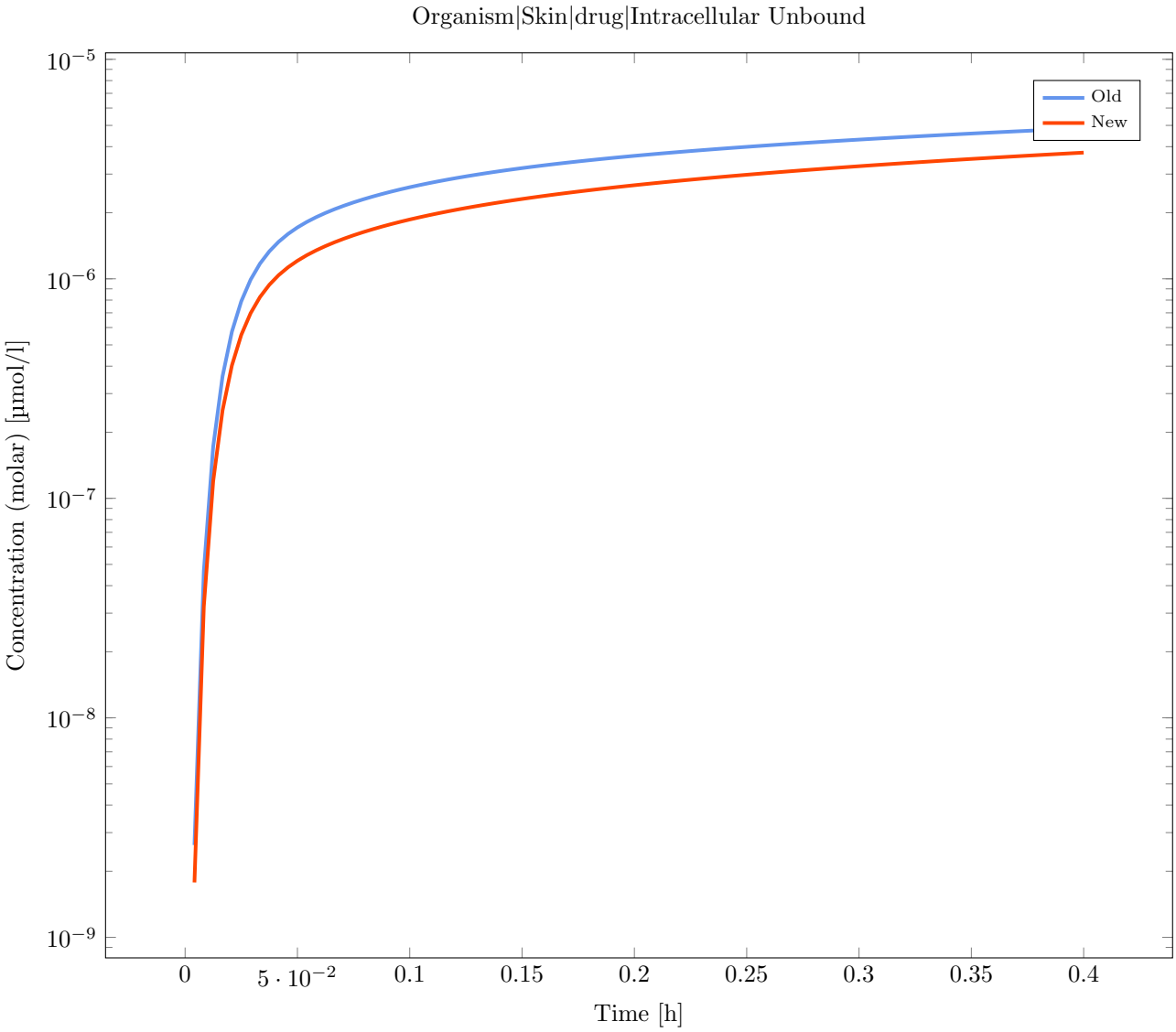
Organism|Skin|drug|Intracellular Unbound
Old
New
Concentration (molar) [µmol/l]
10^{-5}
10^{-6}
10^{-7}
10^{-8}
10^{-9}
0
$5 \cdot 10^{-2}$
0.1
0.15
0.2
0.25
0.3
0.35
0.4
Time [h]

Figure 1.41

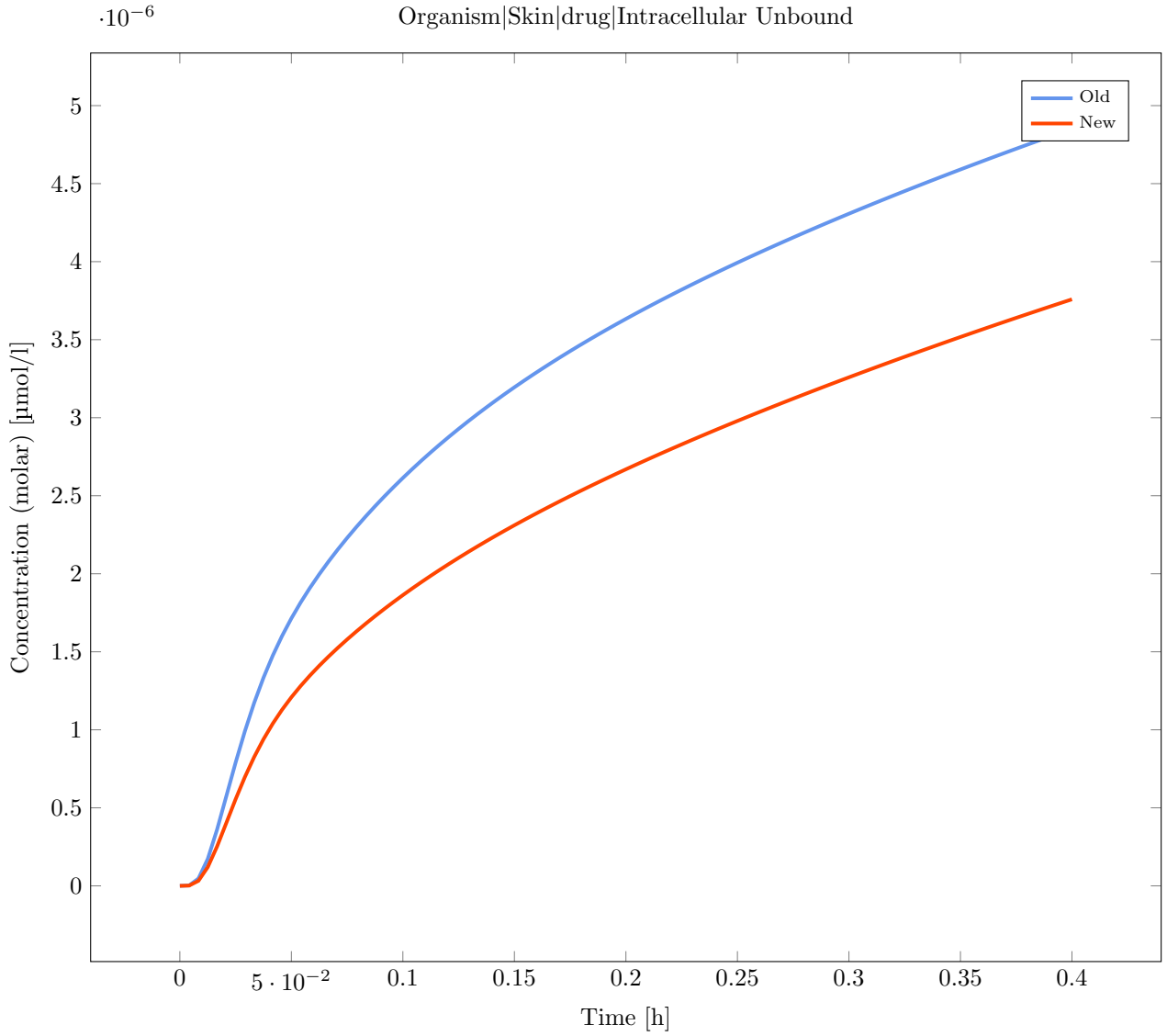

Figure 1.42

Simulation: Human_SingleORAL_PolyParticlesLogNormal_AsSuspention-Human_SingleORAL_-PolyParticlesLogNormal_AsSuspention
Result of the validation: Invalid
Absolute Tolerance: 1.00E-10
Relative Tolerance: 1.00E-5

Output Path: Organism|LargeIntestine|Mucosa|Caecum|Intracellular|drug|Concentration in container
Deviation for 'Organism|LargeIntestine|Mucosa|Caecum|Intracellular|drug|Concentration in container' is 32.92% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.33

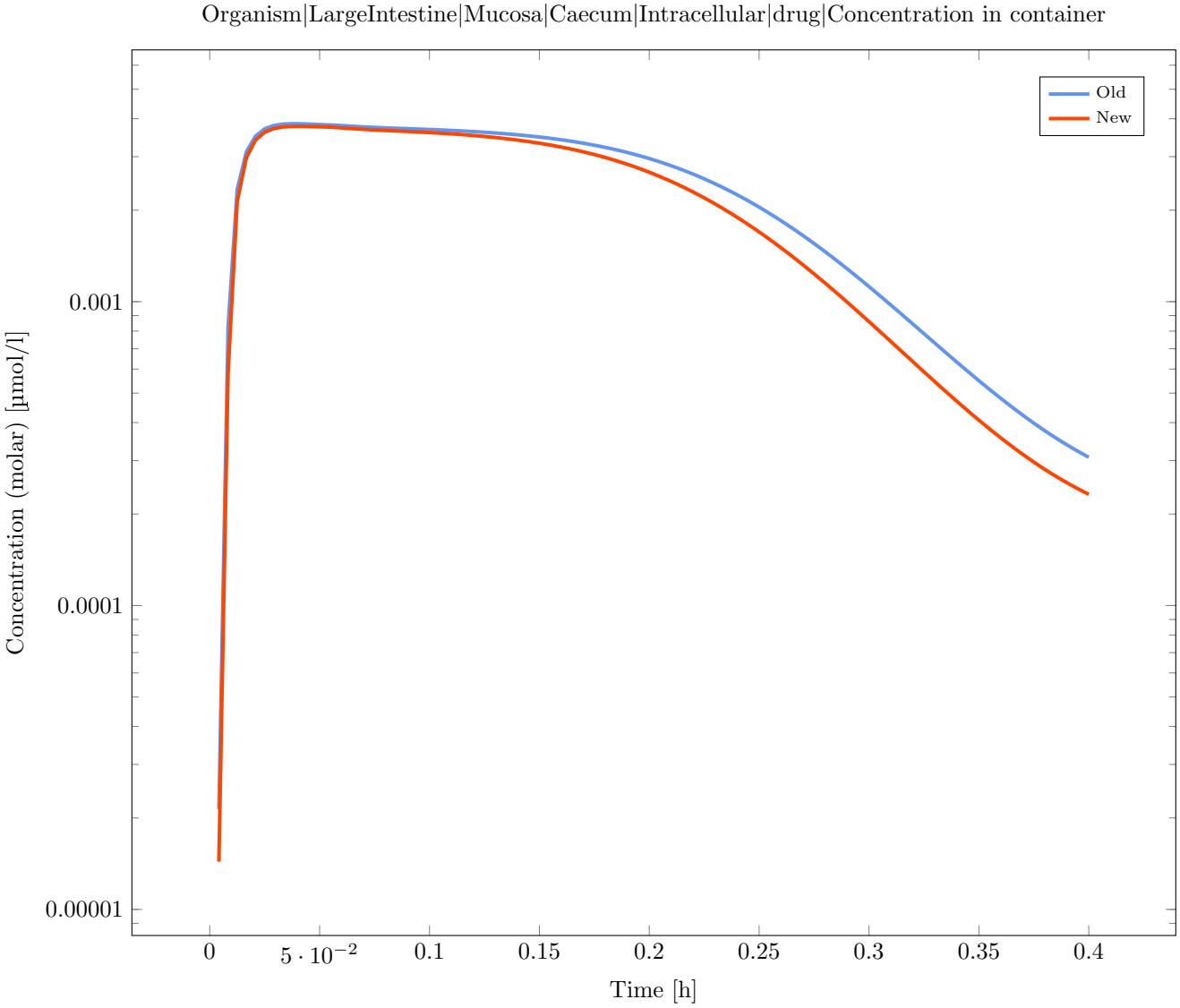
Organism|LargeIntestine|Mucosa|Caecum|Intracellular|drug|Concentration in container
Old
New
Concentration (molar) [µmol/l]
0.001
0.0001
0.00001
0
$5 \cdot 10^{-2}$
0.1
0.15
0.2
0.25
0.3
0.35
0.4
Time [h]

Figure 1.43

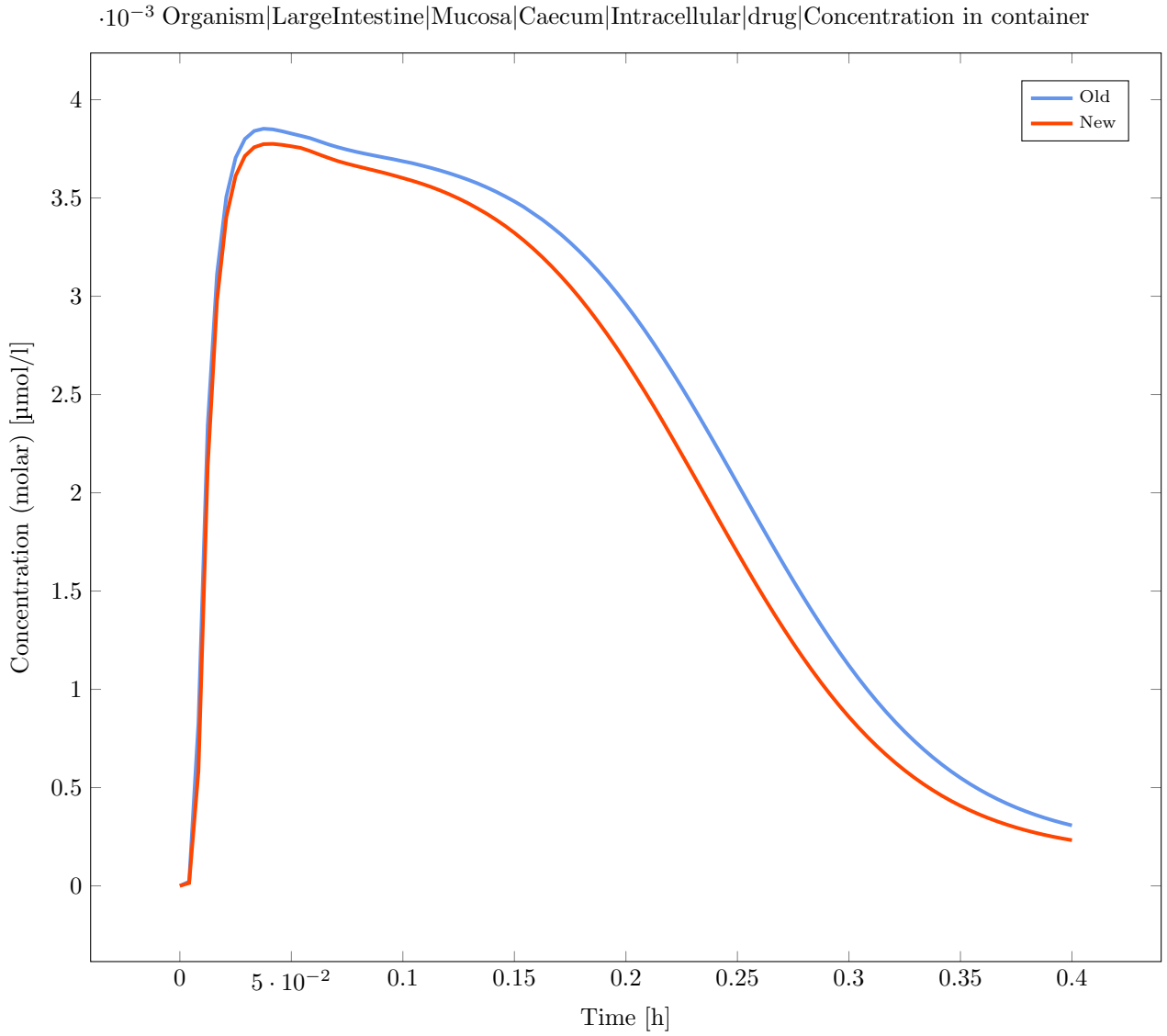

Figure 1.44

Output Path: Organism|SmallIntestine|Mucosa|LowerIleum|Intracellular|drug|Concentration in container
Deviation for 'Organism|SmallIntestine|Mucosa|LowerIleum|Intracellular|drug|Concentration in container' is 32.84% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.33

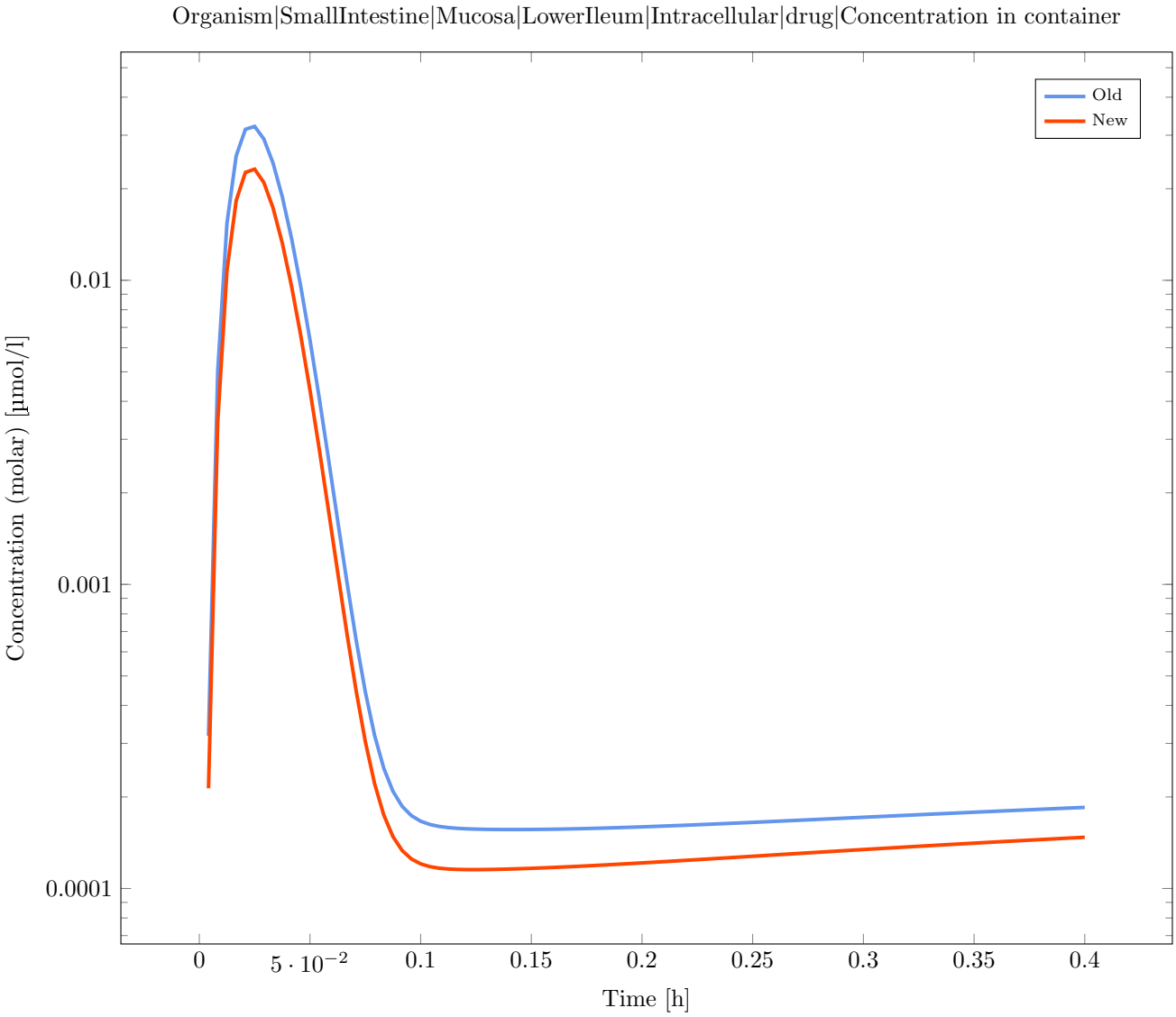
Organism|SmallIntestine|Mucosa|LowerIleum|Intracellular|drug|Concentration in container
Old
New
Concentration (molar) [µmol/l]
0.01
0.001
0.0001
0
$5 \cdot 10^{-2}$
0.1
0.15
0.2
0.25
0.3
0.35
0.4
Time [h]

Figure 1.45

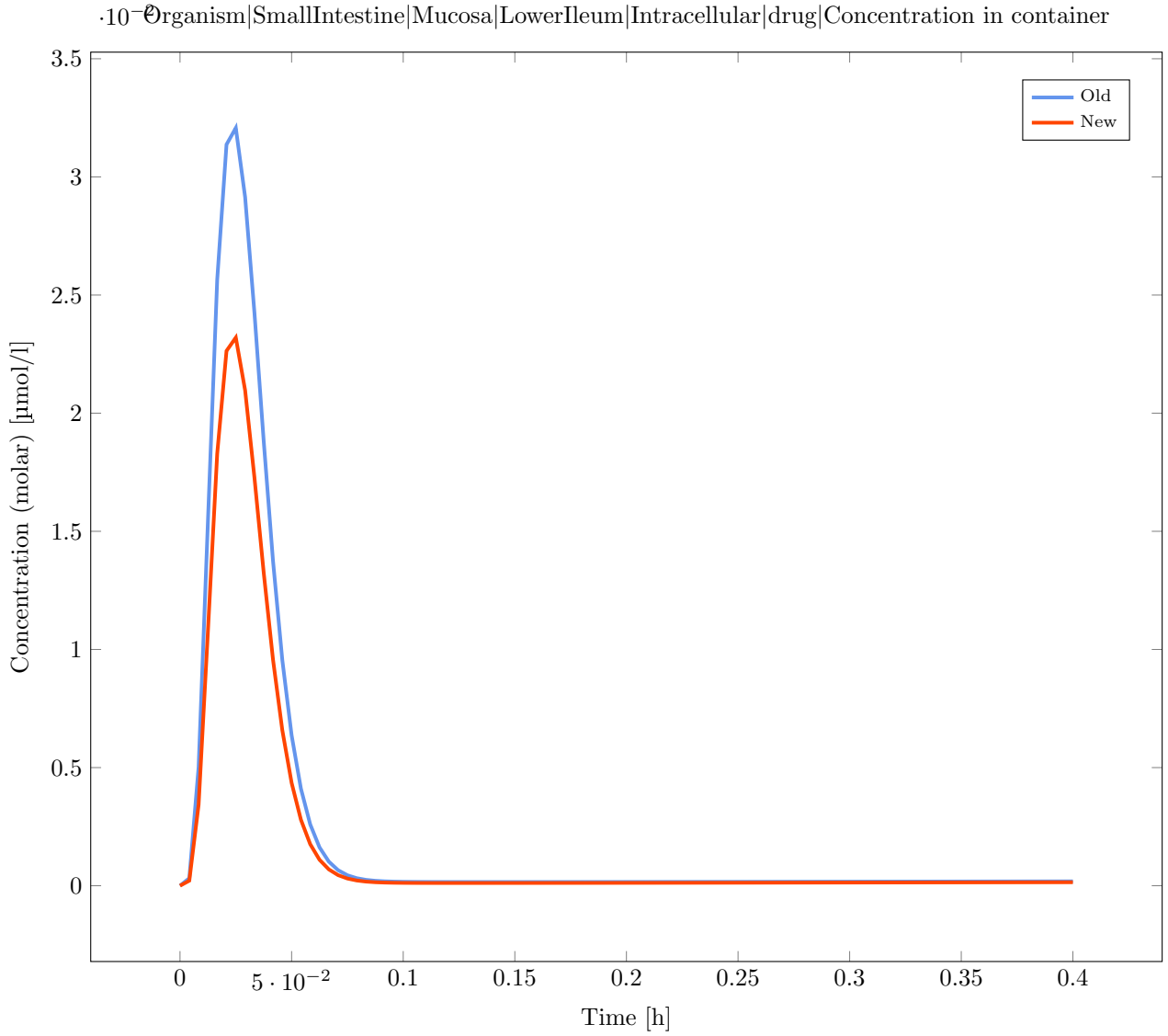

Figure 1.46

Output Path: Organism|SmallIntestine|Mucosa|LowerIleum|Interstitial|drug|Concentration in container
Deviation for 'Organism|SmallIntestine|Mucosa|LowerIleum|Interstitial|drug|Concentration in container' is 32.74% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.33

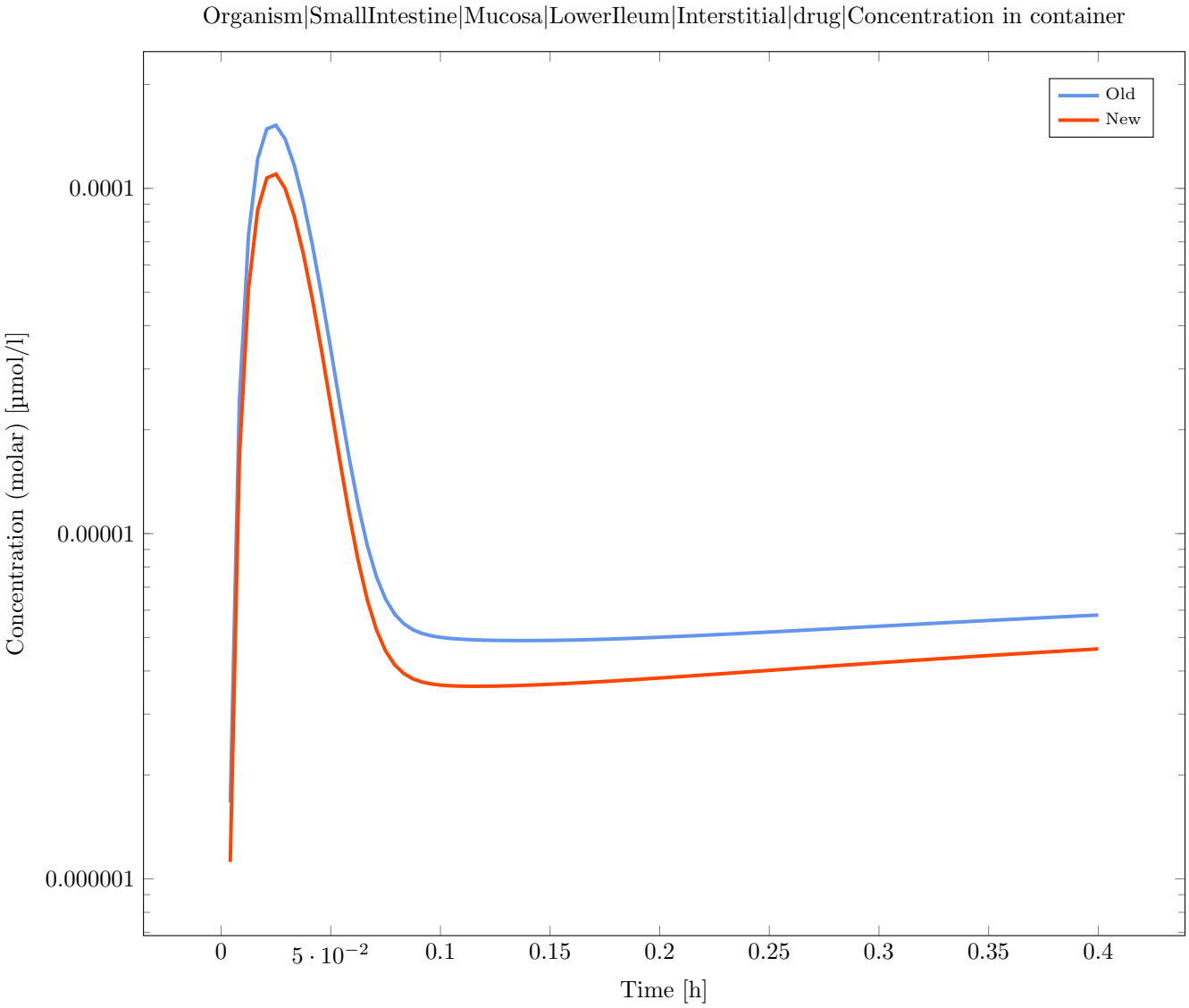
Organism|SmallIntestine|Mucosa|LowerIleum|Interstitial|drug|Concentration in container
Old
New
Concentration (molar) [µmol/l]
0.0001
0.00001
0.000001
0
$5 \cdot 10^{-2}$
0.1
0.15
0.2
0.25
0.3
0.35
0.4
Time [h]

Figure 1.47

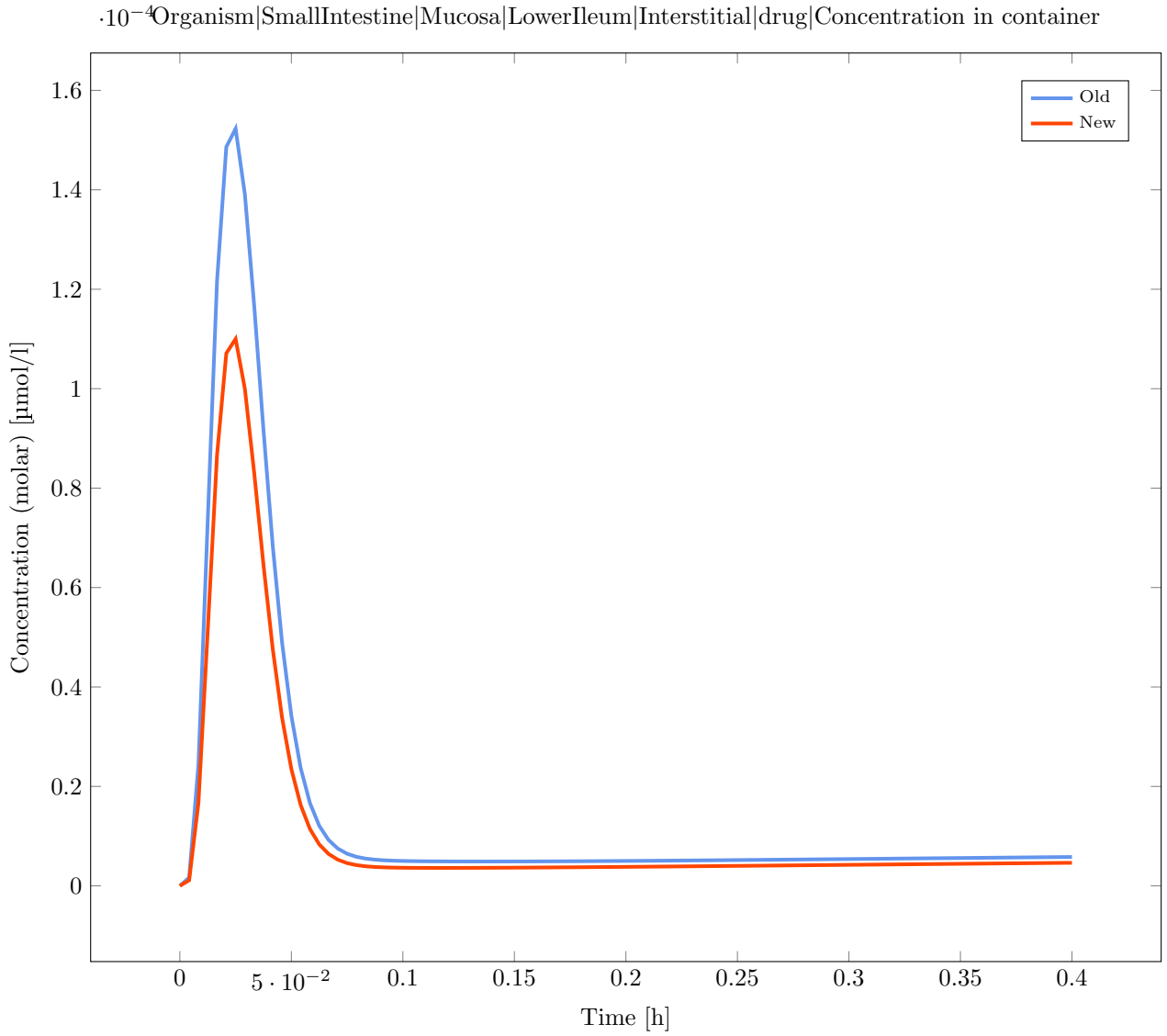

Figure 1.48

Output Path: Organism|SmallIntestine|Mucosa|LowerIleum|Plasma|drug|Concentration in container
Deviation for 'Organism|SmallIntestine|Mucosa|LowerIleum|Plasma|drug|Concentration in container' is 32.74% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.33

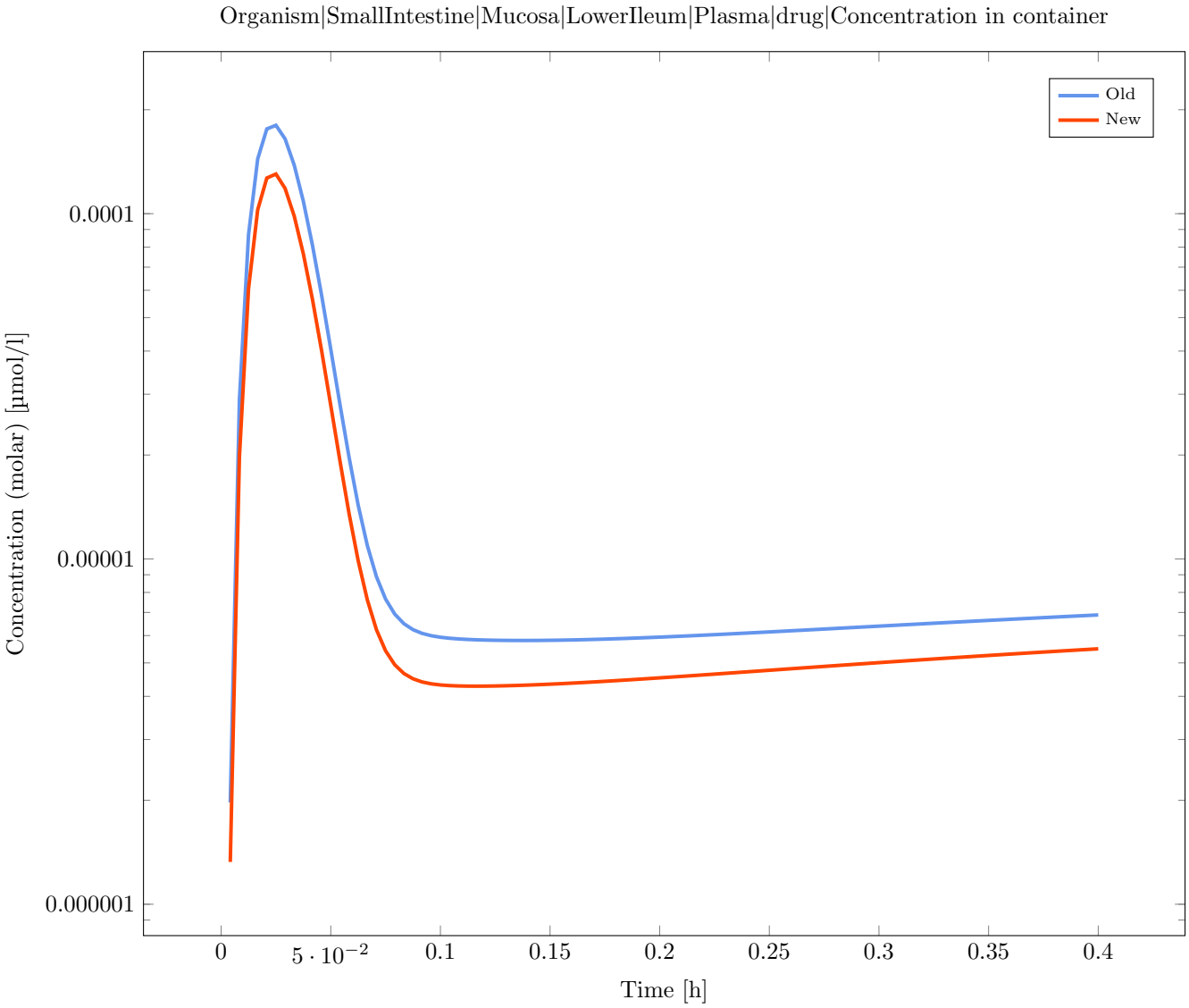
Organism|SmallIntestine|Mucosa|LowerIleum|Plasma|drug|Concentration in container
Old
New
Concentration (molar) [µmol/l]
0.0001
0.00001
0.000001
0
5 · 10−2
0.1
0.15
0.2
0.25
0.3
0.35
0.4
Time [h]

Figure 1.49

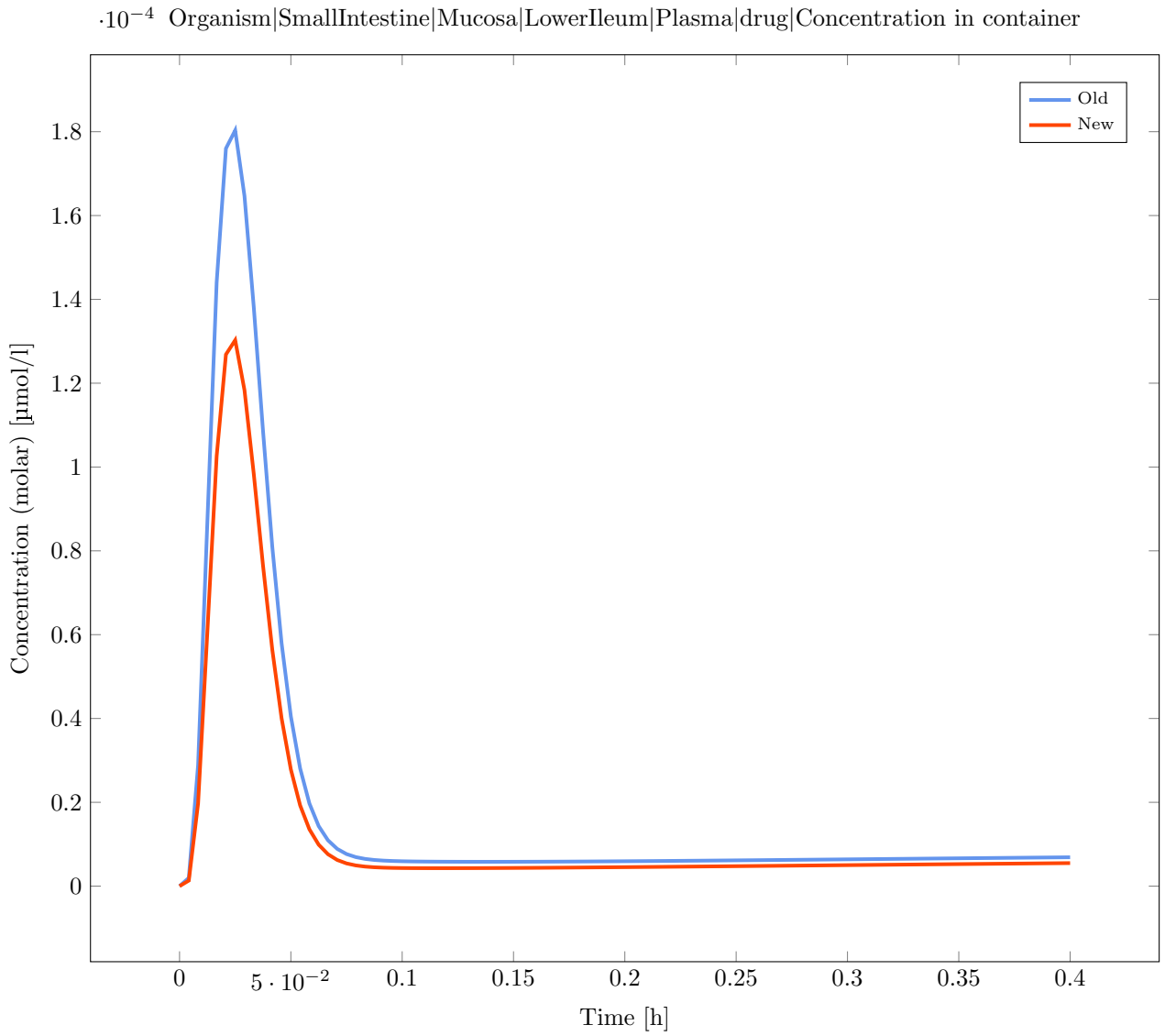

Figure 1.50

Output Path: Organism|Skin|Intracellular|drug|Concentration in container
Deviation for 'Organism|Skin|Intracellular|drug|Concentration in container' is 32.49% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.32

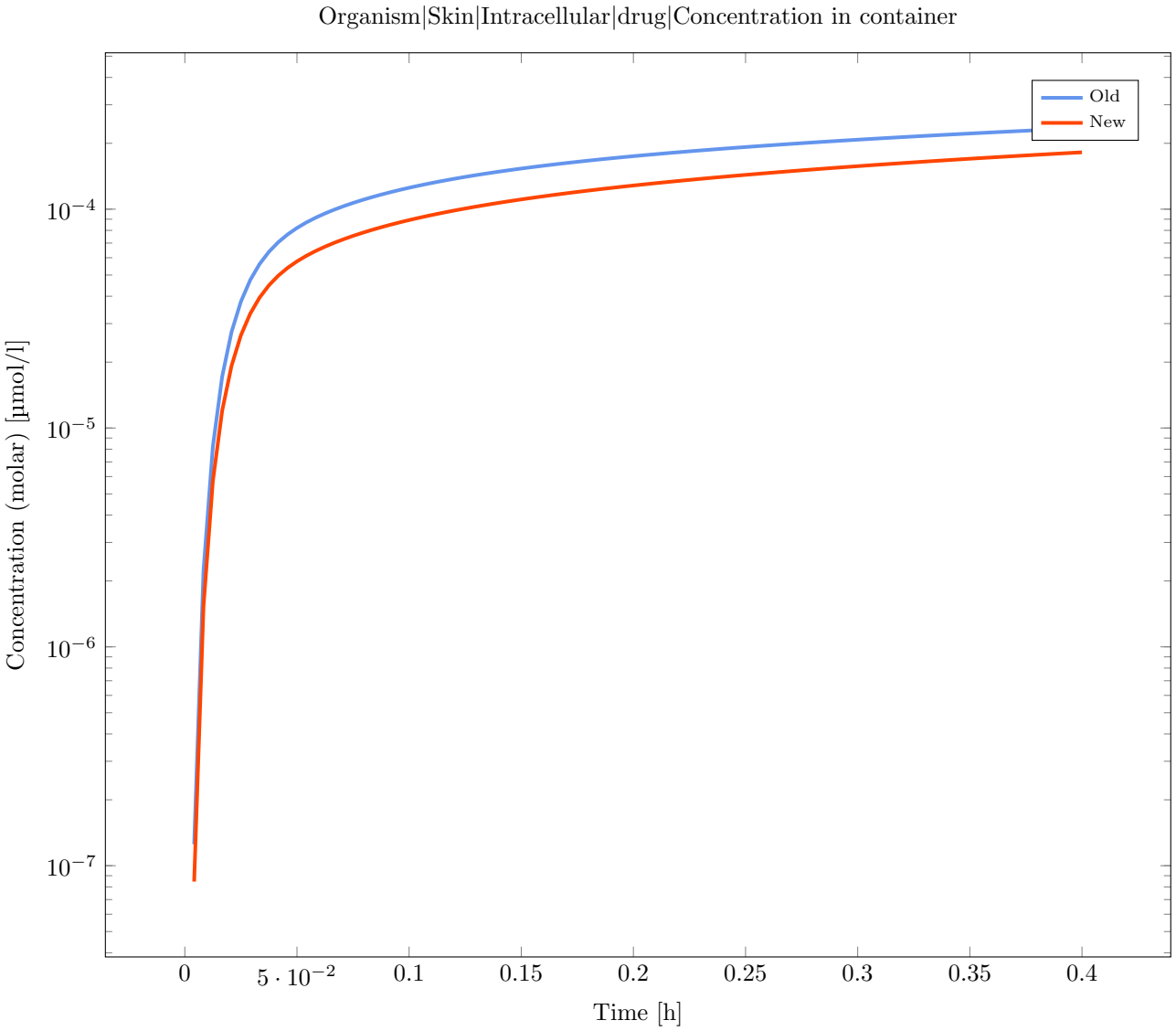
Organism|Skin|Intracellular|drug|Concentration in container
Old
New
Concentration (molar) [µmol/l]
10^{-4}
10^{-5}
10^{-6}
10^{-7}
0
$5 \cdot 10^{-2}$
0.1
0.15
0.2
0.25
0.3
0.35
0.4
Time [h]

Figure 1.51

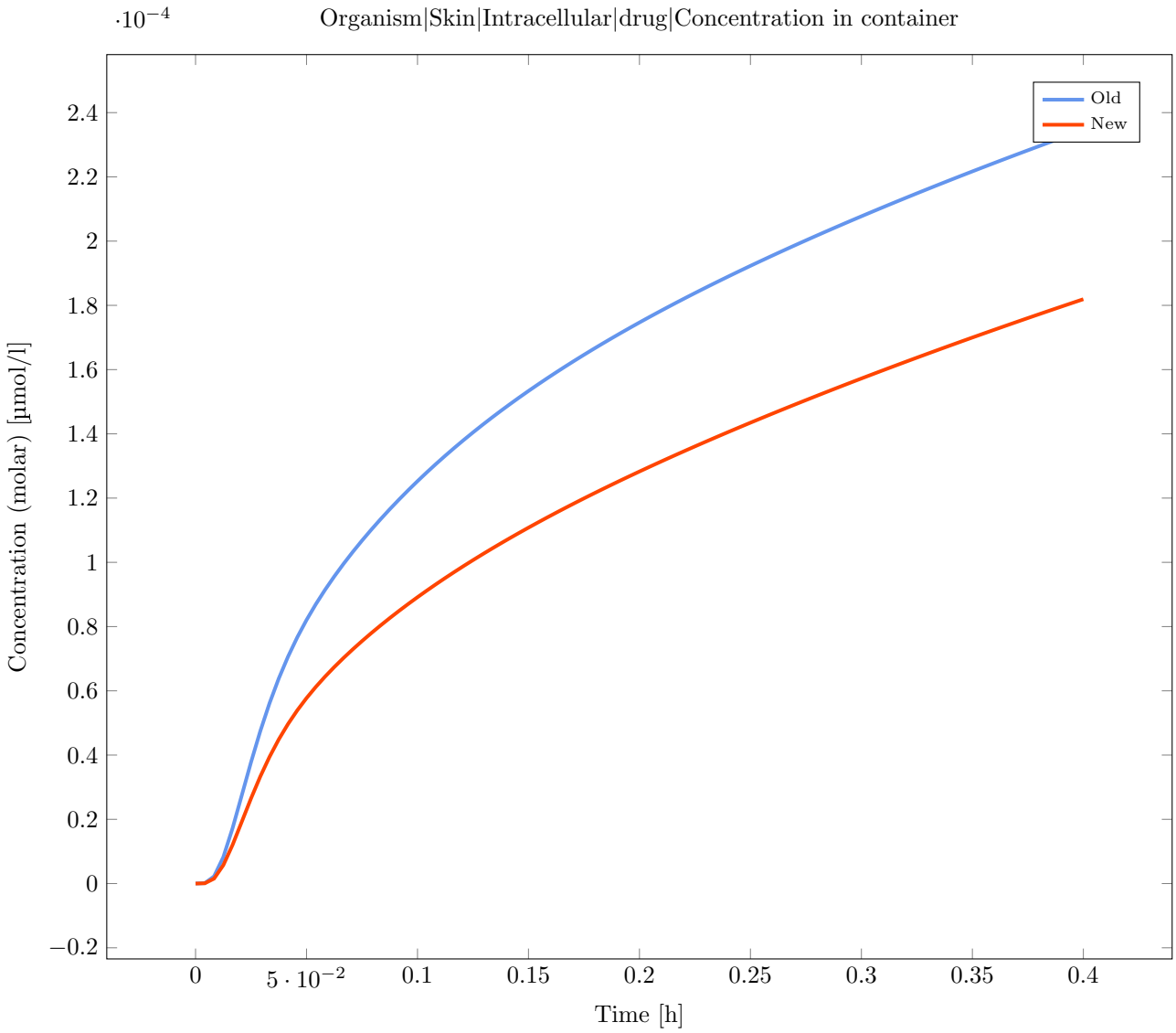

Figure 1.52

Simulation: Human_SingleORAL_PolyParticlesNormal_AsSuspention-Human_SingleORAL_PolyParticlesNo AsSuspention
Result of the validation: Invalid
Absolute Tolerance: 1.00E-10
Relative Tolerance: 1.00E-5

Output Path: Organism|LargeIntestine|Mucosa|Caecum|Intracellular|drug|Concentration in container
Deviation for 'Organism|LargeIntestine|Mucosa|Caecum|Intracellular|drug|Concentration in container' is 32.96% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.33

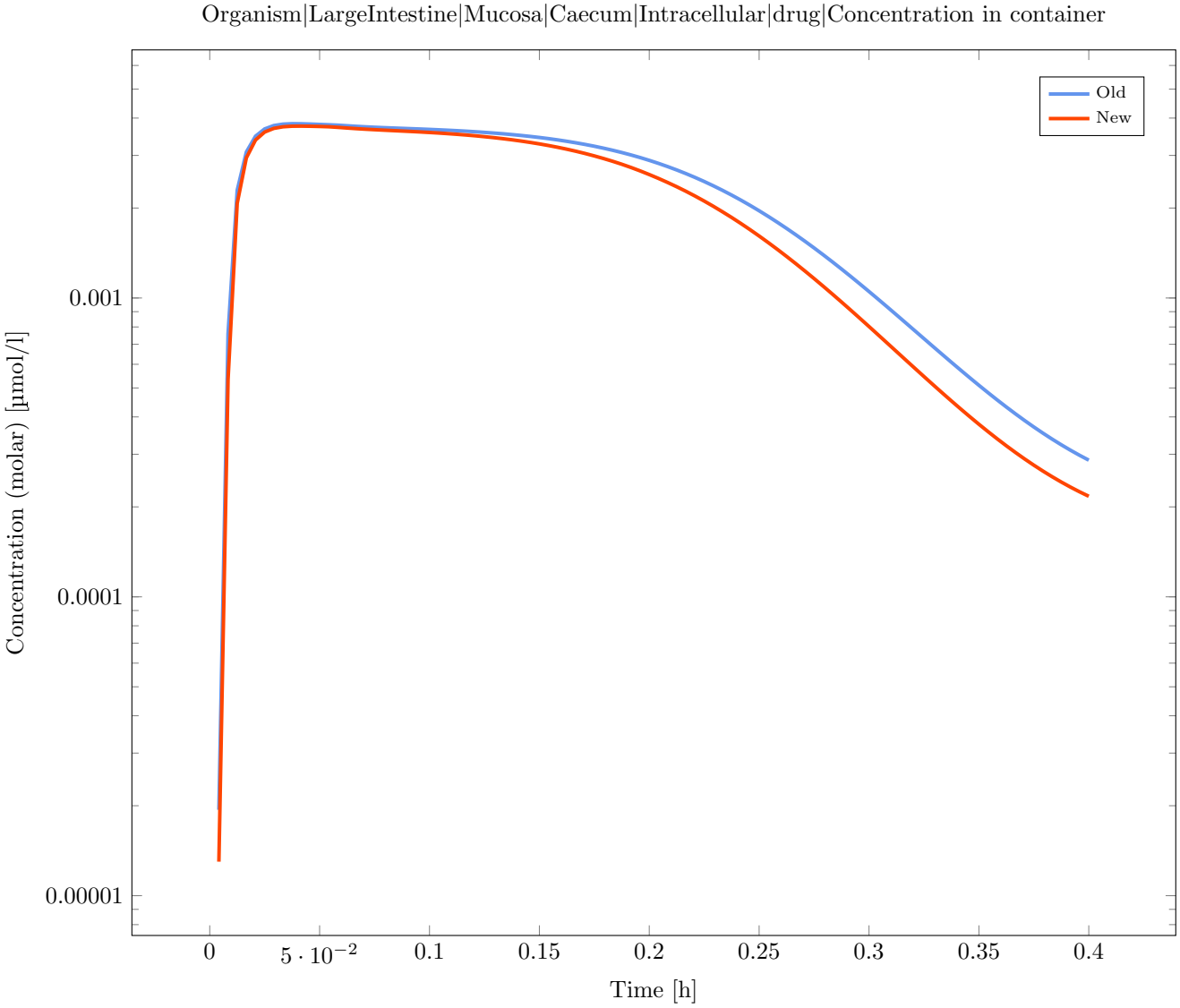
Organism|LargeIntestine|Mucosa|Caecum|Intracellular|drug|Concentration in container
Old
New
Concentration (molar) [µmol/l]
0.001
0.0001
0.00001
0
5 · 10−2
0.1
0.15
0.2
0.25
0.3
0.35
0.4
Time [h]

Figure 1.53

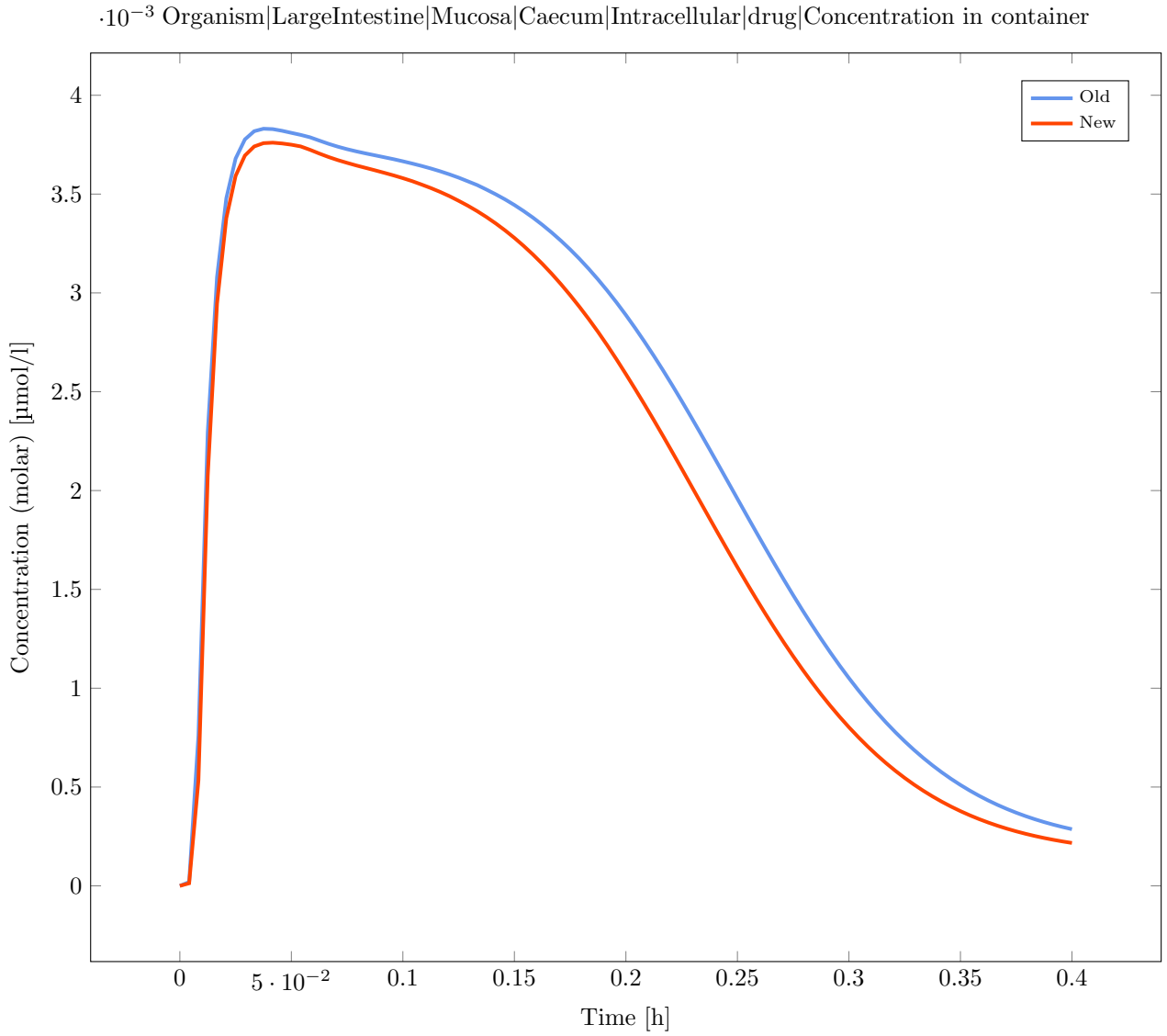

Figure 1.54

Output Path: Organism|SmallIntestine|Mucosa|LowerIleum|Intracellular|drug|Concentration in container
Deviation for 'Organism|SmallIntestine|Mucosa|LowerIleum|Intracellular|drug|Concentration in container' is 32.88% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.33

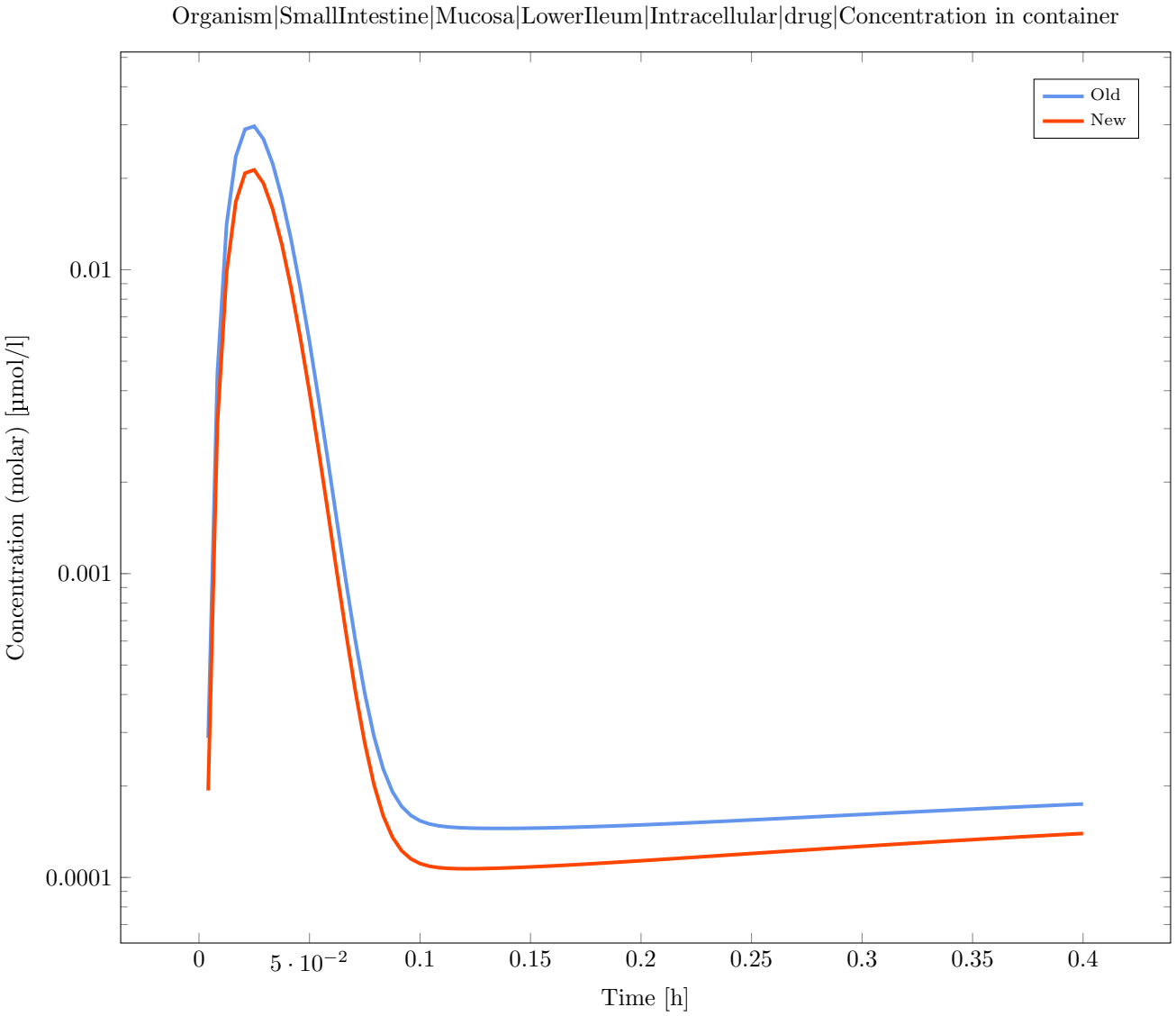
Organism|SmallIntestine|Mucosa|LowerIleum|Intracellular|drug|Concentration in container
Old
New
Concentration (molar) [µmol/l]
0.01
0.001
0.0001
0
$5 \cdot 10^{-2}$
0.1
0.15
0.2
0.25
0.3
0.35
0.4
Time [h]

Figure 1.55

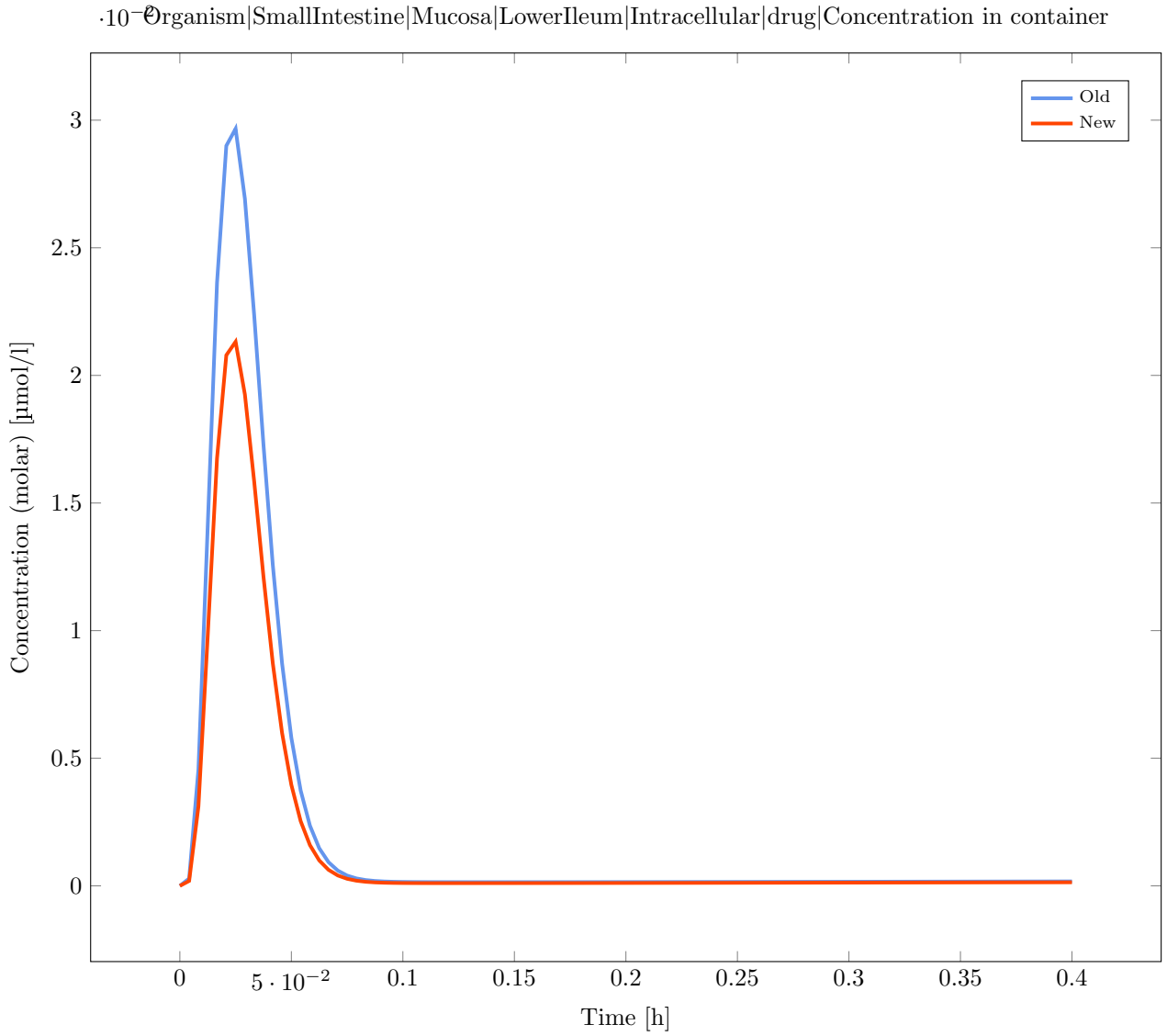

Figure 1.56

Output Path: Organism|SmallIntestine|Mucosa|LowerIleum|Interstitial|drug|Concentration in container
Deviation for 'Organism|SmallIntestine|Mucosa|LowerIleum|Interstitial|drug|Concentration in container' is 32.80% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.33

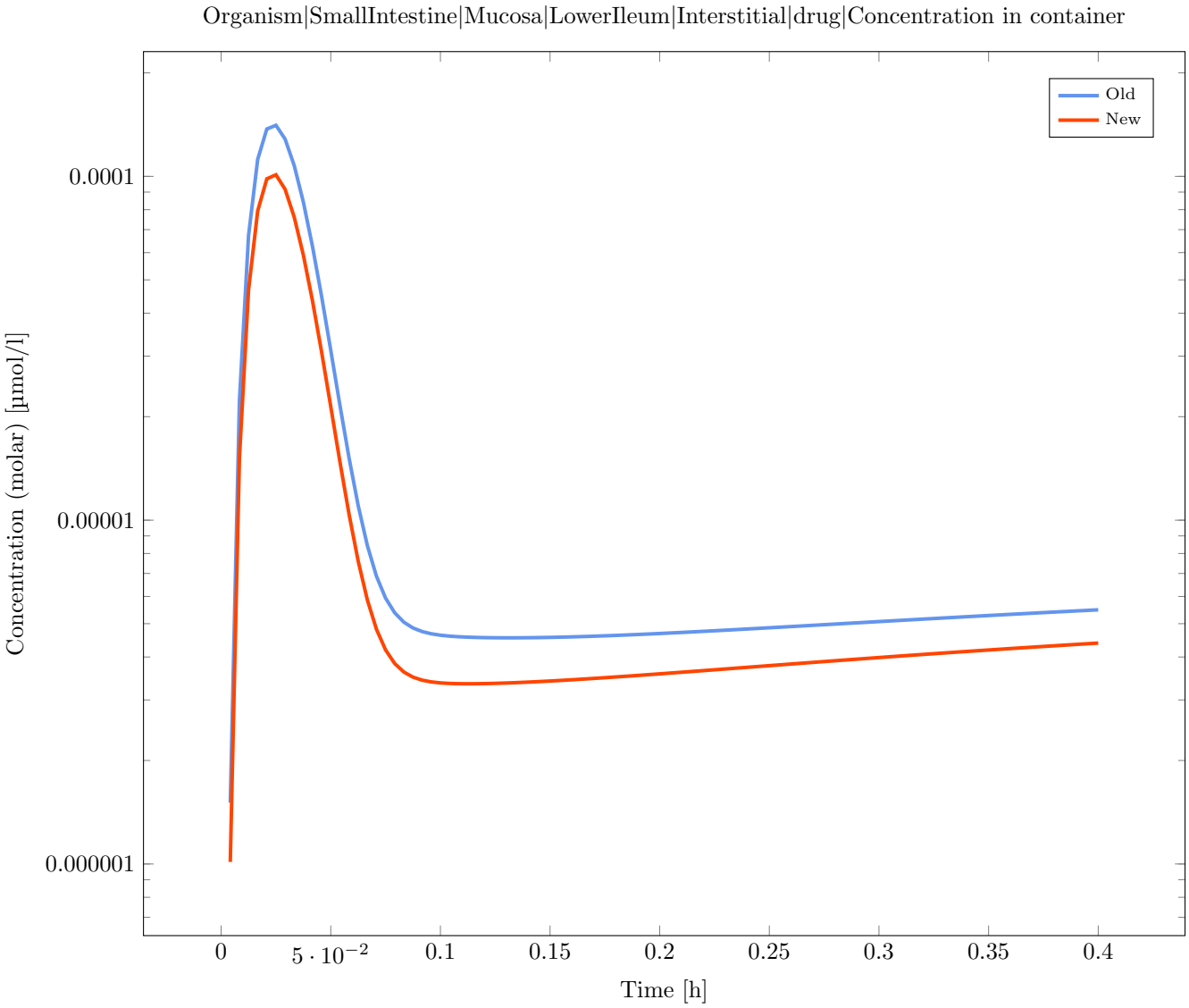
Organism|SmallIntestine|Mucosa|LowerIleum|Interstitial|drug|Concentration in container
Old
New
Concentration (molar) [µmol/l]
0.0001
0.00001
0.000001
0
$5 \cdot 10^{-2}$
0.1
0.15
0.2
0.25
0.3
0.35
0.4
Time [h]

Figure 1.57

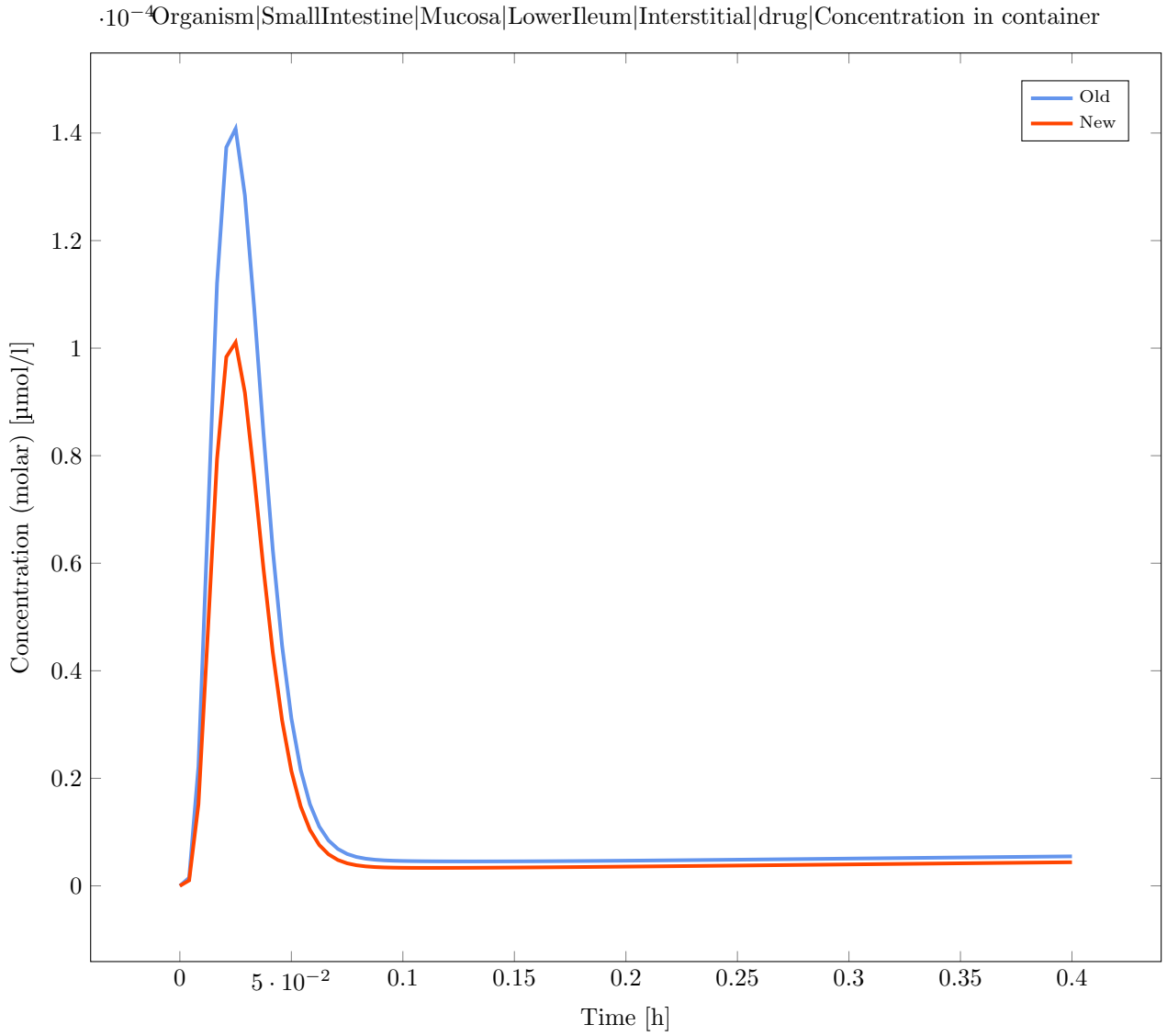

Figure 1.58

Output Path: Organism|SmallIntestine|Mucosa|LowerIleum|Plasma|drug|Concentration in container
Deviation for 'Organism|SmallIntestine|Mucosa|LowerIleum|Plasma|drug|Concentration in container' is 32.80% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.33

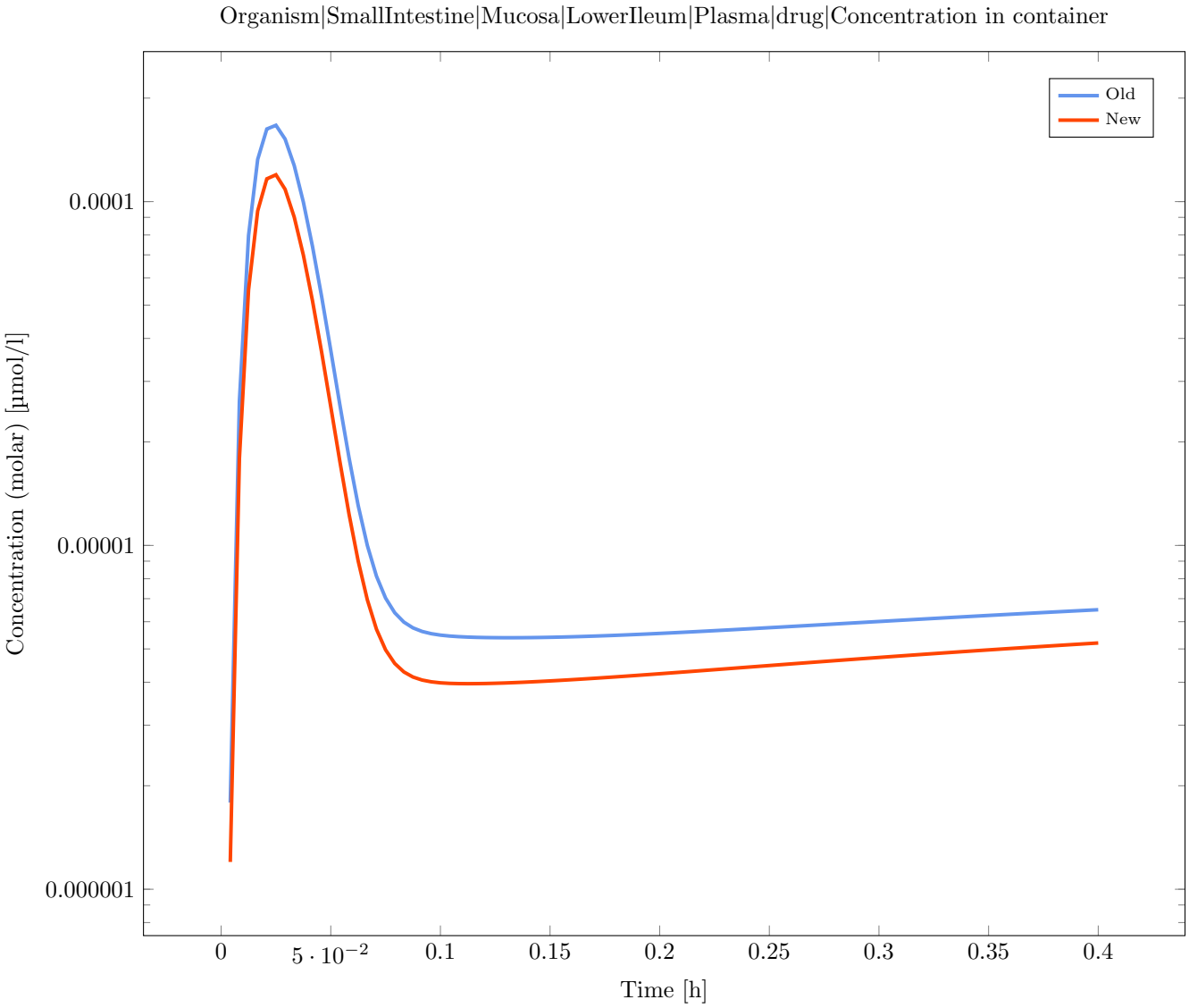
Organism|SmallIntestine|Mucosa|LowerIleum|Plasma|drug|Concentration in container
Old
New
Concentration (molar) [µmol/l]
0.0001
0.00001
0.000001
0
$5 \cdot 10^{-2}$
0.1
0.15
0.2
0.25
0.3
0.35
0.4
Time [h]

Figure 1.59

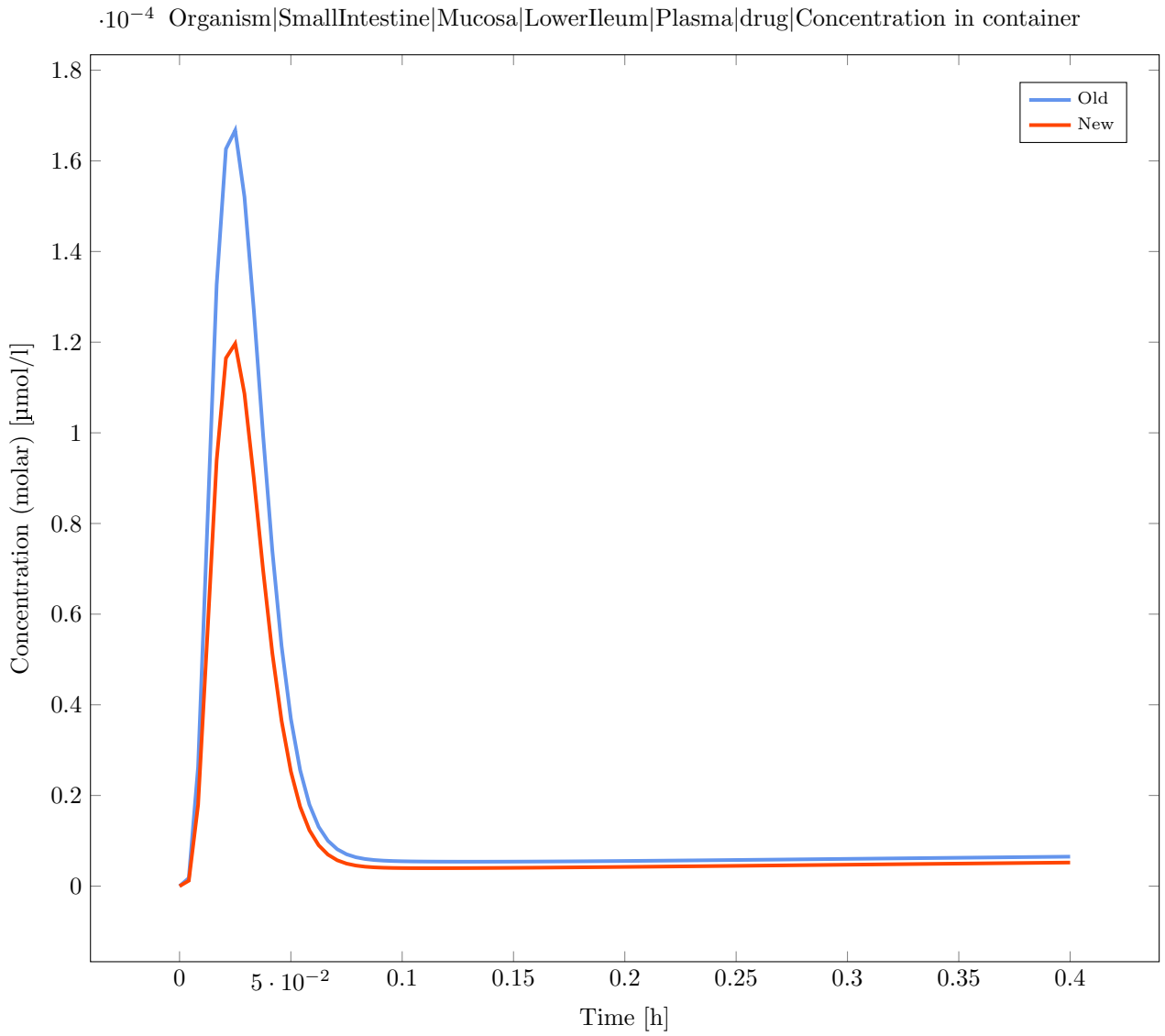

Figure 1.60

Output Path: Organism|Skin|Intracellular|drug|Concentration in container
Deviation for 'Organism|Skin|Intracellular|drug|Concentration in container' is 32.57% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.33

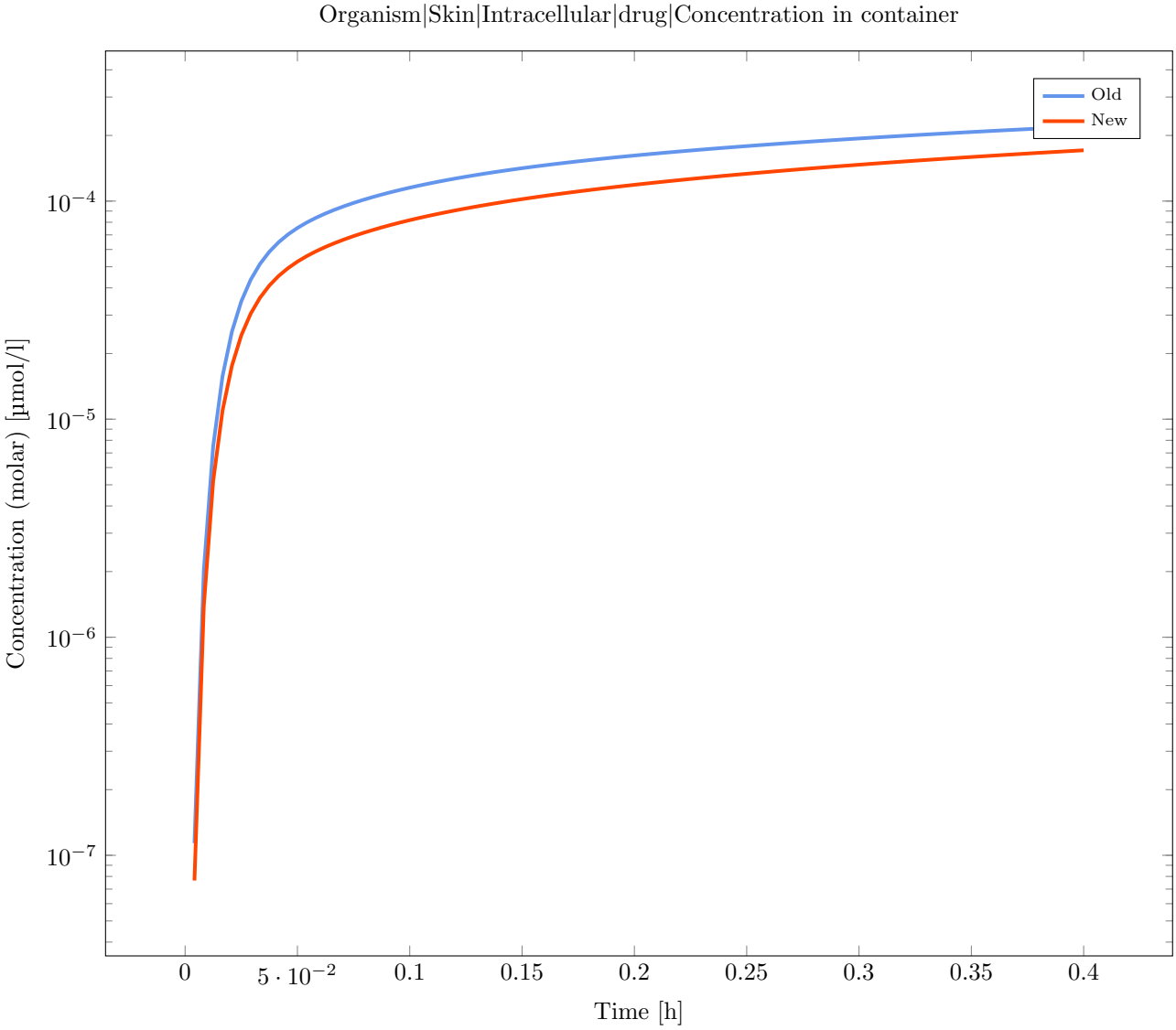
Organism|Skin|Intracellular|drug|Concentration in container
Old
New
Concentration (molar) [µmol/l]
10^{-4}
10^{-5}
10^{-6}
10^{-7}
0
$5 \cdot 10^{-2}$
0.1
0.15
0.2
0.25
0.3
0.35
0.4
Time [h]

Figure 1.61

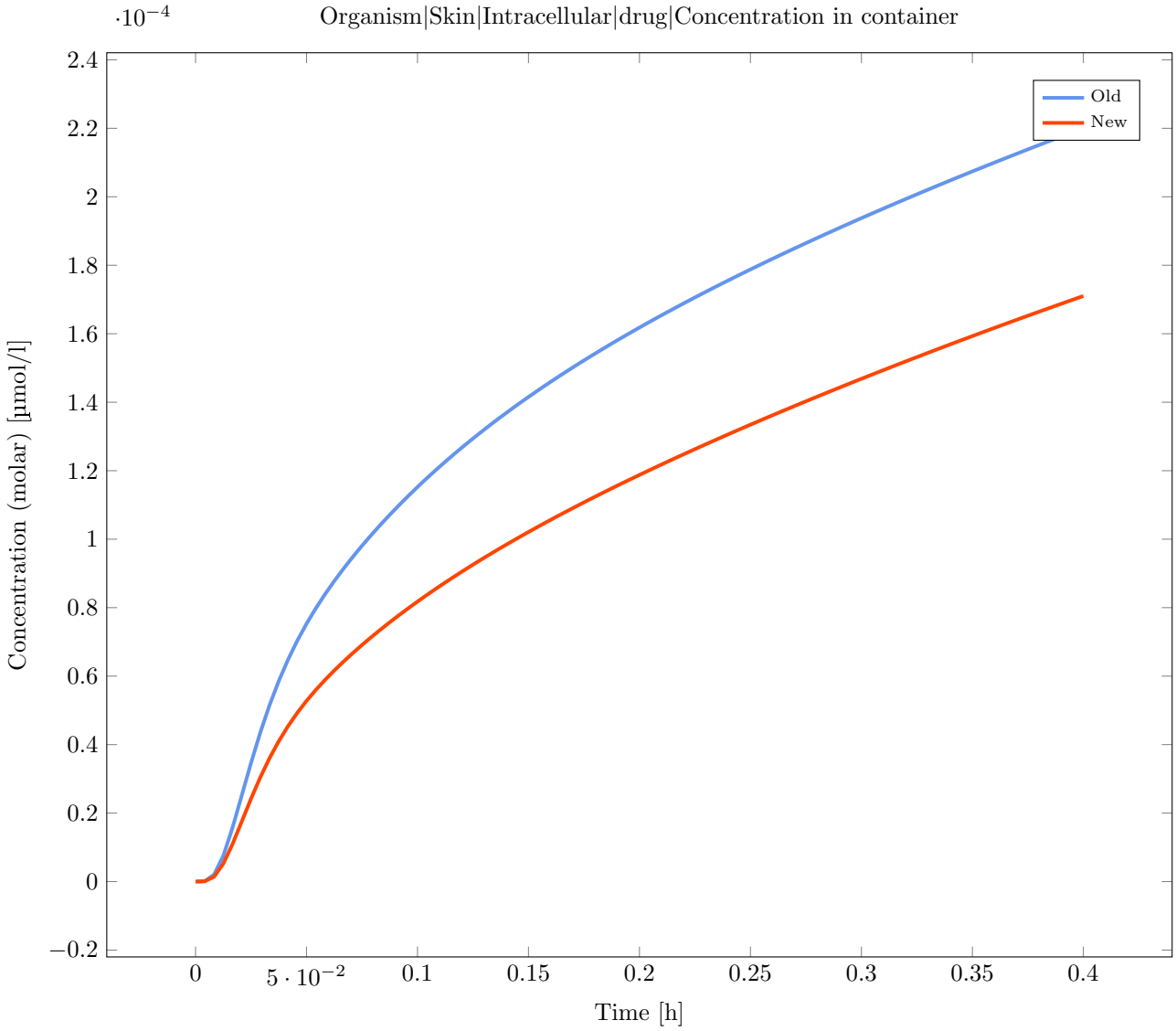

Figure 1.62

Simulation: Human_SingleORAL_PolyParticlesNormal_AsSuspention-Human_SingleORAL_PolyParticlesNo AsSuspention_dissolved_radius
Result of the validation: Invalid
Absolute Tolerance: 1.00E-10
Relative Tolerance: 1.00E-5

Output Path: Organism|SmallIntestine|Mucosa|LowerIleum|Intracellular|drug|Concentration in container
Deviation for 'Organism|SmallIntestine|Mucosa|LowerIleum|Intracellular|drug|Concentration in container' is 29.81% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.30

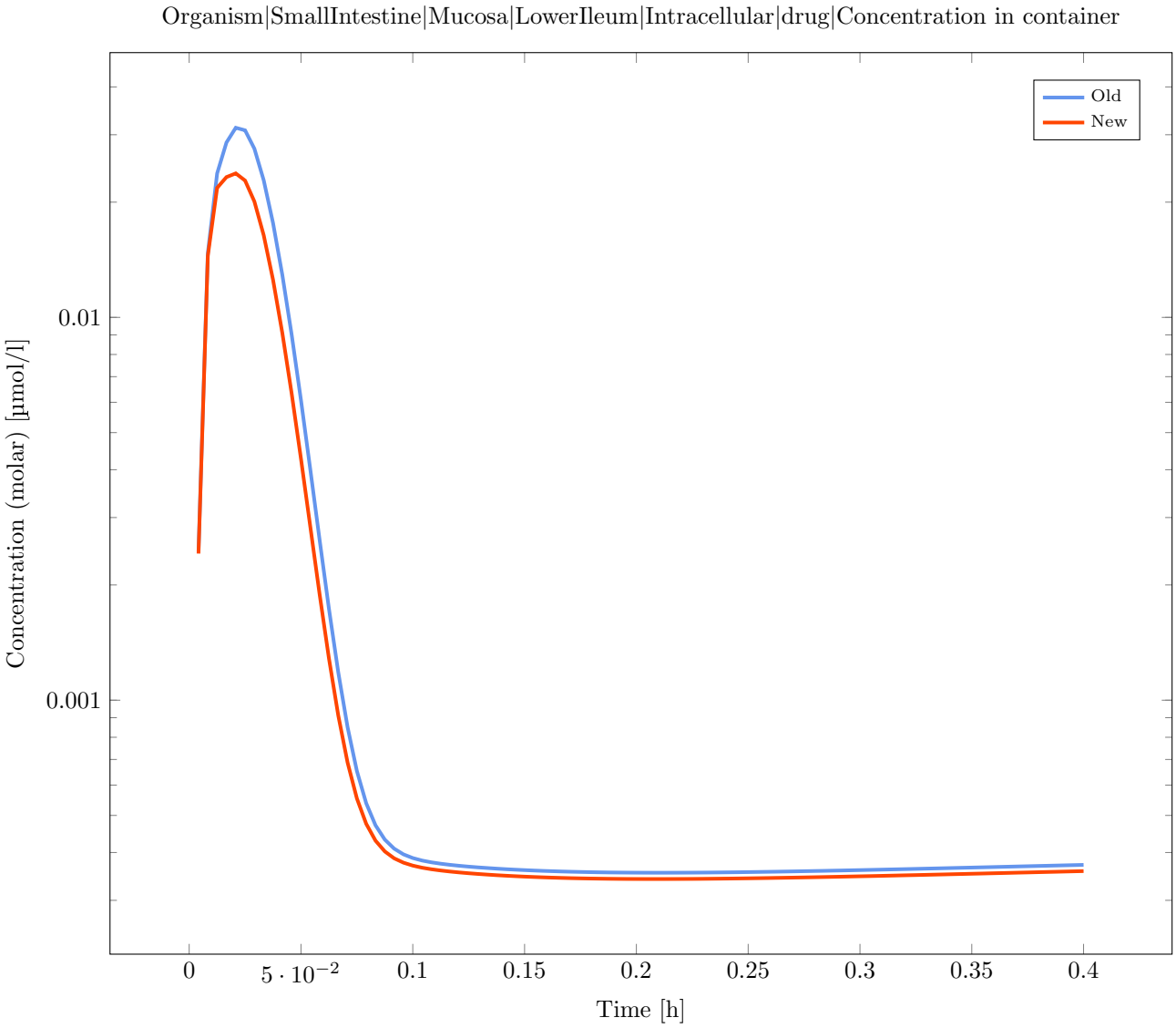
Organism|SmallIntestine|Mucosa|LowerIleum|Intracellular|drug|Concentration in container
Old
New
Concentration (molar) [µmol/l]
0.01
0.001
0
$5 \cdot 10^{-2}$
0.1
0.15
0.2
0.25
0.3
0.35
0.4
Time [h]

Figure 1.63

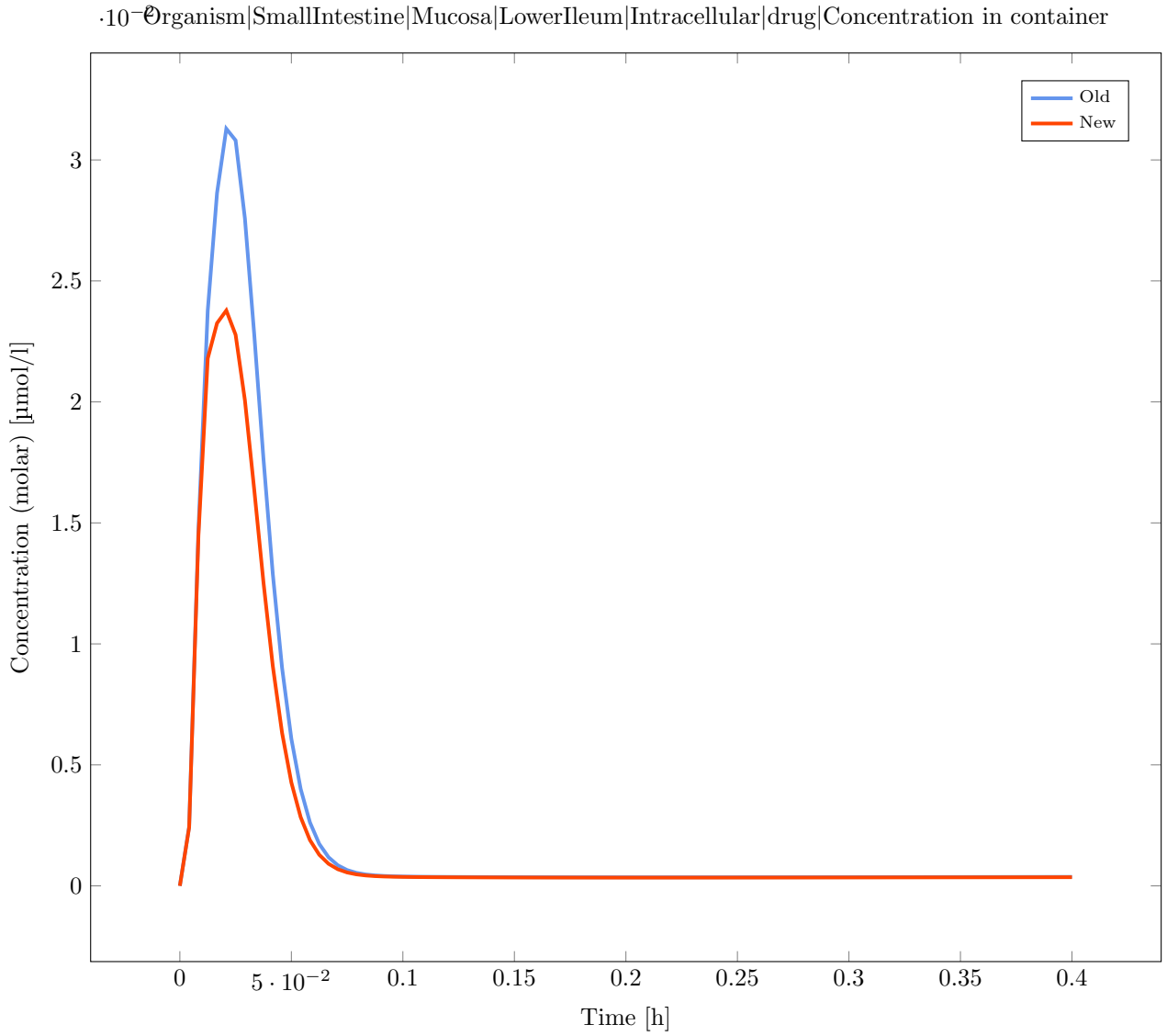

Figure 1.64

Output Path: Organism|Lumen|Feces|drug|Fraction excreted to feces
Deviation for 'Organism|Lumen|Feces|drug|Fraction excreted to feces' is 29.80% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.30

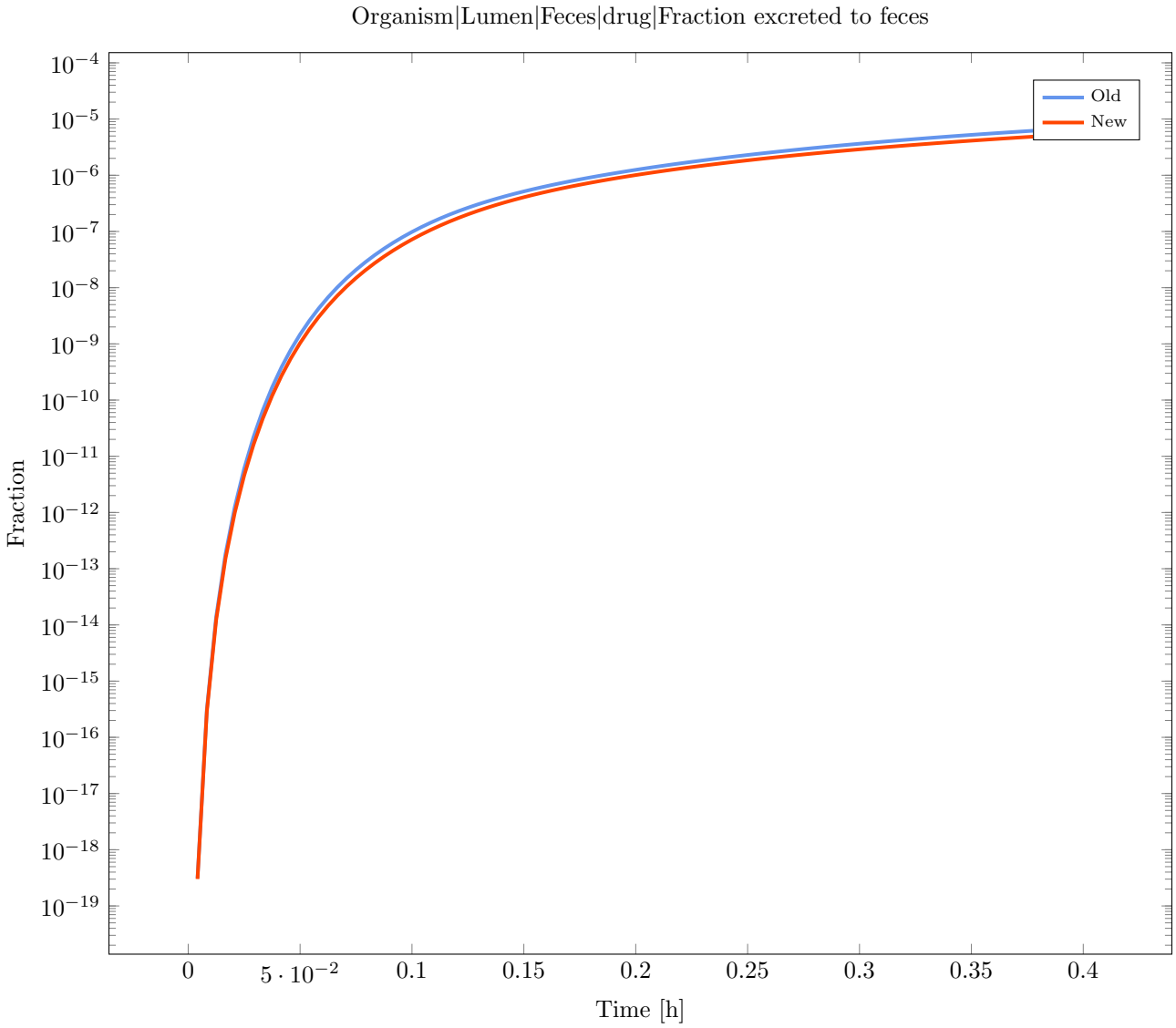
Organism|Lumen|Feces|drug|Fraction excreted to feces
Old
New
Fraction
10^{-4}
10^{-5}
10^{-6}
10^{-7}
10^{-8}
10^{-9}
10^{-10}
10^{-11}
10^{-12}
10^{-13}
10^{-14}
10^{-15}
10^{-16}
10^{-17}
10^{-18}
10^{-19}
0
$5 \cdot 10^{-2}$
0.1
0.15
0.2
0.25
0.3
0.35
0.4
Time [h]

Figure 1.65

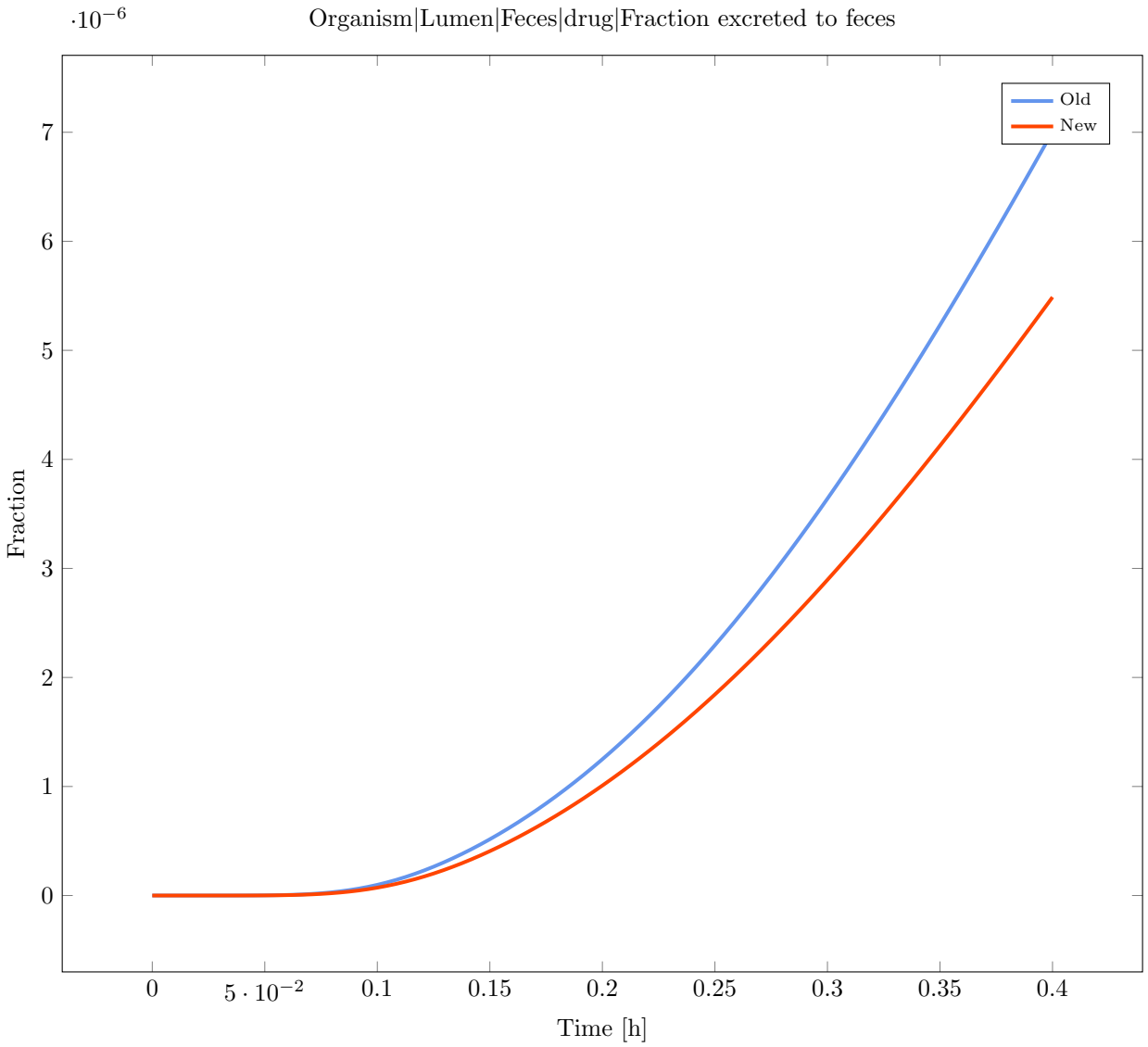

Figure 1.66

Output Path: Organism|Lumen|Feces|drug
Deviation for 'Organism|Lumen|Feces|drug' is 29.80% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.30

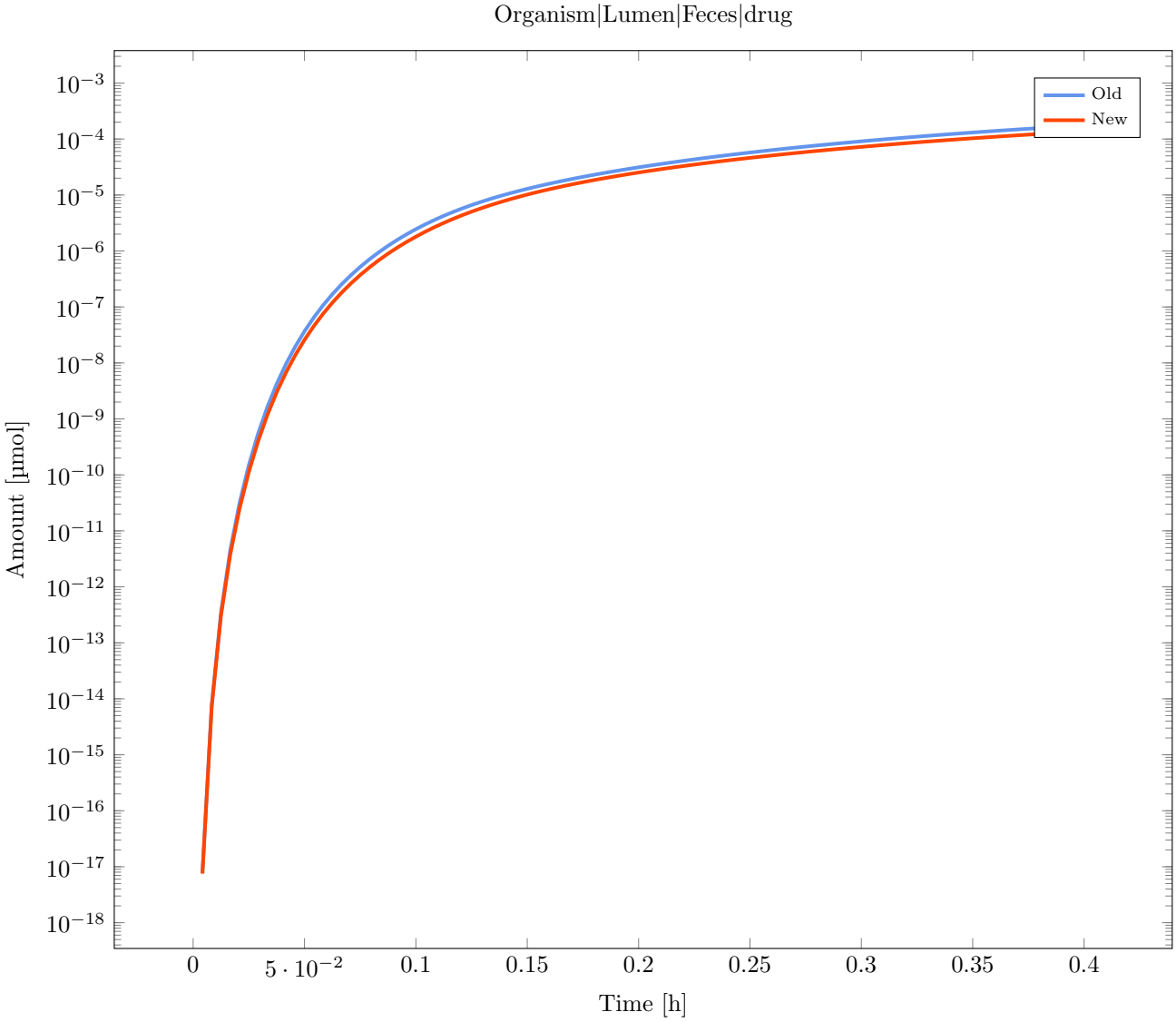
Organism|Lumen|Feces|drug
Old
New
Amount [µmol]
10^{-3}
10^{-4}
10^{-5}
10^{-6}
10^{-7}
10^{-8}
10^{-9}
10^{-10}
10^{-11}
10^{-12}
10^{-13}
10^{-14}
10^{-15}
10^{-16}
10^{-17}
10^{-18}
0
$5 \cdot 10^{-2}$
0.1
0.15
0.2
0.25
0.3
0.35
0.4
Time [h]

Figure 1.67

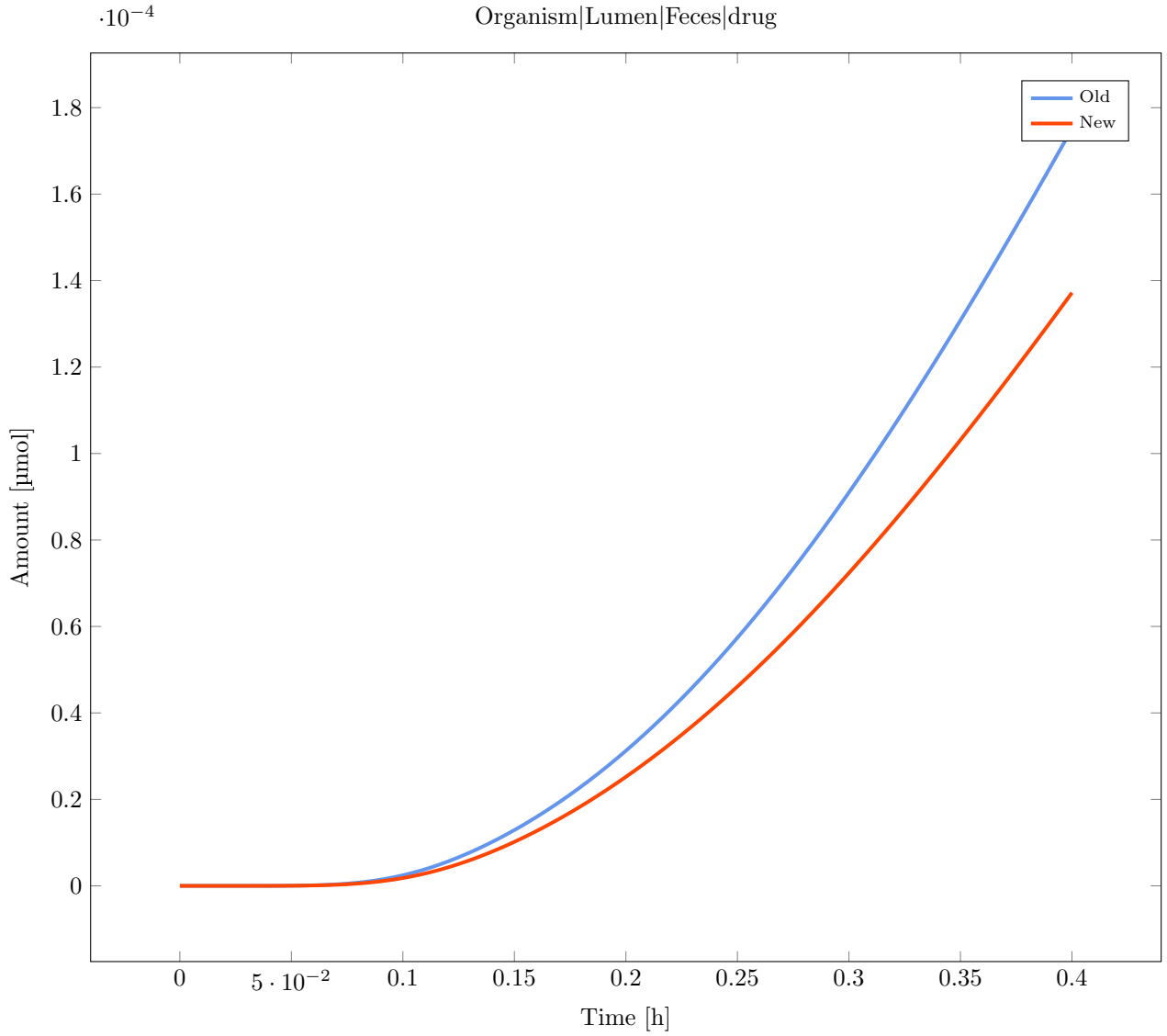

Figure 1.68

Output Path: Organism|Lumen|Feces|drug|Concentration in feces
Deviation for 'Organism|Lumen|Feces|drug|Concentration in feces' is 29.80% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.30

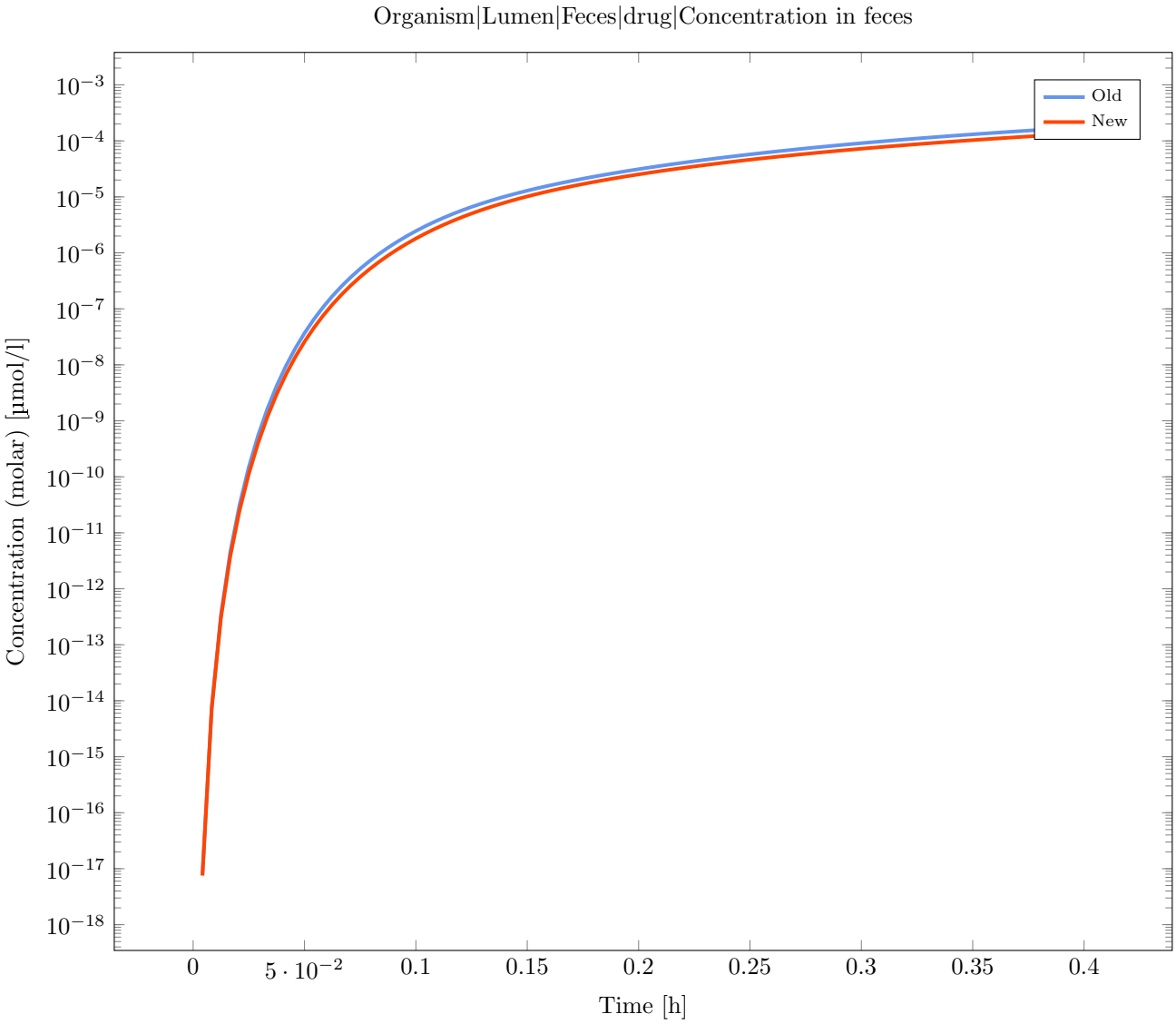
Organism|Lumen|Feces|drug|Concentration in feces
Old
New
Concentration (molar) [µmol/l]
10^{-3}
10^{-4}
10^{-5}
10^{-6}
10^{-7}
10^{-8}
10^{-9}
10^{-10}
10^{-11}
10^{-12}
10^{-13}
10^{-14}
10^{-15}
10^{-16}
10^{-17}
10^{-18}
0
$5 \cdot 10^{-2}$
0.1
0.15
0.2
0.25
0.3
0.35
0.4
Time [h]

Figure 1.69

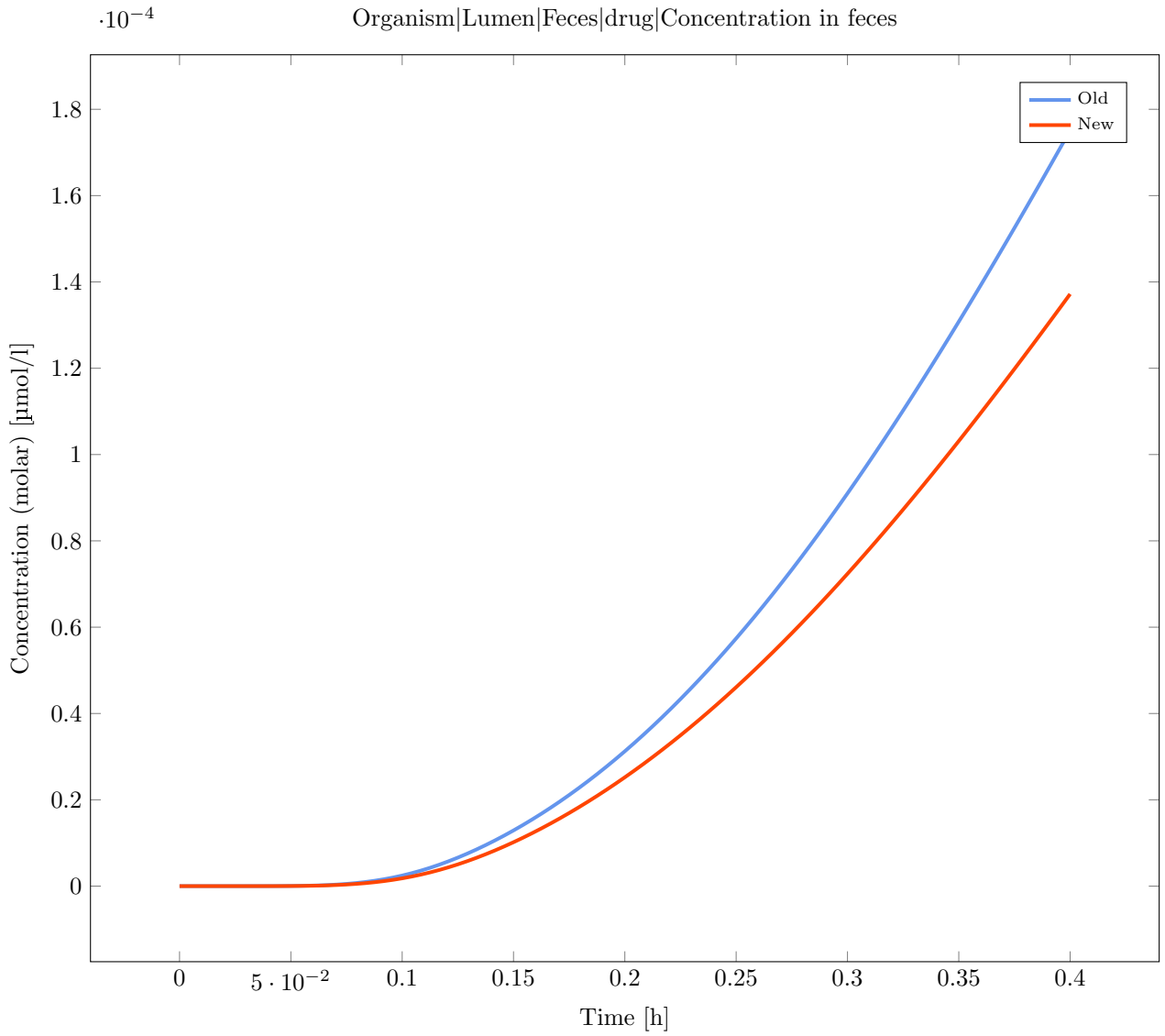

Figure 1.70

Output Path: Organism|SmallIntestine|Mucosa|UpperIleum|Intracellular|drug|Concentration in container
Deviation for 'Organism|SmallIntestine|Mucosa|UpperIleum|Intracellular|drug|Concentration in container' is 28.21% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.28

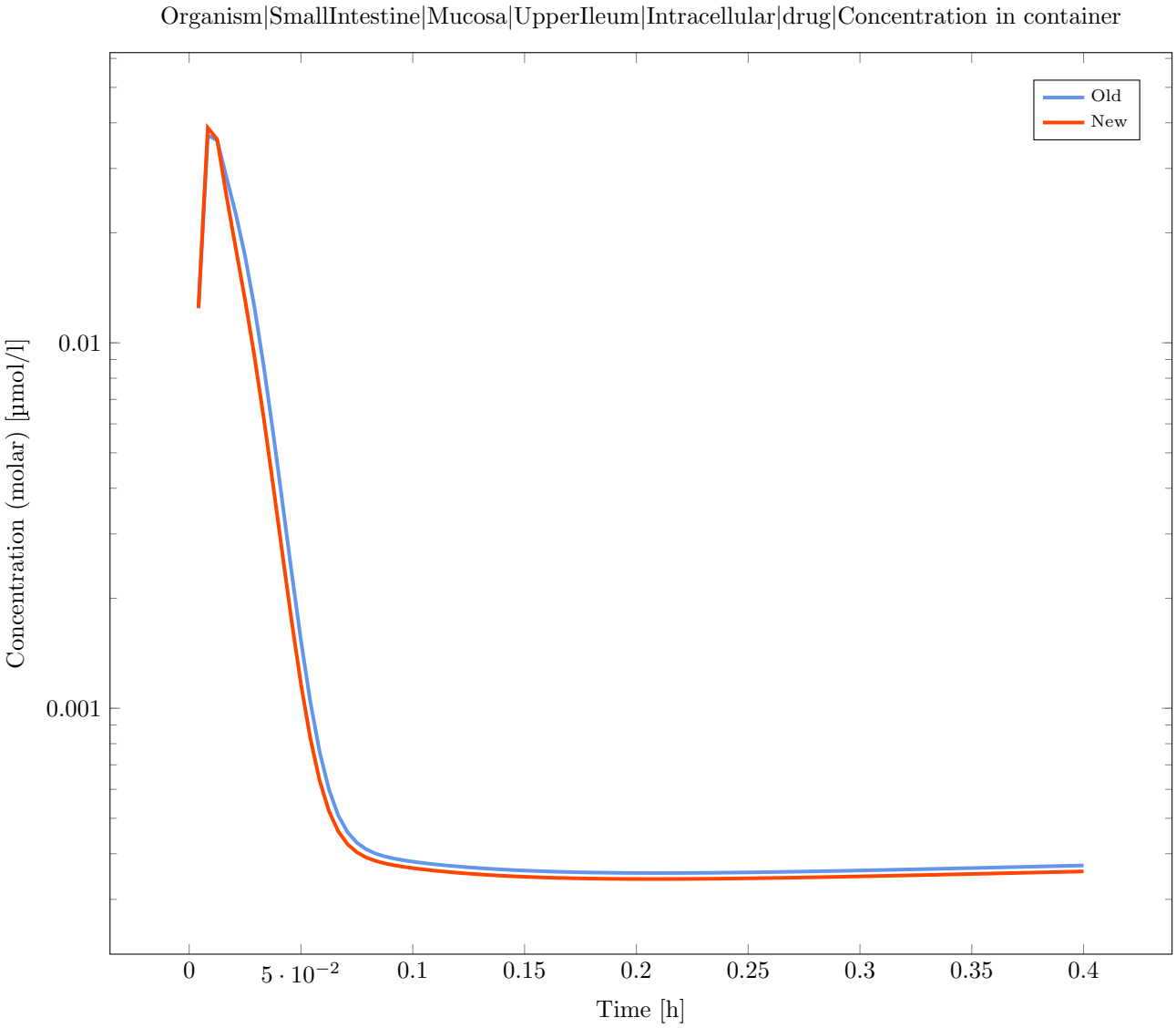
Organism|SmallIntestine|Mucosa|UpperIleum|Intracellular|drug|Concentration in container
Old
New
Concentration (molar) [µmol/l]
0.01
0.001
0
$5 \cdot 10^{-2}$
0.1
0.15
0.2
0.25
0.3
0.35
0.4
Time [h]

Figure 1.71

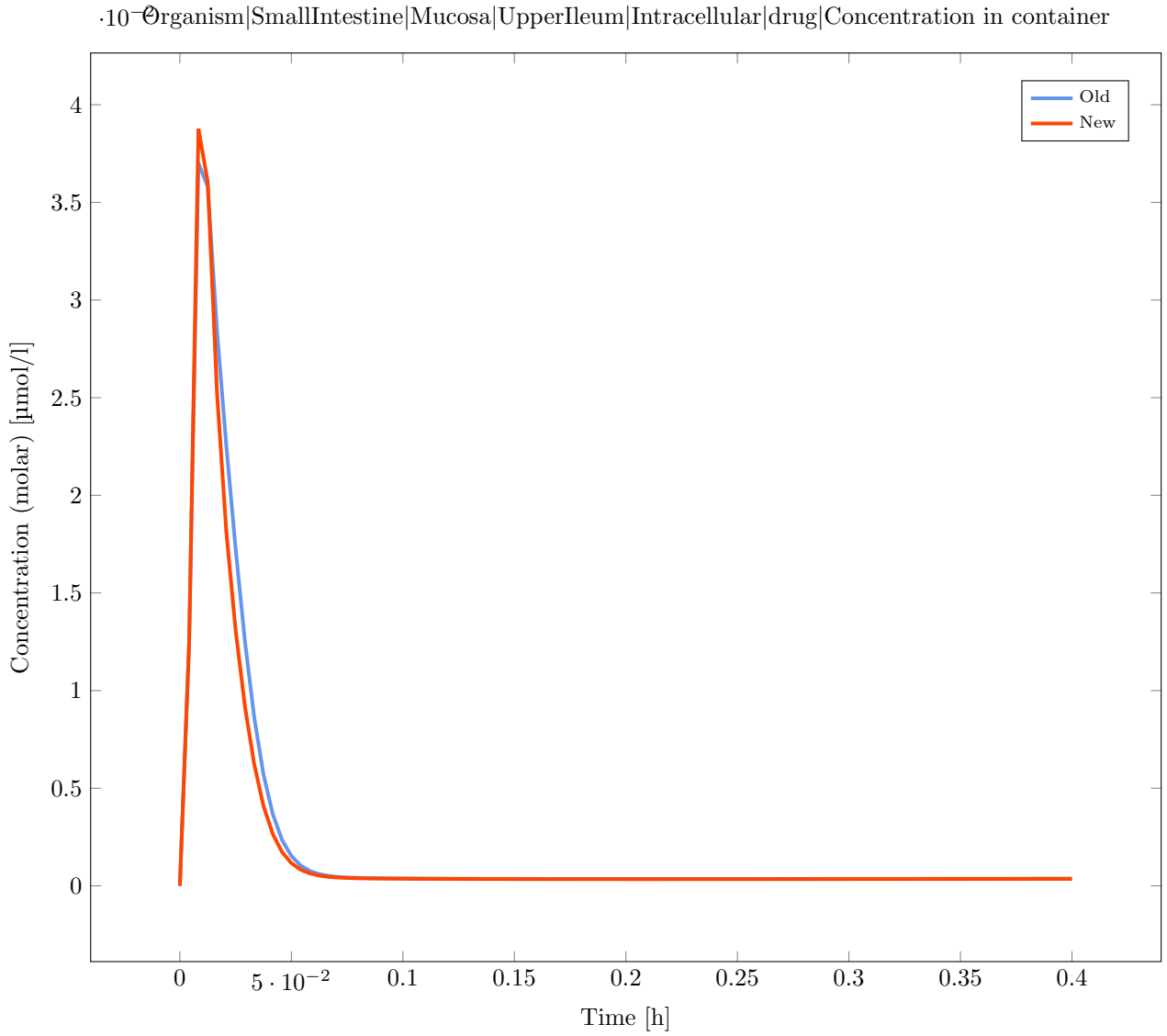

Figure 1.72

Simulation: Human_SingleORAL_PolyParticlesNormal_AsSuspention-Human_SingleORAL_PolyParticlesNo AsSuspention_treat_precipated_drug_as_soluble
Result of the validation: Invalid
Absolute Tolerance: 1.00E-10
Relative Tolerance: 1.00E-5

Output Path: Organism|LargeIntestine|Mucosa|Caecum|Intracellular|drug|Concentration in container
Deviation for 'Organism|LargeIntestine|Mucosa|Caecum|Intracellular|drug|Concentration in container' is 32.96% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.33

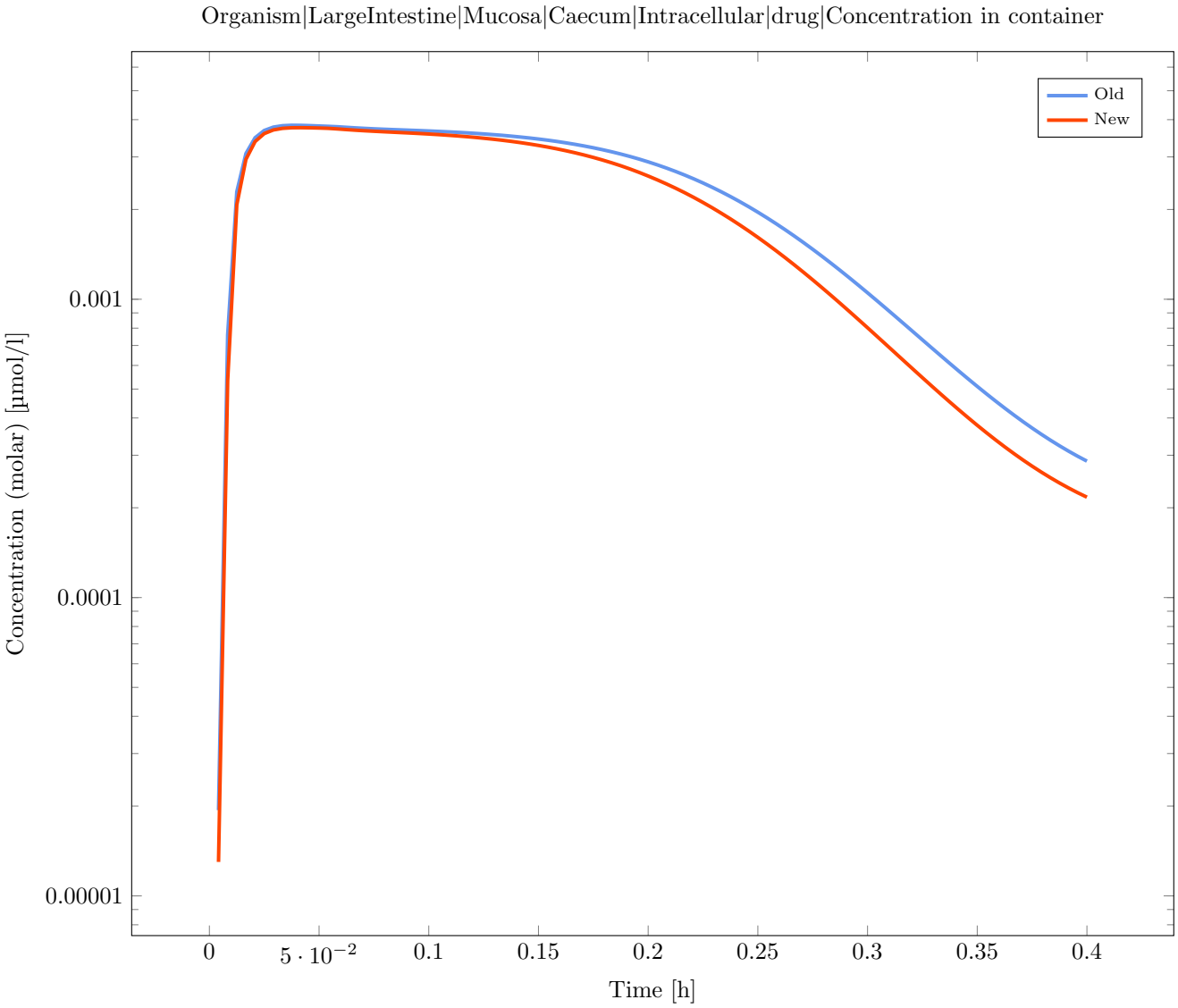
Organism|LargeIntestine|Mucosa|Caecum|Intracellular|drug|Concentration in container
Old
New
Concentration (molar) [µmol/l]
0.001
0.0001
0.00001
0
5 · 10−2
0.1
0.15
0.2
0.25
0.3
0.35
0.4
Time [h]

Figure 1.73

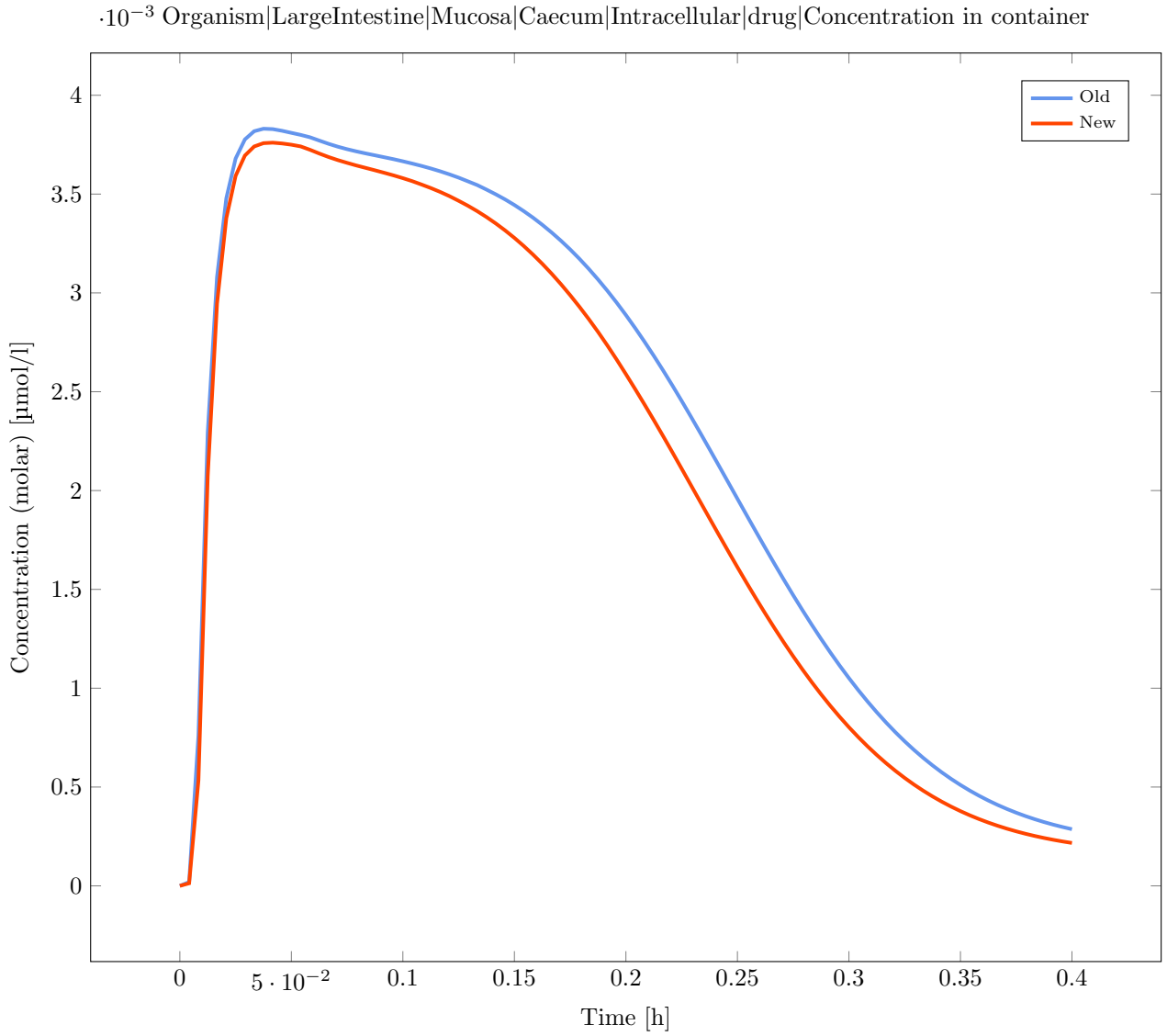

Figure 1.74

Output Path: Organism|SmallIntestine|Mucosa|LowerIleum|Intracellular|drug|Concentration in container
Deviation for 'Organism|SmallIntestine|Mucosa|LowerIleum|Intracellular|drug|Concentration in container' is 32.88% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.33

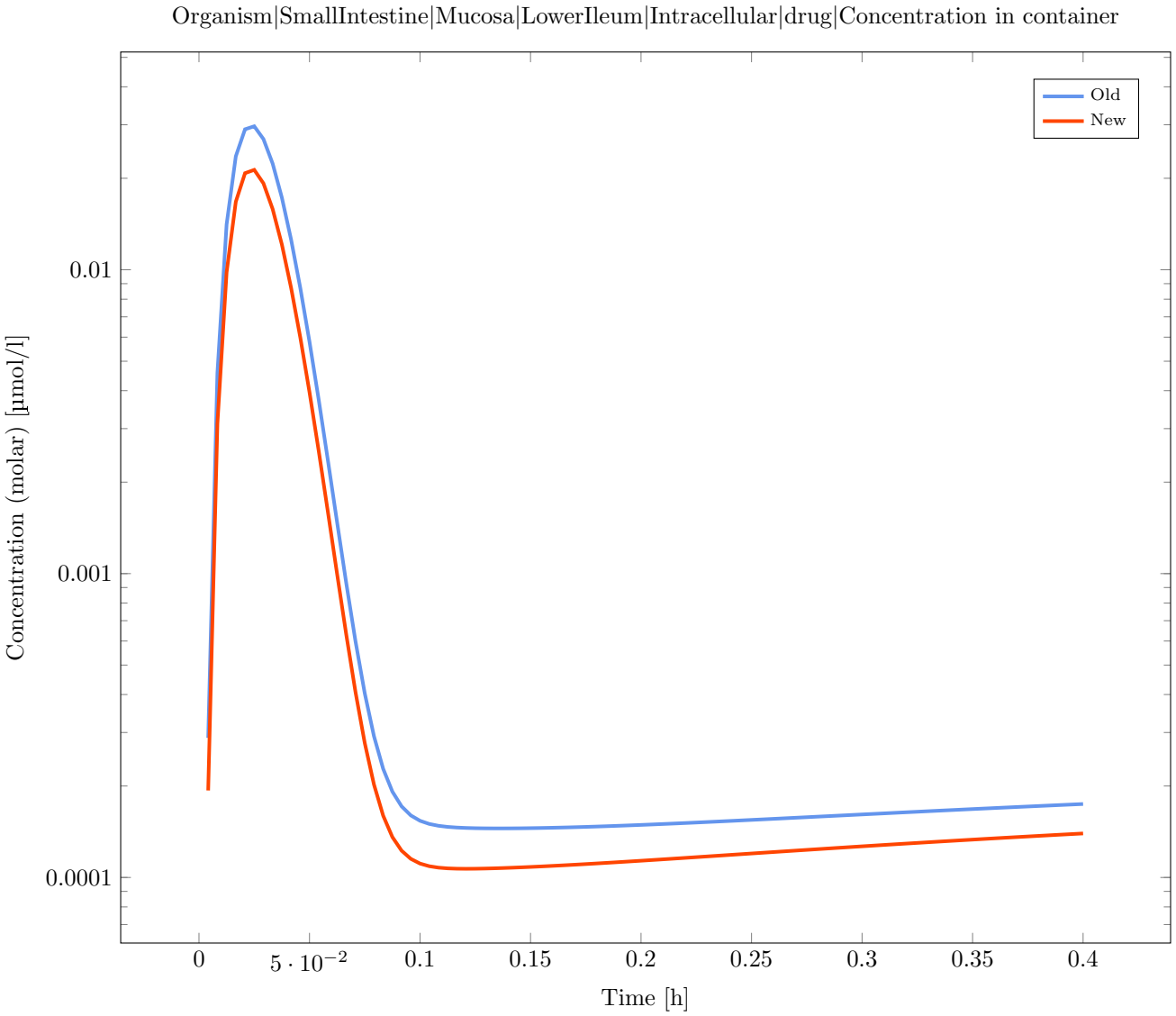
Organism|SmallIntestine|Mucosa|LowerIleum|Intracellular|drug|Concentration in container
Old
New
Concentration (molar) [µmol/l]
0.01
0.001
0.0001
0
$5 \cdot 10^{-2}$
0.1
0.15
0.2
0.25
0.3
0.35
0.4
Time [h]

Figure 1.75

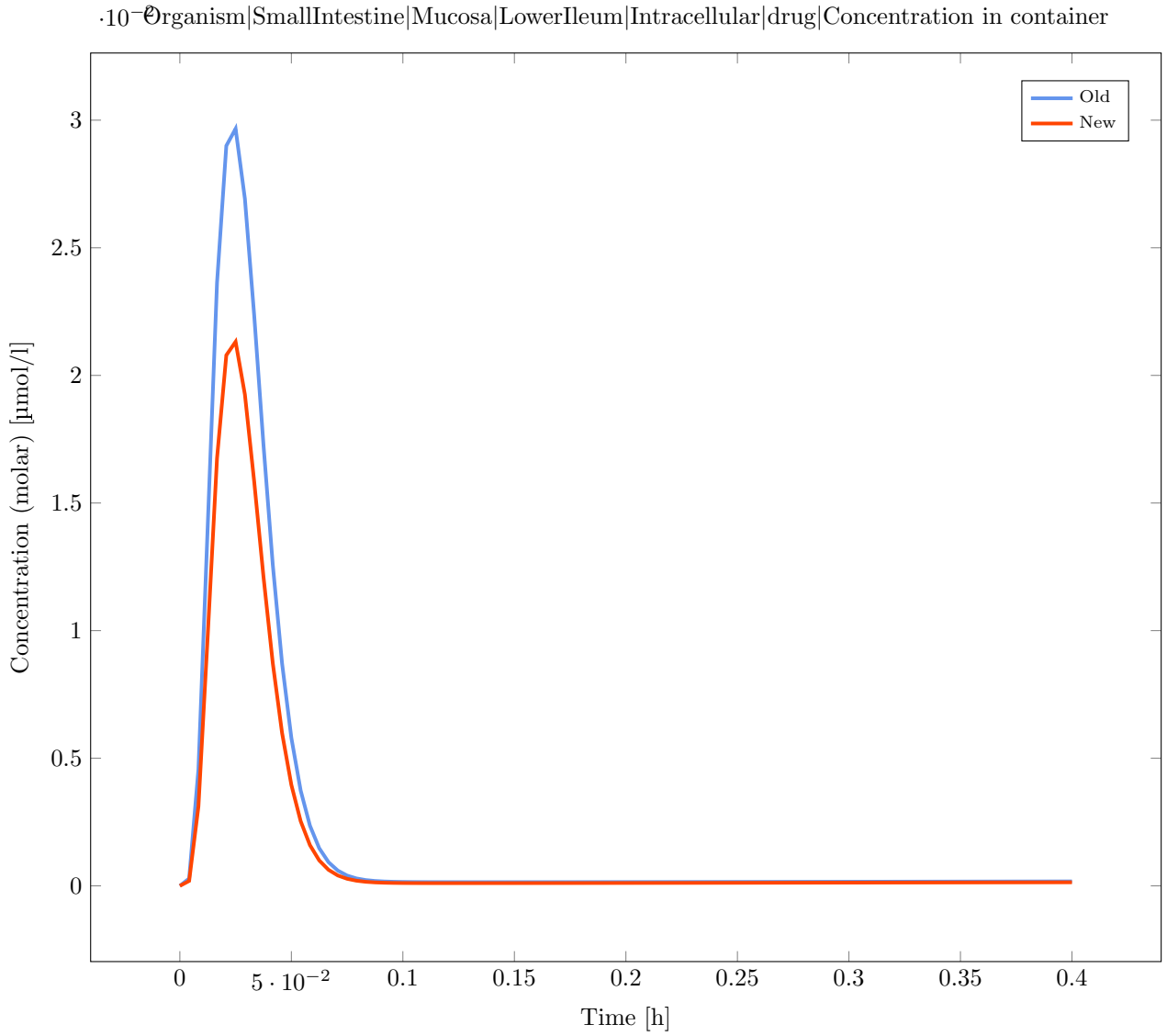

Figure 1.76

Output Path: Organism|SmallIntestine|Mucosa|LowerIleum|Interstitial|drug|Concentration in container
Deviation for 'Organism|SmallIntestine|Mucosa|LowerIleum|Interstitial|drug|Concentration in container' is 32.80% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.33

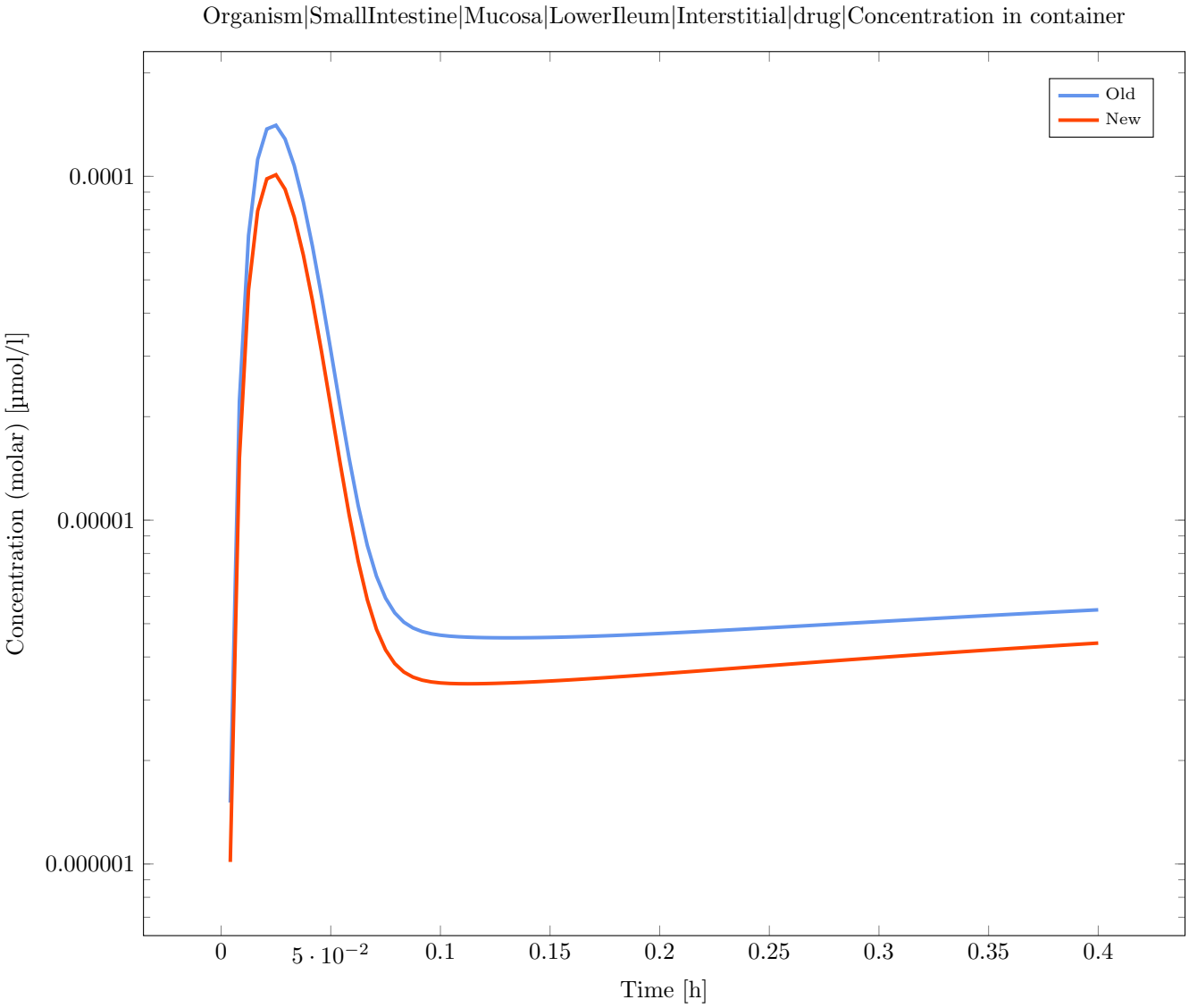
Organism|SmallIntestine|Mucosa|LowerIleum|Interstitial|drug|Concentration in container
Old
New
0.0001
0.00001
0.000001
Concentration (molar) [µmol/l]
0
5 · 10−2
0.1
0.15
0.2
0.25
0.3
0.35
0.4
Time [h]

Figure 1.77

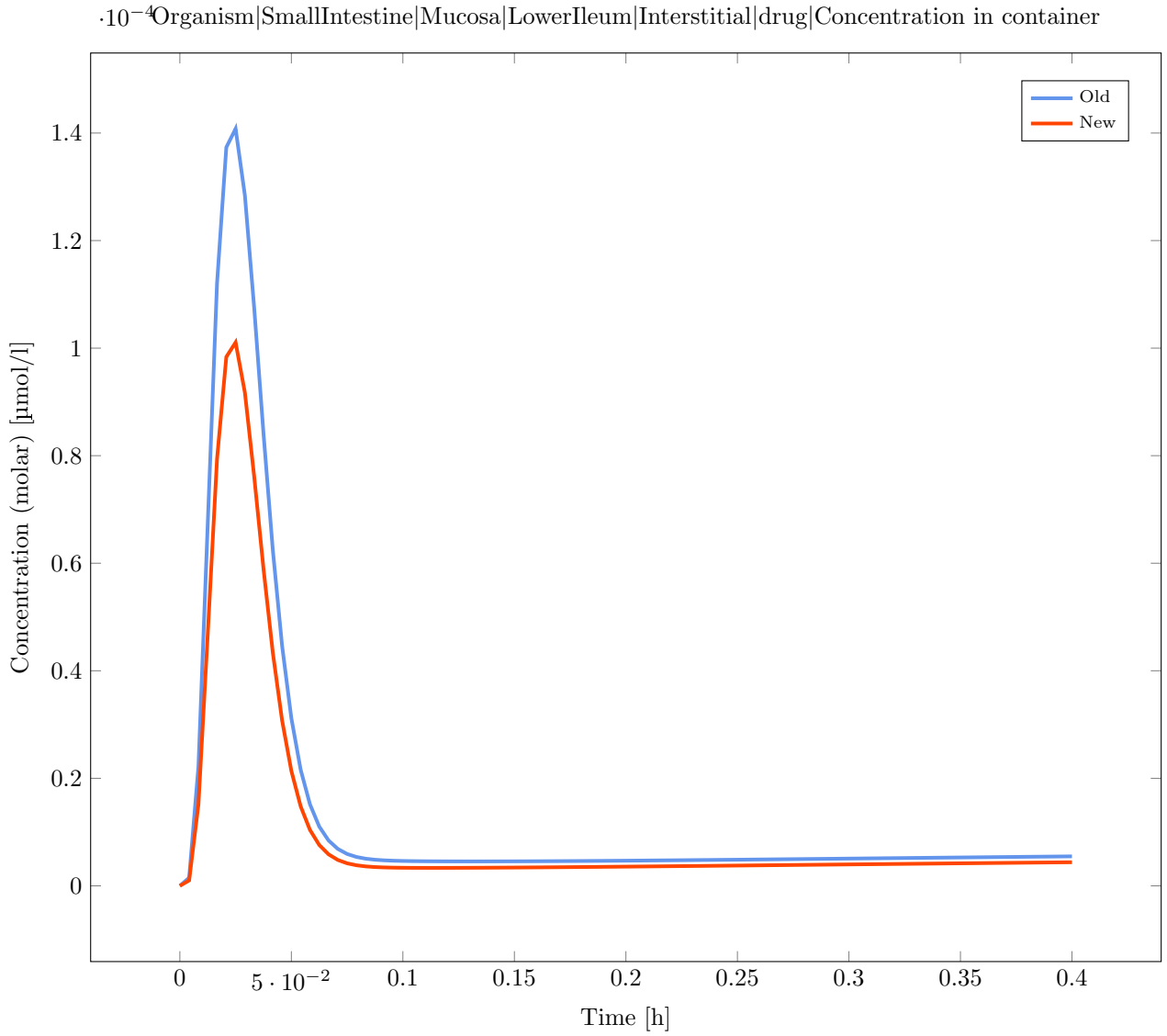

Figure 1.78

Output Path: Organism|SmallIntestine|Mucosa|LowerIleum|Plasma|drug|Concentration in container
Deviation for 'Organism|SmallIntestine|Mucosa|LowerIleum|Plasma|drug|Concentration in container' is 32.80% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.33

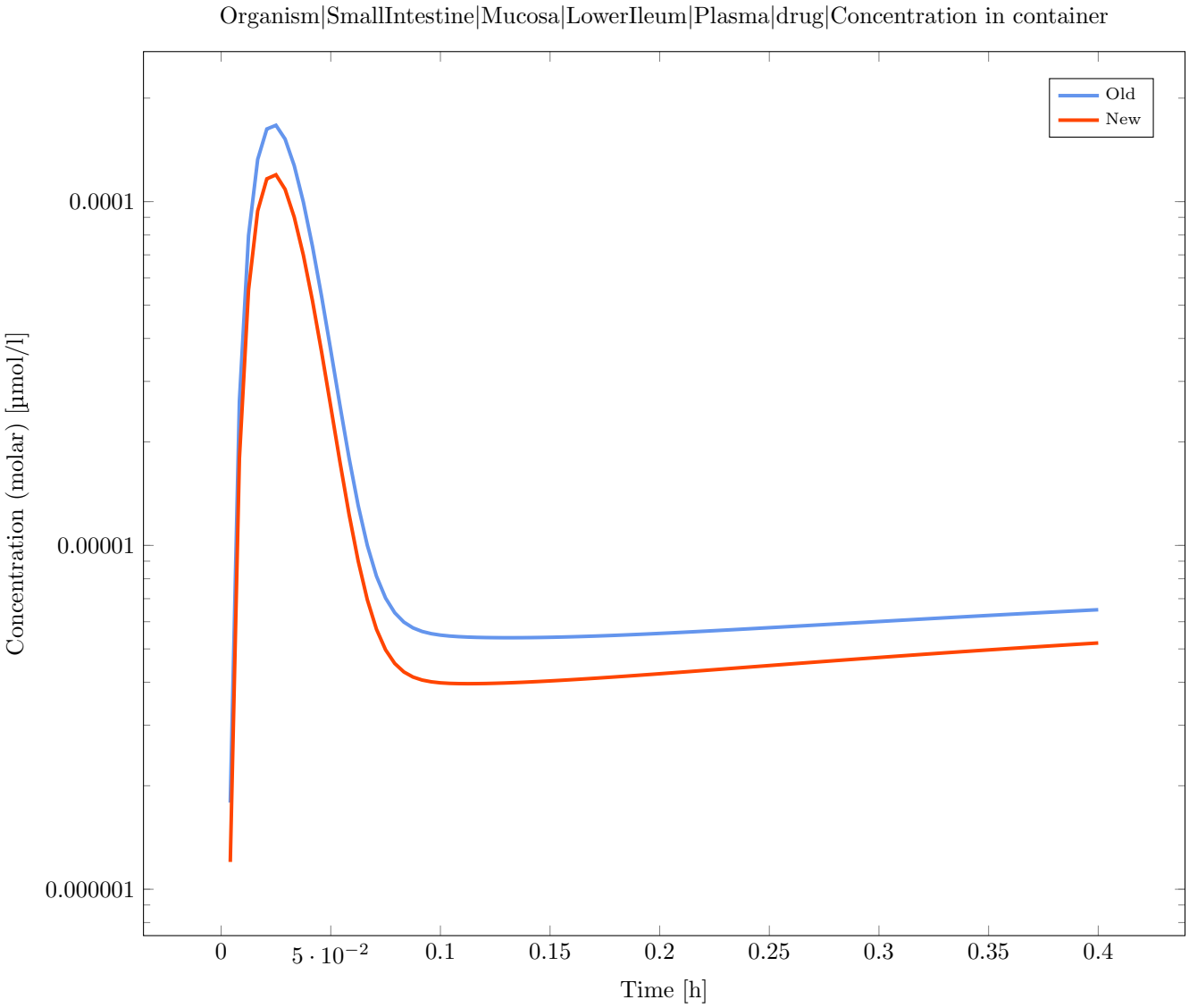
Organism|SmallIntestine|Mucosa|LowerIleum|Plasma|drug|Concentration in container
Old
New
Concentration (molar) [µmol/l]
0.0001
0.00001
0.000001
0
$5 \cdot 10^{-2}$
0.1
0.15
0.2
0.25
0.3
0.35
0.4
Time [h]

Figure 1.79

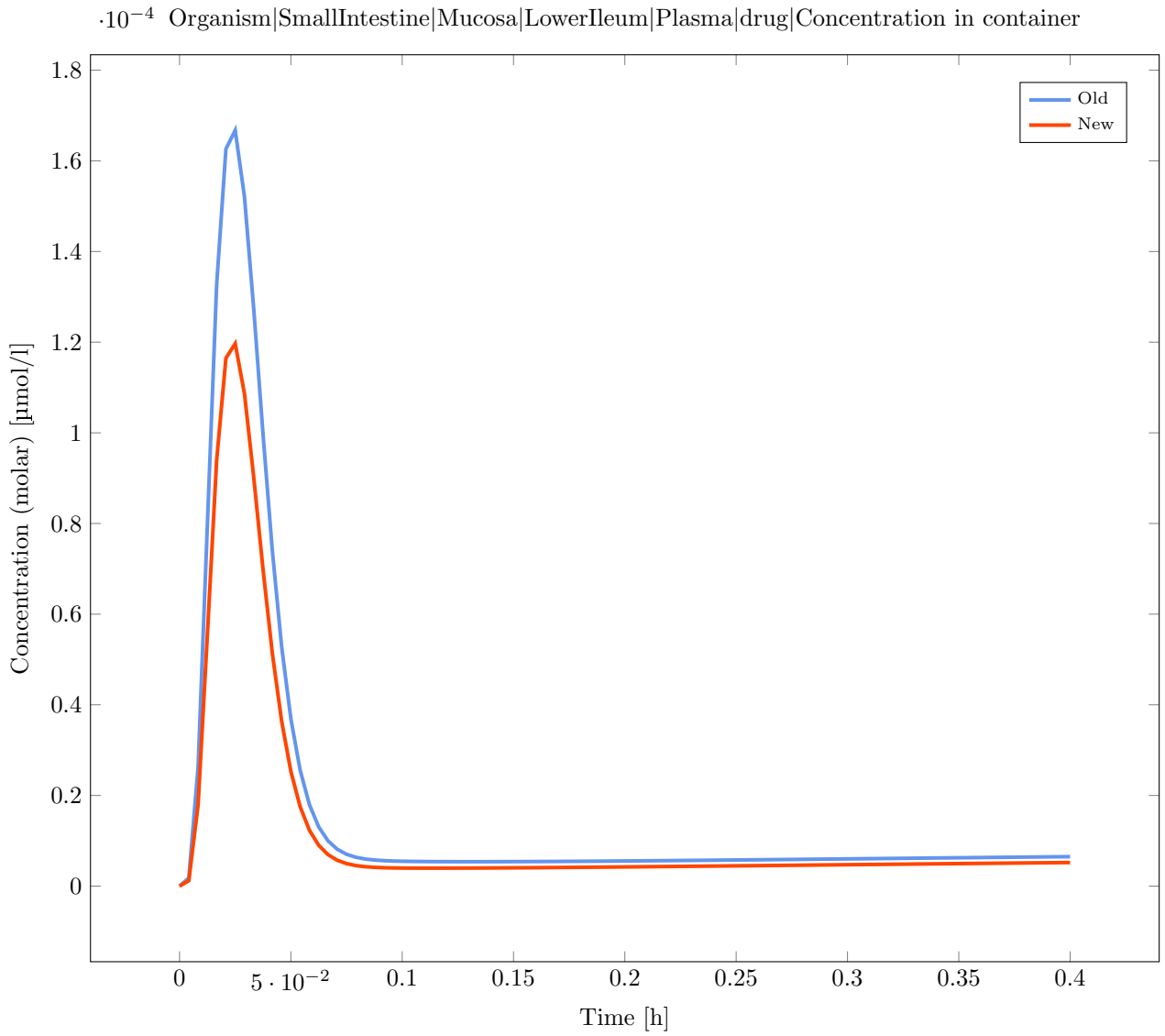

Figure 1.80

Output Path: Organism|Skin|Intracellular|drug|Concentration in container
Deviation for 'Organism|Skin|Intracellular|drug|Concentration in container' is 32.57% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.33

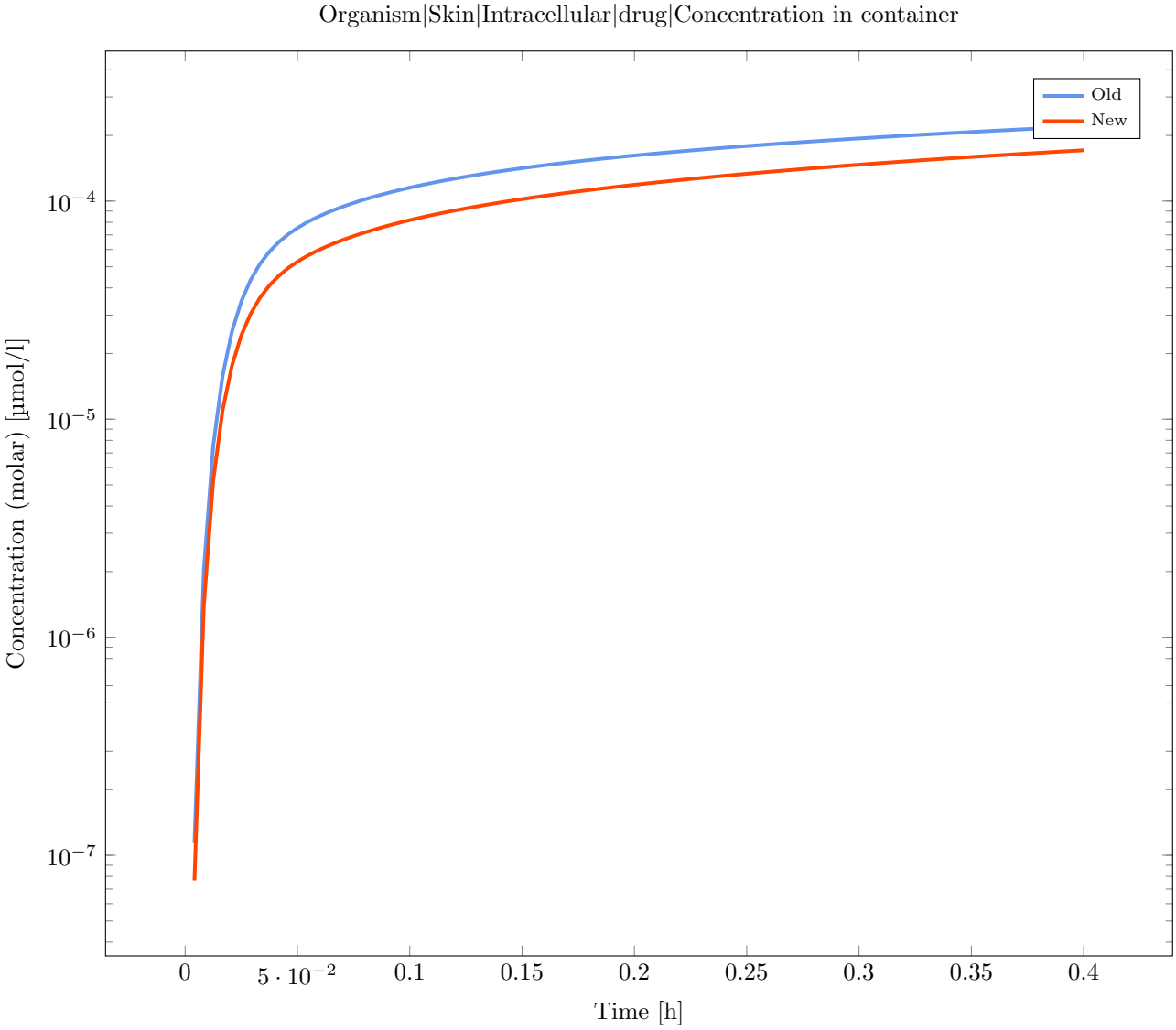
Organism|Skin|Intracellular|drug|Concentration in container
Old
New
Concentration (molar) [µmol/l]
10^{-4}
10^{-5}
10^{-6}
10^{-7}
0
$5 \cdot 10^{-2}$
0.1
0.15
0.2
0.25
0.3
0.35
0.4
Time [h]

Figure 1.81

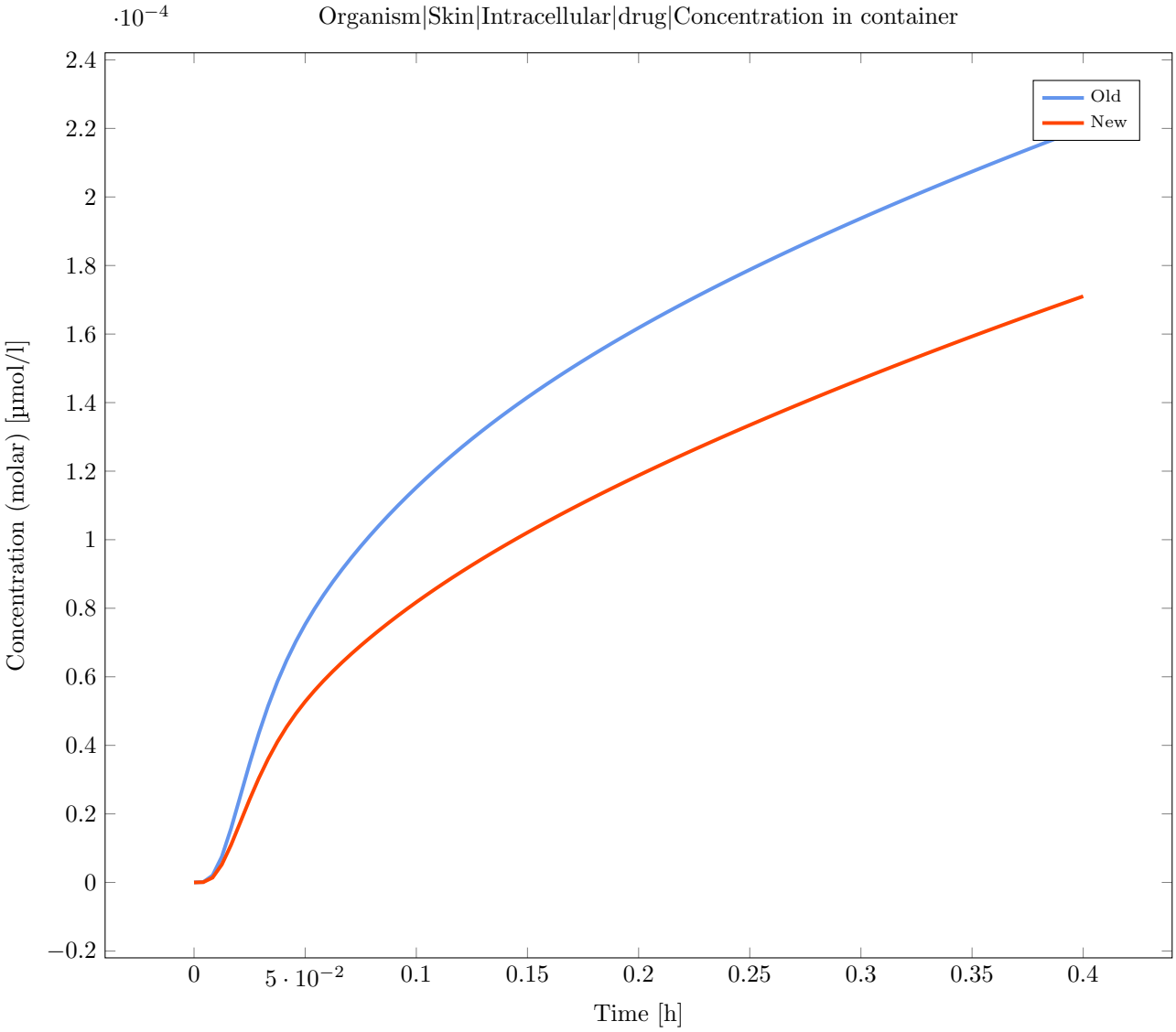

Figure 1.82

Simulation: Human_SingleORAL_Weibull-Human_SingleORAL_Weibull_MW_200_fu_0.2_LogP_5
Result of the validation: Invalid
Absolute Tolerance: 1.00E-10
Relative Tolerance: 1.00E-5

Output Path: Organism|SmallIntestine|Mucosa|LowerJejunum|Intracellular|drug|Concentration in container
Deviation for 'Organism|SmallIntestine|Mucosa|LowerJejunum|Intracellular|drug|Concentration in container' is 3.89% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.04

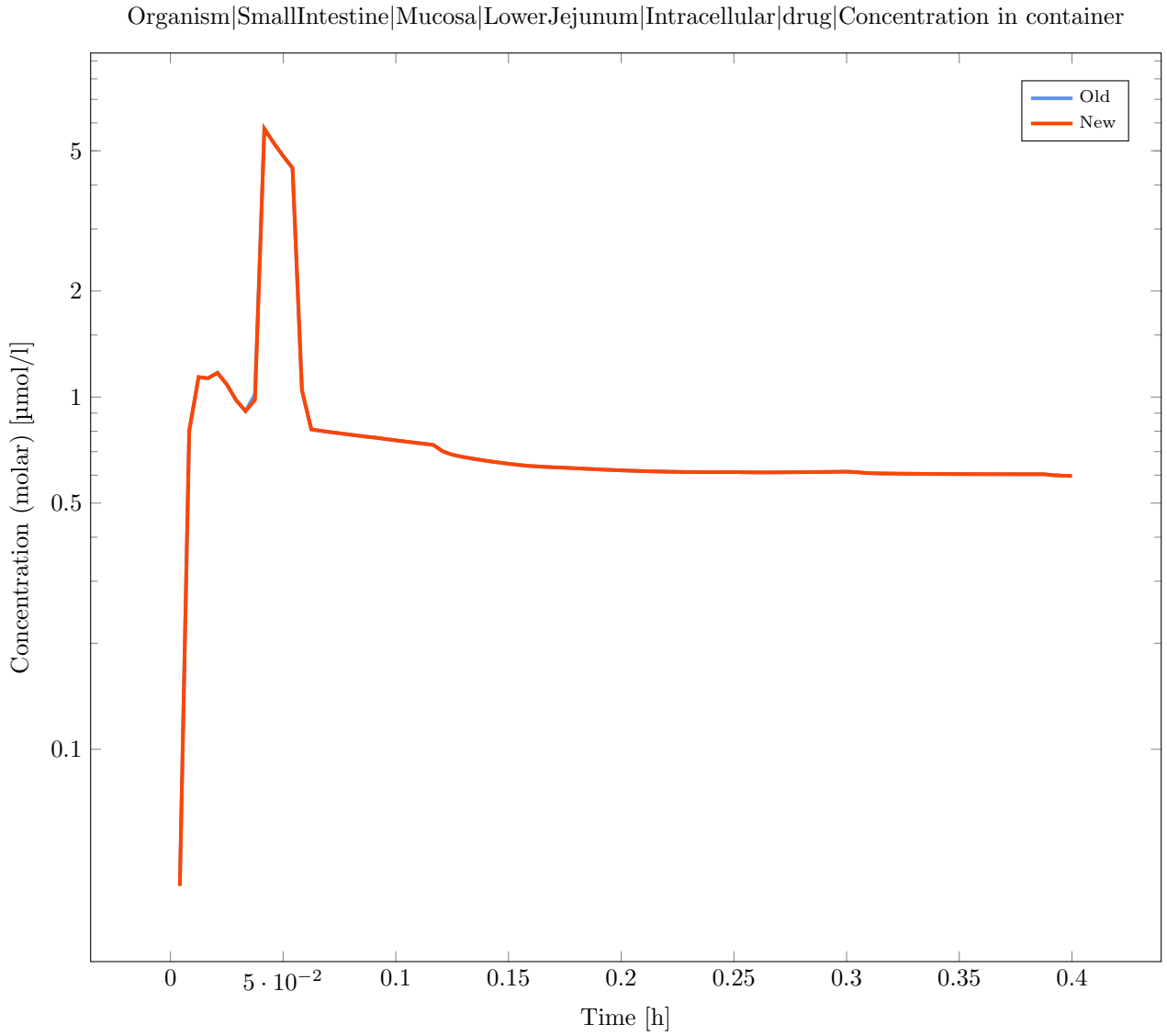
Organism|SmallIntestine|Mucosa|LowerJejunum|Intracellular|drug|Concentration in container
Old
New
Concentration (molar) [µmol/l]
5
2
1
0.5
0.1
0
5 · 10−2
0.1
0.15
0.2
0.25
0.3
0.35
0.4
Time [h]

Figure 1.83

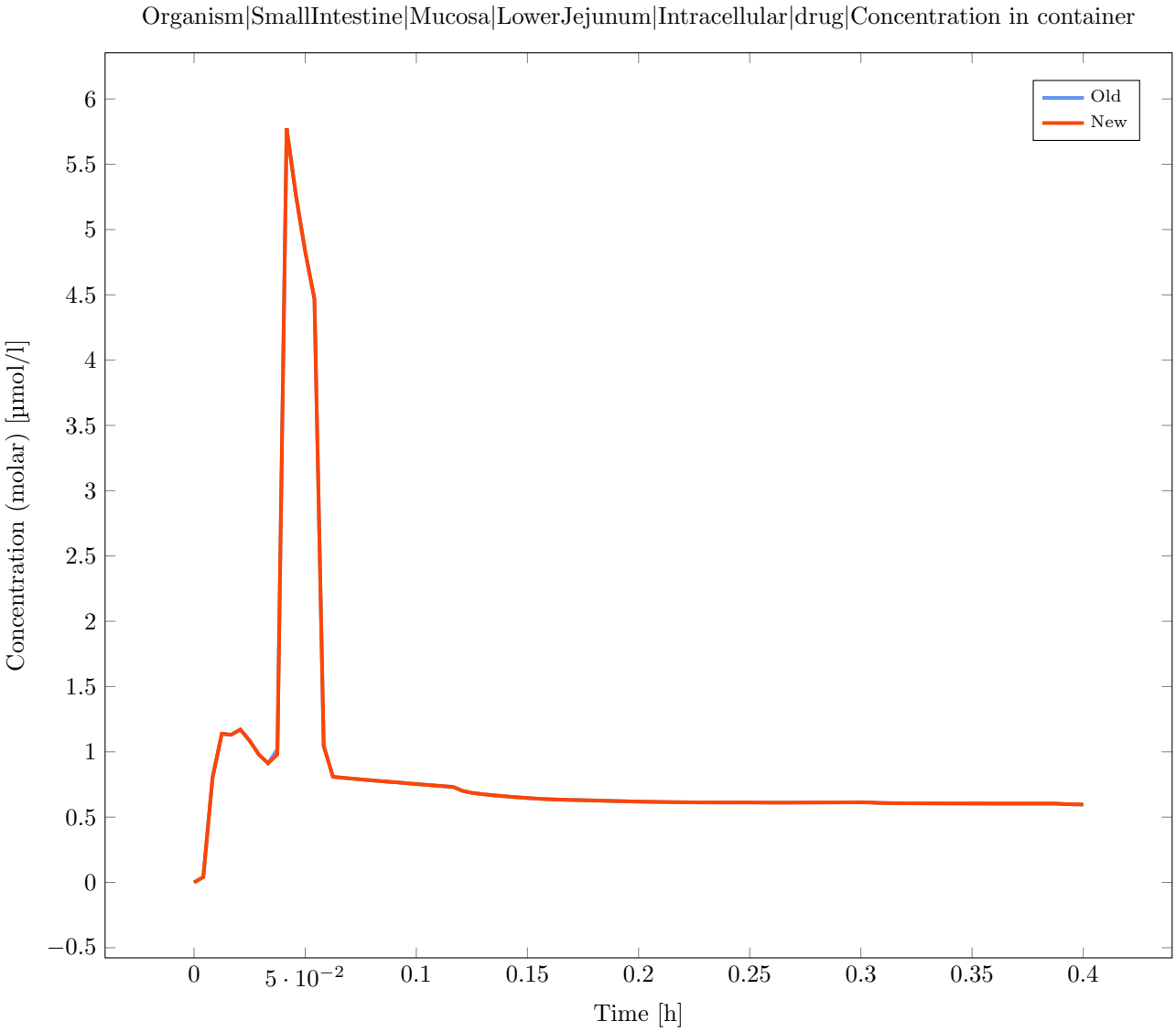

Figure 1.84

Output Path: Organism|SmallIntestine|Mucosa|LowerJejunum|Interstitial|drug|Concentration in container
Deviation for 'Organism|SmallIntestine|Mucosa|LowerJejunum|Interstitial|drug|Concentration in container' is 3.40% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.03

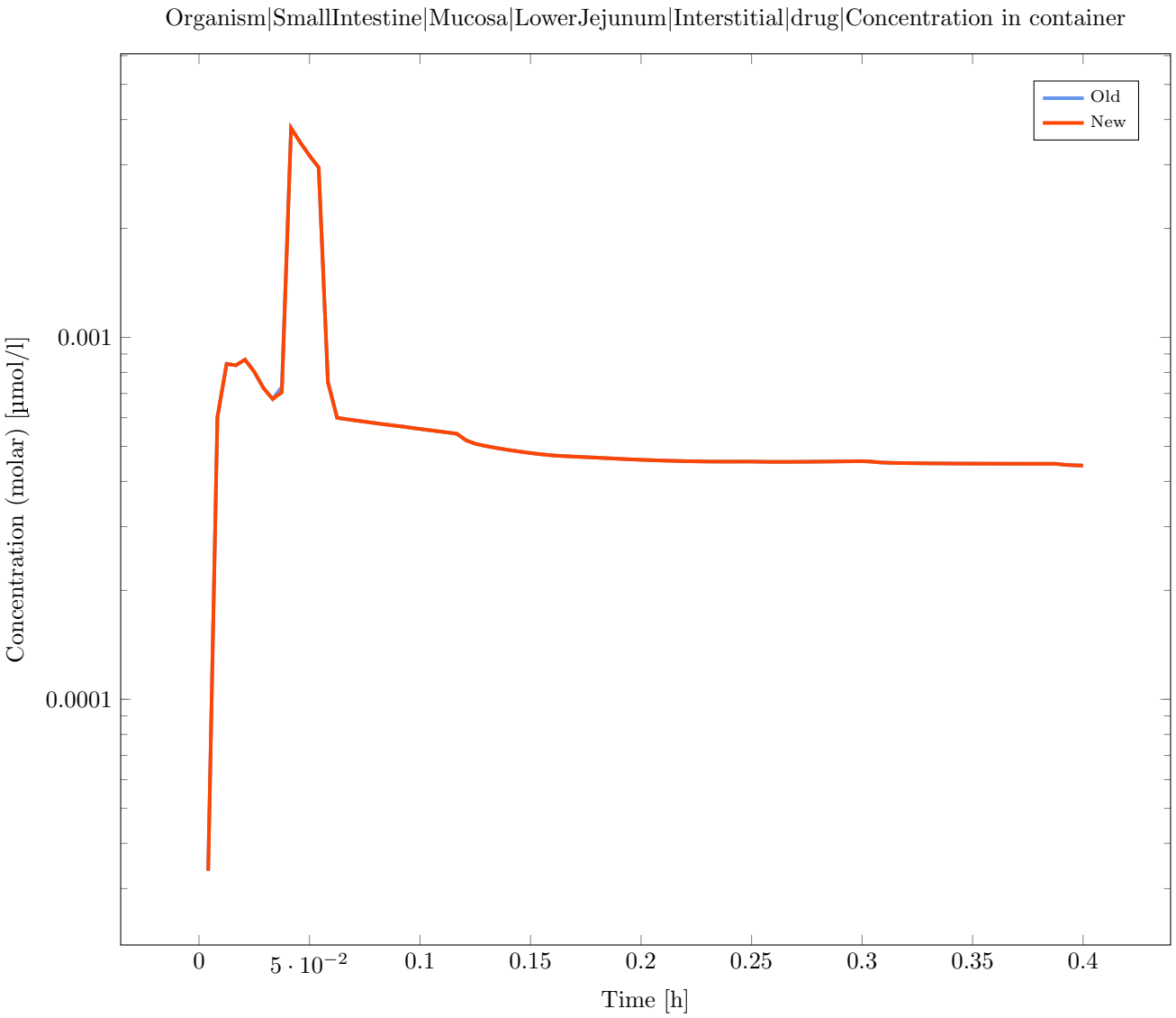
Organism|SmallIntestine|Mucosa|LowerJejunum|Interstitial|drug|Concentration in container
Old
New
Concentration (molar) [µmol/l]
0.001
0.0001
0
5 · 10−2
0.1
0.15
0.2
0.25
0.3
0.35
0.4
Time [h]

Figure 1.85

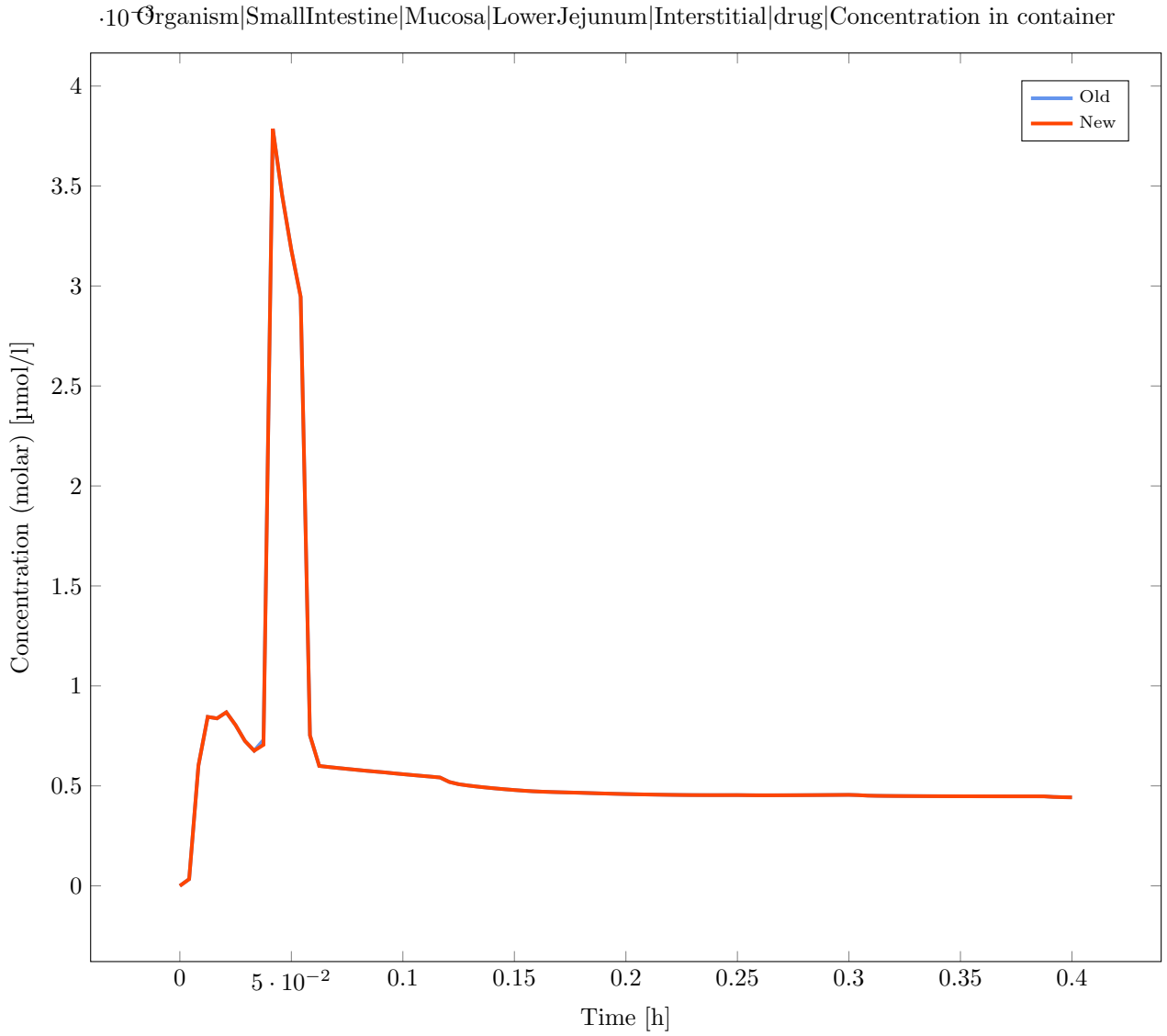

Figure 1.86

Output Path: Organism|SmallIntestine|Mucosa|LowerJejunum|Plasma|drug|Concentration in container
Deviation for 'Organism|SmallIntestine|Mucosa|LowerJejunum|Plasma|drug|Concentration in container' is 3.01% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.03

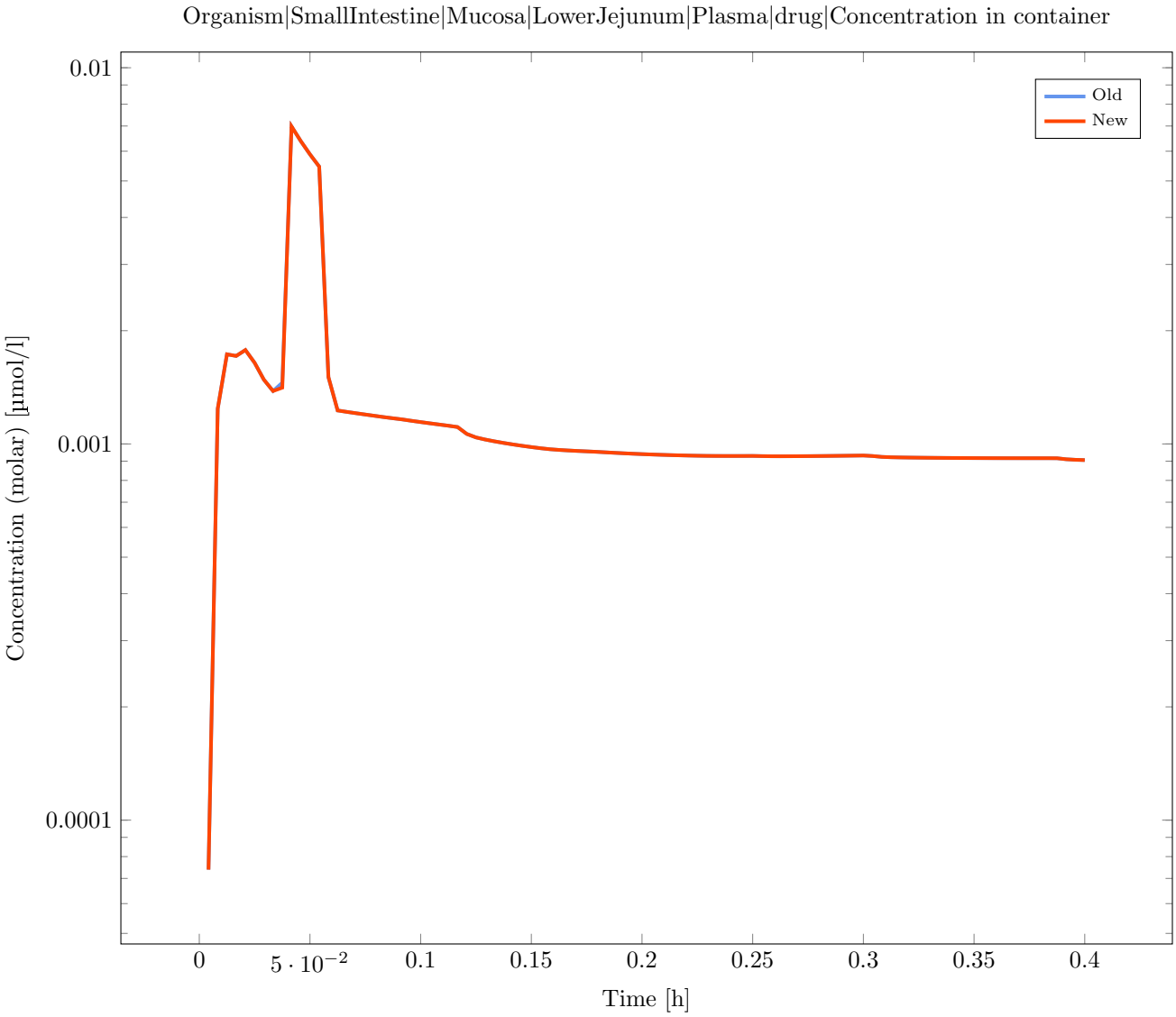
Organism|SmallIntestine|Mucosa|LowerJejunum|Plasma|drug|Concentration in container
Old
New
Concentration (molar) [µmol/l]
0.01
0.001
0.0001
0
5 · 10−2
0.1
0.15
0.2
0.25
0.3
0.35
0.4
Time [h]

Figure 1.87

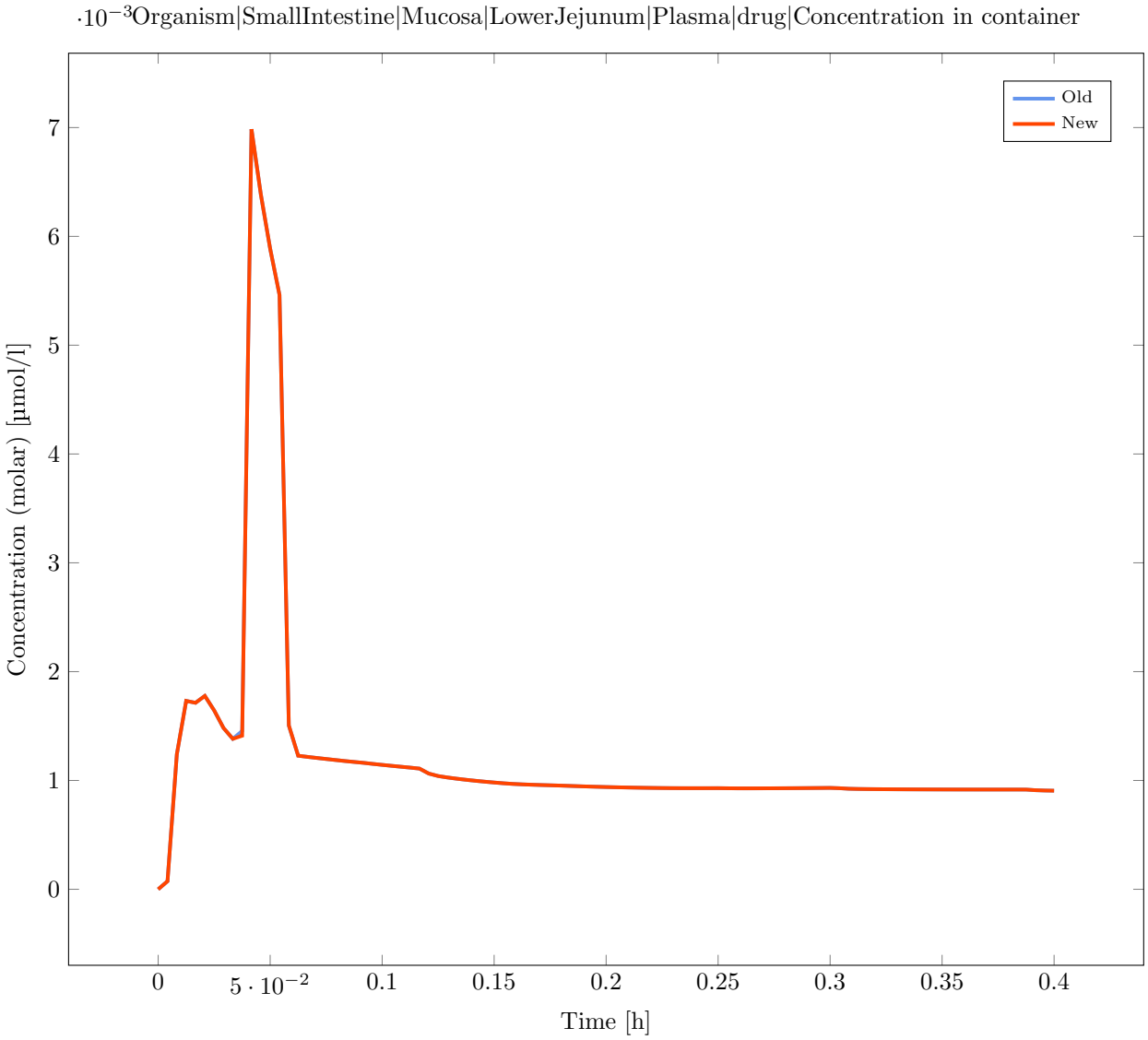

Figure 1.88

Simulation: Rat_MultiORAL_6_6_12_Dissolved-Rat_MultiORAL_6_6_12_Dissolved
Result of the validation: Invalid
Absolute Tolerance: 1.00E-10
Relative Tolerance: 1.00E-5

Output Path: Organism|SmallIntestine|Mucosa|UpperIleum|Intracellular|drug|Concentration in container
Deviation for 'Organism|SmallIntestine|Mucosa|UpperIleum|Intracellular|drug|Concentration in container' is 4.86% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.05

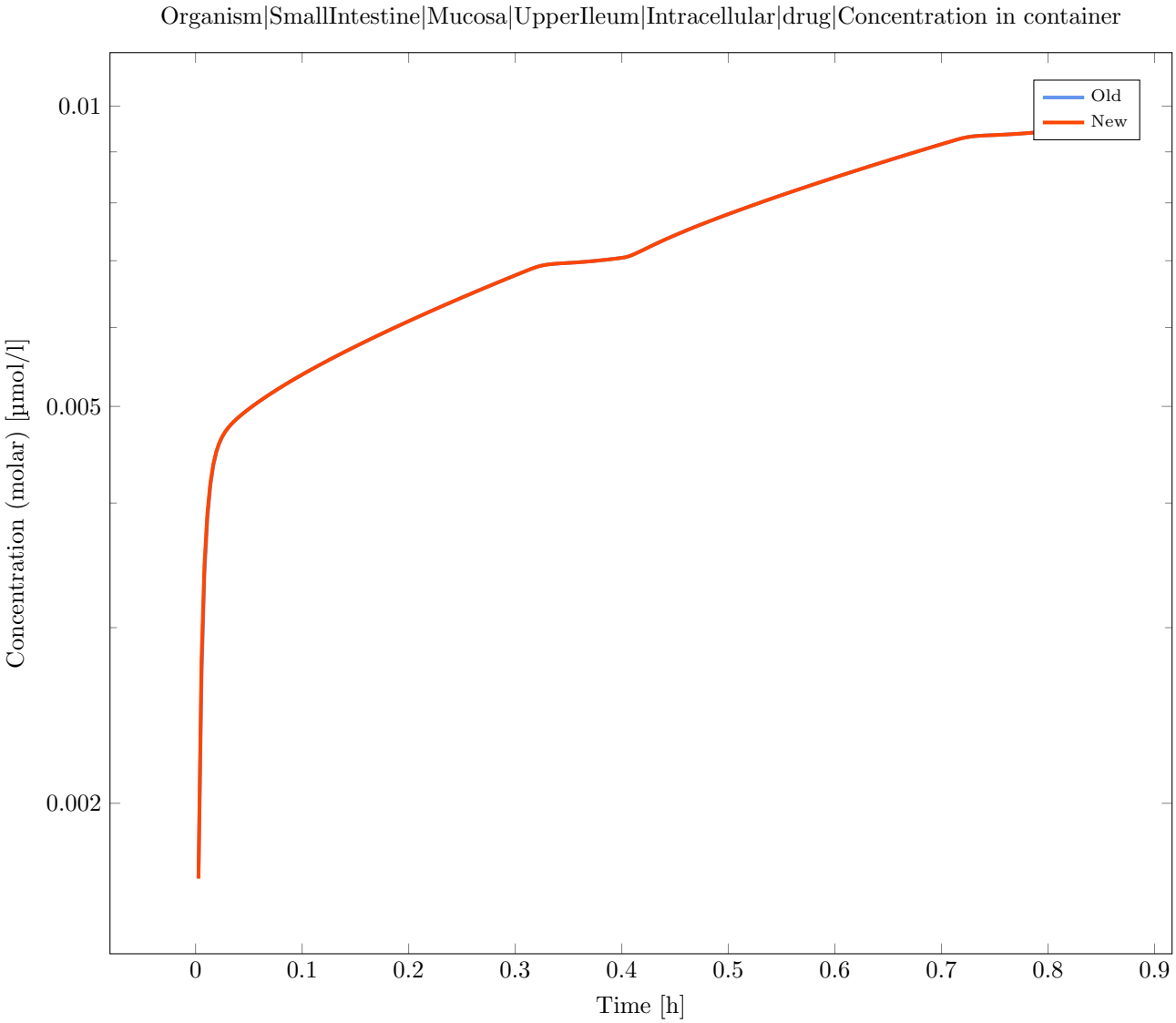
Organism|SmallIntestine|Mucosa|UpperIleum|Intracellular|drug|Concentration in container
Old
New
Concentration (molar) [µmol/l]
0.01
0.005
0.002
0
0.1
0.2
0.3
0.4
0.5
0.6
0.7
0.8
0.9
Time [h]

Figure 1.89

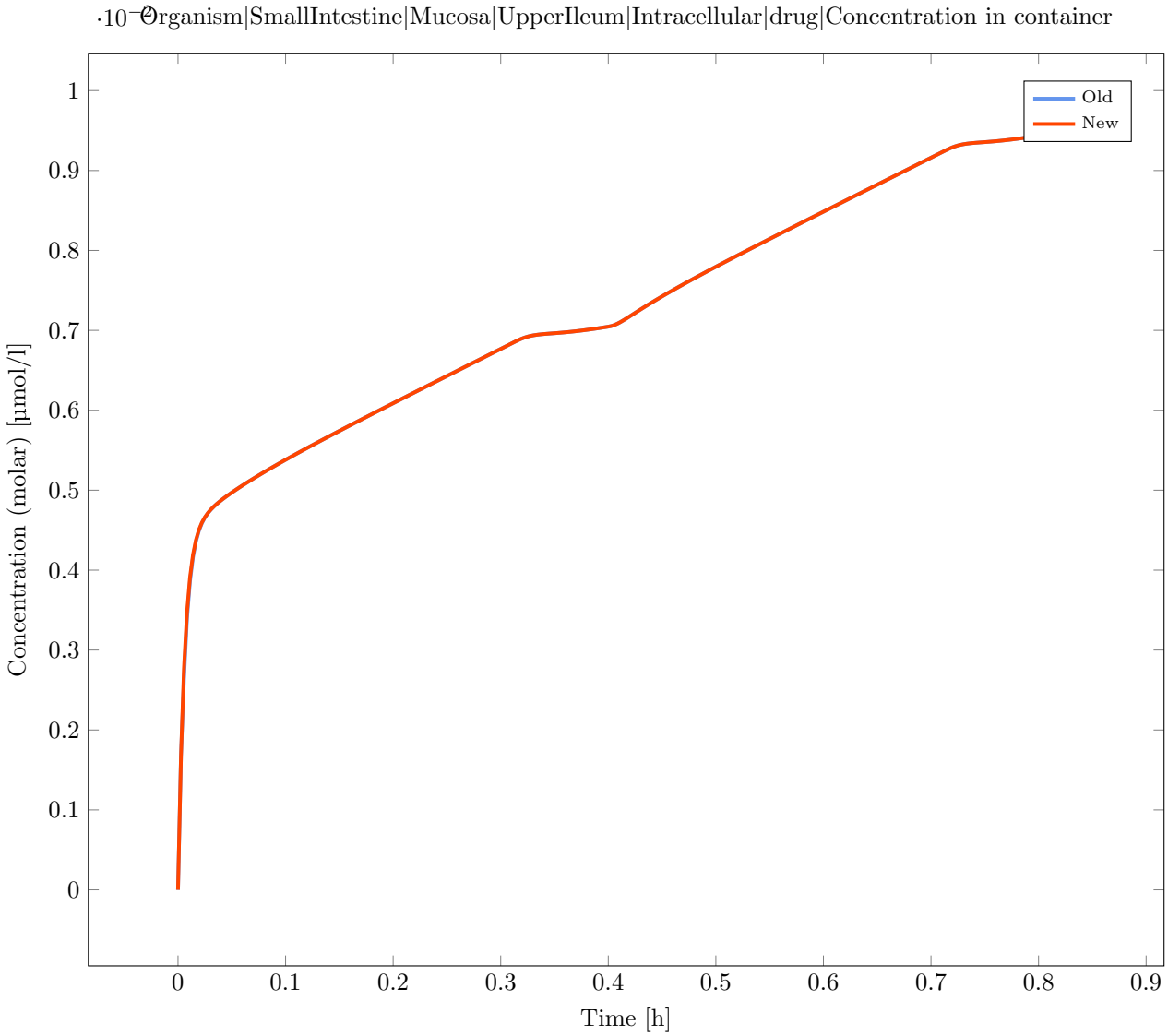

Figure 1.90

Output Path: Organism|SmallIntestine|Mucosa|LowerIleum|Intracellular|drug|Concentration in container
Deviation for 'Organism|SmallIntestine|Mucosa|LowerIleum|Intracellular|drug|Concentration in container' is 4.77% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.05

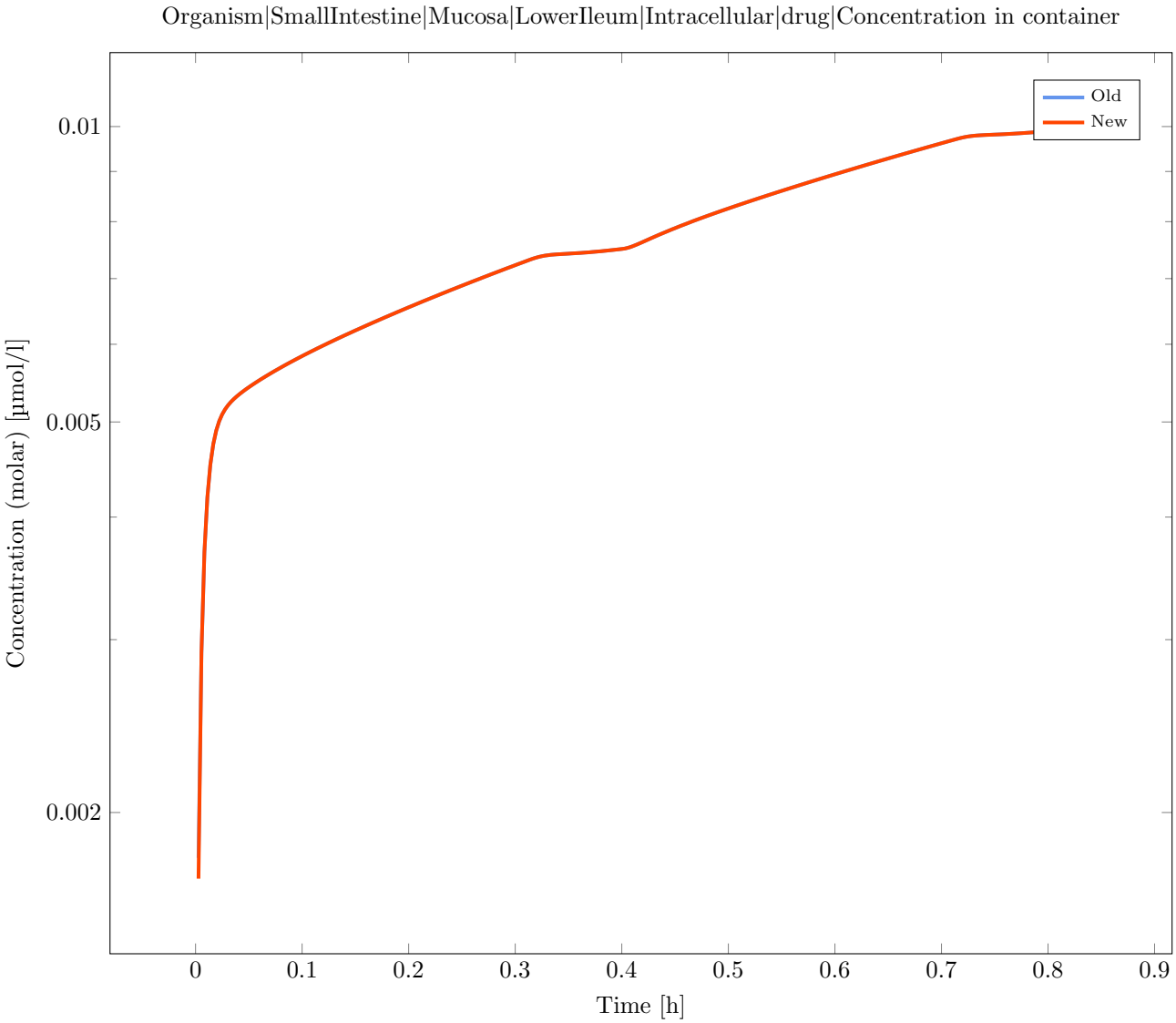
Organism|SmallIntestine|Mucosa|LowerIleum|Intracellular|drug|Concentration in container
Old
New
0.01
0.005
0.002
Concentration (molar) [µmol/l]
0
0.1
0.2
0.3
0.4
0.5
0.6
0.7
0.8
0.9
Time [h]

Figure 1.91

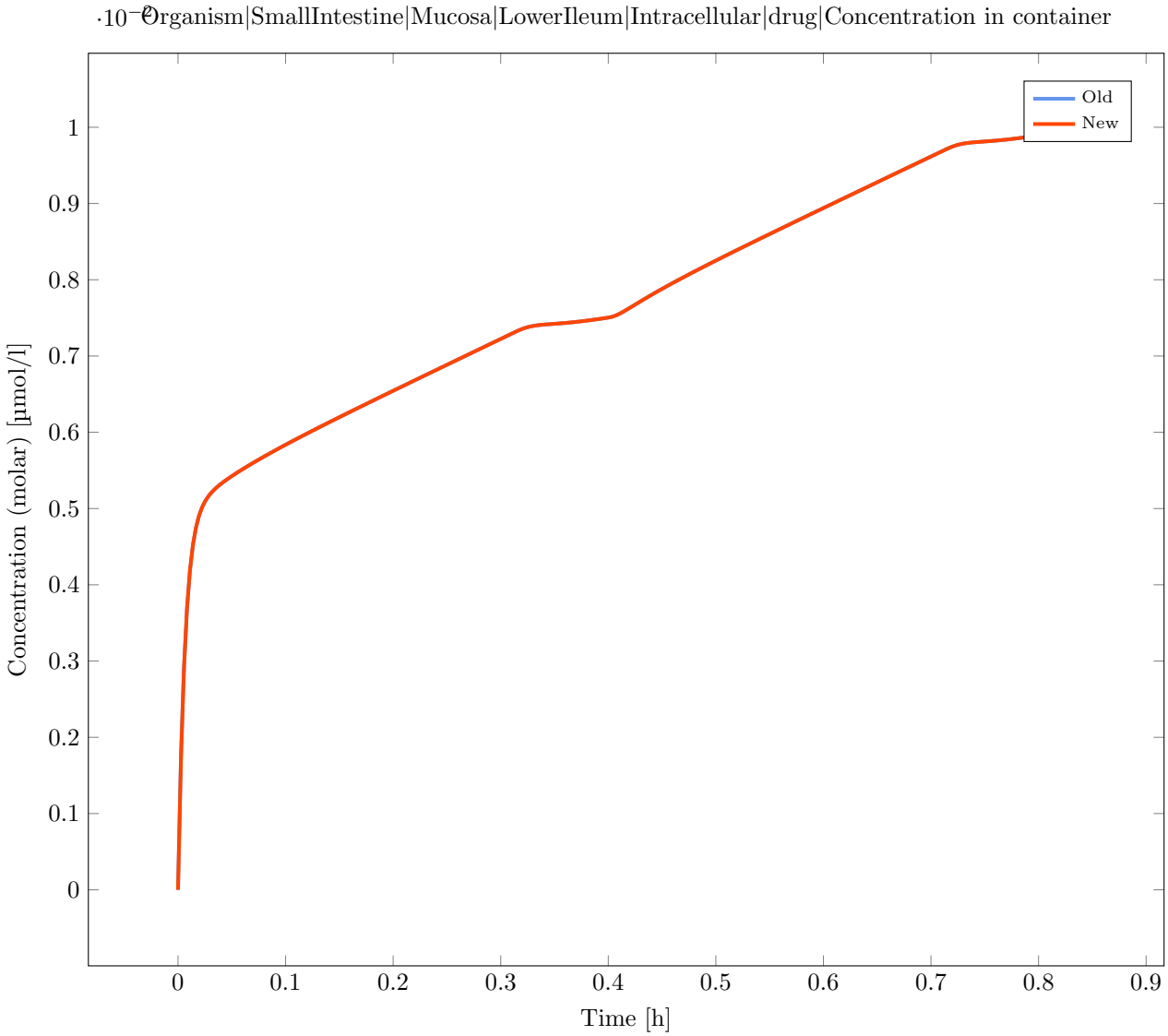

Figure 1.92

Output Path: Organism|LargeIntestine|Mucosa|ColonDescendens|Intracellular|drug|Concentration in container
Deviation for 'Organism|LargeIntestine|Mucosa|ColonDescendens|Intracellular|drug|Concentration in container' is 3.84% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.04

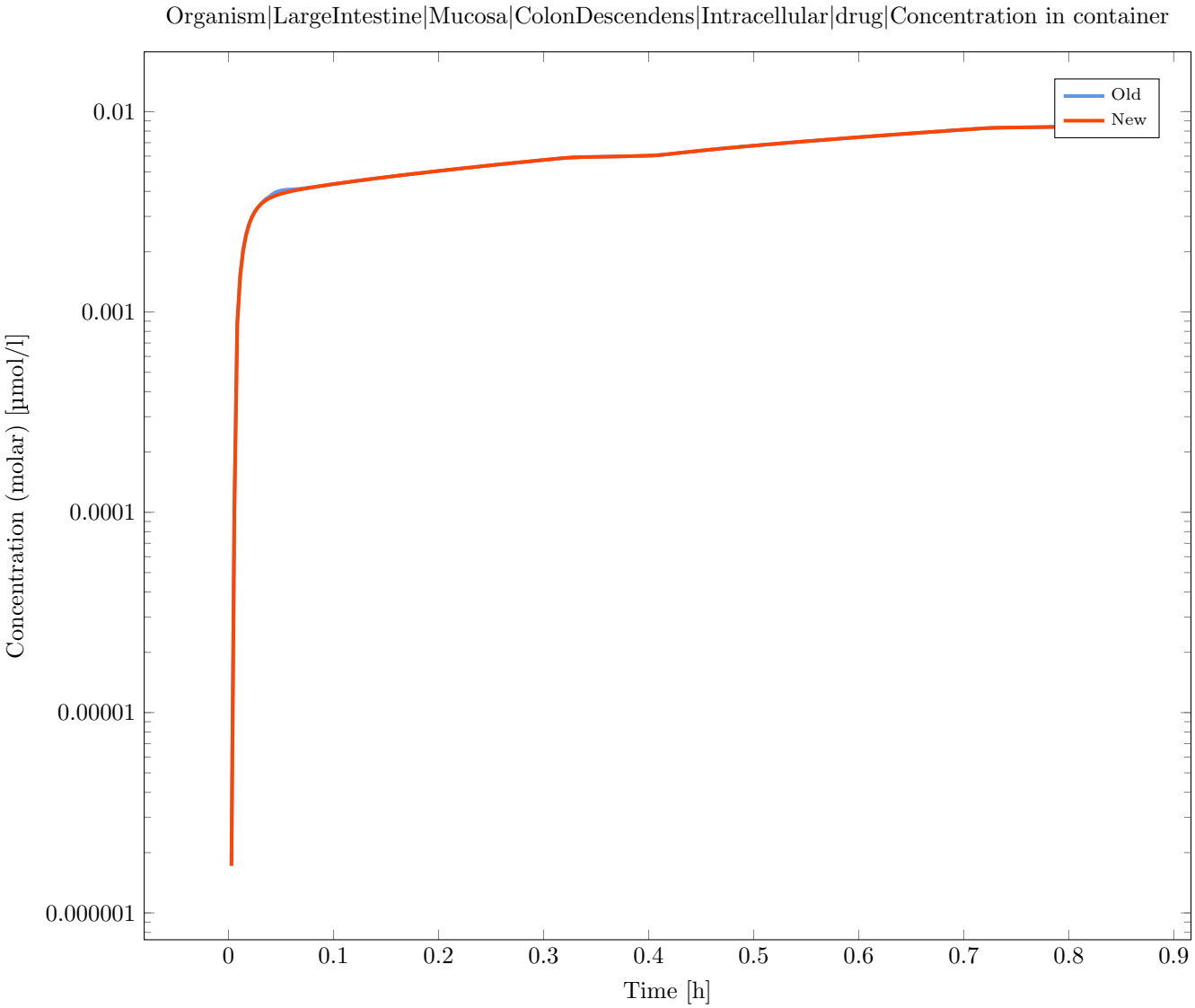
Organism|LargeIntestine|Mucosa|ColonDescendens|Intracellular|drug|Concentration in container
Old
New
Concentration (molar) [µmol/l]
0.01
0.001
0.0001
0.00001
0.000001
0
0.1
0.2
0.3
0.4
0.5
0.6
0.7
0.8
0.9
Time [h]

Figure 1.93

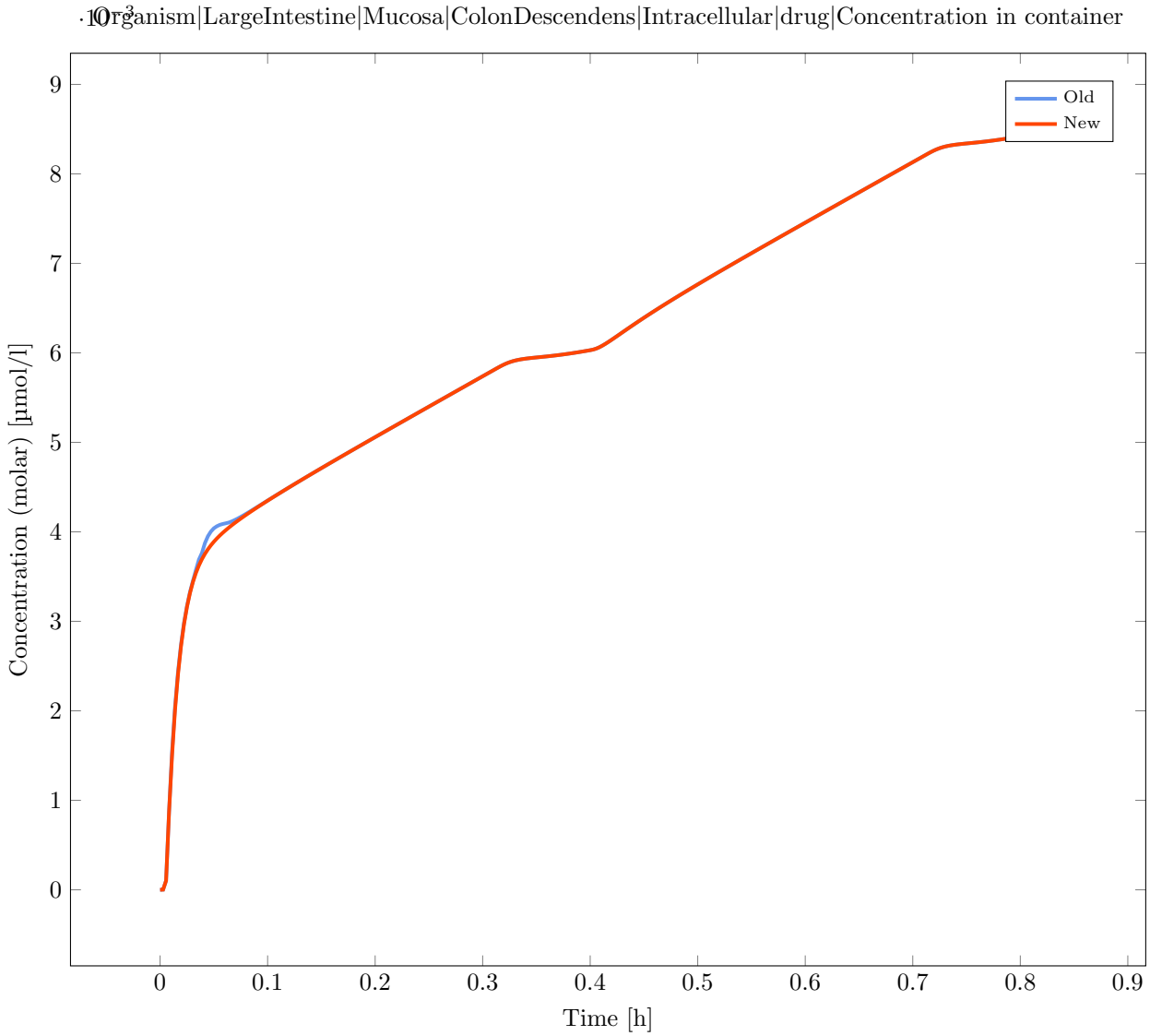

Figure 1.94

Simulation: Rat_MultiORAL_6_6_6_6_Dissolved-Rat_MultiORAL_6_6_6_6_Dissolved
Result of the validation: Invalid
Absolute Tolerance: 1.00E-10
Relative Tolerance: 1.00E-5

Output Path: Organism|SmallIntestine|Mucosa|UpperIleum|Intracellular|drug|Concentration in container
Deviation for 'Organism|SmallIntestine|Mucosa|UpperIleum|Intracellular|drug|Concentration in container' is 4.86% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.05

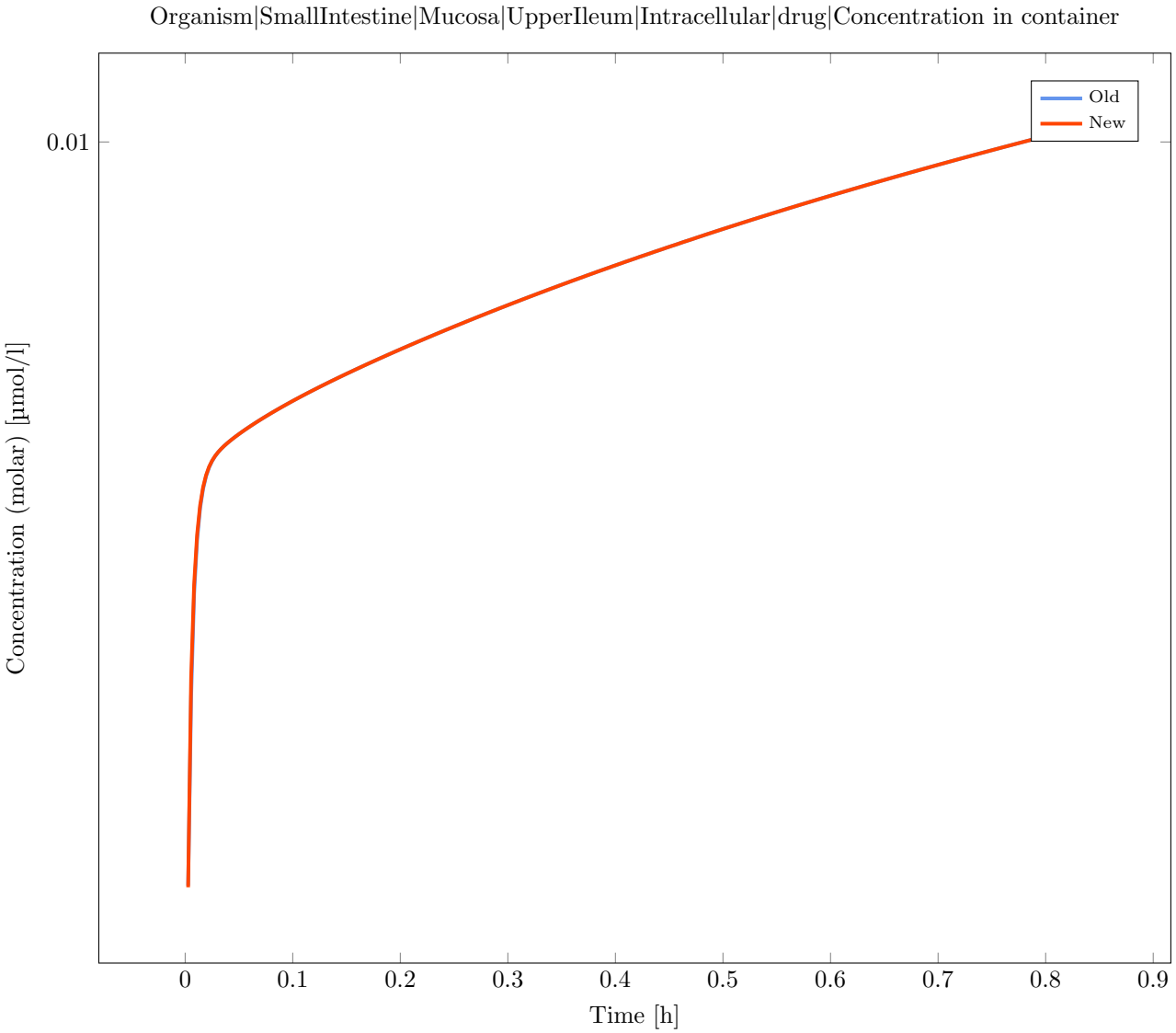
Organism|SmallIntestine|Mucosa|UpperIleum|Intracellular|drug|Concentration in container
Old
New
0.01
Concentration (molar) [µmol/l]
0
0.1
0.2
0.3
0.4
0.5
0.6
0.7
0.8
0.9
Time [h]

Figure 1.95

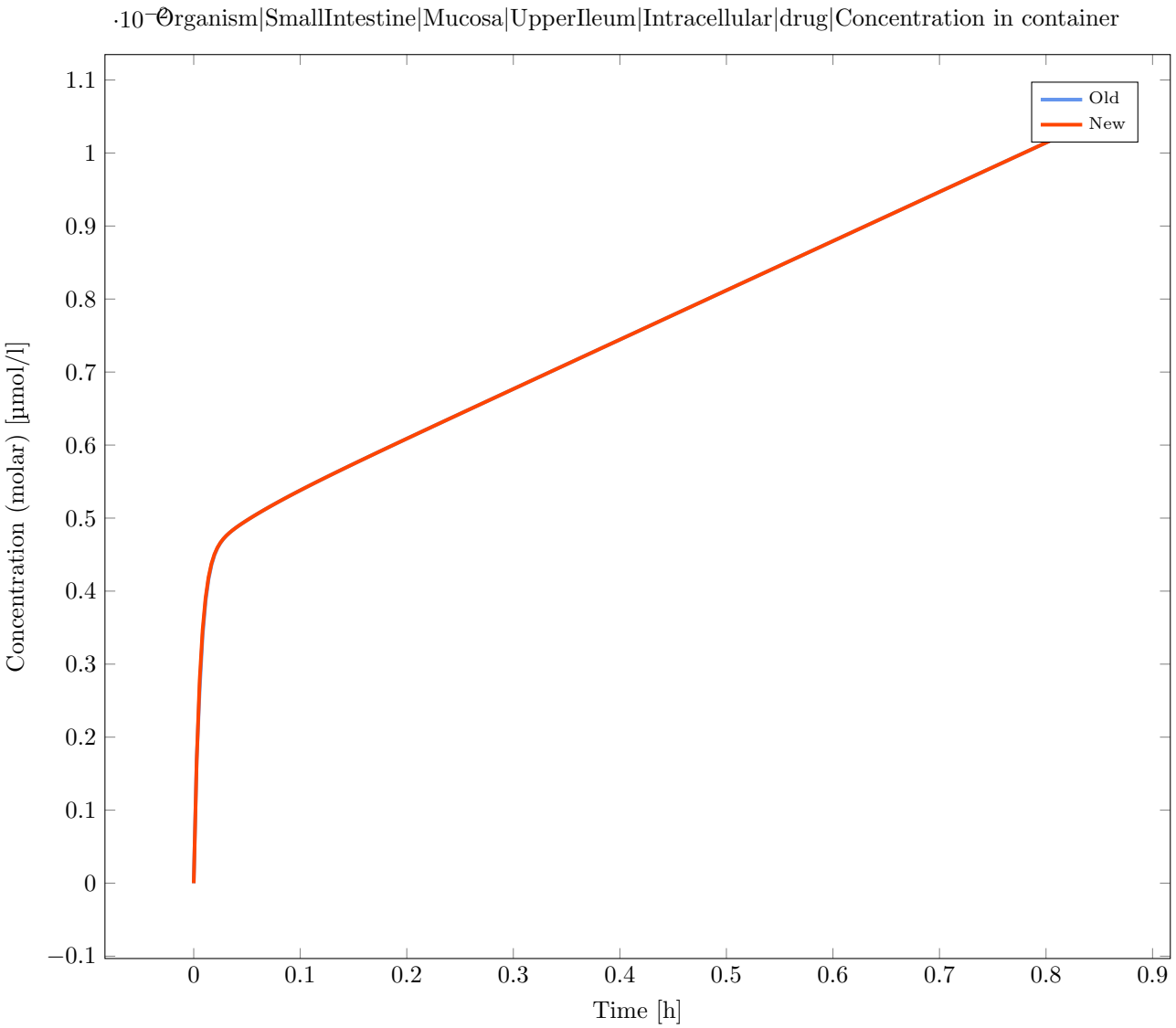

Figure 1.96

Output Path: Organism|SmallIntestine|Mucosa|LowerIleum|Intracellular|drug|Concentration in container
Deviation for 'Organism|SmallIntestine|Mucosa|LowerIleum|Intracellular|drug|Concentration in container' is 4.77% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.05

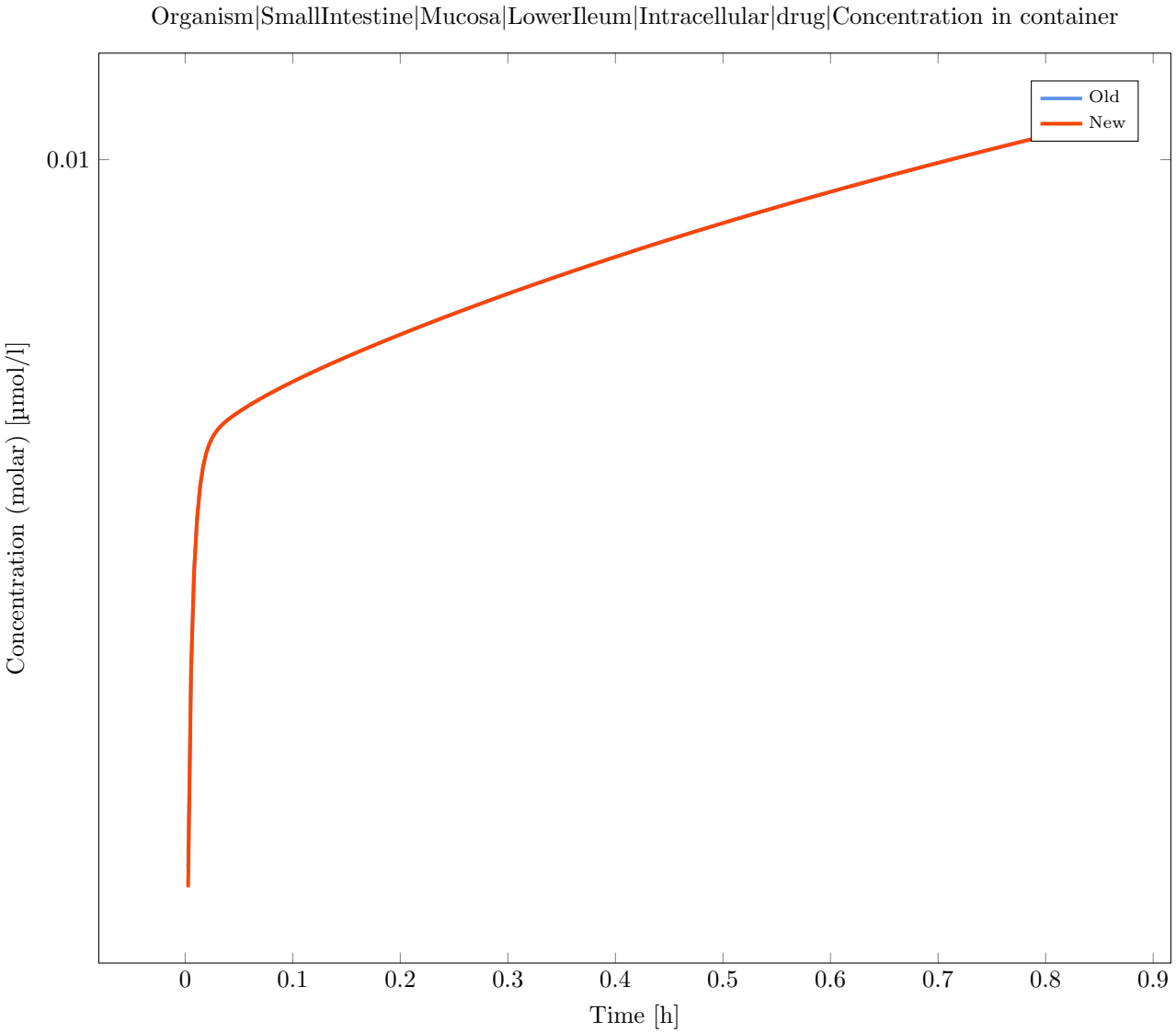
Organism|SmallIntestine|Mucosa|LowerIleum|Intracellular|drug|Concentration in container
Old
New
0.01
Concentration (molar) [µmol/l]
0
0.1
0.2
0.3
0.4
0.5
0.6
0.7
0.8
0.9
Time [h]

Figure 1.97

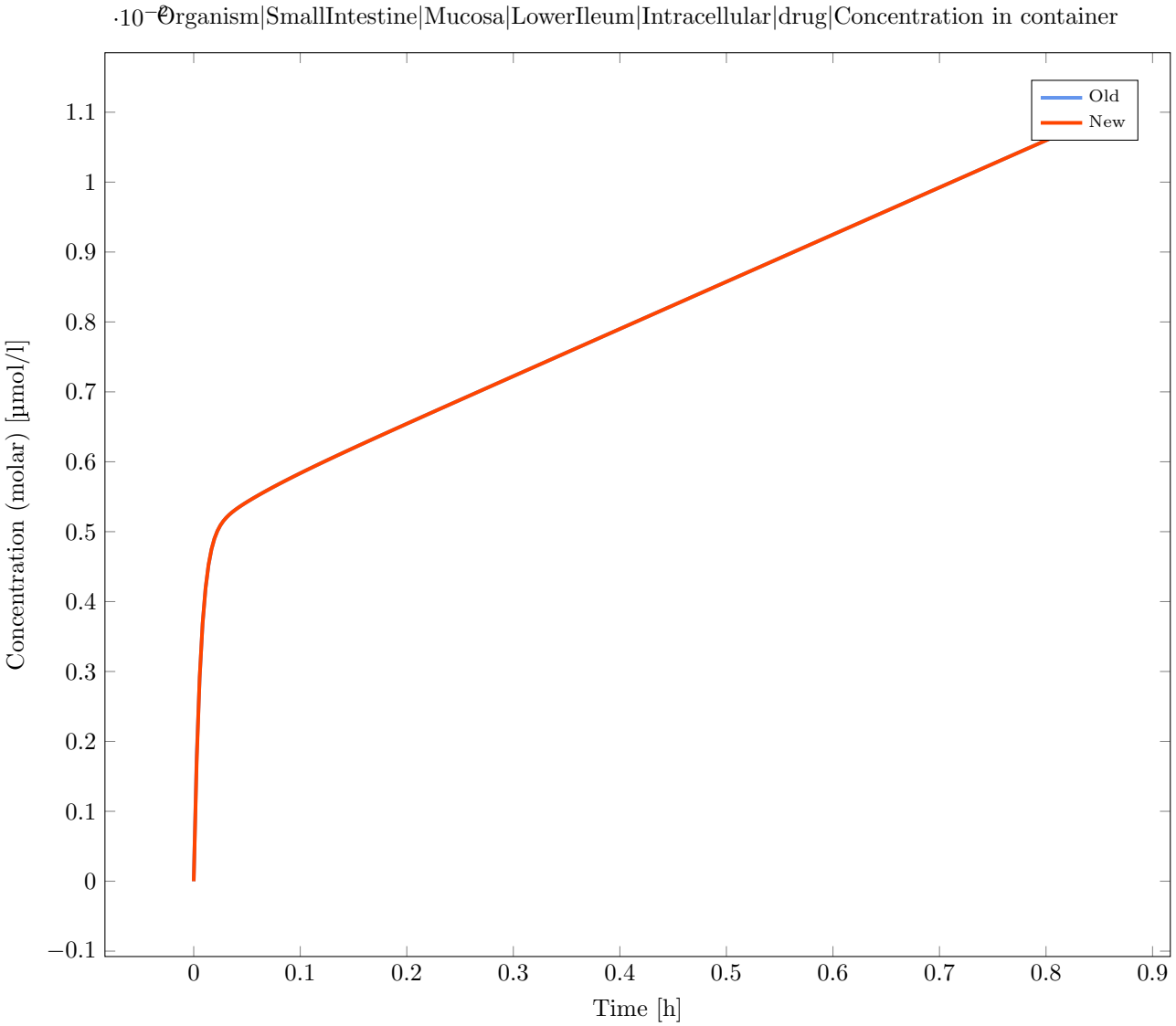

Figure 1.98

Output Path: Organism|LargeIntestine|Mucosa|ColonDescendens|Intracellular|drug|Concentration in container
Deviation for 'Organism|LargeIntestine|Mucosa|ColonDescendens|Intracellular|drug|Concentration in container' is 3.84% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.04

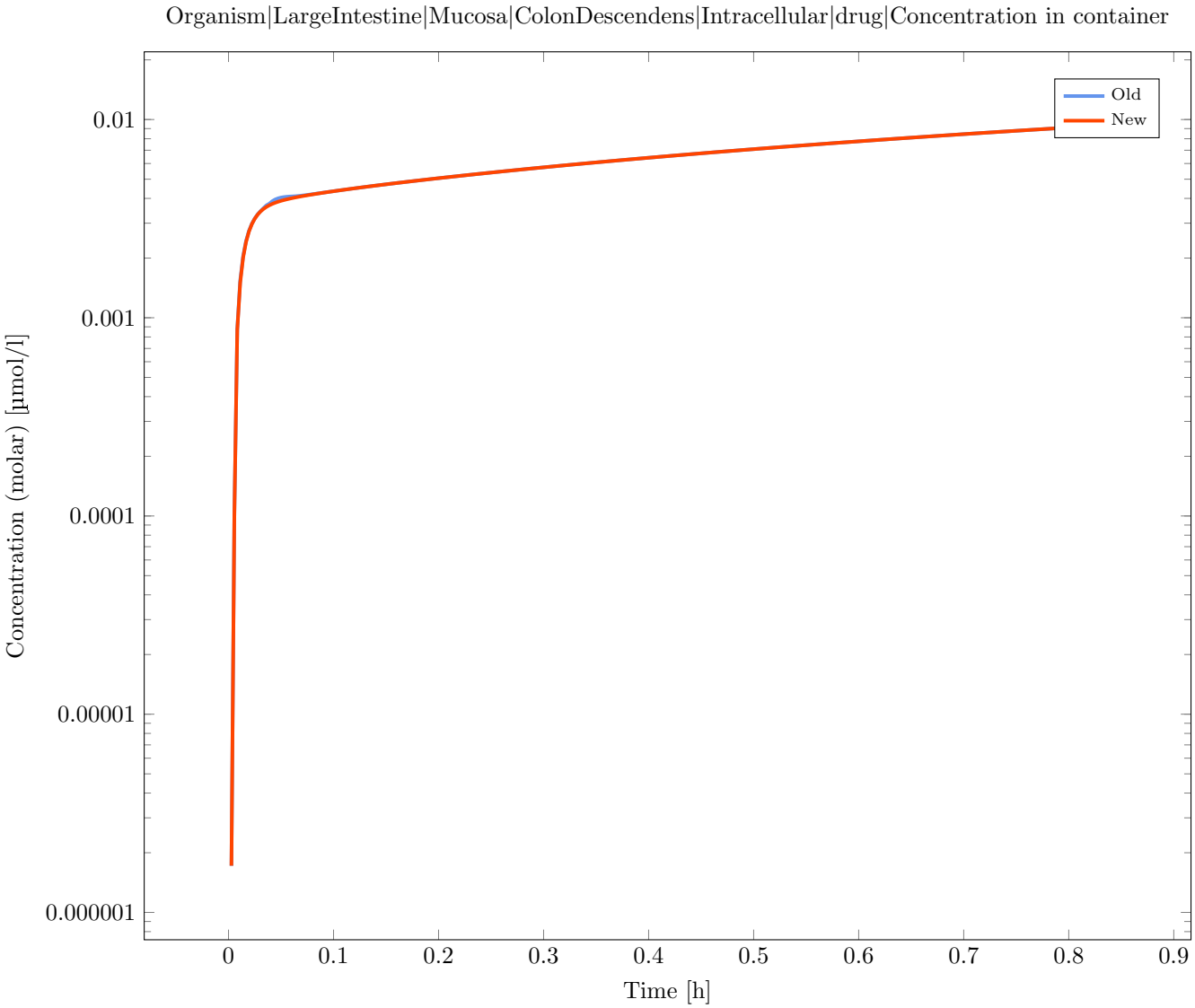
Organism|LargeIntestine|Mucosa|ColonDescendens|Intracellular|drug|Concentration in container
Old
New
Concentration (molar) [µmol/l]
0.01
0.001
0.0001
0.00001
0.000001
0
0.1
0.2
0.3
0.4
0.5
0.6
0.7
0.8
0.9
Time [h]

Figure 1.99

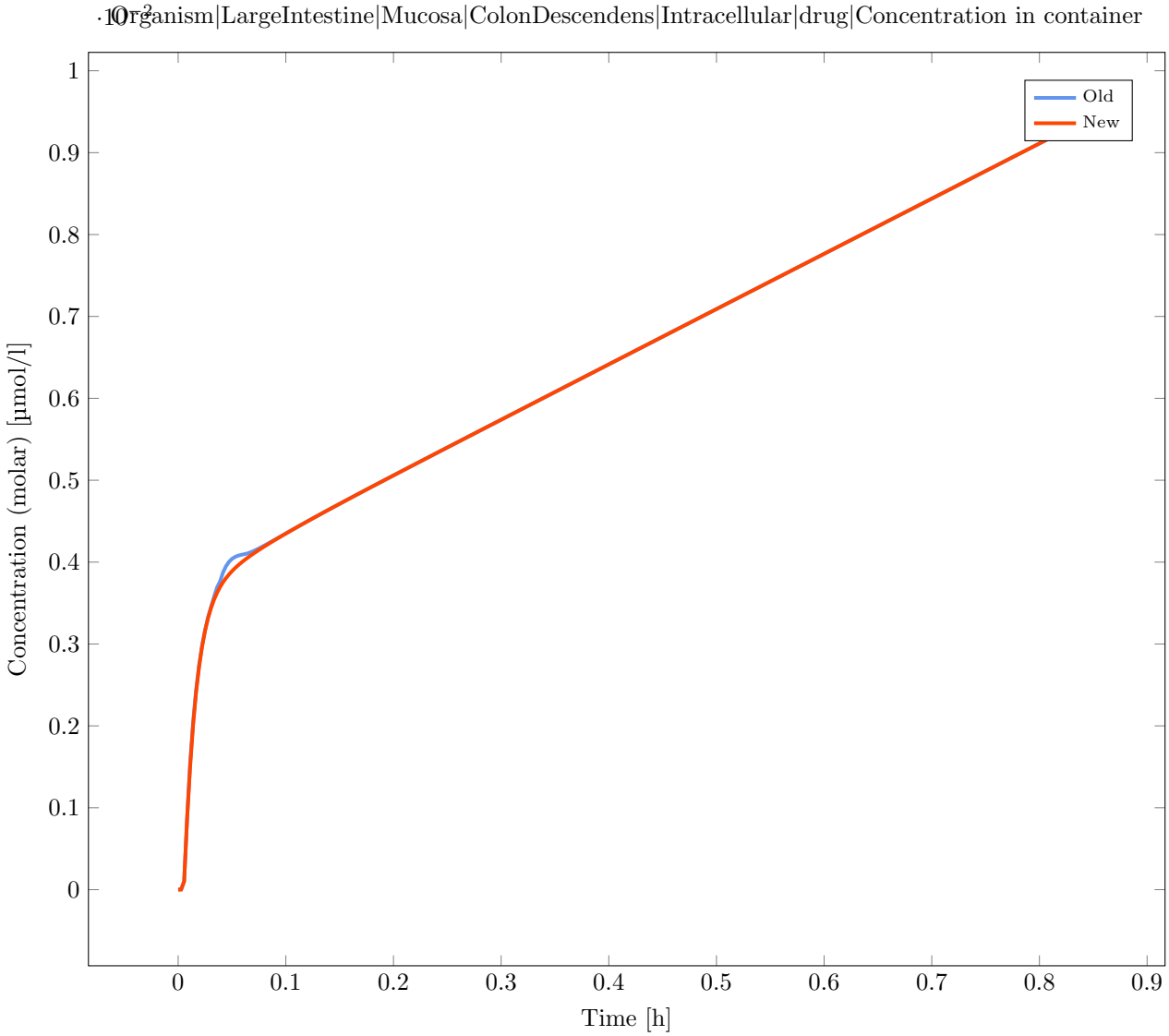

Figure 1.100

Simulation: Rat_MultiORAL_8_8_8_Dissolved-Rat_MultiORAL_8_8_8_Dissolved
Result of the validation: Invalid
Absolute Tolerance: 1.00E-10
Relative Tolerance: 1.00E-5

Output Path: Organism|SmallIntestine|Mucosa|UpperIleum|Intracellular|drug|Concentration in container
Deviation for 'Organism|SmallIntestine|Mucosa|UpperIleum|Intracellular|drug|Concentration in container' is 4.86% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.05

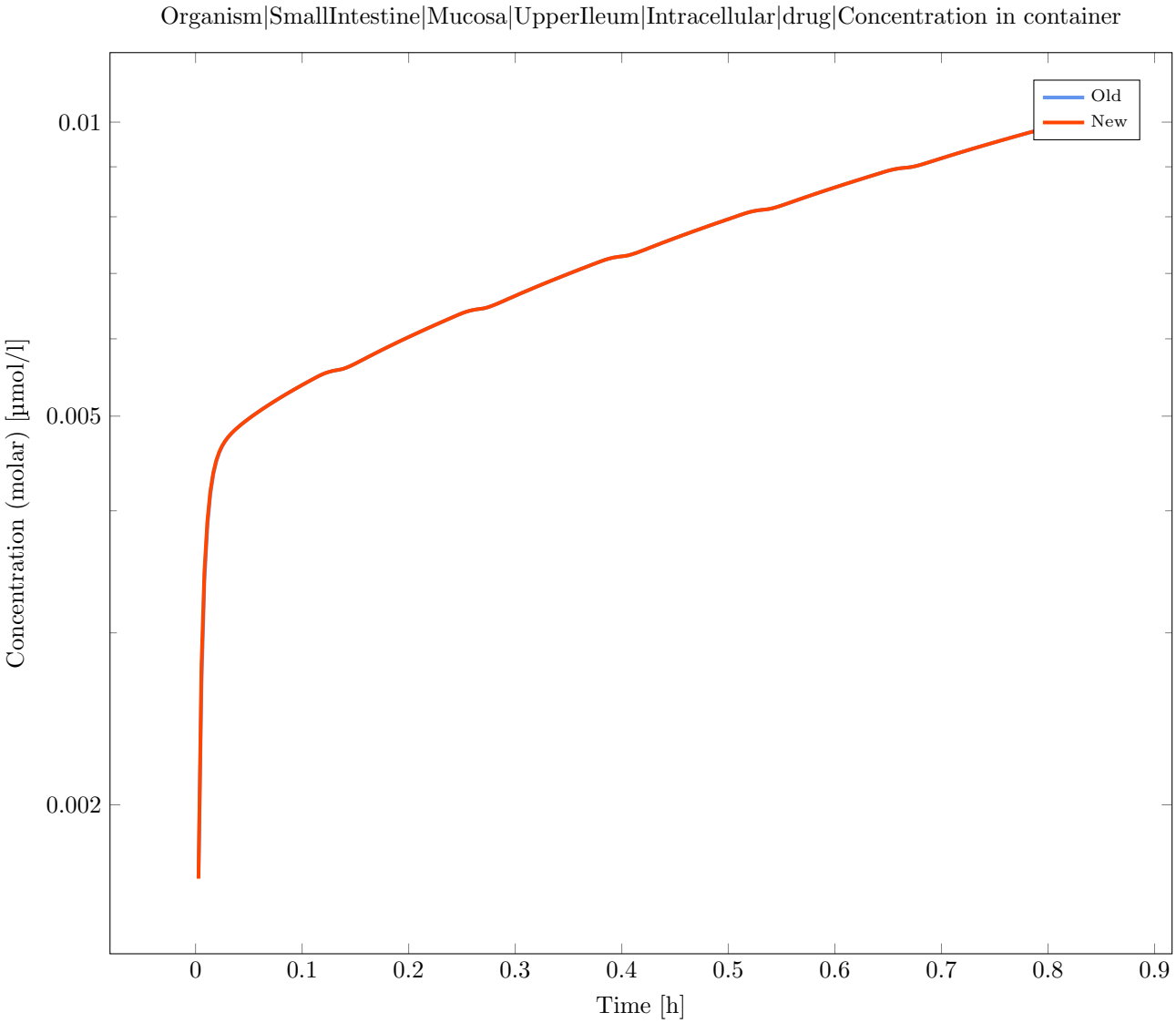
Organism|SmallIntestine|Mucosa|UpperIleum|Intracellular|drug|Concentration in container
Old
New
Concentration (molar) [µmol/l]
0.01
0.005
0.002
0
0.1
0.2
0.3
0.4
0.5
0.6
0.7
0.8
0.9
Time [h]

Figure 1.101

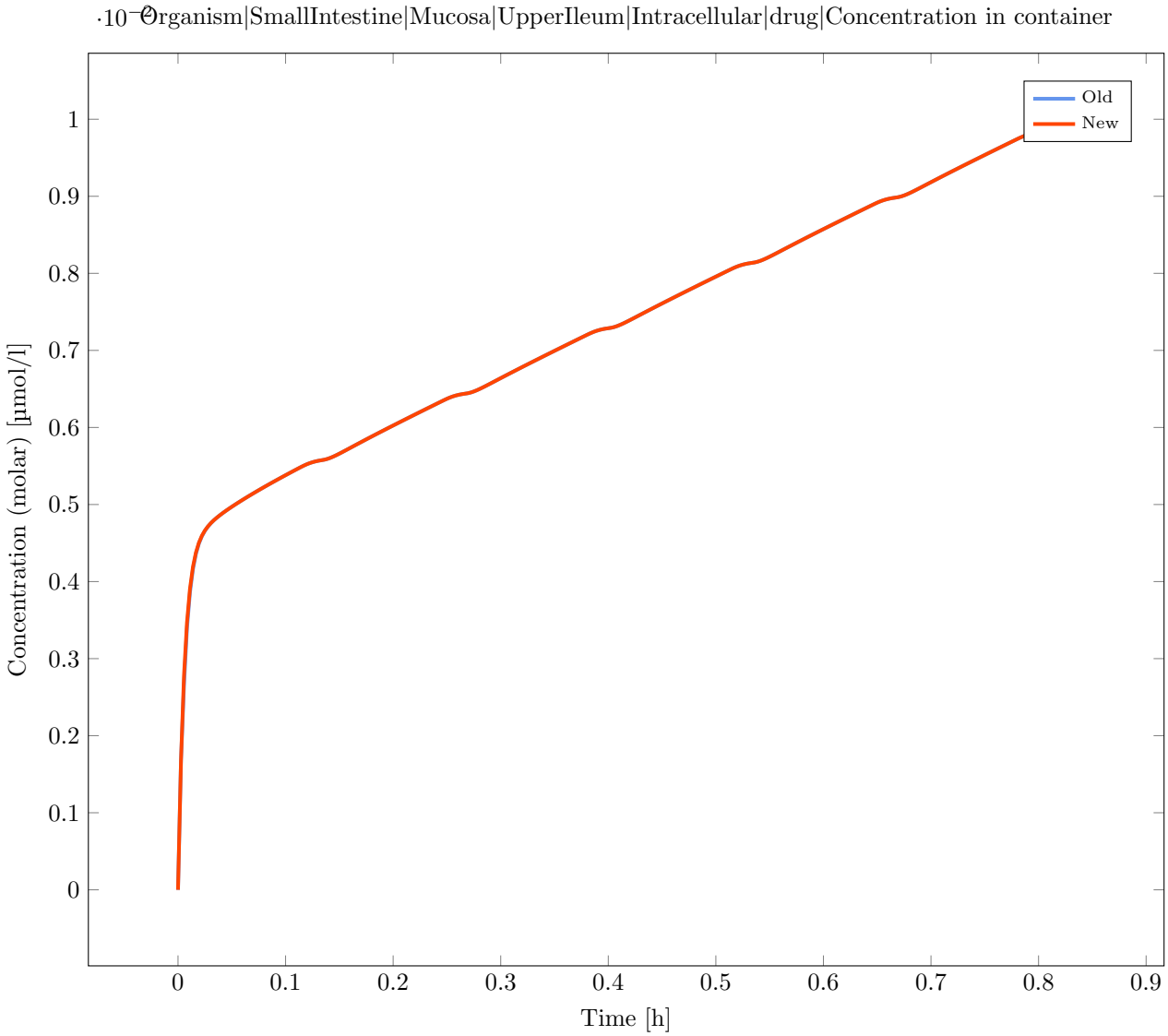

Figure 1.102

Output Path: Organism|SmallIntestine|Mucosa|LowerIleum|Intracellular|drug|Concentration in container
Deviation for 'Organism|SmallIntestine|Mucosa|LowerIleum|Intracellular|drug|Concentration in container' is 4.77% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.05

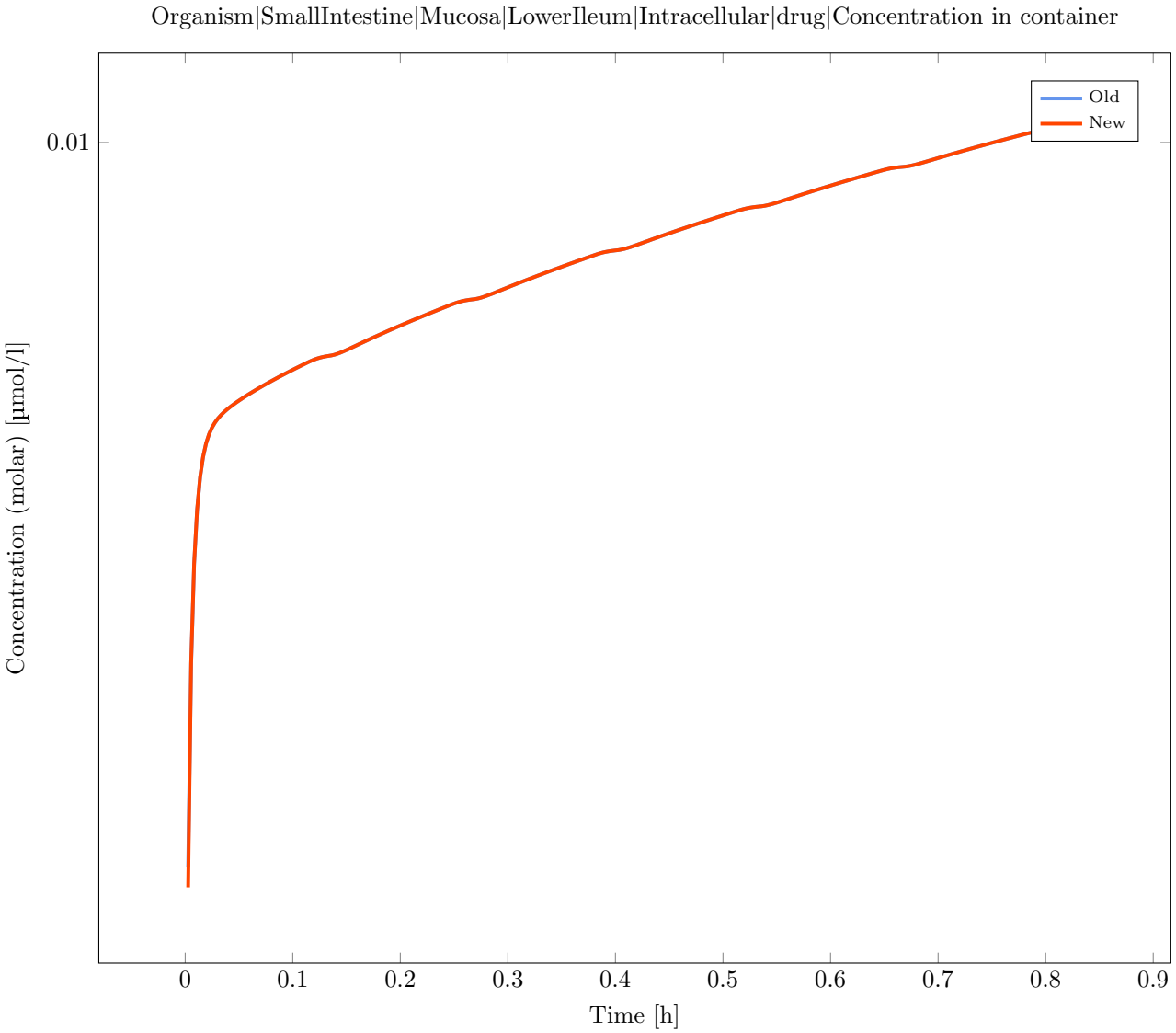
Organism|SmallIntestine|Mucosa|LowerIleum|Intracellular|drug|Concentration in container
Old
New
0.01
Concentration (molar) [µmol/l]
0
0.1
0.2
0.3
0.4
0.5
0.6
0.7
0.8
0.9
Time [h]

Figure 1.103

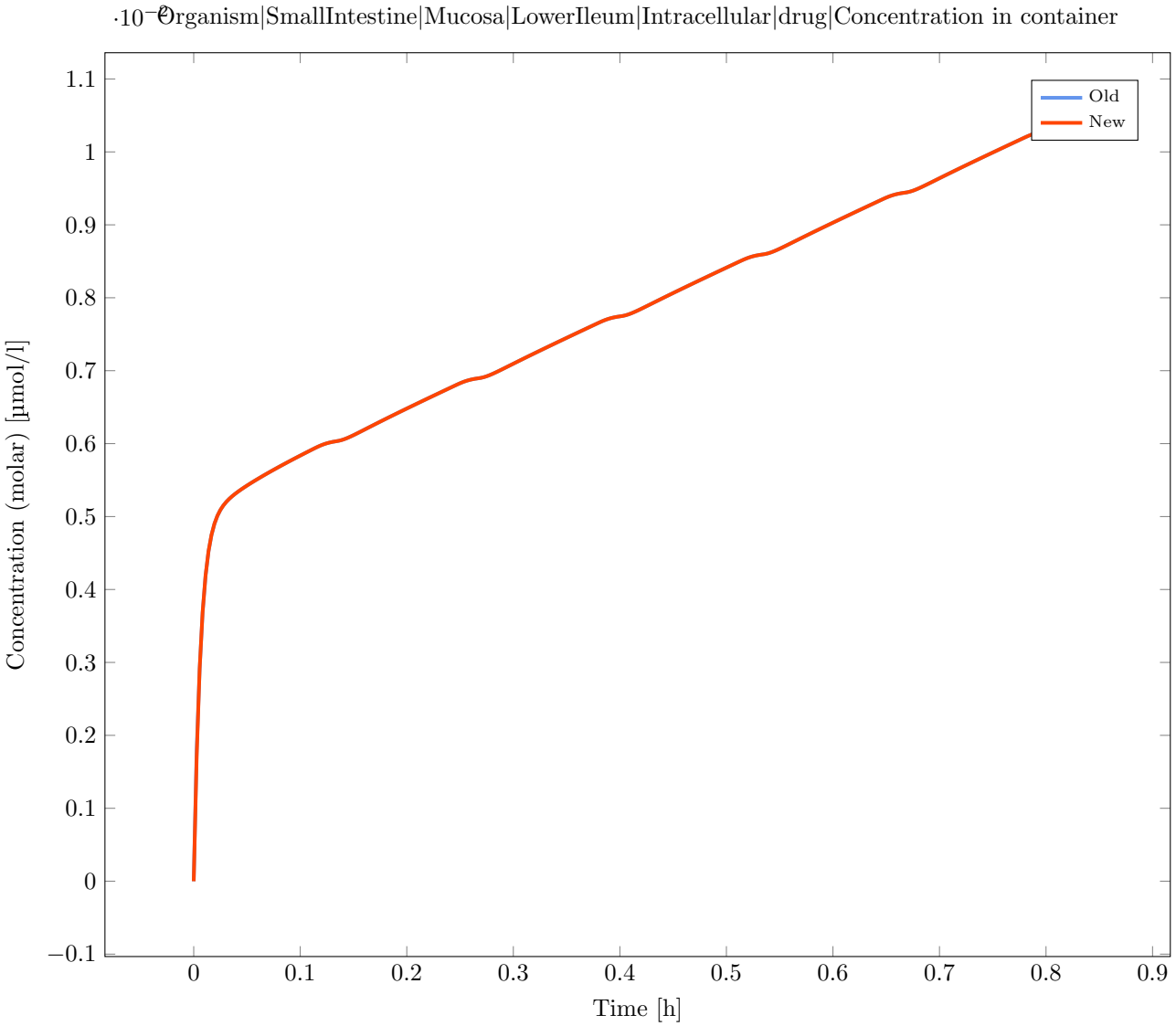

Figure 1.104

Output Path: Organism|LargeIntestine|Mucosa|ColonDescendens|Intracellular|drug|Concentration in container
Deviation for 'Organism|LargeIntestine|Mucosa|ColonDescendens|Intracellular|drug|Concentration in container' is 3.84% and is greater than the allowed max. tolerance of 3.00%
Deviation: 0.04

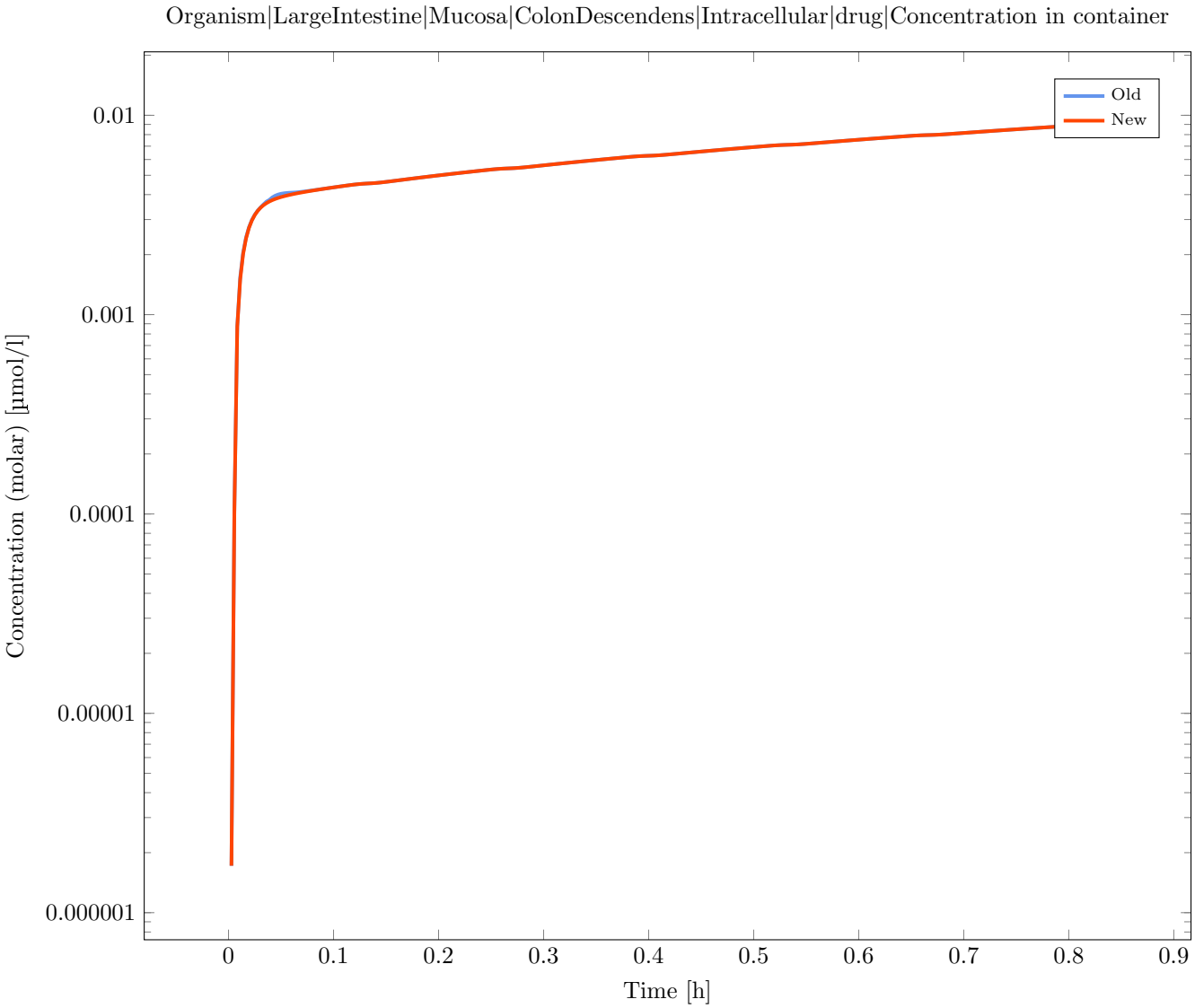
Organism|LargeIntestine|Mucosa|ColonDescendens|Intracellular|drug|Concentration in container
Old
New
Concentration (molar) [µmol/l]
0.01
0.001
0.0001
0.00001
0.000001
0
0.1
0.2
0.3
0.4
0.5
0.6
0.7
0.8
0.9
Time [h]

Figure 1.105

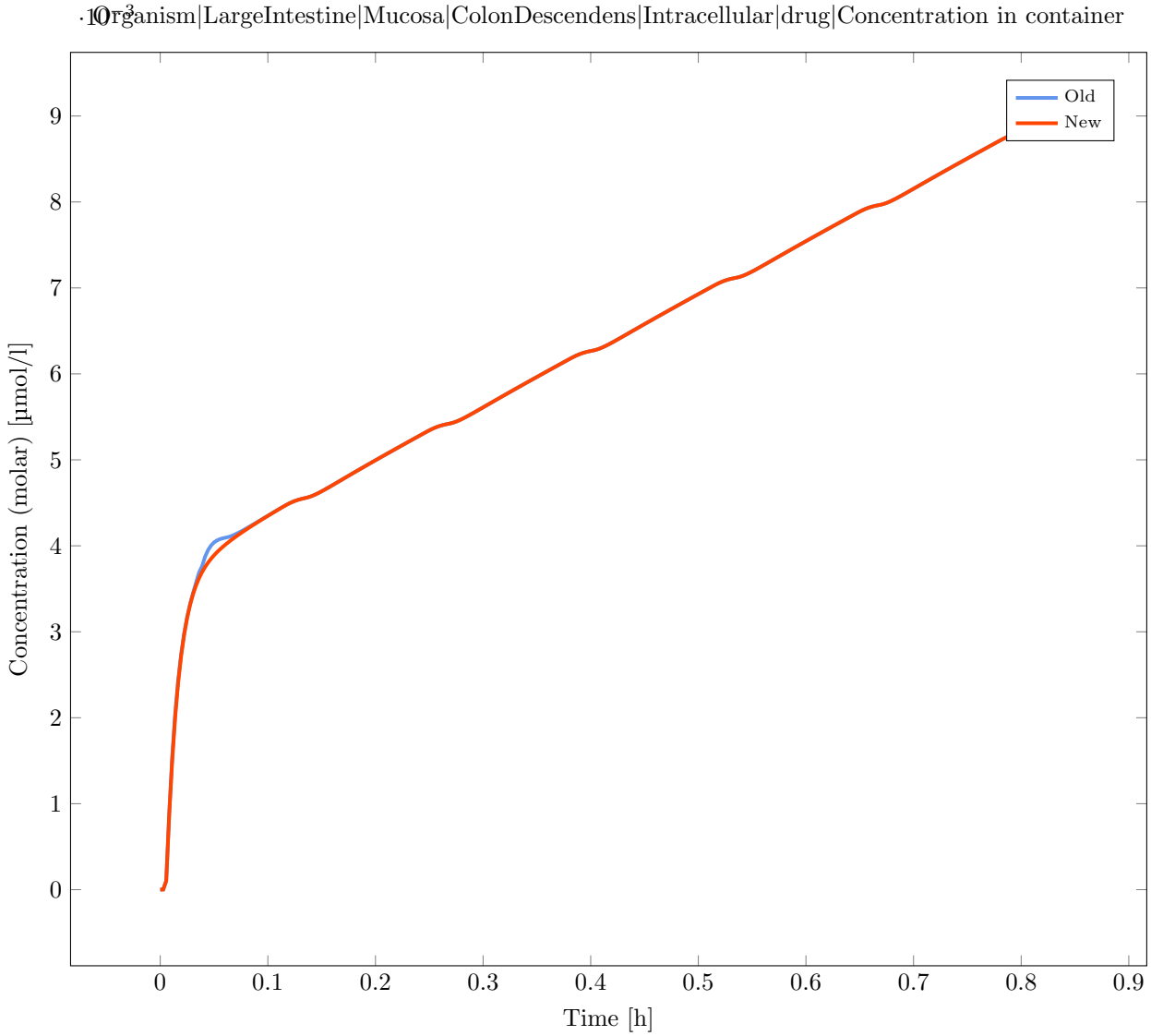

Figure 1.106

1.1.2 Valid Simulations (142/155)

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-05_MM_Mechanismbased_Mechanismbased
Result of the validation: Valid

Simulation: DDI_MultipleCombinations-06_MM_Induction_Induction
Result of the validation: Valid

Simulation: DDI_MultipleCombinations-07_MM_Competitive_Competitive_Mechanismbased_Mechanismbas
Result of the validation: Valid

Simulation: DDI_MultipleCombinations-08_MM_Uncompetitive_Uncompetitive_Mechanismbased_-Mechanismbased
Result of the validation: Valid

Simulation: DDI_MultipleCombinations-09_MM_Noncompetitive_Noncompetitive_Mechanismbased_-Mechanismbased
Result of the validation: Valid

Simulation: DDI_MultipleCombinations-10_MM_Mixed_Mixed_Mechanismbased_Mechanismbased
Result of the validation: Valid

Simulation: DDI_MultipleCombinations-11_MM_Mechanismbased_Mechanismbased_Induction_-Induction
Result of the validation: Valid

Simulation: DDI_MultipleCombinations-12_MM_All_DDI_Types
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: DDI_MultipleCombinations-25_1st_Mechanismbased_Mechanismbased
Result of the validation: Valid

Simulation: DDI_MultipleCombinations-26_1st_Induction_Induction
Result of the validation: Valid

Simulation: DDI_MultipleCombinations-27_1st_Competitive_Competitive_Mechanismbased_Mechanismbased
Result of the validation: Valid

Simulation: DDI_MultipleCombinations-28_1st_Uncompetitive_Uncompetitive_Mechanismbased_-Mechanismbased
Result of the validation: Valid

Simulation: DDI_MultipleCombinations-29_1st_Noncompetitive_Noncompetitive_Mechanismbased_-Mechanismbased
Result of the validation: Valid

Simulation: DDI_MultipleCombinations-30_1st_Mixed_Mixed_Mechanismbased_Mechanismbased
Result of the validation: Valid

Simulation: DDI_MultipleCombinations-31_1st_Mechanismbased_Mechanismbased_Induction_Induction
Result of the validation: Valid

Simulation: DDI_MultipleCombinations-32_1st_All_DDI_Types
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_MultipleIV_Binding-Human_MultipleIV_Binding
Result of the validation: Valid

Simulation: Human_MultipleIV_EffluxBasolateral-Human_MultipleIV_EffluxBasolateral
Result of the validation: Valid

Simulation: Human_MultipleIV_Efflux-Human_MultipleIV_Efflux
Result of the validation: Valid

Simulation: Human_MultipleIV_InfluxBasolateral-Human_MultipleIV_InfluxBasolateral
Result of the validation: Valid

Simulation: Human_MultipleIV_Influx-Human_MultipleIV_ActiveInflux
Result of the validation: Valid

Simulation: Human_MultipleIV_MetabolizmBinding-Human_MultipleIV_MetabolizmBinding
Result of the validation: Valid

Simulation: Human_MultipleIV_Metabolizm-Human_MultipleIV_Metabolizm
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_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_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_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: 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