

Supporting Information

Efficiency Assessment on Single Unit Monomer Insertion Reactions for Monomer Sequence Control: Kinetic Simulations and Experimental Observations

Joris J. Haven,^a Joke Vandenberghe,^a Rafael Kurita,^{a,b} Jonas Gruber,^b Tanja Junkers^{a,c}*

^aPolymer Reaction Design Group, Institute for Materials Research (imo-imomec), Hasselt University, Campus Diepenbeek, Building D, B-3590 Diepenbeek, Belgium

^b Escola Politécnica da Universidade de São Paulo, Avenida Professor Luciano Gualberto, Travessa 3, nº 380, Butantã - São Paulo, 05508-010, Brazil

^cIMEC division IMOMEC, Wetenschapspark 1, B-3590 Diepenbeek, Belgium

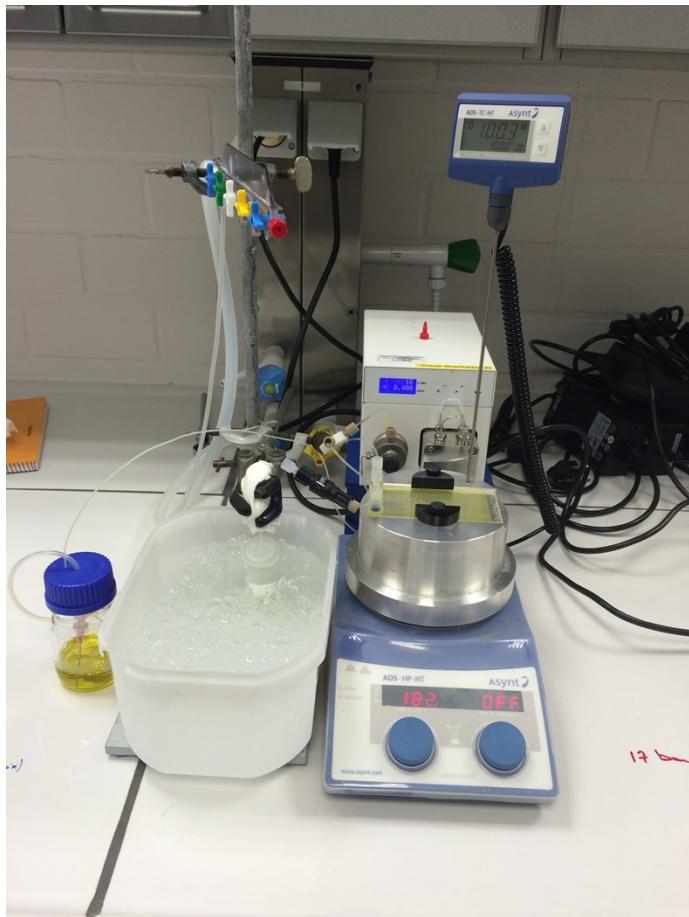


Figure S1. Image of the mesoreactor setup utilized for the upscaled SUMI synthesis.

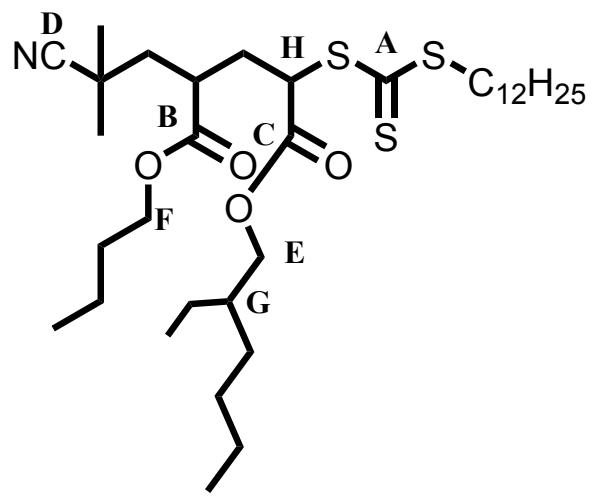


Figure S2. ¹³CNMR peak assignments of the SUMI-2AB product (see Figure S3, S4 and S5).

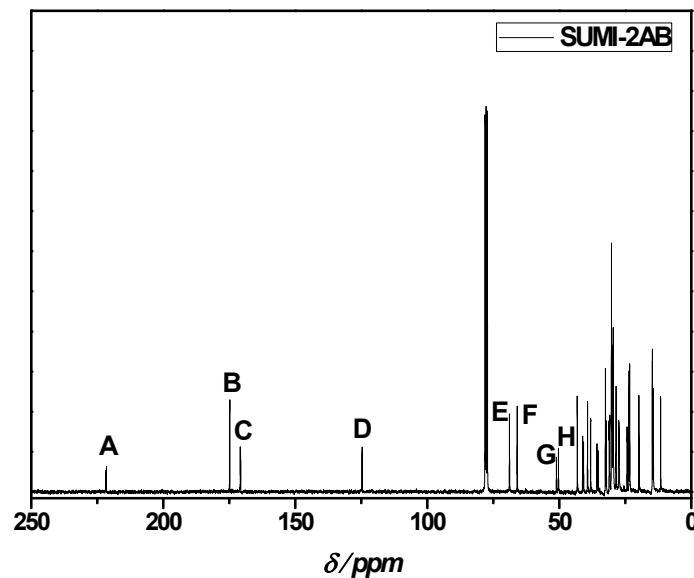


Figure S3. ¹³C NMR spectrum of **SUMI-2AB** (*n*BA-EHA oligo-RAFT agent).

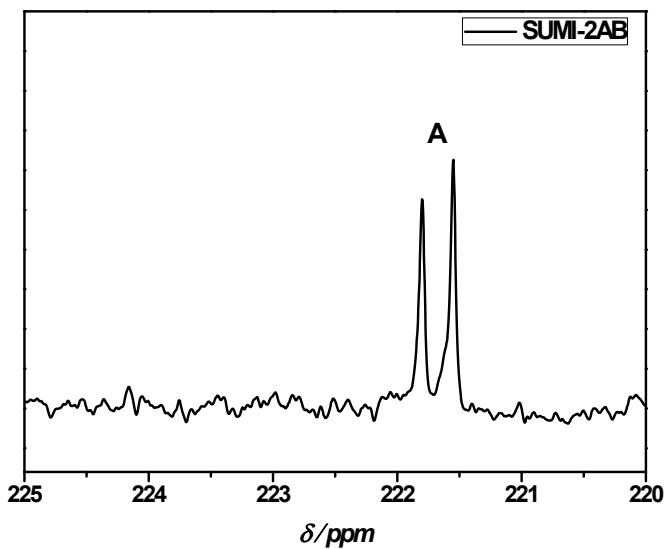


Figure S4. Zoom into ^{13}C NMR spectrum of **SUMI-2AB** (*n*BA-EHA oligo-RAFT agent), The two peaks can be assigned to different diastereomers.

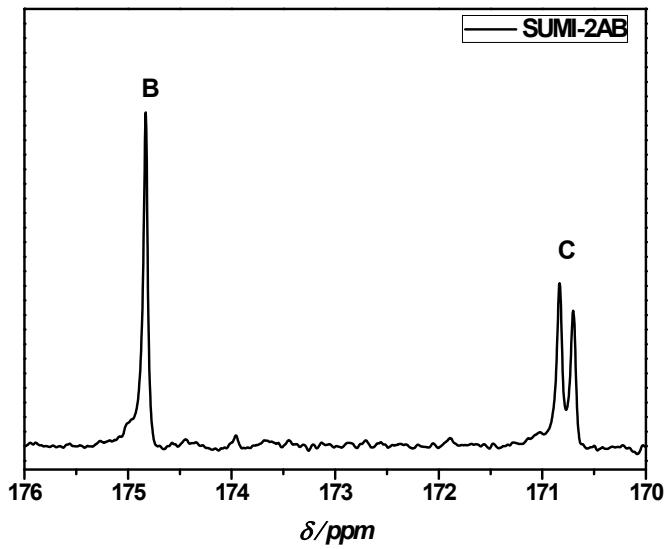


Figure S5. Zoom into ^{13}C NMR spectrum of **SUMI-2AB** (*n*BA-EHA oligo-RAFT agent), the two peaks (C) can be assigned to different diastereomers.

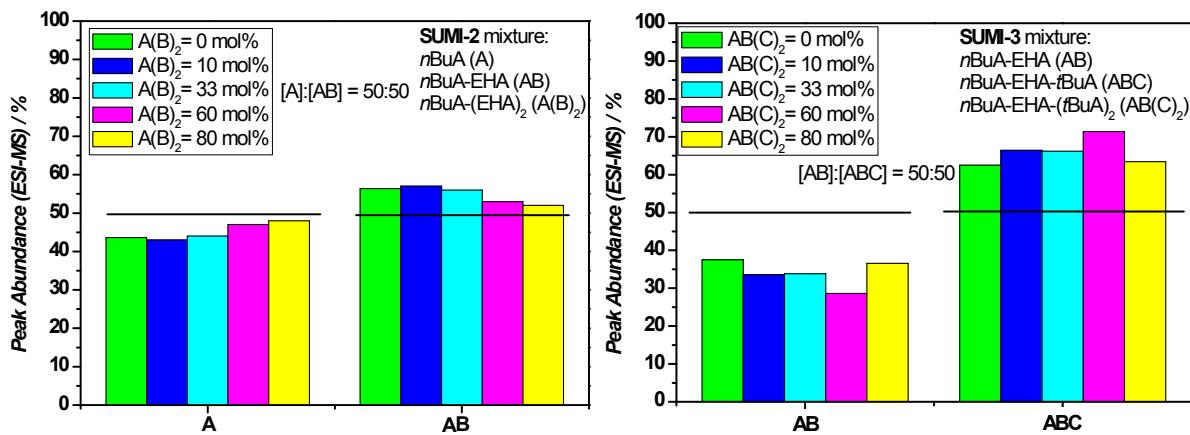


Figure S6. Left: ESI-MS peak abundance of A and AB for a constant A:AB molar ratio in the **SUMI-2** calibration mixture while the moles of A(B)₂ are varied. Right: ESI-MS peak abundance of AB and ABC for a constant AB:ABC molar ratio in the **SUMI-3** calibration mixture while mol% AB(C)₂ is varied.

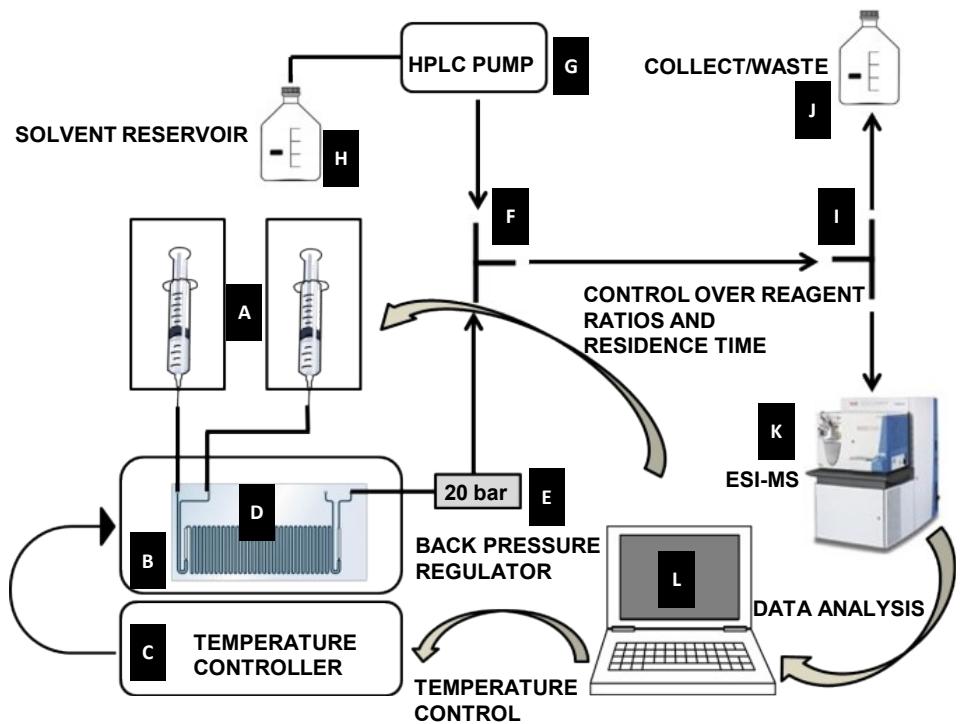


Figure S7. Representation of a flow chart of the on-line setup (top) and image of the on-line setup (bottom).

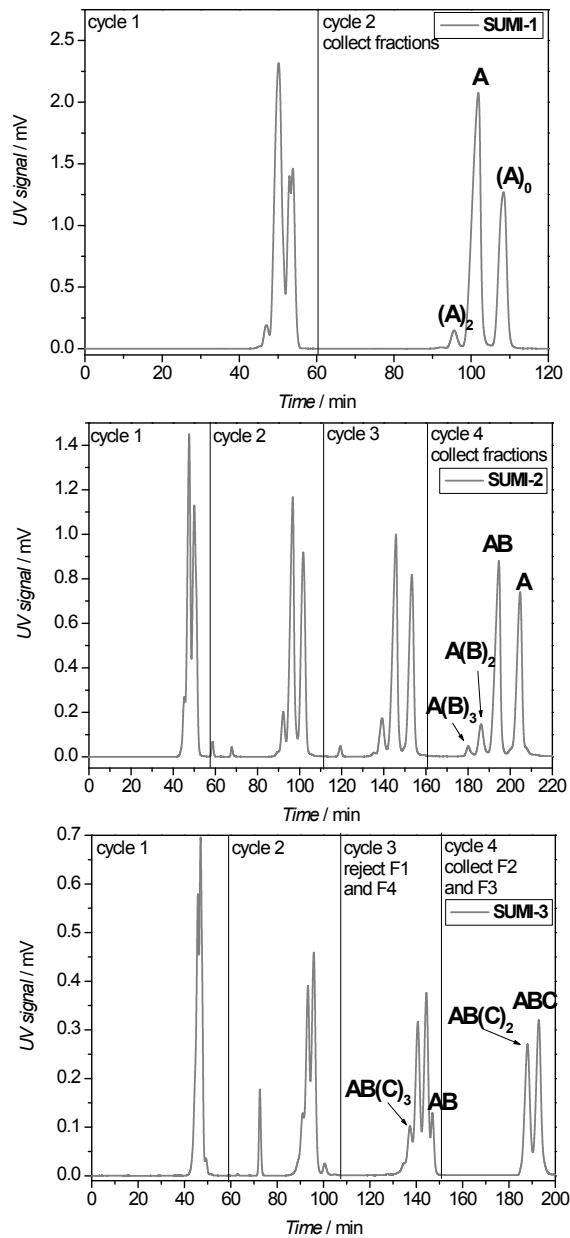
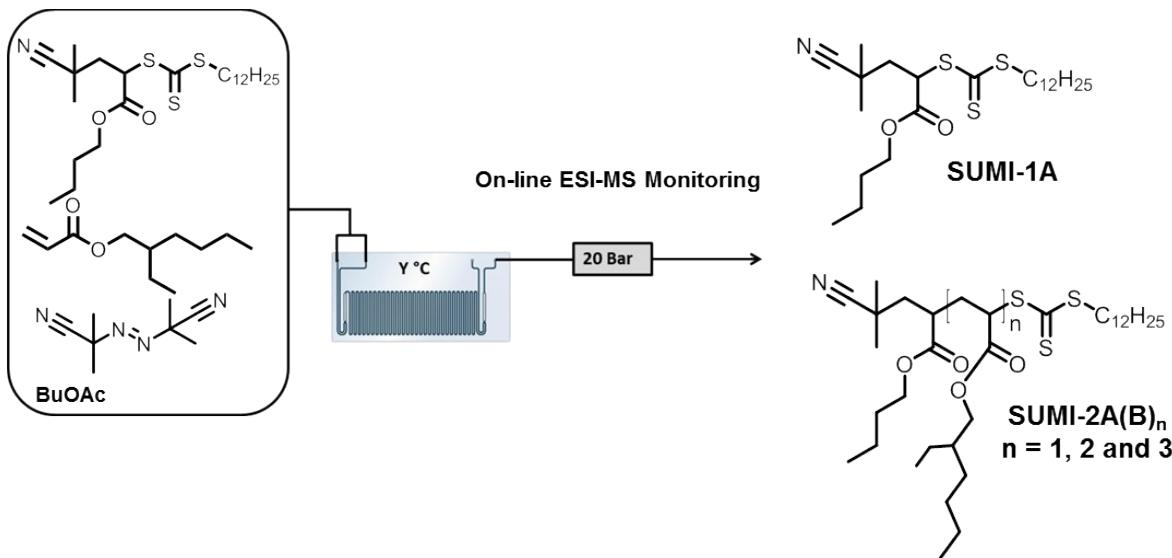


Figure S8. Recycling SEC trace recorded during consecutive purification cycles of **SUMI-1**, **SUMI-2** and **SUMI-3**. (data partially previously published in [2])



Scheme S1. Schematic representation of the reaction products for the microreactor synthesis of **SUMI-2A**. (previously published in [1])

Table S1. Microreactor screening conditions and ESI-MS peak abundances for the synthesis of **SUMI-2AB**. (previously published in [1])

Condition	Temp. (°C)	[EHA] : [SUMI-1A]	Residence Time (min)	A (%)	AB (%)	A(B) ₂ (%)	A(B) ₃ (%)
1	95	5:1	10	50	38	10	2
2	95	10:1	5	41	50	9	0
3	95	10:1	7,5	11	58	27	4
4	100	1:1	5	70	30	0	0
5	100	2:1	5	47	49	4	0
6	100	2:1	10	22	62	16	0
7	100	3:1	8	9	61	27	3
8	100	4:1	6	12	60	25	3
9	100	5:1	2,5	40	54	6	0
10	100	5:1	4	16	62	20	2
11	100	5:1	5	9	61	26	4
12	110	3:1	2,5	19	65	16	0
13	110	3:1	4	9	68	19	4
14	110	2:1	4	28	61	11	0
15	110	2:1	5	20	64	16	0

Table S2. Microreactor screening conditions and molar ratios of all insertion products for the synthesis of **SUMI-2AB**.* (primary experimental results previously published in [1])

Condition	Temp. (°C)	[EHA] : [SUMI-1A]	Residence Time (min)	A (%)	AB (%)	A(B) ₂ (%)	A(B) ₃ (%)
1	95	5:1	10	57	31	10	2
2	95	10:1	5	48	43	9	0
3	95	10:1	7,5	13	56	27	4
4	100	1:1	5	78	22	0	0
5	100	2:1	5	56	40	4	0
6	100	2:1	10	27	57	16	0
7	100	3:1	8	11	59	27	3
8	100	4:1	6	13	59	25	3
9	100	5:1	2,5	48	46	6	0
10	100	5:1	4	20	58	20	2
11	100	5:1	5	11	59	26	4
12	110	3:1	2,5	24	60	16	0
13	110	3:1	4	11	66	19	4
14	110	2:1	4	35	54	11	0
15	110	2:1	5	24	60	16	0

* It is assumed that peak abundances of A(B)₂ and A(B)₃ match the true molar ratio.

Table S3. Theoretical and isolated yields for **SUMI2** and **SUMI3** products, synthesized with different setups. (part of experimental results already previously published in [1] and [2])

Setup	Desired SUMI	Theoretical Yield (%)	Isolated Yield (%)
Batch	SUMI-2AB	50	46
Batch	SUMI-3ABC	30	20
Batch	SUMI-2AC	n.a.	50
Batch	SUMI-3ACB	n.a.	20
Microflow	SUMI-2AB	59	43
Microflow	SUMI-3ABC	34	n.a.
Mesoflow	SUMI-2AB	53	48
Mesoflow	SUMI-3ABC	22	20

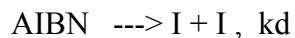
Overview Predici model

Coefficients (CL-independent model)

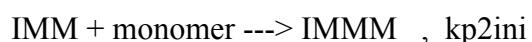
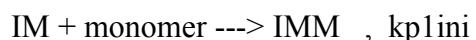
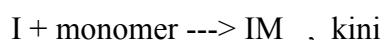
k_t	$1 \cdot 10^9 \text{ L} \cdot \text{mol}^{-1} \cdot \text{s}^{-1}$
k_d	$A = 1.580 \text{e+15 s}^{-1}; E_A/R = 1.551 \text{e+04 J/mol}$
k_{p1}	$7 \cdot 10^4 \text{ L} \cdot \text{mol}^{-1} \cdot \text{s}^{-1}$
k_{p2}	$7 \cdot 10^4 \text{ L} \cdot \text{mol}^{-1} \cdot \text{s}^{-1}$
k_{p3}	$7 \cdot 10^4 \text{ L} \cdot \text{mol}^{-1} \cdot \text{s}^{-1}$
k_{ad1}	$7 \cdot 10^6 \text{ L} \cdot \text{mol}^{-1} \cdot \text{s}^{-1}$
k_{ad2}	$7 \cdot 10^6 \text{ L} \cdot \text{mol}^{-1} \cdot \text{s}^{-1}$
k_{ad3}	$7 \cdot 10^6 \text{ L} \cdot \text{mol}^{-1} \cdot \text{s}^{-1}$
k_{ad4}	$7 \cdot 10^6 \text{ L} \cdot \text{mol}^{-1} \cdot \text{s}^{-1}$
k_{frag1}	$7 \cdot 10^3 \text{ s}^{-1}$
k_{frag2}	$7 \cdot 10^3 \text{ s}^{-1}$
k_{frag3}	$7 \cdot 10^3 \text{ s}^{-1}$
k_{frag4}	$7 \cdot 10^3 \text{ s}^{-1}$
k_{p1ini}	$7 \cdot 10^4 \text{ L} \cdot \text{mol}^{-1} \cdot \text{s}^{-1}$
k_{p2ini}	$7 \cdot 10^4 \text{ L} \cdot \text{mol}^{-1} \cdot \text{s}^{-1}$
k_{p3ini}	$7 \cdot 10^4 \text{ L} \cdot \text{mol}^{-1} \cdot \text{s}^{-1}$

Reaction Modules

Initiator decomposition



Propagation



IMMM + monomer ---> IMMM , kp3ini

RM + monomer ---> RMM , kp1

RMM + monomer ---> RMMM , kp2

RMMM + monomer ---> RMMMM , kp3

SUMI-RAFT equilibria

I + RMTTC ---> INT(ISMR) , kad1

IM + RMTTC ---> INT(ISMR) , kad1

IMM + RMTTC ---> INT(ISMR) , kad2

IMMM + RMTTC ---> INT(ISMR) , kad3

IMMM + RMTTC ---> INT(ISMR) , kad4

I + RMMTTC ---> INT(ISMMR) , kad2

IM + RMMTTC ---> INT(ISMMR) , kad2

IMM + RMMTTC ---> INT(ISMMR) , kad2

IMMM + RMMTTC ---> INT(ISMMR) , kad3

IMMM + RMMTTC ---> INT(ISMMR) , kad4

I + RMMMTTC ---> INT(ISMMMR) , kad3

IM + RMMMTTC ---> INT(ISMMMR) , kad3

IMM + RMMMTTC ---> INT(ISMMMR) , kad3

IMMM + RMMMTTC ---> INT(ISMMMR) , kad4

I + RMMMMTTC ---> INT(ISMMMMR) , kad4

IM + RMMMMTTC ---> INT(ISMMMMR) , kad4

IMM + RMMMMTTC ---> INT(ISMMMMR) , kad4

IMMM + RMMMMTTC ---> INT(ISMMMMR) , kad4

INT(ISMR) ---> RM + Dead , kfrag1

INT(ISMR) ---> RMTTC + Dead , kfrag1

INT(ISMMR) ---> RMM + Dead , kfrag2
 INT(ISMMR) ---> RMMTTC + Dead , kfrag2
 INT(ISMMMR) ---> RMMM + Dead , kfrag3
 INT(ISMMMR) ---> RMMTTC + Dead , kfrag3
 INT(ISMMMMR) ---> RMMMM + Dead , kfrag4
 INT(ISMMMMR) ---> RMMMTTC + Dead , kfrag4
 RM + RMTTC ---> INT(RMSMR) , kad1
 RMM + RMTTC ---> INT(RMMSMR) , kad2
 RMMM + RMTTC ---> INT(RMMMSMR) , kad3
 RMMMM + RMTTC ---> INT(RMMMSMR) , kad4
 INT(RMSMR) ---> RM + RMTTC , kfrag1
 INT(RMMSMR) ---> RMM + RMTTC , kfrag2
 INT(RMMMSMR) ---> RMMM + RMTTC , kfrag3
 INT(RMMMSMR) ---> RMMMM + RMTTC , kfrag4
 INT(RMSMR) ---> RMTTC + RM , kfrag1
 INT(RMMSMR) ---> RMMTTC + RM , kfrag1
 INT(RMMMSMR) ---> RMMTTC + RM , kfrag1
 INT(RMMMSMR) ---> RMMMTTC + RM , kfrag1
 RM + RMMTTC ---> INT(RMSMMR) , kad2
 RMM + RMMTTC ---> INT(RMMSMMR) , kad2
 RMMM + RMMTTC ---> INT(RMMMSMMR) , kad3
 RMMMM + RMMTTC ---> INT(RMMMSMMR) , kad4
 INT(RMSMMR) ---> RMMTTC + RM , kfrag1
 INT(RMMSMMR) ---> RMMTTC + RMM , kfrag2
 INT(RMMSMMR) ---> RMM + RMMTTC , kfrag2
 INT(RMMMSMMR) ---> RMMTTC + RMMM , kfrag3
 INT(RMMMSMMR) ---> RMMTTC + RMMMM , kfrag4

INT(RMSMMR) ---> RMTTC + RMM , kfrag2
 INT(RMMMSMMR) ---> RMMMTTC + RMM , kfrag2
 INT(RMMMSMMR) ---> RMMMTTC + RMM , kfrag2
 RM + RMMMTTC ---> INT(RMSMMR) , kad3
 RMM + RMMMTTC ---> INT(RMMSMMR) , kad3
 RMM + RMMMTTC ---> INT(RMMMSMMR) , kad3
 RMM + RMMMTTC ---> INT(RMMMSMMR) , kad4
 INT(RMSMMR) ---> RMMMTTC + RM , kfrag1
 INT(RMSMMR) ---> RMTTC + RMM , kfrag3
 INT(RMMSMMR) ---> RMMMTTC + RMM , kfrag2
 INT(RMMSMMR) ---> RMMTTC + RMM , kfrag3
 INT(RMMSMMR) ---> RMMMTTC + RMM , kfrag3
 INT(RMMSMMR) ---> RMM + RMMMTTC , kfrag3
 INT(RMMSMMR) ---> RMMMTTC + RMM , kfrag4
 INT(RMMSMMR) ---> RMMMTTC + RMM , kfrag3
 RM + RMMMTTC ---> INT(RMSMMMR) , kad4
 RMM + RMMMTTC ---> INT(RMMSMMMR) , kad4
 RMM + RMMMTTC ---> INT(RMMSMMMR) , kad4
 RMM + RMMMTTC ---> INT(RMMSMMMR) , kad4
 INT(RMSMMMR) ---> RMMMTTC + RM , kfrag1
 INT(RMSMMMR) ---> RMTTC + RMM , kfrag4
 INT(RMMSMMMR) ---> RMMMTTC + RMM , kfrag2
 INT(RMMSMMMR) ---> RMMTTC + RMM , kfrag4
 INT(RMMSMMMR) ---> RMMMTTC + RMM , kfrag4
 INT(RMMSMMMR) ---> RMMMTTC + RMM , kfrag3
 INT(RMMSMMMR) ---> RMMMTTC + RMM , kfrag4
 INT(RMMSMMMR) ---> RMM + RMMMTTC , kfrag4

Termination

I + I ---> Dead , kt

I + RM ---> Dead , kt

I + RMM ---> Dead , kt

I + RMMM ---> Dead , kt

I + RMMMM ---> Dead , kt

IM + RM ---> Dead , kt

IM + RMM ---> Dead , kt

IM + RMMM ---> Dead , kt

IM + RMMMM ---> Dead , kt

IMM + RM ---> Dead , kt

IMM + RMM ---> Dead , kt

IMM + RMMM ---> Dead , kt

IMM + RMMMM ---> Dead , kt

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RM + RM ---> Dead , kt

RM + RMM ---> Dead , kt

RM + RMMM ---> Dead , kt

RM + RMMMM ---> Dead , kt

RMM + RMM ---> Dead , kt

RMM + RMMM ---> Dead , kt

RMM + RMMMM ---> Dead , kt

RMMM + RMMM ---> Dead , kt

RMMM + RMMMM ---> Dead , kt

RMMMM + RMMMM ---> Dead , kt

No efficiency factors have been used and all reactions are considered explicitly.

References

- [1] J. Haven, J. Vandenberghe and T. Junkers, *Chem. Commun.*, 2015, 51, 4611–4614.
- [2] J. Vandenberghe, G. Reekmans, P. Adriaensens and T. Junkers, *Chem. Commun.*, 2013, 49, 10358–10360.