# **Electronic Supporting information**

# Rapid production of the anaesthetic mepivacaine through continuous, portable technology

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# **Reductive amination screening flow set-up**

To find the optimal conditions for the continuous reductive amination the following parameters were targeted: pressure, formic acid equivalents, temperature, and residence time.



Scheme S1 Flow set-up for the reductive amination step

#### Pressure effect

Entry	Residence Time (min)	Formaldehyde (eq.)	Molar Ratio (Pipecolic acid: Formic acid) eq.	Temperature (ºC)	BPR (8 bars)	Conversion (%)*
1			00	No	nd	
2	45	27.0	25	90	Yes	nd
3	45	27.9		120	No	nd
4				120	Yes	3

Table S1 Conditions: Pipecolic acid (1c) 10 mM, formic acid 250 mM. \*Conversion calculated by <sup>1</sup>H-NMR

## Formic acid equivalents

Entry	Residence Time (min)	Formaldehyde (eq.)	Molar Ratio (Pipecolic acid: Formic acid) eq.	Temperature (ºC)	BPR (bars)	Conversion (%)*
			55			
1	45	27.0		120	¥	4
	45	27.9	2650 (Neat)	120	Yes	
2						33
Table S2 Co	nditions: Pinecolic acid	(1c) 10 mM formic aci	d 250 mM *Conversion calculated	by <sup>1</sup> H-NMR		

id (1c) 10 mM, formic acid 250 mM. \*Conversion calculated by <sup>1</sup>H-NMR

## Temperature and residence time effect

Entry	Residence Time	Formaldehyde	Molar Ratio (Pipecolic acid:	Temperature	BPR	Conversion
	(min)	(eq.)	Formic acid) eq.	(ºC)	(bars)	(%)*
1	45	27.9	2650 (Neat)	150	8	>99

2	15	27.9	2650 (Neat)	150	8	80
Table S3 Conditi	ons: Pipecolic a	cid (1c) 10 mM. *Conversion	calculated by <sup>1</sup> H-NMR			

#### Formic acid equivalents vs residence time

Entry	Residence Time (min)	Formaldehyde (eq.)	Molar Ratio (Pipecolic acid: Formic acid) eq.	Temperature (ºC)	BPR (bars)	Conversion (%)*
1	45	27.9	1325 (50% in H2O)	150	8	>99
2	15	27.9	1325 (50% in H2O)	150	8	61
3	45	27.9	265 (10% in H2O)	150	8	84
4	15	27.9	265 (10% in H2O)	150	8	62

Table S4 Conditions: Pipecolic acid (1c) 10 mM. \*Conversion calculated by <sup>1</sup>H-NMR.

# Amide bond formation through acyl fluoride intermediate (batch)



Scheme S2 Amide bond formation trough acyl fluoride intermediate

Entry	PFP (eq.)	Base (eq.)	Amine (eq.)	Activation time (min)	Total reaction time	Conversion* (%)
1	1.1	DIPEA (2)	O-Toluidine (1.0)	30	1.5 days	33
2	1.1	DIPEA (4)	O-Toluidine (1.0)	30	4 days	31
3	1.1	DBU (2)	O-Toluidine (1.0)	30	4 days	22
4	1.1	DIPEA (2)	O-Toluidine (1.5)	30	4 days	44
5	3.0	DIPEA (2)	O-Toluidine (1.0)	30	4 days	73

Table S5 Reaction conditions: 250 mM (1c), 25 °C. \*Conversion measured by HPLC

## Amide bond formation biocatalytic attempt (batch)



Scheme S3 Biocatalytic amine bond formation using Mycobacterium Smegmatis (MsACT)

Entry	Acyl donor	Substrate	Enzyme	Conversion (%)*

1	( <b>4b</b> ) (10 mM)	2,6-Dimethylaniline (250 mM) (25 eq.)	MsAcT (WT)	nd
2			MsAcT-S11C	nd

Table S6 Conditions : [Enzyme] = 2.75 mg/mL, phosphate buffer 100 mM pH 8.0, 45 °C, 48h. \*Conversions calculated by HPLC. Enzymes were obtained following the protocols from the previous literature.<sup>1</sup>

### Process intensification and optimization of the reductive amination (flow)

Entry	Residence Time (min)	Formaldehyde (eq.)	Molar Ratio (methylpipecolinate: Formic acid) eq.	Temperature (ºC)	BPR (bars)	Conversion (%)*
1	45	27.9	53	150	8	>99
2	20	27.9	53	150	8	>99
3	10	27.9	53	150	8	>99
4	1	27.9	53	150	8	39

Conditions screening at 0.10 M of methylpipecolinate (1b) (Residence time effect)

Table S7 Reaction conditions: 0.1 M methylpipecolinate (1b) + 27.9 eq. of formaldehyde (pH adujusted with 15 % v/v solution of acetic acid). 20 % (v/v) solution of formic acid in water. Reactor volume was 10 mL for entries 1-3 and 1.2 mL for entry 4. Reactions were performed at 8 bars. Conversion were calculated by <sup>1</sup>H-NMR.

### Conditions screening at 0.83 M of methylpipecolinate (1b)

Entry	Residence Time (min)	Formaldehyde (eq.)	Molar Ratio (Methylpipecolinate Formic acid) eq.	Temperature (ºC)	BPR (bars)	Conversion (%)*
1	5	16.2	6.4	150	8	> 99
2	10	16.2	6.4	150	8	> 99

Table S8 Reaction conditions: 0.83 M methylpipecolinate (1b) + 16.2 eq. of formaldehyde (pH adjusted with 1 M solution of sodium acetate). 20 % (v/v) solution of formic acid in water. Reactor volume 1.16 mL. Reactions were performed at 8 bars. Conversions were calculated by <sup>1</sup>H-NMR.

#### Conditions screening at 1.6 M of methylpipecolinate (1b)

Entry	Residence Time (min)	Formaldehyde (eq.)	Molar Ratio (Methylpipecolinate: Formic acid) eq.	Temperature (ºC)	BPR (bars)	Conversion (%)*
1	5	8.4	6.4	40	8	n.d.
2	5	8.4	6.4	100	8	28
3	5	8.4	6.4	150	8	>99
4	5	3	3.3	150	8	>99

Table S9 Reaction conditions: 1.6 M methylpipecolinate (1b) + 8.4 eq. of formaldehyde (pH adjusted with 1 M solution of sodium acetate). 20 % (v/v) solution of formic acid in water. Reactor volume 1.5 mL. Reactions were performed at 8 bars. Conversions were calculated by <sup>1</sup>H-NMR.

## Amide Bond formation through Li-amide formation (Flow)

Electrophilic quench semi-continuous mode



Scheme S4 Semi-continuous set-up for the amide coupling reaction.

	Stoichiometry		Collection time in ``STR''		(00)	(00)
Entry	(Amine: n-BuLi)	Residence time (sec)	(min)	volume of electrophile (mc)	remperature ( C)	Conversion* (%)
1		1		20	27 °C	57
2	1:1	10	2	2	27 °C	>99
3		20		2	27 °C	>99

Table S10. <u>Conditions</u>: 0.075 mL of reactor volume, [2,6-dimethylaniline] = 2.5 M, [n-BuLi] = 2.5 M, [Ethyl benzoate] = 0.83 M, dry and degassed THF was used, quench was performed with 1 mL of H<sub>2</sub>O straight after collection was completed. \*Conversions were calculated by <sup>1</sup>H-NMR

#### Telescoped amide bond formation reaction screening:



Scheme S5 Flow set-up for the amide formation

#### Residence time effect

Entry	Stoech (n-Buli: amine)	Stoech (Li-amide: N-methylpipecolinate)	R1 (sec)	R2 (sec)	Conversion (%)
1			1	0.6	26
2	1.1	1 5.1	5	3	37
3	1:1	1.5.1	10	7	52
4			20	15	54

Table S11 Reaction conditions: N-methylpipecolinate (4b) (0.8 M), n-Buli (1.6 M), 2.6-dimethylaniline (1.6 M), reactor volumes 0.075 mL, reaction temperature 23 °C. Conversions were calculated by <sup>1</sup>H-NMR

#### n-BuLi and Li-amide ratio effect

Entry	Stoech (n-Buli: amine)	Stoech (Li-amide: N-methylpipecolinate)	R1 (sec)	R2 (sec)	Conversion (%)
1	1.5:1	2:1		7	57
2		1.5:1	10	6	51
3	2.1	2:1		7	91
4	2.1	5:1		8	60

Table S12 Reaction conditions: N-methylpipecolinate (4b) (0.8 M), n-Buli (1.6 M), 2.6-dimethylaniline (1.6 M), reactor volumes 0.075 mL, reaction temperature 23 °C. Conversions were calculated by <sup>1</sup>H-NMR

Fully telescoped screening

Entry	Stoech (n-Buli: amine)	Stoech (Li-amide: N-methylpipecolinate)	R1 (sec)	R2 (sec)	Conversion (%)
1				7	42
	2:1	2:1	10		
2				225	90

Table S13 Solution of n-BuLi 1.6 M in Hexanes, solution of amine (2) 1.6 M in 2-MeTHF. Amide coupling. Solution of N-methylpipecolinate (4b) 0.8 M in 2-MeTHF. 2-MeTHF was used without additional purification.

#### Fully continuous set-up



Scheme S6 Fully continuous set-up for the synthesis of mepivacaine. Process conditions: **Reductive amination.** Solution of methylpipecolinate **(1b)** 1.6 M in H<sub>2</sub>O pH 4.0 + Formaldehyde 3.0 eq. Formic acid 20 % (v,v) 3.3 eq. **Lithiation**. Solution of n-BuLi 1.6 M in Hexanes 2eq., solution of amine **(2)** 1.6 M in 2-MeTHF 1.0 eq. **Amide coupling.** Solution of N-methylpipecolinate **(4b)** 0.8 M in 2-MeTHF 1.0 eq., stream of Li-amide **(5)** 2.0 eq.

Full coupling of the overall system in continuous was optimal since the gas (butane) generated in the lithiation step caused an intermittent back flow of the organic phase into the membrane separator leading to a very unstable and challenging set-up. Nonetheless, conversions reached ~45%. The placement of a check valve downstream of the membrane separator did not improve the result.

#### **Green metrics**

All the calculations have been performed following the recommendations and assumptions from McElroy et al., 2015.<sup>2</sup> and others.<sup>3,4</sup>

Equation S1 Space time yield (STY)  $Space - time \ yield \ (STY) = \frac{Mass \ of \ product \ (Kg)}{Reaction \ time \ (h) \cdot Reactor \ volume \ (L)}$ Equation S2 E factor  $E \ factor = \frac{Total \ waste \ (Kg)}{Total \ product \ (Kg)}$ Equation S3 Process mass intensity (PMI)  $Process mass intensity (PMI) = \frac{Total raw materials used in process (Kg)}{Total product (Kg)} = \sum [MI]_{solvents, reagents, water, etc}$ Equation S4 Mass intensity (MI)  $Mass intensity (MI) = \frac{Raw material used (Kg)}{Total product (Kg)}$ 

Equation S5 Atom economy (AE)

 $Atom \ economy \ (AE) = \frac{Molecular \ weight_{product}}{\sum Molecular \ weight_{Starting \ materials}} \cdot 100$ 

Comparison between previously existing methods and this work

Metric	This work	Ekenstam <i>et al.,</i> 1957	Suveges <i>et al.,</i> 2017
Space time yield (STY) kg·(L·h) <sup>-1</sup>	0.4	0.037	0.077
E factor (kg waste-(Kg product) <sup>-1</sup> )	18.6	7.5	496.7
Corrected E factor (kg waste·(Kg product) <sup>-1</sup> )*	53.0	36.2	719.2
Process mass intensity (PMI) (Kg total·(Kg product) <sup>-1</sup> )	56.0	39.2	721.2

Table S14 overall metrics for the different analyzed processes. For a more detailed breakdown of each step refer to supplementary excel file. Note that the lower E factors and PMI's reported for the Ekenstam method are not 100% accurate since crucial data was missing for workup and purification stages. Presumably the values would significantly increase. \*Corrected E factor includes the water contribution.

#### Remarks:

- For PMI calculations on Suveges et al., 2017 method, extraction and base volumes for last step are not • reported but, taking into account the final volume of the solution, we assume (at least) the same volumes as for their 1s step.
- For Ekenstam et al., 1957: platinum oxide not considered since it is recovered.
- Wash volumes not known same for recrystallization. .
- For clarity the nomenclature of each molecule is kept as in the original manuscripts. •

#### Additional considerations for green metrics calculations

Work-up, isolation and purification steps are included in the calculations. .

#### Space-Time-Yield (STY)

- STY calculated considering isolated yields (whenever the intermediate/product was isolated)
- For the global STY the lower STY value of the whole process has been taken since it is the bottleneck of the productivity.

#### Process Mass Intensity (PMI)

The following considerations and assumptions were used:

- 25 minutes of production (reductive amination)
- 16 minutes of production for (Amide coupling and Lithiation)
- Total amount of water ``in'' refers to the total amount of water used in each single step.
- Up to 90% of the used solvents can be recycled.
- For the lithiation step the solvent and 2,6-dimethylaniline are fully telescoped into the next step and both are recycled.

E-factor

• Corrected E factor includes the water contribution.

#### Structural Characterization of N-methylpipecolinate (4b)



$$\begin{split} &\delta_{\rm \,H}\,(300~\text{MHz},\text{CDCl}_3)\,3.73\,(3~\text{H},\,\text{s}), 2.96\,(1~\text{H},\,\text{d},\,J\,19.3), 2.76\,(1~\text{H},\,\text{d},\,J\,10.9), 2.25\,(3~\text{H},\,\text{s}), 2.08\,(1~\text{H},\,\text{d},\,J\,25.2), \\ &1.85\,(1~\text{H},\,\text{d},\,J\,20.5), 1.68\,(4~\text{H},\,\text{d},\,J\,55.3), 1.27\,(1~\text{H},\,\text{d},\,J\,45.3). \end{split}$$





High resolution mass spectrometry (4b)



Elemental composition search on mass 158.1172

m/z= 153.1172-163.1172 m/z Theo. Delta Composition Mass (ppm) 158.1172 158.1176 -2.18 C8 H16 O2 N

Figure S1 mass spectrometry (4b)

*m/z* 158.1172 (M<sup>+</sup> + H, 100%)

Delta ppm: 2.18

Type of analysis: +ESI-MS

### **Structural Characterization of mepivacaine (7)**

![](_page_9_Figure_7.jpeg)

<u><sup>1</sup>H-NMR (7)</u>

![](_page_9_Figure_9.jpeg)

$$\begin{split} &\delta_{\rm \,H}\,(400~{\rm MHz},\,{\rm MeOD})\,\,7.17-7.00\,\,(3~{\rm H},\,{\rm m}),\,2.99\,\,(1~{\rm H},\,{\rm dt},\,J\,11.6,\,3.6),\,2.75\,\,(1~{\rm H},\,{\rm dd},\,J\,11.0,\,3.1),\,2.36\,\,(3~{\rm H},\,{\rm s}),\,2.22\,\,(6~{\rm H},\,{\rm s}),\,2.19-2.09\,\,(1~{\rm H},\,{\rm m}),\,2.05-1.95\,\,(1~{\rm H},\,{\rm m}),\,1.92-1.81\,\,(1~{\rm H},\,{\rm m}),\,1.77\,\,(1~{\rm H},\,{\rm m}),\,1.74-1.59\,\,(2~{\rm H},\,{\rm m}),\,1.46-1.28\,\,(1~{\rm H},\,{\rm m}). \end{split}$$

![](_page_10_Figure_0.jpeg)

<sup>13</sup>C NMR (101 MHz, MeOD) δ 174.79, 136.90, 135.17, 129.17, 128.42, 71.05, 56.75, 44.93, 32.04, 26.28, 24.49, 18.67.

High resolution mass spectrometry (7)

![](_page_10_Figure_3.jpeg)

Figure S2 mass spectrometry (7)

*m/z* 247.1812 (M<sup>+</sup> + H, 100%)

Delta ppm: 3.03

Type of analysis: +ESI-MS

### **References**

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