

Supplementary information

Sensitive method to determine melatonin in saliva by automated online in-tube solid-phase microextraction coupled with stable isotope-dilution liquid chromatography-tandem mass spectrometry

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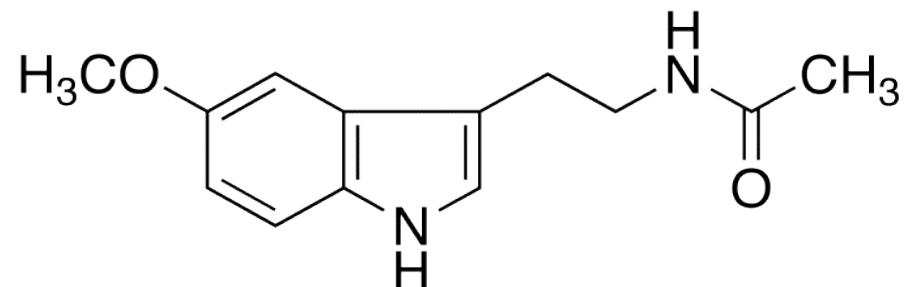
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Supplementary figures

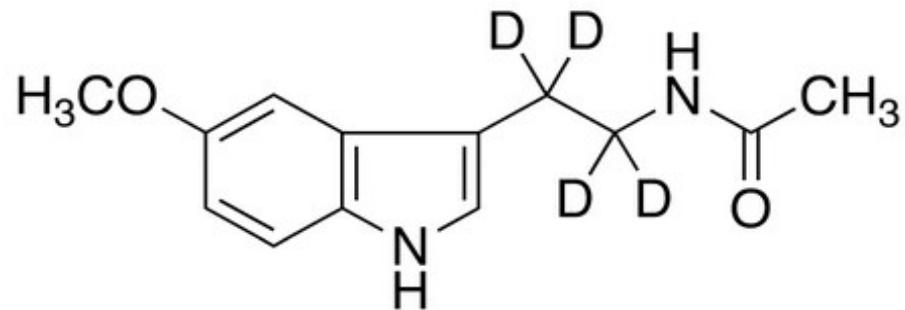
Fig. S1 Structures of MLT and MLT-d₄ used as an internal standard.

Fig. S2 Schematic diagram of the automated online in-tube SPME/LC–MS/MS system.

Fig. S1 Structures of melatonin and melatonin-d₄ used as an internal standard.



Melatonin



Melatonin-d₄

Fig. S2 Schematic diagram of the automated online in-tube SPME/LC-MS/MS system.

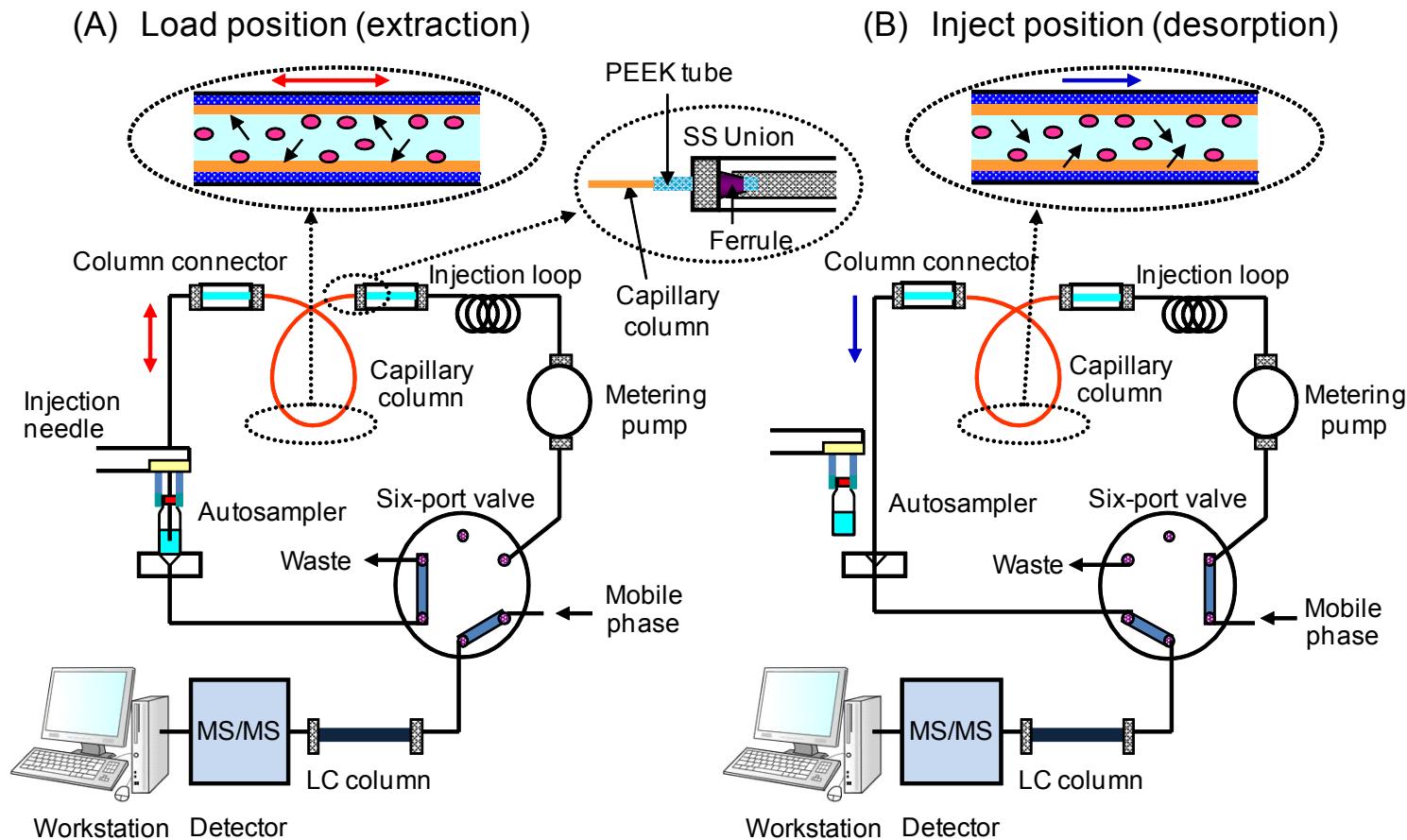


Table S1 MRM conditions for MLT and MLT-d₄.

Compound	Q1 (precursor ion)	Q3 (product ion)	Dwell time (ms)	DP (V)	EP (V)	CE (eV)	CXP (V)
MLT (Quantitation)	233.3	174.3	333	60	15	20	5
MLT (Confirmation)	233.3	159.1	333	60	15	30	15
MLT-d ₄ (Internal standard)	237.3	178.3	333	60	15	20	5

Table S2 Program for the in-tube SPME process.

Sequence	Event	Switching valve	Vial	Draw/eject		
				Cycle ^{a)}	Volume (μL)	Speed (μL min ⁻¹)
1	Conditioning of the capillary	Load	MeOH	D/E (2)	40	200
2	Drawing of air into the capillary	Load	Empty	D (1)	50	200
3	Conditioning of the capillary	Load	Water	D/E (2)	40	200
4	Extraction of analytes into the capillary	Load	Sample	D/E (25)	40	200
5	Needle washing	Load	MeOH	D/E (1)	2	200
6	Desorption of analytes from the capillary	Inject	—	—	—	—
7	HPLC separation of analytes and return to sequence 1	Load	—	—	—	—

^{a)} D: draw, E: ejection.

Table S3 Precision of MLT measurements by in-tube SPME LC–MS/MS.

Compound	Concentration	Precision (RSD %) ^{a)}	
		Intra-day	Inter-day
MLT	1.0 pg mL ⁻¹	4.1	5.8
	10.0 pg mL ⁻¹	3.0	3.6

^{a)} $n = 5$.

Table S4 Recoveries of MLT spiked into saliva samples.

Compound	Concentration (pg mL ⁻¹ saliva)		Recovery (%)
	Spiked	Mean ± SD (n=3)	
MLT	0	3.8 ± 0.1	
	10.0	13.4 ± 1.0	96
	100.0	102.1 ± 3.9	98