



RESEARCH ARTICLE

A FAST SCANNING STRIPPING SQUARE WAVE VOLTAMMETRY ANALYSIS OF ERYTHROMYCIN A IN TILAPIA (*Oreochromis niloticus*) WITH THE DROPPING MERCURY ELECTRODE

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ABSTRACT

A new sensitive analytical approach for determination of erythromycin A in tilapia (*Oreochromis niloticus*) by a fast scanning stripping square wave voltammetry using the dropping mercury electrode (PSA-F) was developed and validated. Optimum parameters for erythromycin A quantification were: Ammonium acetate buffer 0.1 M, pH 8.0 as supporting electrolyte; acetonitrile as the solvents for dissolving erythromycin standard; forward scanning; V_{start} : -400 mV; V_{stop} : -1700 mV; V_{step} : 6 mV; V_{pulse} : 40 mV; T_{dop} : 5000 ms; $V_{\text{electrode}}$: -1100 mV; $T_{\text{electrode}}$: 5s. Peak of erythromycin A appeared at $E_{1/2} = -1430$ mV, separated and distinguished with peaks of other antibiotics. This assay showed high recovery (> 85.07 %), high sensitivity (detection limit 0.52 $\mu\text{g}/\text{kg}$), high precision (RSD, $0.8 \div 2.1$ %), high accuracy (relative error - RE, $85.07 \div 88.56$ %) as well as excellent linearity ($r^2_{\text{adjusted}} = 1.0$). Simpler, reagent-saving and time-saving were other advantages of this assay method. An equivalence of analyzing results between PSA-F and LC-MS/MS could be obviously seen.

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INTRODUCTION

Tilapias are known to have been an important component of fisheries in the Mekong River Delta. The most significant diseases in Nile tilapia (*Oreochromis niloticus*) culture are caused by *Streptococcus iniae*, *Aeromonas hydrophila*, *Trichodina sp.*, *Flexibacter Columnaris*, *Edwardsiella spp.* *Streptococcus spp.*, gram positive bacteria, have become a major problem for

tilapia farmers and there is still no effective commercial vaccine available that can be used to prevent *Streptococcus spp.* in tilapia. They can cause mass death in tilapia farms, and unlike many other tilapia diseases it will affect even large and otherwise healthy fish. The macrolide antibiotic erythromycin has long been the chemotherapeutant of choice to prevent and control *Streptococcus spp.* Erythromycins are broad spectrum antibiotics that exhibit high activity against nearly all Gram positive and Gram-negative bacteria. Erythromycin

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A consists of a polyhydroxylactone and two sugars. The aglycone part of all erythromycin molecules, the erythronolide, is a 14-membered lactone ring (Fig. 1). Depending on the type of erythromycin this lactone ring is substituted via 4-position with a cladinose in case of erythromycin A, erythromycin B, erythromycin E, erythromycin F and with a mycarose in case of erythromycin C and erythromycin D. All erythromycin molecules contain the aminosugar d-desosamine, which is β -glycosidic linked to the 6 position of the lactone ring. The minimum inhibited concentration of erythromycins A, B, C, and D and some of their derivatives were determined against 21 gram-positive and 15 gram-negative microorganisms. Antibacterial activity was confined to gram-positive and very few gram-negative bacteria. Erythromycin B was somewhat less active than erythromycin A, and erythromycin C and D showed about half that activity or even less (Isaac Ongubo Kibwage *et al.*, 1985). Owing to their extensive use in infectious disease therapy, several procedures have been reported for its determination (Table 1). The aim of this work was to determine the feasibility of square wave voltammetry for the direct detection and quantification of erythromycin A in tilapia tissue.

MATERIALS AND METHODS

Reagents and materials

All chemicals and reagents were HPLC grade or p.a. Double-distilled water (DDW) was used throughout the study. The high purity antibiotic standards (>99%) of erythromycin A, erythromycin A, chloramphenicol, furazolidone, florfenicol, ciprofloxacin, colistin, malachite green were purchased from Vietnam Central Institute of Pharmacy.

Apparatus: A fast scanning stripping square wave voltammetry at the slowly dropping mercury electrode was performed in the ANALYZER SQF-505. The mercury dropping electrode was used as a working electrode, silver/silver chloride (saturated KCl) as a reference electrode and a platinum wire as an auxiliary one.

Sample extraction and clean-up procedure

Primary extraction: A 5 g aliquot of a blank or spiked minced tissue sample was mixed with a small volume of erythromycin standard. After a 15-min equilibration period, the tissues were mixed vigorously for 15 min with 25 ml Tris buffer (0.1M; pH 10.5). After a 10-min centrifugation at 3000 g and 4 °C, the supernatant was transferred to a polypropylene tube and the solid residue extracted a second time with 25 ml Tris buffer. Acetic acid (600 μ l) and 5 ml sodium tungstate buffer (0.15M) were added to precipitate the proteins. After equilibration for one hour at 4 °C, the samples were centrifuged at 3000 g for 10 min. The supernatants were further filtered through a plug of glass wool.

Solid phase extraction: The 6-cm³ HLB OASIS extraction cartridges (200 mg) were prepared and conditioned with 10 ml methanol and 10 ml water. The biological samples were placed at the top of the column. Two wash solution volumes were applied before erythromycin elution: 20 ml methanol-water (5:95, v/v) and 5 ml hexane. After the last washing step, the OASIS columns were vacuum-dried for 10 min. Erythromycin was finally eluted with 5 ml methanol-ammonia 30 % (95:5, v/v) and evaporated dry under a nitrogen flow. The extracts were dissolved in 500 μ l NH₄AC-ACN (80/20 v/v), transferred to Eppendorf tubes and centrifuged at 3000 g for 10 min. Aliquots of the supernatant were transferred into the voltammetric cell with 2,500 mL of ammonium acetate 0.1 M, pH 8.0 before being quantified by Analyzer SQF-505 machine in mode stripping square wave voltammetry.

RESULTS AND DISCUSSION

Voltammetric behavior of erythromycin at the slowly dropping electrode

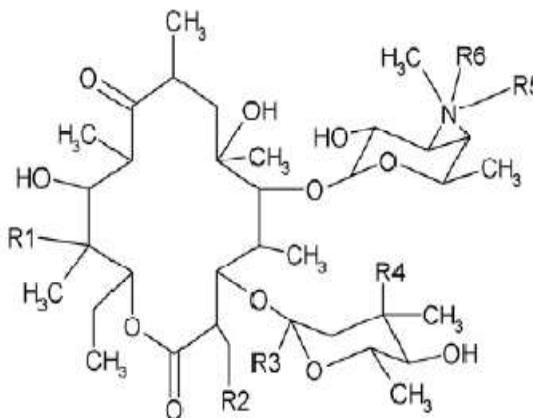
Effect of Supporting Electrolytes and pH values.

The adsorptive peak current of erythromycin has been strongly affected by the type of supporting electrolyte. To study the adsorptive behavior of erythromycin, different supporting electrolytes including sodium acetate, ammonium acetate, citrate-phosphate, borax, Tris buffers were

erythromycin showed the highest peak current and the best peak shape (Table 2 and Fig. 2). The effect of pH of ammonium acetate buffer on the peak current was examined from 7.0 to 10.0. Erythromycin showed highest peak current at pH 8.0 ($E_{1/2} = -1438$ mV, $I = 351.7 \pm 5.7$ nA). Hence ammonium acetate buffer (pH 8.0) was selected for further investigations.

Optimization of measurement conditions

Effect of forward scanning (0 to -1800 mV) and reverse scanning (-1800mV to 0) on the peak current signal was examined. The forward scanning (0 to -1800 mV) showed high peak. Meanwhile, the peak current of the reverse scanning was low. So the forward



Erythromycin	Formula	Molecular mass	R ₁	R ₂	R ₃	R ₄	R ₅
A	C ₃₇ H ₅₇ NO ₁₃	734	OH	H	H	OCH ₃	CH ₃
B	C ₃₇ H ₅₇ NO ₁₂	718	H	H	H	OCH ₃	CH ₃
C	C ₃₈ H ₅₅ NO ₁₃	720	OH	H	H	OH	CH ₃
D	C ₃₆ H ₆₅ NO ₁₂	704	H	H	H	OH	CH ₃
E	C ₃₇ H ₆₇ NO ₁₃	748	OH	-O-		OCH ₃	CH ₃
F	C ₃₇ H ₆₇ NO ₁₄	750	OH	OH	H	CH ₃	CH ₃

Figure 1. Structural formula of erythromycin A and related substances

The effect of the ionic strength of supporting electrolyte was examined at pH 8.0 over the range from 0.05÷0.25 M. Erythromycin showed highest peak current at ammonium acetate 0.1M ($E_{1/2} = -1438$ mV, $I = 254.8 \pm 10.2$ nA). So this value was selected for further studies (Table 3 and Figure 3). Effect of the solvents for dissolving erythromycin A standard on the peak current were examined. Among methanol, acetonitril, ethyl acetate, the peak current increased with a maximum at acetonitril ($E_{1/2} = -1438$ mV, $I = 189.2 \pm 3.5$ nA).

scanning (0 to -1800 mV) was chosen for further investigations. Effect of V_{start} . Mode PSA-F, forward scanning, V_{stop} : -1800 mV, V_{step} : 4 mV, V_{pulse} : 30mV, T_{drop} : 3000ms, $V_{electrolise}$: -700mV, $T_{electrolise}$: 6s, $T_{stabilize}$: 1s. Examining V_{start} from -400mV to -1100mV. V_{start} was optimum at -400mV ($E_{1/2} = -1430$ mV, $I = 106.5 \pm 4.7$ nA; Table 5 and Fig. 5). Effect of V_{stop} . Mode PSA-F, forward scanning, V_{start} : -400 mV, V_{step} : 4 mV, V_{pulse} : 30 mV, T_{drop} : 3000 ms, $V_{electrolise}$: -700 mV, $T_{electrolise}$: 6 s, $T_{stabilize}$: 1 s. Examining V_{stop} from -1700 mV to -

Table 1. Some typical papers published in recently

Author	Sample	Method	LoD (µg/kg)
Zierfels and Petz, 1994	Egg, muscle, milk, liver, kidney of swan	HPLC	<10
Yong-Xi Li <i>et al.</i> , 1998	Human plasma	LC-MS/MS	LoQ 0.5
Kondo <i>et al.</i> , 1999	Human plasma	LC-MS/MS	LoQ 0.05
Dreassi <i>et al.</i> , 2000	Plasma: beef, pork, poultry	HPLC-UV	LoQ 250
	Milk	HPLC-UV	LoQ 25
	Kidney, liver, muscle, gan, fat of beef, pork, poultry.	HPLC-UV	LoQ 125
Huaisheng Wang <i>et al.</i> , 2000	Drug, urine.	ASV-PGCE	5
Carmen Leal <i>et al.</i> , 2001	Chicken	LC-FL	400
Draisci <i>et al.</i> , 2001	Beef	ELISA	0.4
	Muscle and liver of beef	LC-MS/MS	LoQ 50
	Kidney of beef	LC-MS/MS	LoQ 80
Stanley M. Billedeau <i>et al.</i> , 2003	Salmon	LC - ESI/MS	LoD: 5, LoQ: 16
Horie Masakazu <i>et al.</i> , 1999.	Meat and seafood	LC- ESI-MS	10
Michael P. Sche <i>et al.</i> , 2003	Manure	HPLC-MS/MS	0.4-11
Xiao <i>et al.</i> , 2005	Drugs (propionate, base)	HPLC-ESI-MS	1
Deube <i>et al.</i> , 2006	Muscle	LC-MS/MS	0.25
Hui Yun – Hua <i>et al.</i> , 2006	Tilapia	HPLC	400
Jian Wang <i>et al.</i> , 2006	Fresh milk	LC-ESI/MS/MS	0.07
Tang <i>et al.</i> , 2006	Meat	LC-MS/MS	0.1
Deng <i>et al.</i> , 2007	Rat plasma	ECL	0.35
Berrada Houda <i>et al.</i> , 2008	Meat and seafood	LC-ESI/MS	25
Granja <i>et al.</i> , 2009	Honey	LC-MS/MS	LoD 1.27, LoQ 5.0
P. Norouzi <i>et al.</i> , 2009	Human plasma, urine.	CV	LoD 2.4, LoQ 7.0

Table 2. Peak current of erythromycin A was affected by supporting electrolytes, pH values

pH	5.0		6.0		7.0		8.0		9.0		10.0	
	Mean ± SD	RSD (%)	Mean ± SD	RSD (%)	Mean ± SD	RSD (%)	Mean ± SD	RSD (%)	Mean ± SD	RSD (%)	Mean ± SD	RSD (%)
Supporting electrolytes												
Natri acetate	56.67 ^b ± 1.1	1.9	35.5 ^a ± 2.3	6.6	255.5 ^d ± 7.3	2.8	333.4 ^c ± 4.7	1.4	236.2 ^c ± 7.7	3.3	254 ^d ± 4.7	1.9
Ammonium acetate					210.3 ^a ± 5.4	2.6	351.7 ^d ± 5.7	1.6	216.5 ^b ± 2.7	1.2	263.1 ^c ± 1.6	0.6
Citrat-phosphate			207.7 ^d ± 6.0	2.9	168.2 ^a ± 2.3	1.4	194.4 ^b ± 2.6	1.4	229.4 ^c ± 2.1	0.9	200.9 ^c ± 2.0	1.0
Tris					180.0 ^d ± 13.1	7.3	27.5 ^a ± 0.3	1.1	53.4 ^b ± 3.6	6.7	126.4 ^c ± 9.7	7.7
Borax							168.1 ^a ± 1.6	0.9	255.5 ^c ± 6.5	2.6	173.9 ^b ± 1.8	1.0

* Each value was the mean of 5 samples (n = 5)

Effect of V_{step} . Mode PSA-F, forward scanning, V_{start} : -400 mV, V_{stop} : -1700 mV, V_{pulse} : 30 mV, T_{drop} : 3000 ms, $V_{\text{electrolyse}}$: -700 mV, $T_{\text{electrolyse}}$: 6 s, $T_{\text{stabilize}}$: 1 s. Examining V_{step} from 4 mV to 10 mV. V_{step} was optimum at 6.0 mV ($E_{1/2} = -1430$ mV, $I = 214.6 \pm 13.1$ nA) (Table 7 and Fig. 7). Effect of V_{pulse} . Mode PSA-F, forward scanning, V_{start} : -400

optimum at 40 mV ($E_{1/2} = -1430$ mV, $I = 692.6 \pm 14.9$ nA) (Table 8 and Fig. 8). Effect of T_{drop} . Mode PSA-F, forward scanning, V_{start} : -400 mV, V_{stop} : -1700 mV, V_{step} : 6 mV, V_{pulse} : 40 mV, $V_{\text{electrolyse}}$: -700 mV, $T_{\text{electrolyse}}$: 6 s, $T_{\text{stabilize}}$: 1 s. Examining T_{drop} from 1000 ms to 5,000 ms. T_{drop} was optimum at 5,000 ms ($E_{1/2} = -1430$ mV, $I = 381.3 \pm 2.9$ nA)

mV, V_{step} : 6 mV, V_{pulse} : 40 mV, T_{drop} : 5,000 ms, $V_{electrolyse}$: -700 mV, $T_{stabilize}$: 1s. Examining $T_{electrolyse}$ from 3s to 6s. $T_{electrolyse}$ was optimum at 5s ($E_{1/2} = -1430$ mV, $I = 1717.0 \pm 13.7$ nA) (Table 10 and Fig. 10). Effect of $V_{electrolyse}$ Mode PSA-F, forward scanning, V_{start} : -400 mV, V_{stop} : -1700 mV, V_{step} : 6mV, V_{pulse} : 40 mV, T_{drop} : 5,000 ms, $T_{electrolyse}$: 5 s, $T_{stabilize}$: 1 s. Examining $V_{electrolyse}$ from -400 mV to -1400 mV. $V_{electrolyse}$ was optimum at -1100 mV ($E_{1/2} = -1438$ mV, $I = 1863.2 \pm 24.1$ nA) (Table 11 and Fig. 11).

Calibration

Calibration curves and detection Limit: A 25mL supporting electrolyte ammonium acetate 0.1M, pH 8.0 was transferred to the cell and spiked with 5 μ L, 10 μ L, 20 μ L, 30 μ L, 40 μ L, 50 μ L, 60 μ L, 7 μ L, 80 μ L, 90 μ L of stock 250ppm solution of erythromycin in pure acetonitril. The concentrations of erythromycin in the cell were 50 μ g/kg, 100 μ g/kg, 200 μ g/kg, 300 μ g/kg, 400 μ g/kg, 500 μ g/kg, 600 μ g/kg, 700 μ g/kg, 800 μ g/kg, 90 μ g/kg respectively. Mode PSA-F, V_{start} : -400mV, V_{stop} : -1700 mV, V_{step} : 6 mV, V_{pulse} : 40 mV, T_{drop} : 5000 ms, $T_{electrolyse}$: 5s, $V_{electrolyse}$: -1100 mV, $T_{stabilize}$: 1s. A detection limit of 0.57 μ g/kg was obtained for erythromycin. A linear behavior was also observed with a correlation coefficient $r^2_{adjust} = 1.0$ (Fig. 12).

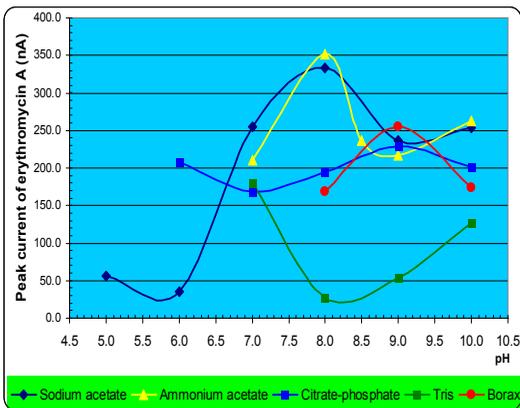


Figure 2. Effect of supporting electrolytes, pH values on peak current of erythromycin A

ions was examined by introducing different

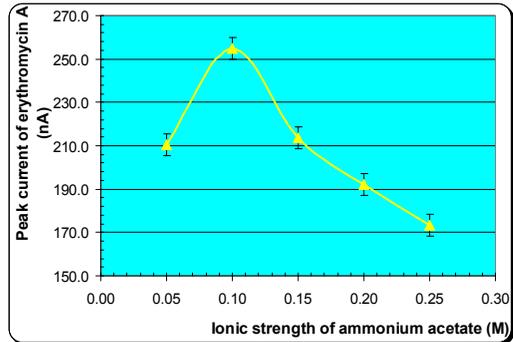


Figure 3. Effect of ionic strength of ammonium acetate on peak current of erythromycin A

Table 4. Peak current of erythromycin A was affected by solvents

Ethyl acetate		Acetonitril		Methanol	
Mean \pm SD	RSD (%)	Mean \pm SD	RSD (%)	Mean \pm SD	RSD (%)
183.6 ^b	0.8	189.2 ^c	1.9	166.3 ^a	3.3

* Each value was the mean of 5 samples (n = 5)

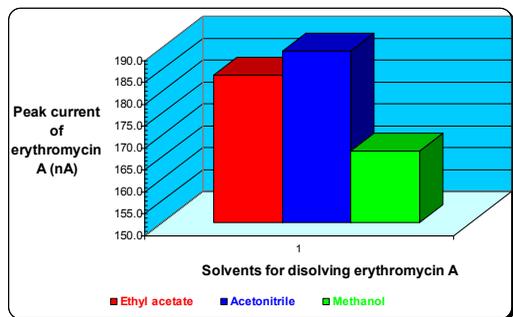
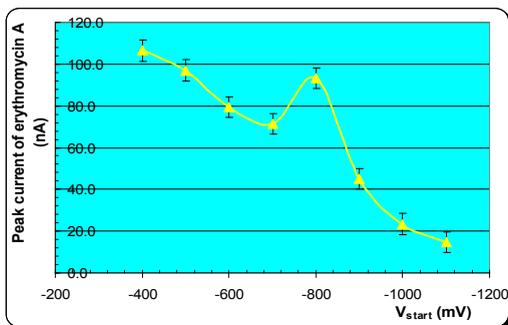


Fig. 4. Effect of solvents for dissolving erythromycin A on peak current



concentrations of some ions to the voltametric cell and recording the corresponding voltammogram using the conditions selected above. It was observed that the additions of 0-5ppm K^+ , Na^+ , Ca^{2+} , Mg^{2+} , Fe^{3+} , Cl^- , SO_4^{2-} , HPO_4^{2-} ions have no effect ($< \pm 5\%$) on the peak response.

Table 5. Peak current of erythromycin A was affected by V_{start}

V_{start} (mV)	-400	-500	-600	-700	-800	-900	-1000	-1100
Mean ± SD	106.5 ^a ± 4.7	97 ^l ± 3.4	79.5 ^c ± 0.8	71.5 ^d ± 4.3	93.3 ^f ± 4.0	45.0 ^c ± 2.6	23.4 ^b ± 1.0	14.8 ^a ± 0.9
RSD (%)	4.4	3.5	1.0	6.0	4.3	5.8	4.3	6.2

* Each value was the mean of 5 samples (n = 5)

Table 6. Peak current of erythromycin A was affected by V_{stop}

V_{stop} (mV)	-1500.0	-1600.0	-1700.0	-1800.0
Mean ± SD	104 ^a ± 1.0	118.2 ^b ± 1.4	136.7 ^c ± 3.9	120.1 ^b ± 1.1
RSD (%)	0.9	1.2	2.9	0.9

* Each value was the mean of 5 samples (n = 5)

Table 7. Peak current of erythromycin A was affected by V_{step}

V_{step} (mV)	4.0	6.0	8.0	10.0
Mean ± SD	162.8 ^a ± 5.8	214.6 ^c ± 13.1	176.9 ^b ± 8.3	165.0 ^{ab} ± 10.9
RSD (%)	3.5	6.1	4.7	6.6

* Each value was the mean of 5 samples (n = 5)

Table 8. Peak current of erythromycin A was affected by V_{pulse}

V_{pulse} (mV)	10	20	30	40
Mean ± SD	230.6 ^a ± 2.7	388.4 ^b ± 12.7	528.0 ^c ± 8.0	692.6 ^d ± 14.9
RSD (%)	1.2	3.3	1.5	2.2

* Each value was the mean of 5 samples (n = 5)

Table 9. Peak current of erythromycin A was affected by T_{drop}

T_{drop} (ms)	1,000	2,000	3,000	4,000	5,000
Mean ± SD	128.3 ^a ± 1.2	197.9 ^b ± 1.9	269.1 ^c ± 12.9	323.2 ^d ± 3.3	381.3 ^e ± 2.9
RSD (%)	0.9	1.0	4.8	1.0	0.8

* Each value was the mean of 5 samples (n = 5)

Table 10. Peak current of erythromycin A was affected by $T_{electrolyse}$

$T_{electrolyse}$ (s)	3	4	5	6
Mean ± SD	1353.6 ^a ± 10.8	1555.4 ^b ± 13.0	1717.0 ^d ± 13.7	1655.4 ^c ± 5.0
RSD (%)	0.8	0.8	0.8	0.3

* Each value was the mean of 5 samples (n = 5)

Table 11. Peak current of erythromycin A was affected by $V_{electrolyse}$

$V_{electrolyse}$ (Mv)	-400.0	-900.0	-1100.0	-1400.0
Mean ± SD	1709.0 ^b ± 17.3	1815.0 ^c ± 3.7	1863.2 ^c ± 24.1	1593.4 ^a ± 13.5
RSD (%)	1.0	0.2	1.3	0.8

* Each value was the mean of 5 samples (n = 5)

Sample analysis

In tilapia samples: Recovery rate: $85.07 \div 88.56\%$, MDL: $0.52 \mu\text{g}\cdot\text{kg}^{-1}$, R_{adjust} : 1.0, RSD: $0.80 \div 2.10 \%$. (Fig. 13). Comparison results between

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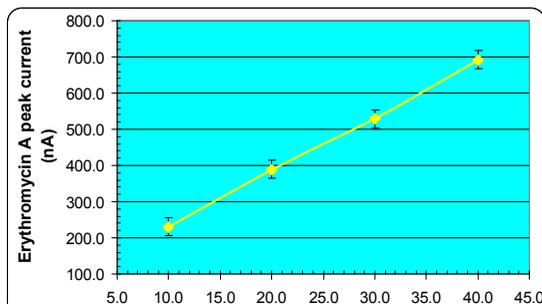


Fig. 8. Effect of pulse width on erythromycin A peak current

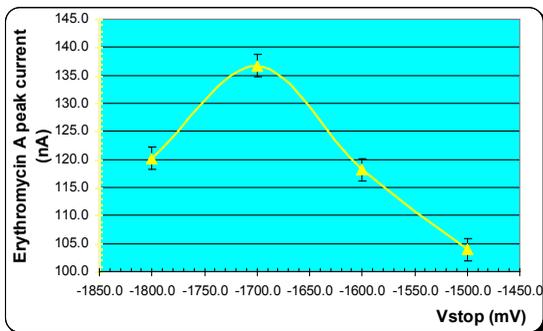


Fig. 6. Effect of V_{stop} on erythromycin A peak current

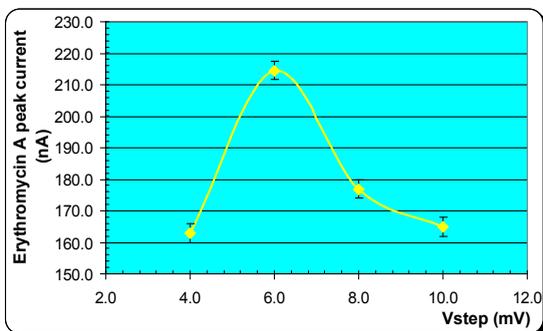


Fig. 7. Effect of V_{step} on erythromycin A peak current

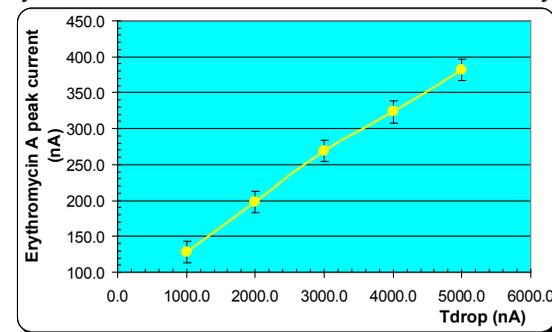


Fig. 9. Effect of T_{drop} on erythromycin A peak current

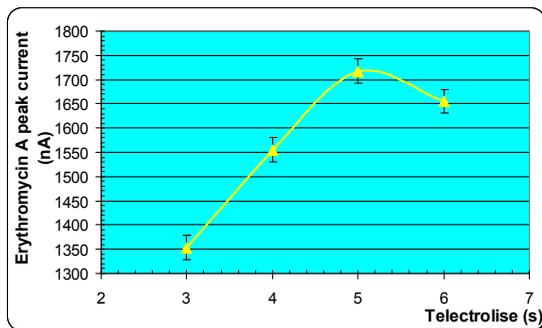


Fig.10. Effect of $T_{\text{electrolise}}$ on erythromycin A peak current

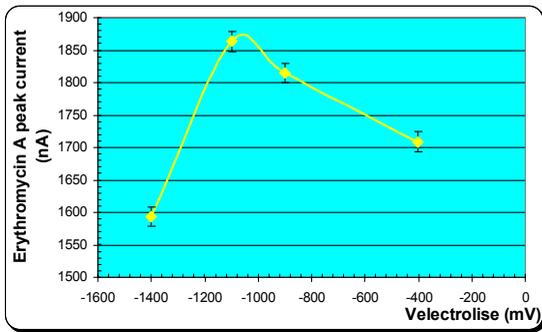


Fig.11. Effect of $V_{\text{electrolise}}$ on erythromycin A peak current

homogeneity could be obviously seen in comparison between two analyzing methods. SWV seems to be superior and more accurate to LC-MS/MS (Table 12)

Validation of quantification method

least square regression method. The calibration curves constructed for erythromycin were linear over the concentration range of $50 \div 400 \mu\text{g/kg}$. Peak areas of erythromycin were plotted versus its concentration and linear regression analysis performed on the resultant curve. A correlation coefficient of $R^2 = 1.0$ with %R.S.D. values

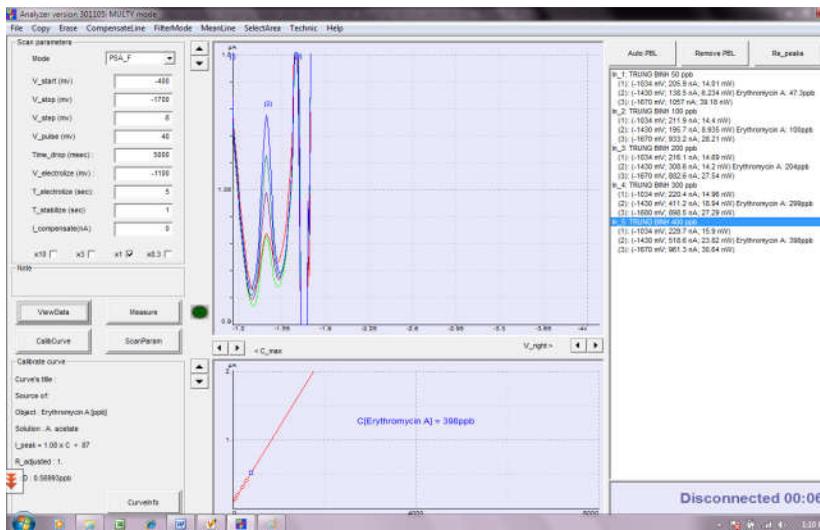
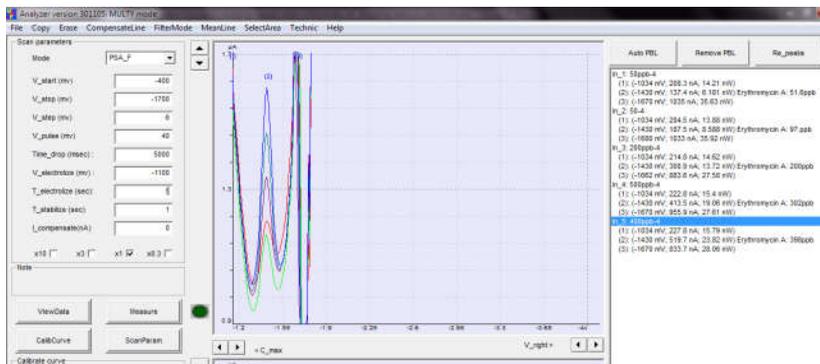


Fig.12. Calibration curve of erythromycin A



range studied were obtained following linear regression analysis (Table 13). Typically, the regression equation for the calibration curve was found to be $Y=1.08*X + 87$.

which is significantly different from that of a blank. Limit of detection was approved by calculations based on the standard deviation of the response (δ) (here the current) which is obtained from blank with 5 replicas and (S) is the slope of the calibration curve according to equation $LOD=3.3(\delta/S)$. The LOD for erythromycin was $0.57 \mu\text{g}/\text{kg}$ (Figure 13).

Precision, Accuracy & Recovery

Precision was investigated by the intra- and inter-

LOD

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analyte that may be detected to produce a response

Table 12. Comparison homogeneity of two analyzing methods

Sample I.D	SWV (LOD = $0.52 \mu\text{g}*\text{kg}^{-1}$)			LC-MS/MS (LOD = $10 \mu\text{g}*\text{kg}^{-1}$)		
	Result ($\text{mg}*\text{kg}^{-1}$)	Mean \pm SD ($\text{mg}*\text{kg}^{-1}$)	R.S.D (%)	Result ($\text{mg}*\text{kg}^{-1}$)	Mean \pm SD ($\text{mg}*\text{kg}^{-1}$)	R.S.D (%)
TL – Blank	N.D	N.A	N.A	N.D	N.A	N.A
TL – Blank	N.D			N.D		
TL – Blank	N.D			N.D		
TL – I	1.31	$1.29^a \pm 0.02$	1.61	1.23	$1.26^a \pm 0.03$	2.10
TL – I	1.27			1.28		
TL – I	1.30			1.27		
TL – II	1.95	$2.02^b \pm 0.07$	3.47	3.14	$2.72^b \pm 0.43$	15.64
TL – II	2.09			2.72		
TL – II	2.02			2.29		
TL – III	2.80	$2.78^c \pm 0.03$	1.25	2.80	$2.80^b \pm 0.01$	0.21
TL – III	2.80			2.80		
TL – III	2.74			2.81		

*LOD: Limit of detection; **N.D: Not detected

Table 13. Linear range in regression analysis of erythromycin

Erythromycin A concentration (ppb)	50.0	100.0	200.0	300.0	400.0
Peak current (Mean \pm SD)	$138.5^a \pm 5.1$	$194.0^b \pm 3.9$	$305.2^c \pm 5.2$	$411.2^d \pm 1.8$	$524.8^e \pm 14.3$
RSD (%)	3.7	2.0	1.6	0.4	2.7

* Each value was the mean of 5 samples

Table 14. Precision (RSD %), accuracy (RE %) and recovery of erythromycin A in tilapia muscles

Day	Spike level ($\mu\text{g}*\text{kg}^{-1}$)	Measured concentration (mean \pm SD, $\mu\text{g}*\text{kg}^{-1}$)	RSD (%)	RE (%)
1	100	$85.07^a \pm 1.26$	1.48	85.07
2	100	$85.34^a \pm 1.79$	2.10	85.34
3	100	$85.31^a \pm 1.24$	1.45	85.31
1	200	$173.25^b \pm 2.34$	1.35	86.63
2	200	$173.03^b \pm 2.09$	1.21	86.51
3	200	$172.82^b \pm 1.39$	0.80	86.41
1	300	$261.62^c \pm 4.53$	1.73	87.21
2	300	$262.57^c \pm 3.42$	1.30	87.52

reproducibility. Repeatability was investigated by injecting six replicate samples of each of the 100, 200, 300 µg/kg standards. Inter-day precision was assessed by injecting the same three concentrations over 3 consecutive days. Accuracy (relative error, RE, %) was calculated by assessing the agreement between measured and nominal concentrations of the fortified samples. Recovery was assessed at erythromycin A, concentrations of 100, 200, 300 µg/kg and the mean value was calculated (Table 14 and Figure 14).

Diffaontiation

the presence of other antibiotic components. The peak response ($E_{1/2}$) of erythromycin A ($E_{1/2} = -1430$ mV) was separated, independent and

monitoring standard solutions of erythromycin A in

($E_{1/2} = -1152$ mV), florfenicol ($E_{1/2} = -78$ mV),

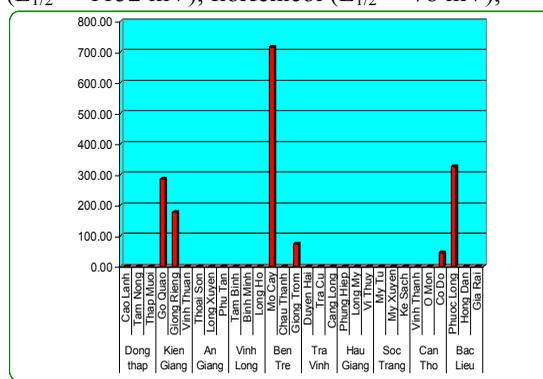
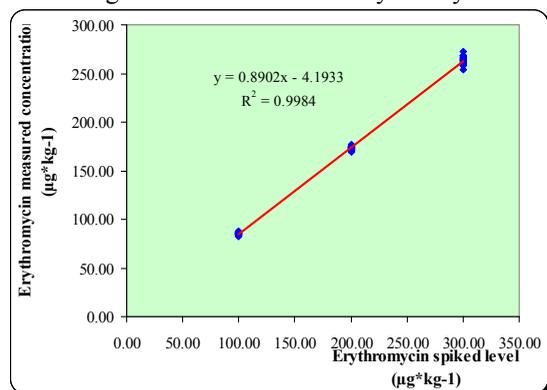


Fig. 14. Erythromycin concentration was detected on fish samples at different times and levels

Fig. 15. Erythromycin surveillance in tilapia aquaculture at ten provinces, three districts in each province of Mekong River Delta, VN

Table 15. Erythromycin residue of tilapia samples from ten provinces, three districts in each province in the Mekong River Delta

No	Province	District	Tilapia					Mean (µg·kg ⁻¹)	RSD (%)	
			M1	M2	M3	M4	M5			
1	Dong Thap	Cao Lanh	N.D	N.D	N.D	N.D	N.D	N.D	N.D	
		Tam Nong	N.D	N.D	N.D	N.D	N.D	N.D	N.D	
		Thap Muoi	N.D	N.D	N.D	N.D	N.D	N.D	N.D	
2	Kien Giang	Go Quao	289.4	287.5	290.3	279.9	280.8	285.6 ^d	1.71	
		Gieng Rieng	175.2	177.9	176.3	175.9	177.8	176.6 ^e	0.67	
		Vinh Thuan	N.D	N.D	N.D	N.D	N.D	N.D	N.D	N.D
3	An Giang	Thoai Son	N.D	N.D	N.D	N.D	N.D	N.D	N.D	
		Long Xuyen	N.D	N.D	N.D	N.D	N.D	N.D	N.D	N.D
		Phu Tan	N.D	N.D	N.D	N.D	N.D	N.D	N.D	N.D
4	Vinh Long	Tam Binh	N.D	N.D	N.D	N.D	N.D	N.D	N.D	
		Binh Minh	N.D	N.D	N.D	N.D	N.D	N.D	N.D	N.D
		Long Ho	N.D	N.D	N.D	N.D	N.D	N.D	N.D	N.D
5	Ben Tre	Mo Cay	713.0	719.2	720.0	715.8	716.9	716.9 ^f	0.39	
		Chau Thanh	N.D	N.D	N.D	N.D	N.D	N.D	N.D	N.D
		Gieng Trom	72.5	73.1	72.8	74.0	72.7	73.0 ^b	0.81	

enrofloxacin, ciprofloxacin ($E_{1/2} = -1336$ mV), colistin ($E_{1/2} = -1120$ mV) malachite green ($E_{1/2} = -1228$ mV). Hence, the determination of

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Application

Tilapia samples from ten provinces in the Mekong River Delta were taken and analyzed to survey the erythromycin residue. Residual results could be obviously seen in the Table 15 and Figure 15.

Conclusion

A new analytical procedure based Square Wave Voltammetry had been developed for determination of erythromycin in tilapia. The proposed method was simple, quick, economical, and sensitive. It should be extensively used for veterinary drug residue screening in food surveillance programs.

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