

## Fosinopril and Carvedilol Reverse Hypertrophy and Change the Levels of Protein Kinase C $\epsilon$ and Components of Its Signaling Complex

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The Publisher regrets that the original article, published in volume 20, number 4, pages 259–271, incorrectly

captured Table 4. This was a typesetting error. The correct Table 4 is shown below.

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**Table 4** Identified results of differential proteins in PKC $\epsilon$  complexes

Protein information <sup>b</sup>	Mr	NO	Sequence coverage (%)	Alterations vs sham		Alterations vs POH-NF	
				POH-NF	POH-F	POH-NF	POH-F
				↓/↑ (fold)	↓/↑ (fold)	↓/↑ (fold)	↓/↑ (fold)
1. Hypothetical protein Loc 28126 UI	21,183	gi 30519925	33	↓4.11±0.75*			
2. Unnamed protein product(37,847/5.55)	37,847	gi 74184434	35	↓4.08±0.72*			
4. Alpha B-Crystallin ( $\alpha$ BC)	19,945	gi 57580	46	↓3.95±0.64*	↓5.08±0.81*	-	-
5. Plectin 1 isoform 6	53,5148	gi 41322931	8			-	-
6. UI							
7. UI							
8. UI							
9. Predicted: similar to mitochondrial ribosomal protein S27(RPS27)	48,131	gi 62642819	34		↑13.34±1.05**		+
10. Tripartite motif protein TRIM34 gamma(Trim34 $\gamma$ )	38,389	gi 12407451	24		↓8.58±0.79**		
12. Myosin regulatory light chain 2 Ventricular/cardiac muscle isoform (MLC2)	18,870	gi 1709058	34		↑4.15±0.65*		↑4.44±0.63 $\Delta$
13. Signal-induced proliferation associated-1(sipa-1 protein)	62,587	gi 27882584	22			+	+
14. UI							
15. Mitotic arrest deficient 1-like 1(Mad1) <sup>a</sup>	60,398	gi 6754602	31				+
16. UI							
17. UI							
18. Lyn A protein tyrosine kinase(Lyn A) <sup>a</sup>	59,295	gi 1198941	32				+
19. UI							+
20. UI							+
21. UI							+
22. UI							+
23. UI							+
24. Transforming growth factor, beta receptor II, isoform 1(TGF- $\beta$ RII-1)	65,630	gi 27363476	23				-
25. UI							
26. UI							↓10.32±0.86***
							↓10.35±0.93***

UI, protein unidentified or having a negative identified result; ↑, increased expression; ↓, decreased expression; +, newly appeared protein; -, disappeared protein; Mr, theoretical molecular mass; NO, accession number of identified protein in the web, <http://www.matrixscience.com/>

<sup>a</sup>The verified proteins by immunoblot

<sup>b</sup>The numbers in protein information indicate the spot numbers on 2-D gels  
Co-IP was performed on 3 samples for 9 times in each group. Each co-IP product was run 3 different gels. Data are represented as means±SD of at least 5 different experiments  
\* $P<0.05$  vs. sham; \*\* $P<0.01$  vs. sham; \*\*\* $P<0.001$  vs. POH-NF;  $\Delta P<0.05$  vs. POH-NF