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Flat C, 23/F., Lucky Plaza,
315-321 Lockhart Road, Wan Chai, Hong Kong, China

ESPS Peer-review Report

Name of Journal: World Journal of Biological Chemistry

ESPS Manuscript NO: 7729

Title: Functional analysis of human Na⁺/K⁺-ATPase FHM2 mutations expressed in *Xenopus* oocytes (re-submission of No. 7676)

Reviewer code: 01004042

Science editor: Zhai, Huan-Huan

Date sent for review: 2013-11-30 23:17

Date reviewed: 2013-12-03 02:51

CLASSIFICATION	LANGUAGE EVALUATION	RECOMMENDATION	CONCLUSION
<input type="checkbox"/> Grade A (Excellent)	<input type="checkbox"/> Grade A: Priority Publishing	Google Search:	<input type="checkbox"/> Accept
<input type="checkbox"/> Grade B (Very good)	<input checked="" type="checkbox"/> Grade B: minor language polishing	<input type="checkbox"/> Existed	<input type="checkbox"/> High priority for publication
<input type="checkbox"/> Grade C (Good)	<input type="checkbox"/> Grade C: a great deal of language polishing	<input type="checkbox"/> No records	<input checked="" type="checkbox"/> Rejection
<input type="checkbox"/> Grade D (Fair)	<input type="checkbox"/> Grade D: rejected	BPG Search:	<input checked="" type="checkbox"/> Minor revision
<input checked="" type="checkbox"/> Grade E (Poor)		<input type="checkbox"/> Existed	<input type="checkbox"/> Major revision
		<input type="checkbox"/> No records	

COMMENTS TO AUTHORS

The aim of this work is Functional characterization of ATP1A2 mutations that are related to Familial Hemiplegic Migraine (FHM). The approach is to express mutants of this protein in oocytes and examine the location and function of this protein. In general the work is straight forward and interesting. There are minor errors in writing. However, there is a major flaw in this work. "To reduce ouabain sensitivity, the mutations Q116R and N127D were introduced in the α 2-subunit. This construct is herein referred to as WT." Thus WT is already mutated. In the judgement of this reviewer, one cannot conclude that the consequences of the mutations would have been the same if the control was not already mutated even though the initial mutations are in a different domain of the protein. This raises doubt on validity of any of the conclusions of this work.



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Title: Functional analysis of human Na⁺/K⁺-ATPase FHM2 mutations expressed in *Xenopus* oocytes (re-submission of No. 7676)

Reviewer code: 00197104

Science editor: Zhai, Huan-Huan

Date sent for review: 2013-11-30 23:17

Date reviewed: 2013-12-10 21:09

CLASSIFICATION	LANGUAGE EVALUATION	RECOMMENDATION	CONCLUSION
<input type="checkbox"/> Grade A (Excellent)	<input type="checkbox"/> Grade A: Priority Publishing	Google Search:	<input checked="" type="checkbox"/> Accept
<input type="checkbox"/> Grade B (Very good)	<input checked="" type="checkbox"/> Grade B: minor language polishing	<input type="checkbox"/> Existed	<input type="checkbox"/> High priority for publication
<input checked="" type="checkbox"/> Grade C (Good)	<input type="checkbox"/> Grade C: a great deal of language polishing	<input type="checkbox"/> No records	<input type="checkbox"/> Rejection
<input type="checkbox"/> Grade D (Fair)	<input type="checkbox"/> Grade D: rejected	BPG Search:	<input type="checkbox"/> Minor revision
<input type="checkbox"/> Grade E (Poor)		<input type="checkbox"/> Existed	<input type="checkbox"/> Major revision
		<input type="checkbox"/> No records	

COMMENTS TO AUTHORS

In continuation to previous work, the authors have studied the functional impact of 7 new mutations in the alpha2 subunit of the NaK ATPase. They show that these mutations that are related to familial and sporadic hemiplegic migraine have different effects on enzyme activity and trafficking to the plasma membrane. The technical quality of the work is good, and the data are convincing and well discussed. Minor comments: - A careful rereading is necessary to correct a few typos (for example familial instead of familiar in familiar hemiplegic migraine, aim and introduction sections). - In the paragraph "Functional consequences" of the discussion ref 26 has been used instead of ref 15 (about P979L mutation)



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Title: Functional analysis of human Na⁺/K⁺-ATPase FHM2 mutations expressed in *Xenopus* oocytes (re-submission of No. 7676)

Reviewer code: 00503000

Science editor: Zhai, Huan-Huan

Date sent for review: 2013-11-30 23:17

Date reviewed: 2014-02-18 23:01

CLASSIFICATION	LANGUAGE EVALUATION	RECOMMENDATION	CONCLUSION
<input type="checkbox"/> Grade A (Excellent)	<input type="checkbox"/> Grade A: Priority Publishing	Google Search:	<input type="checkbox"/> Accept
<input checked="" type="checkbox"/> Grade B (Very good)	<input checked="" type="checkbox"/> Grade B: minor language polishing	<input type="checkbox"/> Existed	<input type="checkbox"/> High priority for publication
<input type="checkbox"/> Grade C (Good)	<input type="checkbox"/> Grade C: a great deal of language polishing	<input type="checkbox"/> No records	<input type="checkbox"/> Rejection
<input type="checkbox"/> Grade D (Fair)	<input type="checkbox"/> Grade D: rejected	BPG Search:	<input type="checkbox"/> Minor revision
<input type="checkbox"/> Grade E (Poor)		<input type="checkbox"/> Existed	<input type="checkbox"/> Major revision
		<input type="checkbox"/> No records	

COMMENTS TO AUTHORS

The authors study the functional consequences of seven mutations related to familial and sporadic hemiplegic migraine, in the alpha2 subunit of the NA, K ATPase. This is an interesting work that complements previous works of the authors. I consider that some issues needs to be resolved before publication. Major comments: -Due to the methodological strategy employed, the authors are studying mutations in an already mutated protein. To strengthen their conclusions, the author need to provide some evidence that the mutations introduced in the WT have no further consequences in the protein and the mutants they are studying -Western blot analyses lack a loading control - The authors must provide statistical analyses of their data. Minor comments: -In the title, the abbreviation should be changed for the complete name of the disease -In the whole manuscript, familiar must be changed for familial



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Title: Functional analysis of human Na⁺/K⁺-ATPase FHM2 mutations expressed in *Xenopus* oocytes (re-submission of No. 7676)

Reviewer code: 00504439

Science editor: Zhai, Huan-Huan

Date sent for review: 2013-11-30 23:17

Date reviewed: 2014-02-28 16:14

CLASSIFICATION	LANGUAGE EVALUATION	RECOMMENDATION	CONCLUSION
<input type="checkbox"/> Grade A (Excellent)	<input type="checkbox"/> Grade A: Priority Publishing	Google Search:	<input type="checkbox"/> [Y] Accept
<input type="checkbox"/> [Y] Grade B (Very good)	<input type="checkbox"/> [Y] Grade B: minor language polishing	<input type="checkbox"/> [] Existed	<input type="checkbox"/> [] High priority for publication
<input type="checkbox"/> [] Grade C (Good)	<input type="checkbox"/> [] Grade C: a great deal of language polishing	<input type="checkbox"/> [] No records	<input type="checkbox"/> [] Rejection
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<input type="checkbox"/> [] Grade E (Poor)		<input type="checkbox"/> [] Existed	<input type="checkbox"/> [] Major revision
		<input type="checkbox"/> [] No records	

COMMENTS TO AUTHORS

The authors provide a firm and well performed studies on WT and mutant pumps, I have very few comments, since the authors did not number the MS I will try to inform about the location of my points. 1- Introduction page, the authors mention that K binding stimulates dephosphorylation and conformational change E2 to E1 without mentioning the role of cytoplasmic ATP in inducing the E1 form. 2-Results, under electrogenic Na,Na exchange. and not "und" 3-From the abstract I have got the impression that K1003E has no impact on pump function, yet the mutation produces a clear shift in Na,Na exchange, because the affinity of the third Na site can be changed independently, it would be more precise to say that K1003 had no impact on K interaction,