Reviewer's report

Title: Clinical Features and Phylogenetic Tree of Coxsackievirus A9 in Northern Taiwan in 2011

Version: 2 Date: 4 September 2012

Reviewer: Teruo Yamashita

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Major Compulsory Revisions:
1. Page 8 (Viral identification): Authors use RNA from throat swabs for RT-PCR. However, Oberste et al. use cell culture supernatant in reference 9. I guess the method is not suitable for amplifying the virus nucleotides from throat swabs of patients.

Page 8, last sentence: Accession nos. are need for 19 formerly reported CV-A9 sequences and isolates in this study.

2. Page 10 (Results), line 7: Patients no. of age <1 and 1 are more than age 2. Age 9 and 10 are more than age 8. Why aged 2 and 8 years old?

3. Page 10 (Results), line 3 from bottom: What sre influenza-like viral illnesses in table 1?

Minor Essential Revisions:
1. Page 5(Introduction), line 5: “VP2” is incorrect. VP1 region is used for typing.

2. Page 7(Methods), 6: “real-time” should be removed.

3. Page 11, line 4, Page 14, line 3 from bottom: The most commonly seen rash pattern?

4. Page 14, line 5 from bottom: 61 should be Sixty one.

5. Page 15, line 8: 96 should be Ninety six.

6. Page 16, line 5: usually is ?

7. In references: There are no titles or bold characters in reference 2, 5, 10,

8. Table 1: Age (year) --- 4.6(3.4) Need asterisk mark. Male/female (ratio) ---- 65/35(1.9)* Remove asterisk mark.

Level of interest: An article whose findings are important to those with closely related research interests

Quality of written English: Needs some language corrections before being published

Statistical review: No, the manuscript does not need to be seen by a statistician.