

## ***Interactive comment on “MP CBM-Z V1.0: design for a new CBM-Z gas-phase chemical mechanism architecture for next generation processors” by Hui Wang et al.***

### **Anonymous Referee #2**

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My previous review had some formatting issues where 3 of the lines were truncated, and I'm sorry for any confusion caused. I include these comments below.

#### **General Comments**

5. CBM-Z output is plotted for 10 model hours, and within this time the relative error introduced by the optimisations is less than 0.05% (just), as seen in Figure 4 and mentioned throughout the manuscript. However, it is clear that this error is increasing for some species (e.g. H<sub>2</sub>O<sub>2</sub>, SO<sub>2</sub>, and H<sub>2</sub>SO<sub>4</sub> as presented, and possibly others not shown). If these simulations were run for longer than 10 hours, would these errors still be below 0.05%? For confidence in the improvements described I would expect to see

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that the errors remain low for the length of a typical CTM simulation, which could be years depending on how the author's CTM is used. If the conditions used are more realistic (see point 4 above) do these errors increase?

#### **Specific Comments and Technical Corrections**

P1L19 - I question whether the <0.05% figure is correct (see General Comments)

P6L17 - as I have mentioned in the General Comments, how robust is the error of range <0.05%?

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