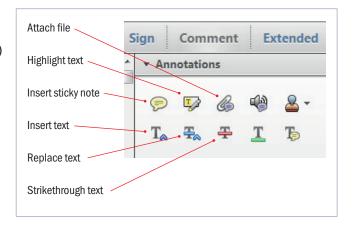
# Making corrections to your proof

Please follow these instructions to mark changes or add notes to your proof. You can use Adobe Acrobat Reader (download the most recent version from https://get.adobe.com) or an open source PDF annotator.

For Adobe Reader, the tools you need to use are contained in **Annotations** in the **Comment** toolbar. You can also right-click on the text for several options. The most useful tools have been highlighted here. If you cannot make the desired change with the tools, please insert a sticky note describing the correction.

Please ensure all changes are visible via the 'Comments List' in the annotated PDF so that your corrections are not missed.

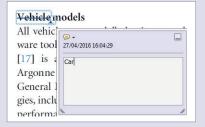


## Do not attempt to directly edit the PDF file as changes will not be visible.



## Replacing text

To replace text, highlight what you want to change then press the replace text icon, or right-click and press 'Add Note to Replace Text', then insert your text in the pop up box. Highlight the text and right click to style in bold, italic, superscript or subscript.





## **Inserting text**

Place your cursor where you want to insert text, then press the insert text icon, or right-click and press 'Insert Text at Cursor', then insert your text in the pop up box. Highlight the text and right click to style in bold, italic, superscript or subscript.





## **Deleting text**

To delete text, highlight what you want to remove then press the strikethrough icon, or right-click and press 'Strikethrough Text'.

## Vehicle models

All vehicles are modelled using comple ware tools developed with industry inpo-[17] is a vehicle simulation model Argonne National Laboratory in con General Motors to analyse specific vehgies, including the cost, fuel economy ar performance of particular ICEV and BE



## Highlighting text

To highlight text, with the cursor highlight the selected text then press the highlight text icon, or right-click and press 'Highlight text'. If you double click on this highlighted text you can add a comment.



## **QUERY FORM**

JOURNAL: Journal of Physics B: Atomic, Molecular and Optical Physics

AUTHOR: A M P Mendez et al

TITLE: Scaling rules for the ionization of biological molecules by highly charged ions

ARTICLE ID: ab9c36

Your article has been processed in line with the journal style. Your changes will be reviewed by the Production Editor, and any amendments that do not comply with journal style or grammatical correctness will not be applied and will not appear in the published article.

The layout of this article has not yet been finalized. Therefore this proof may contain columns that are not fully balanced/matched or overlapping text in inline equations; these issues will be resolved once the final corrections have been incorporated.

- Q1: Please check that the names of all authors as displayed in the proof are correct, and that all authors are linked to the correct affiliations.

  Please also confirm that the correct corresponding or has been indicated. Note that this is your last opportunity to review and amend this information before your article is published.
- Q2: If an explicit acknowledgment of funding is required, please ensure that it is indicated in year ticle. If you already have an Acknowledgments section, please check that the information there is complete and correct
- Q3: Please check that the funding information below is correct for inclusion in the article metadata. Secretaria de Ciencia y Tecnica, Universidad de Buenos Aires 20020170100727BA Consejo Nacional de Investigaciones Científicas y Técnicas PIP 2014 Agencia Nacional de Promoción Científica y Tecnológica 2017-2945
- Q4: We have been provided with ORCID iDs for the authors as below. Please confirm whether the numbers are correct.

  A M P Mendez 0000-0003-3568-7730

  C C Montanari 0000-0002-7325-6125
- Q5: Please be aware that the colour figures in this article will only appear in colour in the online version. If you require colour in the printed journal and have not previously arranged it, please contact the Production Editor now.
- Q6: Please specify the corresponding author and provide his/her email address.
- Q7: Please provide the title of the book in reference [4].

J. Phys. B: At. Mol. Opt. Phys. 0 (2020) 000000 (5pp)

https://doi.org/10.1088/1361-6455/ab9c36

## Scaling rules for the ionization of biological molecules by highly charged ions

## A M P Mendez<sup>®</sup>, C C Montanari<sup>®</sup> and J E Miraglia

Instituto de Astronomía y Física del Espacio, Consejo Nacional de Investigaciones Científicas y Técnicas-Universidad de Buenos Aires, Pabllón IAFE, 1428 Buenos Aires, Argentina

Received 6 March 2020, revised 26 May 2020 Accepted for publication 12 June 2020 Published XX XX XXXX



We investigate scaling rules for the ionization cross sections of multicharged ions on molecules of biological interest. The cross sections are obtained using a methodology presented in [Mendez et al J. Phys. B (2020)], which considers distorted-wave calculations for atomic targets combined with a molecular stoichiometric model. We examine ions with nuclear charges Z from +1 to +8 impacting on five nucleobases –adenine, cytosine, guanine, thymine, uracil-, tetrahydrofuran, pyrimidine, and water. We investigate scaling rules of the ionization cross section with the ion charge and the number of active electrons per molecule. Combining these two features, we define a scaling law for any ion and molecular target, which is valid in the intermediate to high energy range, i.e., 0.2–5 MeV amu<sup>-1</sup> for oxygen impact. Thus, the forty ion-molecule systems analyzed here can be merged into a single band. We confirm the generality of our independent scaling law with several collisional systems.

Keywords: ionization, scaling, molecules, charged-ions, DNA, multicharged ions

(Some figures may appear in colour only in the online journal)

## 1. Introduction

The interest in the ionization of biological molecules by multicharged ions has increased due to medical and environmental implementations [1], including medical treatments [2–4] and contaminant recognition in biological materials [5, 6]. Many semiempirical [7] and theoretical efforts are currently being undertaken [8-13] to get reliable values for the ionization cross sections of these molecular systems.

In recent work [8], we combined the continuum distortedwave calculations (CDW) for atoms and the simple stoichiometric model (SSM) to approximate the ionization cross sections of complex molecular targets by the impact of charged ions. The molecular ionization cross section  $\sigma_{\rm M}$  was expressed as a linear combination of atomic CDW calculations  $\sigma_A$ , weighted with the number of atoms for each specie  $n_A$ , i.e.,  $\sigma_M = \sum_A n_A \sigma_A$ . The CDW-SSM approximation showed consistent results for over a hundred of biologically relevant ion-molecule systems. As expected, in the high energy range (i.e., above 5 MeV amu<sup>-1</sup>), the ionization cross sections of the molecular systems follow the  $Z^2$  dependence predicted by the first Born approximation. However, at intermediate energies, the dependence with Z is not straightforward since non-perturbative models are mandatory.

This contribution constitutes a follow-up to our previous work [8]. We introduce here a two-folded scaling rule for the ionization cross sections of complex molecules by charged ions. Our approach considers the dependence of the cross section with the ion charge Z and incorporates the scaling of the ionization with the number of active electron  $n_e$  of the molecular targets. Scaling rules are generally very useful since they can be used as first-order approximations in experimental measurements and multipurpose codes.

## 2. Scaling rules

## 2.1. Scale with the ion charge

In the development of our scaling rule, we examine forty collisional systems. The target-ion systems are composed of eight targets: the DNA and RNA nucleobases -adenine, cytosine, guanine, thymine, uracil-, tetrahydrofuran (THF), pyrimidine, and water; and five ion species: H+, He+2, Be+4, C+6, and  $O^{+8}$ . We consider these systems as a benchmark for the present

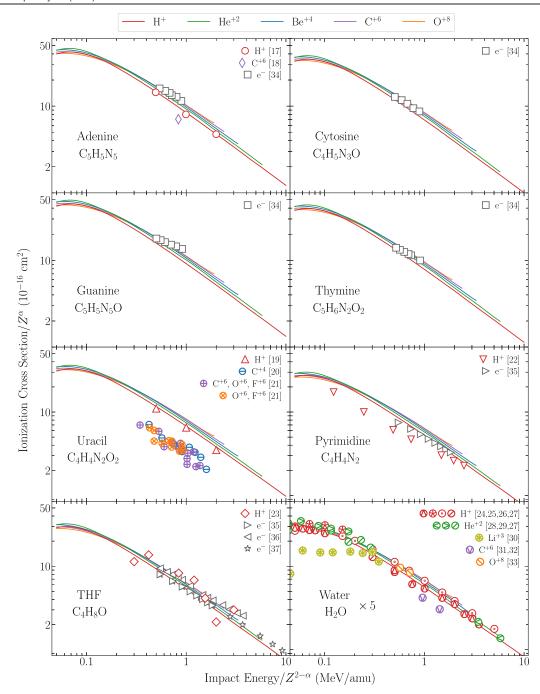


Figure 1. Scaled ionization cross section  $\sigma/Z^{\alpha}$  as a function of the ion impact energy  $E/Z^{2-\alpha}$  with  $\alpha=1.2$ . Colors are associated with the incident ion labeled on top of the figure. Curves: present CDW-SSM theoretical results. Symbols: experimental data [17–33]. Electron impact ionization values [34–37] are included with the corresponding equi-velocity conversion.

**Table 1.** Number of active electrons per target at intermediate to high energies obtained from the CDW calculations [8].

Molecule	$n_{\rm e}$	Molecule	$n_{\rm e}$	Molecule	ne
H <sub>2</sub> O	6	$CO_2$	12	C <sub>4</sub> H <sub>5</sub> N <sub>3</sub> O	37
$N_2$	8	$C_4H_8O$	28	$C_5H_6N_2O_2$	42
$O_2$	8	$C_4H_4N_2$	28	$C_5H_5N_5$	45
CH <sub>4</sub>	8	$C_4H_4N_2O_2$	36	$C_5H_5N_5O$	49

We found two types of Z-scaling laws in the literature applicable to the intermediate impact energy range. The rule

suggested by Janev and Presnyakov [14] considers  $\sigma/Z$  versus E/Z to be the *natural* reduced form of the ionization cross section  $\sigma$  and the incident ion energy E. More recently, Montenegro and co-workers [15, 16] suggested an alternative scaling by taking into account that the cross section is a function of  $Z^2/E$  at high energies. Their scaling, given by

$$\sigma/Z^{\alpha} = f(E/Z^{2-\alpha}),\tag{1}$$

keeps the  $Z^2/E$  relationship for any value of the parameter  $\alpha$ . The authors proposed  $\alpha = 4/3$  for the ionization of He and H<sub>2</sub> by differently charged ions [15].

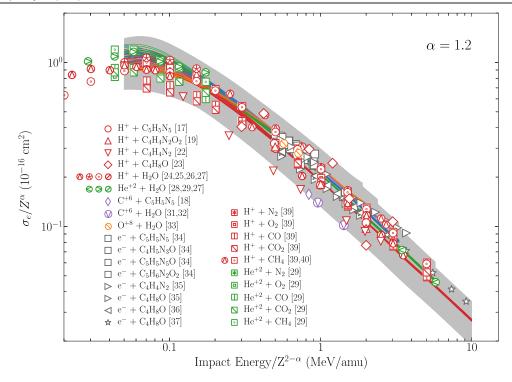


Figure 2. Ionization cross section reduced with the ion charge Z and scaled with number of active electrons per molecule  $n_e$ , given by equation (3) with  $\alpha = 1.2$ . Curves: present CDW-SSM theoretical results. Symbols: experimental data [17–19, 22–29, 31–33, 39, 40]. Electron impact ionization values [34–37] are included with the corresponding equi-velocity conversion.

Following the work of Montenegro and collaborators, we found that the parameter that best converges the CDW–SSM cross sections of the forty collisional systems over the broadest energy range is  $\alpha=1.2$ . The validity of this particular scaling is evident in figure 1, where–for each target–the CDW–SSM curves corresponding to different ions lay one over the other. It is worth noting that our theoretical results are valid for impact energies above the maximum of the cross sections, which corresponds to an impact energy range from 50 keV for H $^+$  to 250 keV amu $^{-1}$  for O $^{+8}$ .

We also examined the experimental data available for the forty ion-target systems [17–2 $\frac{1}{\sqrt{3}}$ –33] with the  $Z^{\alpha}$ -scaling rule. For targets with none or little experimental data, we included electron impact ionization results [34–37] at high velocity with the corresponding equivelocity conversion. As can be noted, most of the data in figure 1 confirm the present scaling, even for  $O^{+8}$  in water [33]. Only two data sets are off our predictions: the ionization cross section of uracil by swift C, O, and F ions from references [20, 21], and the values for  $Li^{+3}$  in water from reference [30] for E < 600 keV amu<sup>-1</sup>. In the case of uracil, recent CTMC calculations by Sarkadi [38] are also above the experimental values by Tribedi and collaborators [20, 21].

## 2.2. Scale with the molecular target

The good results obtained in the scaling with the ion charge encouraged us to further investigate a scaling law that could predict values for ionization cross sections of any ion in any molecule. To this end, we considered the number of active electrons in each molecule  $n_e$  proposed in reference [8] and combined it with the  $Z^{\alpha}$ -scaling from section 2.1.

In our previous work, we noticed that the CDW ionization cross sections  $\sigma_A$  of atomic targets H, C, N, and O scale with the number of active electrons per atom  $\nu_A$ , as  $\sigma_e = \sigma_A/\nu_A$ , where  $\nu_A$  is 1 for H and 4 for C, N, O, i.e.,

$$\frac{\sigma_{\rm H}}{1} \sim \frac{\sigma_{\rm C}}{4} \sim \frac{\sigma_{\rm N}}{4} \sim \frac{\sigma_{\rm O}}{4}.$$
 (2)

By means of the SSM, we define the number of active electrons per molecule as  $n_e = \sum_A n_A \nu_A$ . The  $n_e$  values for the molecular targets considered throughout this work are displayed in table 1. The scaling with the molecular number of active electrons proved to give excellent results, as shown in figure 6 of reference [8].

## 2.3. Scale with the ion charge and the molecular target

By incorporating the  $Z^{\alpha}$  reduction and the scaling with the number of active electrons, we introduce the scaled and reduced ionization cross section of molecules  $\tilde{\sigma}$ , which is expressed as a function of  $E/Z^{2-\alpha}$ , and it is given by

$$\tilde{\sigma} = \frac{\sigma_{\rm e}}{Z^{\alpha}} = \frac{\sigma_{\rm M}/n_{\rm e}}{Z^{\alpha}},\tag{3}$$

where  $\sigma_{\rm M}$  is the ionization cross section for the molecular target,  $n_{\rm e}$  is the number of active electrons per molecule displayed in table 1, and the parameter is  $\alpha=1.2$ . Figure 2 shows the theoretical and experimental values of  $\tilde{\sigma}$  (given by equation (3)) for all the systems displayed in figure 1. As can be noted, the scaling works very well and is independent of the ion charge or the complexity of the molecular target. Our theoretical curves lay in a narrow band valid for any charged

ion (reduced with  $Z^{\alpha}$ ) in any molecule (scaled with the number of active electrons) with a dispersion of about  $\pm 20\%$ . If we consider the experimental data, the uncertainty of our scaling grows to  $\pm 30\%$ , which is schematized in figure 1 with a gray area. It is worth noting that we did not include in this figure the data for uracil from references [20, 21], and Li<sup>+3</sup> on water [30]. The discussion about these experimental values exceeds the present work.

We consider the present scaling robust enough to be valid for different ion-molecule combinations. We tested the generality of our model by including in figure 1 several data sets of molecular targets not considered previously, such as the measurements by Rudd *et al* [29, 39] for  $H^+$  and  $He^{+2}$  in  $N_2$ ,  $O_2$ ,  $CH_4$ , CO and  $CO_2$ , and recent values by Luna *et al* [40] for  $H^+$  in  $CH_4$ .

The good agreement shown in figure 2 summarizes the main result of this work and holds the validity of the present scaling for different ions and targets. Although the theoretical CDW-SSM results are valid for energies above the maximum of the cross sections, the scaling of the experimental data extends even to lower impact energies, as can be noted in figure 2. New measurements for other ions and molecules are expected to reinforce the present proposal.

## 3. Conclusions

We present scaling rules for the ionization cross sections of highly charged ions in biological targets. The first scaling reduces the nature of the projectile by scaling the cross section with the ion charge,  $Z^{\alpha}$ , as a function of the reduced impact energy  $E/Z^{2-\alpha}$ , with  $\alpha=1.2$ . The second scaling considers the molecular description of the target by taking into account the number of active electrons per molecule,  $n_{\rm e}$ . The last scaling law combines the  $Z^{\alpha}$ -reduction with the  $n_{\rm e}$ -scaling of the cross section, and it becomes independent of the ion charge and the molecular target. The scalings are obtained by means of CDW–SSM calculations for five different charged ions in eight targets and tested with the available experimental data. The generality of our independent scaling is proved to be valid in a wide energy range by considering a significant number of experimental data sets for other collisional systems.

## **Acknowledgments**

This work was finantially supported by Consejo Nacional de Investigaciones Científicas y Técnicas (PIP 2014), Agencia Nacional de Promoción Científica y Tecnológica (PICT 2017-2945), and Universidad de Buenos Aires (UBACyT 20020170100727).

## **ORCID iDs**

A M P Mendez https://orcid.org/0000-0003-3568-7730 C C Montanari https://orcid.org/0000-0002-7325-6125

## References

- [1] Liamsuwan T and Nikjoo H 2013 *Phys. Med. Biol.* **58** 641–72
- [2] Mohamad O, Sishc B J, Saha J, Pompos A, Rahimi A, Story M D, Davis A J and Kim D N 2017 Cancers 9 66

- [3] Solov'yov A V, Surdutovich E, Scifoni E, Mishustin I and Greiner W 2009 Phys. Rev. E 79 011909
- [4] Denifl S, Märk T D and Scheier P 2012 ed G García Gómez-Tejedor and M Fuss (Dordrecht: Springer)
- [5] Gafur N A, Sakakibara M, Sano S and Sera K A 2018 Water 10 1507
- [6] Benedetti D, Nunes E, Sarmento M, Porto C, Iochims dos Santos C E, Ferraz Dias J and da Silva J 2013 Mutation Research/Genetic Toxicology and Environmental Mutagenesis 752 28–33
- [7] de Vera P, Garcia-Molina R, Abril I and Solov'yov A V 2013 Phys. Rev. Lett. 110 148104
- [8] Mendez A M P, Montanari C C and Miraglia J E 2020 J. Phys. B: At. Mol. Opt. Phys. 53 055201
- [9] Quinto M A, Monti J M, Tachino C A, Weck P F, Fojón O A, Champion C and Rivarola R D 2020 Radiat. Phys. Chem. 167 108337
- [10] Lüdde H J, Horbatsch M and Kirchner T 2019 J. Phys. B: At. Mol. Opt. Phys. 52 195203
- [11] Lüdde H J, Horbatsch M and Kirchner T 2018 Eur. Phys. J. B 91 99
- [12] Lüdde H J, Achenbach A, Kalkbrenner T, Jankowiak H-C and Kirchner T 2016 Eur. Phys. J. D 70 82
- [13] Champion C et al 2012 J. Phys.: Conf. Ser. 373 012004
- [14] Janev R K and Presnyakov L P 1980 J. Phys. B: At. Mol. Opt. Phys. 13 4233
- [15] DuBois R D, Montenegro E C and Sigaud G M 2013 AIP Conf. Proc. 1525 679
- [16] Montenegro E C, Sigaud G M and DuBois R D 2013 Phys. Rev. A 87 012706
- [17] Iriki Y, Kikuchi Y, Imai M and Itoh A 2011 Phys. Rev. A 84 052719
- [18] Bhattacharjee S, Bagdia C, Chowdhury M R, Mandal A, Monti J M, Rivarola R D and Tribedi L C 2019 Phys. Rev. A 100 012703
- [19] Itoh A, Iriki Y, Imai M, Champion C and Rivarola R D 2013 Phys. Rev. A 88 052711
- [20] Agnihotri A N et al 2012 Phys. Rev. A 85 032711
- [21] Agnihotri A N et al 2013 J. Phys. B: At. Mol. Opt. Phys. 46 185201
- [22] Wolff W, Luna H, Sigaud L, Tavares A C and Montenegro E C 2014 J. Chem. Phys. 140 064309
- [23] Wang M, Rudek B, Bennett D, de Vera P, Bug M, Buhr T, Baek W Y, Hilgers G and Rabus H 2016 Phys. Rev. A 93 052711
- [24] Luna H et al 2007 Phys. Rev. A 75 042711
- [25] Bolorizadeh M A and Rudd M E 1986 Phys. Rev. A 33 888
- [26] Rudd M E, Goffe T V, DuBois R D and Toburen L H 1985 *Phys. Rev.* A 31 492
- [27] Toburen L H, Wilson W E and Popowich R J 1980 *Radiat. Res.* 82 27–44
- [28] Ohsawa D, Sato Y, Okada Y, Shevelko V P and Soga F 2005 Phys. Rev. A 72 062710
- [29] Rudd M E, Goffe T V and Itoh A 1985 Phys. Rev. A 32 2128
- [30] Luna H, Wolff W, Montenegro E C, Tavares A C, Ludde H J, Schenk G, Horbatsch M and Kirchner T 2016 Phys. Rev. A 93 052705
- [31] Dal Cappello C, Champion C, Boudrioua O, Lekadir H, Sato Y and Ohsawa D 2009 Nucl. Instrum. Methods Phys. Res. B 267 781–90
- [32] Bhattacharjee S, Biswas S, Monti J M, Rivarola R D and Tribedi L C 2017 Phys. Rev. A 96 052707
- [33] Bhattacharjee S et al 2016 J. Phys. B: At. Mol. Opt. Phys. 49 065202
- [34] Rahman M A and Krishnakumar E 2016 Electron ionization of DNA bases J. Chem. Phys. 144 161102
- [35] Bug M U, Baek W Y, Rabus H, Villagrasa C, Meylan S and Rosenfeld A B 2017 Radiat. Phys. Chem. 130 459 -79
- [36] Wolff W, Rudek B, da Silva L A, Hilgers G, Montenegro E C and Homem M G P 2019 J. Chem. Phys. 151 064304

- [37] Fuss M, Muñoz A, Oller J C, Blanco F, Almeida D, Limão-Vieira P, Do T P D, Brunger M J and García G 2009 *Phys. Rev.* A **80** 052709
- [38] Sarkadi L 2016 J. Phys. B: At. Mol. Opt. Phys. 49 185203
- [39] Rudd M E, DuBois R D, Toburen L H, Ratcliffe C A and Goffe
- T V 1983 *Phys. Rev.* A **28** 3244 [40] Luna H, Wolff W, Montenegro E C and Sigaud L 2019 *Phys.* Rev. A 99 012709