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1 Interplay between ionizing radiation effects and aging in *C. elegans*

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Abstract

Living species are chronically exposed to environmental ionizing radiations from sources that can be overexpressed by nuclear accidents. In invertebrates, reproduction is the most radiosensitive studied endpoint, likely to be connected with aging. Surprisingly, aging is a sparsely investigated endpoint after chronic ionizing radiation, whereas understanding it is of fundamental interest in biology and medicine. Indeed, aging and aging-related diseases (e.g., cancer and degenerative diseases) cause about 90 % of deaths in developed countries. Therefore, glp-1 sterile Caenorhabditis elegans nematode was used to assess the impact of chronic gamma irradiation on the lifespan. Analyses were performed, at the individual level, on aging and, in order to delve deeper into the mechanisms, at the molecular level, on oxidative damage (carbonylation), biomolecules (lipids, proteins and nucleic acids) and their colocalization. We observed that ionizing radiation accelerates aging (whatever the duration (3 to 19 days)/dose (0.5 to 24 Gy)/dose rate (7 and 52 mGy.h⁻¹) tested) leading to a longevity value equivalent to that of wt nematode (~25-30 days). Moreover, the level of protein oxidative damage (carbonylation) turned out to be good cellular biomarker of aging, since it increases with age. Conversely, chronic radiation treatments reduced carbonylation levels and induced neutral lipid catabolism whatever the dose rate and the final delivered dose. Finally, under some conditions a lipidprotein colocalization without any carbonyl was observed; this could be linked to yolk accumulation in glp-1 nematodes. To conclude, we noticed through this study a link between chronic gamma exposure, lifespan shortening and lipid level decrease associated with a decrease in the overall carbonylation.

31 1. Introduction

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As nature provided environmental chronic ionizing radiations that can be overexpressed by nuclear accidents, e.g., Fukushima [1], environmental species are chronically exposed to ionizing radiation. Reported effects on gamma ray exposure are heterogeneous, depending on the mode of irradiation, and suffer from a lack of knowledge regarding the impact of chronic gamma irradiation. For many invertebrates, reproduction is the most radiosensitive studied parameter, after both acute and chronic irradiation [2-4]. This also includes roundworm Caenorhabditis elegans (C. elegans), where reduction of the progeny number is observed after both acute and chronic irradiation, whereas the hatching success decreases after acute irradiation [5-8]. While germline ablation has been shown to impact lifespan, a reproduction impairment could also influence the aging process and, finally, the longevity. Aging is considered to be a complex process involving both several molecular pathways and stochastic events for its modulation [9]. Energy resources are of crucial importance for biomolecular fitness, which slows down the aging process. There is a constant trade-off between investment in somatic maintenance and reproduction. Indeed, in organisms such as C. elegans, the reproduction has been proven to be an energetically expensive process with profound metabolic effect on the lifespan [10]. Moreover, germline ablated nematodes showed a significantly prolonged lifespan of 60 %, and some of these conditional mutants are well suited to study the biochemistry of aging [11]. C. elegans is one of the premier metazoan models extensively used to investigate the molecular mechanisms of aging [12]. Despite the apparent complexity of aging, experiments performed on the C. elegans model have helped to outline the few key mechanisms that affect lifespan [13]: germline ablation [14], mTOR pathway [15] and IIS pathway [16]. Some of the transcription factors that are involved in lifespan extension by germline elimination are DAF-2, DAF-12 and DAF-16-, which are also involved in the IIS pathway [17]. Moreover, another gene-block is known to regulate worm development, phenotype and lifespan, e.g., clock genes (clk-1, clk-2, clk-3 and gro-1) [18]. Numerous mechanisms, like autophagy and proteostasis control [19], modulate aging and influence these genes. Finally, various genetic factors, e.g., the mammalian tumour suppressor p53 ortholog CEP-1, were reported to regulate aging and induce an adaptive response in C. elegans [20]. The most popular theory of aging, called the "free-radical theory of aging" (FRTA), is based on free-radical induced oxidative damage. Reactive Oxygen Species (ROS) appear to be an important signalling intermediate, whereas mitochondria are crucial organelles for life and aging by hosting numerous metabolic processes, particularly electron transport and ATP-production [21]. Various biomolecular targets have been recognized to be involved in the molecular processes that regulate aging. Recently, it was shown that epigenetic drift, mediated by various PTMs such as DNA methylation and histone modification, impacts gene expressions and biomolecular interactions, which in turn finally affects aging [22]. Moreover, as the aging proceeds, massive metabolome and proteome changes that include decreased abundance of proteins acting for transcription termination, mRNA

degradation, mRNA stability, protein synthesis, and proteasome function [23] occur. Furthermore, a decrease in protein turnover has been identified as one of the biomarkers of aging with no relation to the protein abundance over time [24]. In the course of aging, there is an exponential increase of oxidative damage in proteins of human fibroblasts between the ages of 17 and 80 [25], while lipid membrane modifications have been proposed to play a major role in the process of aging [26]. Elevation of lipid droplets (highly conserved neutral lipid storage), can cause various metabolic syndromes, including accelerated aging. One of the most studied oxidative damages to proteins is carbonylation [27]. Protein carbonylation is an irreversible post-translational modification, and is also an aging and oxidative stress biomarker [28]. Protein carbonylation is not an enzymatic process and can be induced via four chemical mechanisms [29]. Since the FRTA theory of aging and gamma ionizing radiation are mainly mediated through ROS generation, they can share some similar mechanisms. An old theory [30] recalled recently [31], stated that ionizing radiation accelerates the process of aging by sharing ROS intermediates and similar mechanisms [32]. Given that aging is a poorly investigated endpoint after chronic exposure to ionizing radiations, a conditional sterile glp-1 C. elegans mutant (a dedicated model for lifespan studies) was used to assess the impact of chronic gamma irradiation on its lifespan without any influence of either reproduction or progeny. Glp-1 regulates both somatic and germline growth and also differentiation in C. elegans, whereas the glp-1 sterile mutant facilitates the study of radiation effects on a C. elegans cohort that does not reproduce. To test the impact of the irradiation period, the dose, and the dose rate, chronic gamma radiation exposure was performed at 2 different dose rates, i.e., 7 and 52 mGy.h⁻¹, during 2 different periods, i.e., from eggs to young adult stage (64 h), and from eggs to old stage (19 days), with recovery conditions. This was accompanied by non-irradiated control, in order to follow the normal aging process. The lifespan, the levels of carbonyls, i.e., carbonylation of proteins, and of biomolecules like proteins, lipids and nucleic acids, were monitored under all of these conditions. Conversely, despite already known radioinduced apoptosis [33, 34]. DNA damage was not analyzed here mainly because DNA repair, which is a rapid process, and apoptosis are expected to be low in germline-less glp-1 C. elegans.

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2. Results

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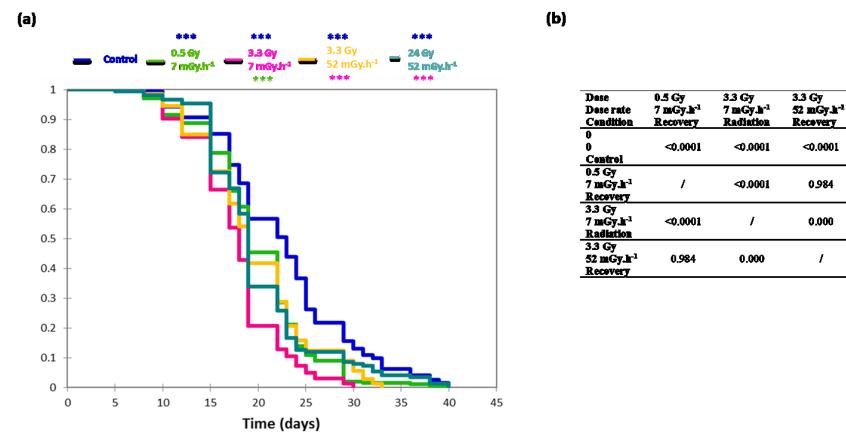
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2.1. Impact of low chronic gamma irradiation on the lifespan

- We used the *glp-1* mutant strain of *C. elegans*, which are sterile at a restrictive temperature (25°C).
- 97 Together with control experiments, we performed chronic gamma ray irradiations (as described in the
- 98 Materials and Methods section, summary in Figure S1) with two different dose rates of 7 and 52
- 99 mGy.h⁻¹ and during two different periods of life, as follows:
- Gamma irradiation from eggs to young adult stage (i.e., 64 hours of development time, equivalent to the 1st day of adulthood);
 - Gamma irradiation from eggs to old stage (i.e., 19 days, which is equivalent to the 17th day of adulthood).

The aim of the study was, firstly, to compare the effects induced by these 2 irradiation dose rates (given that chronic exposure conditions are being used) for a same period of exposure (eggs to L4, given 2 different doses) and also for a same dose, i.e., 3.3Gy (at the old stage, with one exposure ending earlier, yielding recovery groups). As shown in Figure 1, we noticed a highly significant lifespan shortening (Log-Rank p<0.0001) upon low chronic gamma irradiation in comparison with non-irradiated controls, which showed an expected lifespan as previously reported [11]. Moreover, we also noticed that the highly significant lifespan shortening was observed whatever the dose rate, the final dose or the duration of irradiation (Fig. 1). More precisely, the irradiation condition of 7 mGy.h⁻¹ differed significantly from all of the other irradiation conditions assessed (Fig. 1) and showed a maximum lifespan shortening effect. This result indicates that the "long term irradiation" treatment (which spans 19 days) with a dose of 7 mGy.h⁻¹ may induce the maximal shortening effect on the animal's lifespan expectancy and differs significantly from the other irradiation treatments. Furthermore, we observed that a higher dose did not result in a worsening effect on the lifespan reduction. Accordingly, as shown in Fig. 1a and b, a "long-term irradiation" with a higher dose rate like 52 mGy.h⁻¹ did not result in a greater shortening of the lifespan expectancy. To complete the irradiation effect on the worm population, we performed a "recovery treatment", which involves gamma ray irradiation from eggs to young adult stage (over a development period spanning 64 h) followed by a recovery period, until animal death. We then observed that the animals of the "recovery treatment" groups, irradiated with either 7 or 52 mGy.h⁻¹, were not significantly different from each other (Fig. 1b). Altogether, these results could indicate that gamma ray irradiation can induce a phenomenon with a binary response without any dose response relationship, i.e., as soon as C. elegans have been irradiated at the egg stage, the lifespan has been shortened and nearly equals that of N2 wild type C. elegans (Figure S2). To compare, the lifespan of N2 C. elegans exposed to chronic gamma irradiation from eggs to L4 stage with the dose rate of 52 mGy.h⁻¹ did not show a significant difference relative to the unirradiated control ones (Kaplan-Meier analysis, P-value = 0.075) (Fig. S2). To

- investigate the underlying mechanisms in order to explain our results, we monitored the assay at the
- molecular levels.



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Figure 1 – (a). Kaplan-Meier estimation of the *glp-1* mutant lifespan after gamma irradiation. Various conditions were considered as individual statistical groups: Control (blue line)- *C. elegans* without irradiation treatment; 0.5 Gy (green line)- Gamma irradiation from eggs to L4 stage with 7 mGy.h⁻¹ and recovery until death occurs; 3.3 Gy (yellow line)- Gamma irradiation from eggs to L4 stage with 52 mGy.h⁻¹ and recovery until death occurs; 24 Gy (turquoise line)- Gamma irradiation from eggs to old stage with 52 mGy.h⁻¹ and recovery until death occurs. The experiment was performed in triplicate. Treatments with stars are significantly different (Log-Rank, $\alpha = 0.01$, p<0.0001***). Blue stars depict a significant difference between control and all treatments; green stars depict a significant difference between 0.5 Gy and 3.3 Gy irradiations with a same dose rate (7 mGy.h⁻¹); pink stars depict a significant difference between 3.3 Gy cumulative irradiation (with 7 mGy.h⁻¹) and treatments with 52 mGy.h⁻¹ (with cumulative doses of 3.3 and 24 Gy). (b) Statistical analysis summary (XLSTAT software, Addinsoft, New York, NY, USA) of the results plotted in Figure 1-(a), showing significant degree differences between survivals calculated with the Log-Rank test with 99% confidence.

24 Gy

52 mGy.h¹

Radiation

< 0.0001

0.9

0.000

0.484

2.2. Biomolecular characterization of aging, radiation and recovery processes

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In situ visualization of C. elegans carbonylation and biomolecular colocalization were performed by a recently published method [35]. Figure 2a gives an example of carbonyl and biomolecule visualization under our conditions. As shown in Figure 2, we were able to uncover the evolution of both the carbonylation and biomolecule levels in the course of the normal aging and as a response to contrasted ionizing radiation conditions. Figures 2b and f (carbonyl panel) reveal a significant carbonylation increase, 2.5-3 fold, between 1 (young adult worm group) and 17 days (elderly worm group) of adulthood. Interestingly, we also noticed that irradiated groups did show a comparable increase in carbonylation. Conversely, a chronic gamma ray irradiation induced a detectable reduction of the carbonyl staining (Figs. 2b and f, carbonyl panel) in both young (2-3 fold change) and old worm (1.5-5 fold change) groups. We also observed that this carbonylation decrease did not depend on the dose rate, the final dose or the duration of the gamma irradiation (Figs. 2b and f). On the basis of the results presented above, we can conclude that a chronic gamma irradiation decreases the level of carbonylation, whereas aging can increase the carbonylation levels in both unirradiated and irradiated nematodes. All of these results have been confirmed by protein carbonylation measurement in vitro on extracted proteins using new methodology recently developed [36] (data not shown).

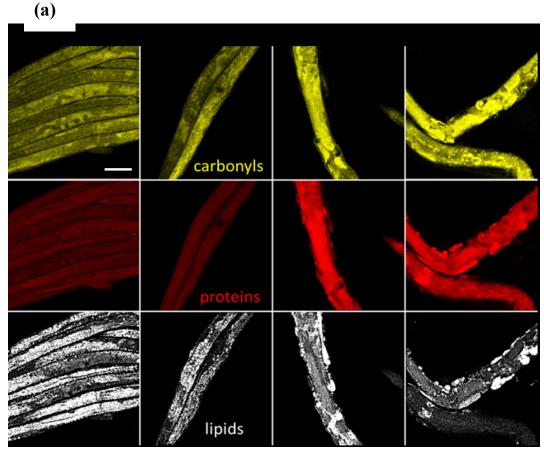
Interestingly, lipid level trends after irradiation are similar to carbonylation ones. Namely, irradiation decreases lipid levels at the young (3.75-7.5 fold) and old stages (3-13 fold) (Figs. 2c and g, lipid panel). However, no significant changes in the lipid levels have been observed with aging. These results have been confirmed by lipid extraction and thin layer chromatography (HPTLC) measurement (data not shown) to avoid erroneous conclusions due to a methodology bias; e.g., loss of labelling signal due to the disruption of fat storage organelles [37]. In addition, the lipid distribution along the nematode body appears to be impacted by aging and irradiation. Those observations are simplified in Scheme S1. However, only a few worms were observed per condition and need to be verified through visualization of lipids and their distribution; e.g., with electron microscopy.

The protein level is not significantly changed either by aging or by irradiation (Figs. 2e and h, protein panel). Beyond this result, the stable level of protein constituted a labelling control, showing that old worms are not easier to permeabilize than young ones are. A complementary assay for protein measurement has been done by extraction and BCA assay measurement. Protein concentrations vary between 1 and 1.52 mg.ml⁻¹ over all conditions (young and old worms, irradiated and unirradiated controls); protein concentrations are not significantly different between conditions (Kruskal-Wallis test, P-value = 0.1257) (Supplementary data, Figure S3).

172 On the other hand, chronic exposure of N2 C. elegans to gamma rays (with 3 different dose rates: 50,

75 and 100 mGy.h⁻¹) did not significantly alter the levels of carbonylation and biomolecules (protein, 173

- lipid and DNA) [8] compare to the controls, except at 6.8 Gy where the lipid levels decreased
- dramatically (Figure S4).



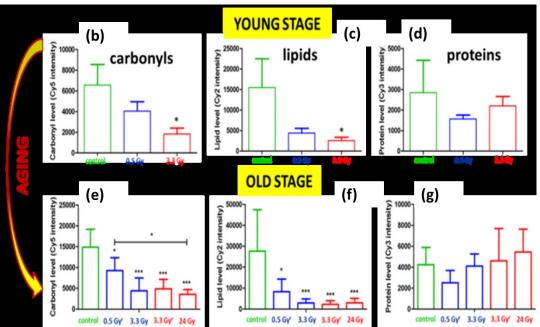


Figure 2 – (a). Impact of aging, radiation and recovery after different radiation treatments on biomolecule levels of *C. elegans*, at young (3rd day of nematode life) and old (19th day of nematode life) stages. Biomolecules were visualized *in situ* with confocal microscope. Mean ± SE, n=15. Scale bar is 10 μm. Confocal images were visualized by using a colour-blind friendly palette. Those images are examples of young and old worms (control and irradiated) with the labeling of carbonyls (yellow), proteins (red) and lipids (white). Irradiated worms represented here had been subjected to a dose rate of 52 mGy.h⁻¹ cumulating the dose of 3.3 Gy and 24 Gy at the young and old stages, respectively. Biomolecular levels of carbonyls (**b, e**), of lipids (**c, f**), and of proteins (**d, g**) were measured *in situ* and quantified with ImageQuantTL at two stages; i.e., the young (**b, c, d**) and old stages (**e, f, g**). Mean ± SE, n=3, 3, 6 (young: control, 0.5 and 3.3 Gy), n=9, 6, 6, 9, 6 (old: control, 0.5 Gy, 3.3 Gy, 3.3 Gy recovery and 24 Gy). Treatments with stars are significantly different (Tukey and Kruskal-Wallis tests; * < 0.05; **<0.01; ***<0.01.

To highlight which biomolecule holds carbonyls, the biomolecule colocalizations were assessed from the young stage (3rd day of nematode life) up until the old stage (19th day of nematode life) using a dedicated software [38], and following a methodology as depicted in **Scheme S1**. Therefore, biomolecule colocalization under all conditions during adulthood was compared (**Figure 3**).

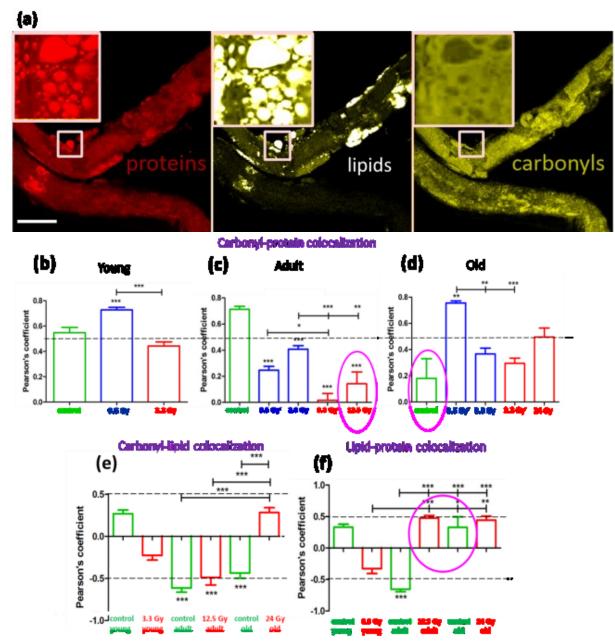


Figure 3 – (a). Biomolecules were visualized *in situ* with a confocal microscope. This confocal image is an example of old worms that display the colocalization between lipids and proteins with partial exclusion of carbonyls, with the higher magnification in the white zoom box. Confocal images were visualized by using a colour-blind-friendly palette. Mean ± SE, n=15. Scale bar is 10 μm. **(b-f)** Impact of aging, radiation and recovery after irradiation on colocalization between biomolecules in young (3rd day of nematode life), adult (10th day of adulthood) and old (19th day of nematode life) *C. elegans*. Colocalization was measured via Pearson's coefficient in ImageJ using the JACoP plugin. The colour green represents the controls in different stages, blue represents irradiation with a dose rate of 7 mGy.h⁻¹, and red represents irradiation with 52 mGy.h⁻¹. r stands for recovery after irradiation that took place during the development from eggs to young stage. Carbonyl-protein colocalization was calculated at the young, adult and old stage in Graphs b, c and d, respectively. Carbonyl-lipid colocalization is calculated in Graph e, while protein-lipid colocalization is depicted in Graph f. Pink circles represent the samples that were finally analyzed for lipid and protein colocalization. Treatments with stars are significantly different (Tukey and Kruskal-Wallis tests; * < 0.05; **<0.01; ***<0.001). The dashed line indicates limits of Pc coefficient-significance.

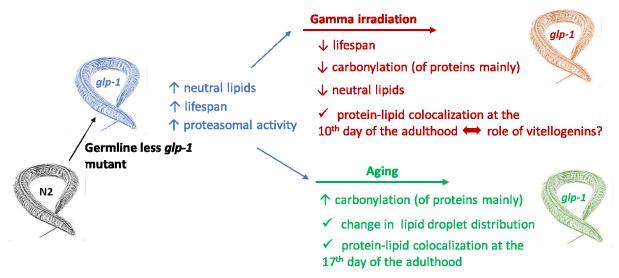
Positive and negative colocalizations are considered to be significant if the value of Pearson's coefficient (Pc) ranges between 0.5 and 1 and between -0.5 and -1, respectively. Values in between -0.5 and 0.5 were then considered as non-significant and values close to zero mean that there is no correlation. Results are shown in **Figures 3b-f**. Significant differences between Pc, illustrating colocalization between two macromolecules, were observed at the young, adult and old stage; i.e., 1st day, 10th day and 17th day of adulthood. Colocalization of carbonyls and proteins was firstly analyzed (**Figures 3b-d, first row**). Under most conditions, carbonyls colocalize mainly with proteins. However, at the young stages, the high dose rate (52 mGy.h⁻¹) irradiation tends to display a lower colocalization between carbonyls and proteins than the low dose rate (7 mGy.h⁻¹) irradiation (**Figure 3b**); it is the same at the 10th day of nematode adulthood, under all tested irradiations compared to the control group (**Figure 3c**). However, at the same adulthood stage (10th day), high dose rate (52 mGy.h⁻¹, in recovery and long-radiation groups) irradiated worms manifested something never observed yet; namely, no colocalization between carbonyls and proteins (**Figure 3c**). This phenomenon is also observed in control old nematodes (19 days), at the 17th day of adulthood; that is, with a delay of 7 days compared to the adult stage irradiated group (52 mGy.h⁻¹) (**Figure 3d**).

Although carbonyls do not colocalize with proteins, they might colocalize with other biomolecules, such as nucleic acids or lipids. Namely, carbonyls colocalize positively with nucleic acids only in young stage irradiated worms (0.5 Gy), under the same condition in which nucleic acids also display a colocalization with proteins (data not shown). On the other hand, carbonyls and lipids do not colocalize in young-stage nematodes for control (Pc <0.5) (Figure 3e). Conversely, 52 mGy.h⁻¹ radiation seems to induce a negative correlation (still weak as <-0.5) between carbonyls and lipids, except for old-stage worms, in which a non-significant positive correlation is observed. This negative correlation is enhanced and becomes significant at the adult stage (in both control and irradiated worms) and at the old stage (in control worms) (Fig. 3e).

Finally, the Pc between lipids and proteins has been calculated (**Figure 3f**). We observed a similar pattern to that found for carbonyls and lipids at the young stage (both control and irradiated), for adult controls (showing a negative correlation) and for irradiated old worms (**Fig. 3e**). Conversely, at the adult stage (10th day of adulthood) after chronic irradiation at 52 mGy.h⁻¹ and for the old control, the Pc pattern is totally opposite to that between lipids and carbonyls, as it shows a slight positive colocalization between lipids and proteins (**Fig. 3f**). Positive colocalization between proteins and lipids that partially exclude carbonyl groups can also be visualized from the confocal image (**Figure 3a**).

To conclude with regard to these biomolecular colocalization results, proteins and carbonyls mostly colocalize at the young stage of *C. elegans*, as published before [35], while there is only a weak colocalization between proteins and lipids. This can be due to the level change of carbonyls, which

makes possible the colocalization measurement with more or less accuracy. Conversely, we observed that adult irradiated nematodes and old control worms show negative colocalization between carbonyls and proteins associated with a positive lipid and protein colocalization. A scheme has been included to sum up the results (Scheme 1).



Scheme 1. Illustrations of summarized conclusions are displayed here upon aging and chronic ionizing radiation of sterile *C. elegans*.

3. Discussion

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Understanding aging is of fundamental interest in biology and medicine [17], since aging and aging-related diseases (e.g., cancer and degenerative diseases) cause about 90% of deaths in developed countries. However, questions remain about i) the specificity of aging-related mechanisms, and ii) the influence of organism environment on aging.

A low-dose gamma irradiation can modulate the lifespan of various organisms [39]. With regard to C. elegans, only acute gamma irradiations above 1000 Gy have already been observed to reduce the lifespan [40], but no results have been found after chronic irradiations. We showed herein that the C. elegans sterile glp-1 mutant had a significantly shortened lifespan after all irradiation treatments. Furthermore, different radiation treatments, i.e., different dose, dose rate and duration of exposure, did not result in a significant difference between treated groups. It is possible that our experimental conditions resulted in an increased protective mechanism with a threshold reached by the first delivered dose to organisms, independently of the duration and dose rate, and then led to a similar lifespan after irradiation. As some developmental stages are more sensitive to radiations than others [40], shorter radiation exposure but with the same dose rates, over specific developmental stages, e.g., diakinesis, would be interesting to perform in order to test for modulation effects on the lifespan. Our work has been done on a conditional sterile mutant that has prolonged lifespan due to germline mutation. Therefore, noted changes in longevity cannot be simply associated to metabolic trade-offs between reproduction and longevity, but rather are instead consequences of distinct, pro- and antilongevity signals originating from various regions of the gonad [41]. In addition, irradiation effects on the lifespan can then be more emphasized in this mutant as the period of observation is more extended than for wild worms.

A biochemical study of radiation-induced accelerated aging was performed with emphasis on protein damage monitoring; i.e., protein carbonylation, a molecular modification that has been previously known to be irreversible and impacted by aging and ionizing radiation [42, 43]. In fact, carbonylation turned out to be a good biomarker of aging, since old nematodes display higher carbonylation than young ones. These results are in accordance with previous reports on aging [25, 44]. Moreover, a decrease in protein turnover has been identified as one of the biomarkers of aging with no relation to protein abundance over time [24]. Such a phenomenon perhaps explains the higher susceptibility to carbonylation and aggregation of proteins in old worms. Otherwise, radiation treatment gave a rather unexpected result regarding carbonylation level detection compared to its control counterpart; i.e., radiation reduces carbonylation regardless of the dose or duration of the irradiation. The decrease in carbonylation could be partially explained by chronic irradiation induction of chymotrypsin-like proteasomal activity [8], whereas proteasome activity of germline ablated *glp-1* worm mutants has already highly increased caspase, trypsin, and chymotrypsin-like proteasome activities in comparison

to wild worms [45]. In addition, 26S proteasome regulators, like rpn-6.1, that reduce proteotoxic stress have already been reported to be upregulated following a constant mild-heat stress, such as 25°C [45]. Our experiments also involved a temperature of 25°C under all our working conditions, including the control ones. In *wt* nematodes, the protein carbonylation level has been found to be quite stable after chronic irradiation, with an increasing proteasomal activity [8]. As some hypotheses stated that carbonylation partially takes place in mitochondria [46], the difference between both models could be due to the mitochondria number, which is higher in N2 than in *Glp-1* mutants (without germ cells).

With regard to lipids, their level in glp-1 is higher than in N2 wt nematodes, which is in agreement with a previous report [14]. Moreover, our experimentation showed that radiation also induced neutral lipid catabolism whatever the dose or the dose rate. Our results indicate that lipid catabolism induced by radiation correlates with a reduced longevity, which is finally equivalent to the longevity value of wt nematodes (~25-30 days). Lipids are very important signalling molecules that constitute the cell membrane and form comprehensive energy storage. Lipid homeostasis is in charge of endocrine signalling of longevity. However, regulating mechanisms of lipids in C. elegans are not fully understood and are very complex [47]. It was already reported that germline ablation in worms alters lipid metabolism and affects longevity. With our methodology, the neutral lipids were mainly labelled in this study. In C. elegans, lipids are stored in 2 types of organelles, lipid droplets and yolk. Lipids were observed at the young stage in a small dotted pattern, which is consistent with previous research where LDs appear as bead organelles highly enriched in Tri Acyl Glycerol (TAGs) [48, 49]. One of the assumptions for lipid catabolism after irradiation can be that irradiated worms tend to have a lower bacterial food intake after irradiation, resulting in a reduced uptake of lipids, which is not the case in the normal aging process [24]. Another possibility can be that a higher rate of TAG molecule degradation would increase the release of free-fatty acids and the production of energy [47].

With regard to biomolecular colocalization during radiation and aging, especially between lipids, proteins and carbonyls, explanations are unclear at this time, particularly for the lipid-protein colocalization partially excluding carbonyls. One category of protein that could correspond to this "lipoprotein" description of positive colocalization between lipids and proteins is vitellogenin. Indeed, vitellogenin is the most abundant lipoprotein of the yolk, which is highly accumulated in the aged glp-1 sterile mutant nematodes [50] and overexpressed after gamma chronic irradiation of wt C. elegans (Dubois et al, Submitted 2018). In addition, this protein is not very prone to carbonylation due to its constitutive amino-acids [51]. To confirm this hypothesis, additional experiments need to be performed, such as triplex labelling between lipids, proteins and vitellogenin. This phenomenon could constitute a marker of both irradiation and aging, but requires further investigations to be fully elucidated.

In C. elegans, it is believed that IIS (insulin/IGF1 signalling) and germline originating signals can modify DAF-16 transcription factor activity, which can, in turn, modulate the lifespan expectancy [16]. By recruiting different signalling pathways (DAF-16, SKN-1/Nrf, oleic acid, lipase, etc.), germline ablation can enhance immune defence, proteostasis and stress resistance. Notably, fat metabolism, i.e., fatty acid desaturation and thus fat composition, can be profoundly modified [52]. Indeed, through the mutation of the glp-1 gene, C. elegans overexpress triglyceride lipase Lipl-4 and fatty acid desaturases, which induces autophagy, lipophagy and desaturation into higher but healthier lipid status [53]. Lipids can be beneficial but also detrimental for longevity [14]. After gamma radiation exposure, Glp-1 individuals seem to lose their properties; it seems that the decrease of lipids is associated to a reduction of the lifespan. To fight against chronic irradiation, N2 individuals stimulate the 20S proteasome complex [8] and conversely seem to maintain their lipid content at the same level for the reproductive function, despite an overexpression of lipid transporters (Dubois et al, Submitted 2018). However, a decrease occurs at 6.8 Gy. Two different compensatory mechanisms could occur during gamma ray exposure: one increasing the level of lipids (with an increase of the lipid transporter expression), and one decreasing the level of lipids. Recently, a nuclear protein, SKN-1, was reported to drive two opposite responses in C. elegans, thus promoting longevity by a mechanism other than that set for protection against oxidative damage [54]. In addition, SKN-1 plays a direct role in maintaining lipid homeostasis [14]. Irradiation yields oxidative stress, e.g., protein oxidation and lipid peroxidation, and then triggers protection against oxidative damage. SKN-1, in this context, could then be implied in the response to radiation and trigger lipid catabolism. However, lipid catabolism can also be induced by several other pathways and the question of the link between lipids, reproduction and lifespan is far from elucidated [41, 47].

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To conclude, this study contributes to assert that fat catabolism is not necessarily beneficial, as we showed that the lifespan reduction after chronic gamma exposure is associated with a lipid level decrease. In addition, and by the same way, a decrease in carbonyl content cannot be considered as a reliable marker of health because it appears to be linked with a lifespan decrease after chronic gamma irradiation. A proteolysis increase induced by a commitment in fighting oxidative stress induced by irradiation can probably yield energy consumptions that are of negative impact for organism health. Finally, whereas some phenomena, i.e., colocalization of lipids and proteins in some developmental stages, seem shared by aging and irradiation processes, other mechanisms, such as lipid metabolism and oxidative protein level, seem different as organisms react in opposite ways. These results enable to conclude that aging and irradiation processes undergo different mechanisms under our conditions. Numerous cascades of events can yield this result. Some specific tests on SKN-1 and downstream proteins would be necessary to address this question and give evidence of the specific mechanisms that take place during irradiation exposure and aging.

4. Materials and methods

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4.1. Strain and maintenance

- 357 The *glp-1* sterile *C. elegans* strain (Bristol N2) used in this study was provided by the Caenorhabditis
- 358 Genetics Center (MN, USA). Nematode growth medium (NGM) and M9 medium were prepared by
- standard protocols [55]. The nematodes were cultured at 15 °C, 70 % relative humidity in darkness, on
- NGM agar seeded with Escherichia coli OP50 strain, according to the standard method [55]. Bacteria
- were prepared following the standard protocol [56]. Seeded plates were exposed to UVC (20 min,
- Bio-LinkCrosslinker; 254 nm, 200 μWm2) to suppress bacterial activity [57].

4.2. Double-synchronized population for ionizing radiation

- 364 Samples for ionizing radiation were prepared in 3 steps. All those steps were developed at 15 °C and
- 365 70 % relative humidity in darkness, which was constantly controlled by HOBO data logger [58]. All
- NGM plates that were used for this experiment were seeded with 10 x concentrated bacteria (OP 50) to
- prevent starvation and metabolic changes. The first step was to transfer 100 gravid worms with a
- 368 worm picker onto 10 NGM large plates (9-cm diameter plates) followed by 5 days incubation at 15 °C
- 369 to stimulate reproduction. The second step represents 1st synchronization on the fifth day, in which
- those plates were rinsed with M9 buffer and the worms were cleansed 3 times in M9 buffer in glass
- tubes with a centrifuge speed of 400 g for 2 minutes. Cleaned samples were subjected to bleaching (20
- 372 % NaOH, 10 % hypochlorite and 70 % H₂0) for 5 minutes in order to release the eggs from the
- worms' uterus. After this, fast centrifuge is necessary to prevent egg degeneration (800 g for 1 minute
- followed by the removal of the maximum amount possible of supernatant). Cleaning of the samples
- was the same as above (3 times). The number of eggs was counted under the binocular in 1 µl of M9.
- From there, the volume that contained 5000 eggs was transferred onto 20 NGM large plates followed
- by 5 days of incubation, as mentioned above. The last and third day involved synchronizing these
- 378 plates. The rinsing and washing were performed as in the second step. Samples were passed through
- 379 the saccharose gradient (3 and 7%), in order to separate only gravid worms whose eggs from the
- 380 uterus should be synchronized over 3 hours (internal communication). Bleaching was performed on
- 381 those gravid worms and the eggs were counted as in the second step. This time, the volume
- corresponding to 3000 eggs was transferred onto 50 6-cm NGM plates because their volume fitted the
- 383 ¹³⁷Cs source. Furthermore, 50 eggs were transferred to 15 6-cm NGM plates to follow the life span.

4.3. Ionizing radiation

- A Monte Carlo simulation of the dose was performed before irradiation to determine by modelling the
- correct position of the petri dish. The same homogeneous ¹³⁷Cs source was used for both irradiations,
- varying the distance only to yield 7 or 52 mGy.h⁻¹. Plates were arranged perpendicularly to the ¹³⁷Cs
- source, with 1 to 3 RPL (Radio photo luminescent) dosimeters placed on the back of the container to

perform a posterior metrology analysis of dosimeters and check the stability of the dose rate during the experiment. Irradiation was conducted at 25 °C and 70 % relative humidity in darkness. IR began from eggs at different dose rates in one incubator, where the temperature and humidity were controlled by data logger [58]. Supplementary data, **Figure S1a** shows the experimental device, as well as a summary of these results **S1b**. Homogeneous data have been obtained for all of the experiments.

There were four irradiation schemes with two different time exposures and two different dose rates: the first irradiation groups were from eggs to L4 stage (during 64 hours) with 7 and 52 mGy.h⁻¹ reaching 0.5 and 3.3 Gy, respectively; the second irradiation groups were from eggs to "old" stage (during 19 days) with the same dose rates reaching 3.3 Gy and 24 Gy, respectively. Due to the limitation of the maximum dose rate (52 mGy.h⁻¹) from the ¹³⁷Cs gamma irradiation source (**Sup data**, **Fig S1a**), there was no possibility of obtaining a cumulative dose of 24 Gy at the young stage (L4, 64 h of irradiation). After each completed treatment, samples were washed 3 times with M9 and quickly frozen in a grinder tube with the minimal volume. Samples were stored at -80 °C until use.

4.4. Experimental design for following biomolecular levels during radiation and aging

Although the study was focused on the young and old stage of worms during radiation and aging, the interest was also to follow their biomolecule levels in between the young and old stage. The biomolecules of interest were: carbonyls, proteins, lipids and nucleic acids. Various effects were monitored every second or third day, when also HOBO data were undertaken. On those days plates were washed 3 times with M9 as mentioned above. After washing the worms, they were transferred onto fresh 6-cm NGM plates with 10 times concentrated bacteria to avoid starvation. The samples for biomolecular labelling were collected from all replicates; i.e., 10 worms from 10 replicates were taken and quickly frozen in 100 µl of M9. Aging, radiation and recovery effects were followed from the young stage to the old stage, whereas dose and dose rate effects were observed within the same stage. The aging effect represents a normal control without any radiation treatment. The radiation effect represents long-term radiation (from eggs to old stage) with either of the chosen dose rates, 7 and 52 mGy.h⁻¹. However, the recovery effect represents short-term radiation, where nematodes were irradiated during the developmental period from eggs to the young stage with both dose rates (7 and 52 mGy.h⁻¹), and after letting them recover up until the old stage. The dose rate effect compares two different dose rates but the same final dose; i.e., 7 and 52 mGy.h⁻¹ at the old stage. Finally, the dose effect represents all of the different doses at the same stage, including control which represents 0 dose.

4.5. Life span

There were approximately 50 worms on a 6-cm plate to follow their life span. Life span was monitored in biological triplicate for every condition. Furthermore, 4 different radiation treatments were performed at least for 64 hours in radiation incubators. All of the effects mentioned above were

- followed: normal aging (control), cumulative radiation (from eggs to old stage with 7 and 52 mGy.h⁻¹)
- and recovery effect (radiation from eggs to the young stage and recovery). The time points for
- following the life span under the binocular were the same as in the previous section; i.e., every second
- or third day during radiation and after every day except on weekends. The important strategy to notice
- here is that, when radiations were finished and plates were transferred to control incubators without
- radiation, they were separated from the control plates by lead and shell to prevent a bystander effect.
- The radiation-induced bystander effect is the phenomenon occurring in non-irradiated cells displaying
- damaged phenotype received from nearby irradiated cells [59].
- 4.6. Multiplex *in situ* labelling
- 4.6.1. Labelling of carbonyls, proteins and lipids
- Chemoporation and labelling were performed in 40 % isopropanol as in the previous protocol [35];
- 434 i.e., carbonyls with 4 μg.ml⁻¹ Cy5 Hz (CyTM5 Mono Hydrazide, GE Healthcare Life Sciences,
- Buckinghamshire, UK); proteins with 0.5 μg.ml⁻¹ Cy3 NHS ester (Cyanine3 NHS ester minimal dye,
- 436 Interchim, Montluçon, France) and lipids with 2 μg.ml⁻¹ Bodipy 505/515 (Bodipy 505/515, Molecular
- 437 Probes ThermoFisher Scientific, Eugene, Oregon, USA).
- 4.6.2. Labelling of carbonyls, proteins and nucleic acids
- The protocol was the same as above, but the last labelling was for nucleic acids with 5X Sybr Green II
- in 70 % ethanol (Sybr® Green II Nucleic Acid Stain, Lonza, Basel, Switzerland) for 15 minutes.
- Sybr green II labels DNA and RNA in situ [35]. There is significantly higher quantity of RNA
- compared to DNA generally, but particularly here because there is less DNA in sterile nematodes
- compared to wild ones, since the gonad is empty (no cells nor embryos). To conclude, most of the
- signal obtained here is from RNA.
- 4.7. Slide preparation, confocal microscopy imaging and fluorescence quantification
- Images were collected with an LSM 780 confocal microscope (Carl Zeiss, France) by using a 10× dry
- N.A. 0.45 and 20× dry (N.A. 0.80) or 63× oil (N.A. 1.4) oil immersion objective, with a pinhole
- setting of 1.31 A.U. and 8 or 12-bit images. Imaging of multiplex labelled samples and fluorescence
- quantification was performed according to the previous protocol [35].
- 450 4.8. Colocalization
- 451 Biomolecular colocalization was measured across the whole body of *C. elegans* with Pearson's
- coefficient, as depicted with a blue square in **Scheme S1**. Namely, we sampled 15 different points
- across the body of C. elegans (Scheme S1, yellow numeric points in a blue square) and tested them on
- different colocalizations: protein-carbonyl, carbonyl-lipid and protein-lipid colocalization. These 15

different points were quite homogeneous in terms of the colocalization parameter and they were averaged in the end. 4.9. Statistical analyses Statistical analysis significance was calculated in GraphPad Prism. Before each analysis, the normality (Shapiro-Wilk test) and homogeneity of data variance (Levene's test) were tested. When these assumptions were met, then parametric tests were used; i.e., one-way ANOVA with post-hoc Tukey test. Otherwise, nonparametric tests were used; i.e., Kruskal-Wallis with Dunn's multiple comparison test. The lifespan statistical analyses were done with XLSTAT software (Addinsoft, New York, NY, USA). Survival data by the Kaplan-Meier method and differences between survivals were calculated using the Log-Rank test with 99% confidence. Three statistical parameters are presented: Log-rank, Wilcoxon and Tarone-Ware.

474 Authors' contribution

- 475 Mira Kuzmic participated in the experimental design, performed data acquisition, analysis and
- interpretation and most of the writing of the article.
- 477 Simon Galas performed statistical analysis of the lifespan, critical analysis, associated figures and
- captions, and contributed to the writing of the article.
- 479 Catherine Lecomte contributed to setting up the global experimental design, and to the breeding and
- 480 maintenance of the nematodes with her expertise in *C. elegans*, as well as to writing the article.
- 481 Cecile Dubois participated in carbonlyation quantification in the Oxy-DIGE method.
- Nicolas Dubourg contributed to the ionizing radiation experiment and the dosimetry calculation.
- Sandrine Frelon contributed to the global experimental design throughout the entire project, and she
- also performed signal integration, data acquisition, analysis and interpretation, and participated in the
- writing of the article.

486 Conflict of interest

- 487 No benefit of any kind will be received from this paper either directly or indirectly by any of the
- 488 authors.

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496 5. References

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