Response to manuscript by Pfaff *et al.*: Evidence against a role of DJ-1 in methylglyoxal detoxification

DOI 10.1074/jbc.L117.797464

Gilbert Richarme

From the Université Paris Descartes-Sorbonne Paris Cité, UMR 8601, 75270 Paris, France

Edited by Gerald W. Hart

The manuscript by Pfaff et al. (1) claims that DJ-1 doesn't prevent glycation in Drosophila and that its deglycase activity is a Tris artifact. I found the following points in Ref. 1 to be troubling. (i) Pfaff et al. didn't provide final Tris concentrations. (ii) The presence of Tris in our DJ-1 preparation is not reported (2). DJ-1, prepared in Tris or phosphate buffer, displays similar activities (3), and its active site mutant C106S is inactive (2). (iii) In the study by Pfaff et al., DJ-1 samples displayed massive protein aggregation and potential protein inactivation. (iv) Pfaff et al. (1) didn't consider that lactate formation implicates a deglycase activity that couldn't result from a Tris effect that would have only displaced the cysteine/methylglyoxal/hemithioacetal equilibrium toward methylglyoxal formation. Moreover, stable lysine/arginine methylglyoxal adducts wouldn't have been affected by Tris (2, 3). (v) They also didn't take into account that the apparent glyoxalase III activity of DJ-1 reflects its deglycase activity and should have been investigated (2-4). (vi) Consequently, their paper implies that glyoxalase III activities reported by several groups would be artifactual (3, 4). (vii) They didn't observe that, in their publication (Fig. 1E), DJ-1- and glyoxalase-deficient cells displayed similar increases in protein glycation levels. (viii) Protein glycation levels increase by a factor of 3 to 10 in deglycase-deficient cells (reviewed in Ref. 3). (ix) Deglycases prevent acrylamide formation by degrading Maillard adducts (5). (x) In addition to protein repair, DJ-1/Park7 and its bacterial homologs perform nucleotide and DNA repair (6). Because Maillard adducts are their substrates, we renamed them "DJ-1 family Maillard deglycases" (3, 6). Ref. 3 provides a methodical rebuttal to the paper by Pfaff *et al* (1).

References

- Pfaff, D. H., Fleming, T., Nawroth, P., and Teleman, A. A. (2017) Evidence against a role for the Parkinsonism-associated protein DJ-1 in methylglyoxal detoxification. *J. Biol. Chem.* **292**, 685–690
- Richarme, G., Mihoub, M., Dairou, J., Bui, L. C., Leger, T., and Lamouri, A. (2015) Parkinsonism-associated protein DJ-1/Park7 is a major protein deglycase that repairs methylglyoxal- and glyoxal-glycated cysteine, arginine, and lysine residues. *J. Biol. Chem.* **290**, 1885–1896
- 3. Richarme, G., and Dairou, J. (2017) Parkinsonism-associated protein DJ-1 is a *bona fide* deglycase. *Biochem. Biophys. Res. Commun.* **483**, 387–391
- Lee, J. Y., Song, J., Kwon, K., Jang, S., Kim, C., Baek, K., Kim, J., and Park, C. (2012) Human DJ-1 and its homologs are novel glyoxalases. *Hum. Mol. Genet.* 21, 3215–3225
- Richarme, G., Marguet, E., Forterre, P., Ishino, S., and Ishino, Y. (2016) DJ-1 family Maillard deglycases prevent acrylamide formation. *Biochem. Biophys. Res. Commun.* 478, 1111–1116
- Richarme, G., Liu, C., Mihoub, M., Abdallah, J., Leger, T., Joly, N., Liebart, J. C., Jurkunas, U., Nadal, M., Bouloc, P., Dairou, J., and Lamouri, A. (2017) Guanine glycation repair by DJ-1/Park7 and its bacterial homologs. *Science*, 10.1126/science.aag1095

The author declares that he has no conflicts of interest with the contents of this article.

¹To whom correspondence should be addressed. E-mail: richarme@ paris7.jussieu.fr.