

PublisherInfo		
PublisherName	:	BioMed Central
PublisherLocation	:	London
PublisherImprintName	:	BioMed Central

Old flies oxidize

ArticleInfo		
ArticleID	:	3851
ArticleDOI	:	10.1186/gb-spotlight-20001205-01
ArticleCitationID	:	spotlight-20001205-01
ArticleSequenceNumber	:	288
ArticleCategory	:	Research news
ArticleFirstPage	:	1
ArticleLastPage	:	2
ArticleHistory	:	RegistrationDate : 2000-12-05 OnlineDate : 2000-12-05
ArticleCopyright	:	BioMed Central Ltd2000
ArticleGrants	:	
ArticleContext	:	130591111

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In the December 5 [Proceedings of the National Academy of Sciences](#), Zou *et al.* find that only some of the processes of [aging](#) in the fly can be explained by increased oxidative stress (*Proc Natl Acad Sci USA* 2000, **97**:13726-13731). Zou *et al.* analyze expression profiles of both aging flies and young flies exposed to the free-radical generator paraquat, using microarrays of approximately 8000 expressed sequence tags (ESTs) that cover 30-40% of the *Drosophila* genome. Of these ESTs, 43 are upregulated with age, and 89 are downregulated. There are reductions in transcripts involved in reproduction, metabolism and protein turnover. Some genes encoding detoxification agents and chaperones are upregulated, while others are downregulated. Many of these changes are similar to those seen in the [mouse](#), although the specific genes involved tend to differ. One third of all age-regulated genes show significant changes in response to oxidative stress. This suggests that free radicals are not the only causal factor in aging.

References

1. *Proceedings of the National Academy of Sciences*, [<http://www.pnas.org/>]
2. Molecular biology of aging.
3. Gene expression profile of aging and its retardation by caloric restriction.