promised development is a relief. However, the future may be more complex than commonly envisioned by positivistic reductionists.

Studies involving complex disease or disease-like test systems may not be reduced to simple integrative or 'translational' (ref. 1) research that merely seeks confirmation of in vitro cell and molecular data/concepts. For example, "translational" research would negate the fact that cells in vitro may exhibit features (physiology, pharmacology, and ultrastructure) distinct from those recorded in vivo3. Clinical and other in vivo studies of new bioactive molecules should include the active search for off-label compound actions, as these often lead to innovative treatments³⁻⁵. Thus, complex biosystems investigators remain a driving force behind the discovery of drugs and essential disease mechanisms³⁻⁵. This development, which is complementary to molecular medicine, is in keeping with the fallacy of the paradigm of genetic determinism for many common diseases.

A reappraisal of 'wholistic' in vivo research as both a validation and a discovery activity will be of tremendous importance to reductionists who now may have to adopt (explore and explain) incorrect or insufficiently researched textbook ideas of physiology and histopathology. Scientists trained in exploratory in vivo approaches will also speed up critical evaluations of in vitro research standards, to reduce the risk that they become 'segregative'. Collaborations giving equal merit to in vitro and in vivo research will vouch for the lack of delay in exploring the mechanisms involved in original in vivo discoveries (not forgetting iconoclastic observations). Encouraging scientists to study complex biosystems, using ever-improved physiological and histopathological methods as well as adopting the molecular techniques, is urgently needed. Indeed, the failure to understand the importance of exploratory *in vivo* approaches may profoundly slow discovery in medicine³⁻⁵.

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Oxidative DNA damage and embryo development

To the editor-We read with interest the findings of Parman and colleagues regarding the etiology of thalidomide-induced embryopathy¹. The authors suggest that excess reactive oxygen species (ROS) may be involved in the teratogenic process of thalidomide. Their conclusions are based on increased levels of DNA damage in thalidomide-exposed rabbit embryos, concomitant with induction of embryonic developmental damage. Pre-treatment of the mother with the anti-oxidant α -phenyl-N-t-butylnitrone was shown to reduce embryonic dysmorphogenesis. In earlier studies, the authors presented evidence suggesting that excess ROS may be involved in the induction of phenytoin-associated congenital malformations², and others have suggested a role for ROS in ethanol-induced embryopathy3.

Excess ROS may also be important in the maldevelopment seen in the offspring of diabetic mothers. This idea, has been substantiated by in vivo demonstrations of diminished dysmorphogenesis in the offspring of diabetic rodents given dietary antioxidative agents, such as butylated hydroxytoluene⁴, vitamin E (ref. 5), vitamin C (ref. 6) and lipoic acid⁷. Pregnant diabetic mice transgenic for Sod1 show fewer embryo malformations than non-transgenic pregnant diabetic controls8. It has been proposed further that oxidative stress in embryos exposed to a diabetic environment may be the result of increased levels of the isoprostane 8-epi-PGF₂ in the embryos⁹. Also, the fact that high-amplitude mitochondrial swelling in embryonic neuroectoderm of these embryos¹⁰ is diminished by antioxidative treatment of the mother¹¹ suggests the presence of an embryonic ROS imbalance, with conceivable consequences for the rate of apoptosis in susceptible cell lineages in the embryo¹².

Finally, just as in thalidomide teratogenicity, the fetuses and embryos of diabetic rodents have increased rates of DNA damage^{13,14}, suggesting the possibility of a common teratological pathway involving altered expression of genes under the control of transcription factors sensitive to oxidative stress. Investigations of diabetic pregnancies have identified candidate genes, including catalase¹⁵ and cyclooxygenase-2 (ref.9).

These intriguing genetic and biochemical relationships associated with teratological process(es) demand further study. The effect of excessive ROS on embryogenesis may emerge as a more general mechanism in teratogenesis than previously thought.

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