Several clinical trials of bone marrow stem cell therapy for myocardial infarction are ongoing, but the mechanistic basis for any potential therapeutic effect is currently unclear. A growing body of evidence suggests that the potential improvement in cardiac function is largely independent of cardiac muscle regeneration. A study by Fazel et al. in this issue of the JCI provides evidence that bone marrow-derived c-kit+ cells can lead to an improvement in cardiac function in mutant hypomorphic c-kit mice that is independent of transdifferentiation into either cardiac muscle or endothelial cells, but rather is associated with the release of angiogenic cytokines and associated neovascularization in the infarct border zone (see the related article beginning on page 1865). These findings suggest the potential therapeutic effect of specific paracrine pathways for angiogenesis in improving cardiac function in the injured heart.

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Paracrine angiogenic pathways for cardiac repair by bone marrow precursors

In the current issue of the JCI, Fazel and coworkers report on their use of genetically engineered mouse models toward this goal (16) (for brief diagrammatic overview, see Figure 1). Utilizing hypomorph-c-kit mutant mice (compound heterozygote Kit<sup>Wt</sup>/Kit<sup>Wt</sup> mice) that harbor an intrinsic defect in hematopoietic stem cell mobilization, the authors noted an accelerated progression to dilated cardiomyopathy following acute myocardial infarction versus control mice, which is associated with a decrease in neoangiogenesis in the injured myocardium. By reconstituting the bone marrow of irradiat-
ed wild-type mice with GFP-tagged cells, the authors determined that there are only rare c-kit+ cells in the heart at baseline and the vast majority of the cells that are resident in the heart following acute myocardial infarction are bone marrow derived, suggesting that it might be possible to rescue the cardiac defect in the mutant mice via bone marrow reconstitution with wild-type bone marrow cells. The authors go on to document this rescue of the heart phenotype with wild-type bone marrow precursors derived from wild-type mice. (C) Coronary ligation results in an acute myocardial infarction in the mutant mice, with the marked migration of the wild-type GFP-tagged c-kit+ cells into the infarct border zone. Compared with mutant c-kit mice that have not had reconstitution with wild-type bone marrow precursors, there is a markedly smaller extent of chamber dilation and associated dysfunction in the wild-type transplanted mutant c-kit mice. The beneficial effect is not associated with transdifferentiation into cardiac myocytes or vascular cells but rather is associated with the release of angiogenic cytokines and an increase in neovascularization.

**Experimental and clinical implications**

On the experimental front, the study (16) raises several critical issues. First and foremost is whether these studies in a mutant mouse model system have any direct bearing on ongoing human clinical trials. It will become critical to rigorously quantitate changes in specific paracrine factors following cell therapy in small and large animal models of myocardial infarction. Ascribing any therapeutic benefit to an effect on neovascularization may require conditionally knocking down (or out) specific cytokines in the transplanted cell population, where one would lose the presumed therapeutic effect on cardiac function as well as on neangiogenesis. Second, if it is indeed a paracrine effect, an examination of the relative efficacy of the intramyocardial delivery of angiogenic cytokines in experimental model systems of acute myocardial infarction appears warranted. If indeed the effects of cell therapy in the acutely ischemic heart are not related to transdifferentiation, it could become valuable to define which autologous cell type has the optimal profile of angiogenic paracrine factors to drive the desired neovascularization end point. It will also become of interest to determine if there is a distinct functional difference and role for the previously resident c-kit+ cells in the heart (17) versus the influx of c-kit+ cells from the bone marrow. In this regard, further characterization, purification, and functional analyses of specific cell subsets could be informative.

On the clinical front, these studies raise the larger question as to whether the mechanistic basis for cardiovascular cell therapy following acute myocardial infarction might be more closely related to the biology of wound healing than regenerative medicine. The short-term effects of infiltrating or transplanted cell types from bone marrow might be partially based on the triggering of an inflammatory response, with the secondary release of cytokines and nonspecific angiogenic cues. In a similar manner, homing of these cells into the injured myocardium might represent the migration of inflammatory cells across a breakdown in the vascular permeability barrier that accompanies many forms of tissue injury. Interestingly, one of the clear messages from the placebo-controlled trials is that there can be a statistically significant, partial recovery of cardiac function in control groups (5). In the only long-term study to date, the control group eventually reached the same extent of functional recovery seen in the treated groups (2), which might relate to the acceleration of cardiac scar formation by inflammatory pathways. Inflammation is known to be linked to atherogenesis, and recent studies suggest luminal loss in the infarct-related artery following intracoronary delivery of bone marrow precursors (18). Accordingly, careful clinical monitoring for these long-term effects seems warranted.
A parable for regenerative medicine

In this ongoing parable of regenerative medicine, the initial hope was that we were in an advantageous position to take a “shot on goal” by moving quickly to clinical studies. The rationale was noble, clear, and compelling, particularly given the unmet clinical need and the robust results of early scientific studies. However, it now is becoming increasingly clear that we may not have the optimal cell type in hand, let alone a clear understanding of other key variables such as in vivo delivery, efficiency of grafting, and suppression of alternative, unwanted cell phenotypes (e.g., pacemaker cell formation in the midst of cardiac muscle regeneration). Cardiovascular stem cell biology still remains one of the most intriguing fields of scientific inquiry in the cardiovascular field and holds great long-term potential. Perhaps, given our growing understanding of the complexity of cell therapy for heart disease, the time has come to “move the ball down the field” by first trying to understand the mechanistic basis of this potential therapy.

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The leak stops here: platelets as delivery vehicles for coagulation factors

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Gene therapy is an attractive approach for the treatment of hemophilia, as continuous expression of donated clotting factor VIII (FVIII) DNA would ensure clotting factor replacement at constant circulating levels rather than at the peaks and troughs that characterize the current protein infusion therapeutic approach. In this issue of the JCI, Shi et al. describe an interesting variant of a gene transfer approach for hemophilia (see the related article beginning on page 1974). They show that targeted expression of FVIII in megakaryocytes, with storage in the α-granules of platelets, has the advantage of delivering clotting factors directly to the site of an injury, where platelets accumulate in large numbers and undergo activation accompanied by release of granule contents. Earlier clinical experience with gene transfer into hematopoietic cells highlighted the potential safety risks of this approach, but an F8 transgene may represent a lower risk than transgenes for growth factors or their receptors.

Gene transfer for the treatment of genetic disease remains one of the most compelling ideas in modern molecular medicine. In genetic disease, where the therapeutic objective is long-term expression of a specific protein, there are — broadly speaking — two potential strategies: gene transfer into a stem cell via an integrating vector, so that all daughter cells contain the genetic modification, and gene transfer with a nonintegrating vector into long-lived postmitotic cells (such as skeletal or cardiac muscle, hepatocytes, or cells in the central nervous system), so that long-lasting expression is achieved even without vector integration. Both of these strategies have been used to achieve long-term expression of a clotting factor, with phenotypic correction, in large animal models of hemophilia, one by use of a retroviral vector in neonatal (rapidly dividing) hepatocytes (1), and the other by use of an adeno-associated virus vector introduced into mature hepatocytes (2–4) (Figure 1A).

One of the advantages of hemophilia as a model for gene therapy is that tissue-specific expression of the donated gene is not

Nonstandard abbreviations used: FVIII, factor VIII.
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