Diabetic foot ulcers (DFUs), a leading cause of amputations, affect 15% of people with diabetes. A series of multiple mechanisms, including decreased cell and growth factor response, lead to diminished peripheral blood flow and decreased local angiogenesis, all of which can contribute to lack of healing in persons with DFUs. In this issue of the JCI, Gallagher and colleagues demonstrate that in diabetic mice, hyperoxia enhances the mobilization of circulating endothelial progenitor cells (EPCs) from the bone marrow to the peripheral circulation (see the related article beginning on page 1249). Local injection of the chemokine stromal cell–derived factor–1α then recruits these EPCs to the cutaneous wound site, resulting in accelerated wound healing. Thus, Gallagher et al. have identified novel potential targets for therapeutic intervention in diabetic wound healing.

Over 100 known physiologic factors contribute to wound healing deficiencies in individuals with diabetes (Figure 1). These include decreased or impaired growth factor production (6–8), angiogenic response (8, 9), macrophage function (10), collagen accumulation, epidermal barrier function, quantity of granulation tissue (8), keratinocyte and fibroblast migration and proliferation, number of epidermal nerves (11), bone healing, and balance between the accumulation of ECM components and their remodeling by MMPs (12). Wound healing occurs as a cellular response to injury and involves activation of keratinocytes, fibroblasts, endothelial cells, macrophages, and platelets. Many growth factors and cytokines released by these cell types are needed to coordinate and maintain healing.

Molecular analyses of biopsies from the epidermis of patients have identified pathogenic markers that correlate with delayed wound healing. These include overexpression of c-myc and nuclear localization of β-catenin (13). Coupled with a reduction in and abnormal localization of EGFR and activation of the glucocorticoid pathway, keratinocyte migration is inhibited (13, 14). At the nonhealing edge (callus) of DFUs, keratinocytes show an absence of migration, hyperproliferation, and incomplete differentiation (13, 14). Fibroblasts demonstrate a phenotypic change as well as decreased migration and proliferation.

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In contrast, cells from an adjacent non-ulcerated area display the appearance of a normal phenotype but are still physiologically impaired. However, they are able to respond to administration of additional growth factors or cellular therapy. Microarray analyses of patient biopsies have confirmed these clinical findings by showing that the transcription profiles of epithelial cells from the two locations (callus and adjacent nonulcerated skin) are distinct and recognizable (14).

**Molecular surgery: integrative approach to healing**

Proper debridement, defined as the removal of hyperkeratotic, infected, and nonviable tissue from a wound, is essential as it accelerates wound healing (15, 16) via a multitude of mechanisms (15, 17). It is essential that pathological diagnosis is used to assure elimination of the hyperkeratotic epithelium and evaluate the extent of debridement (Figure 2). Molecular markers (13) may be utilized in the future development of rapid pathology tests for the guidance of effective debridement procedure. DFU debridement should be done in a sequential fashion until no callus or hyperkeratotic tissue is seen in the periphery of the wound and no scar or infection is present in the base of the wound.

**Bone marrow progenitors in diabetic wound healing**

Bone marrow–derived endothelial progenitor cells (EPCs) play a significant role in the process of neovascularization in response to...
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Molecular basis of debridement. A typical foot ulcer in a person with diabetes is shown at top. (i) The nonhealing edge (callus) containing ulcerogenic cells with molecular markers indicative of healing impairment. (ii) Phenotypically normal but physiologically impaired cells, which can be stimulated to heal. Despite a wound’s appearance after debridement, it may not be healing and may need to be evaluated for the presence of molecular markers of inhibition and/or hyperkeratotic tissue (e.g., c-myc and β-catenin). We expect more such molecular markers will be identified in the future. Once a wound is debrided, pathology analyses along with immunohistochemistry should determine whether the extent of debridement was sufficient. If the extent of debridement was not sufficient (lower left diagram), cells positive for c-myc (green) and nuclear β-catenin (purple) will be found, indicating the presence of ulcerogenic cells, which will prevent the wound from healing and indicate that additional debridement is necessary. Lack of healing is also demarcated by a thicker epidermis, thicker cornified layer, and presence of nuclei in the cornified layer. If the debridement was successful (lower right diagram), no staining for c-myc or β-catenin will be found, indicating an absence of ulcerogenic cells and successful debridement. These markers of inhibition are useful, but the most important goal is actual healing as defined by the appearance of new epithelium, decreased area of the wound, and no drainage. This information should be stored electronically in the Wound Electronic Medical Record (WEMR), which provides an objective analysis coupled with pathology and microbiology reports.

Debridement

Monitor outcomes by WEMR

Which wound will heal?

Evaluation by pathology

No healing

Healing

EPC mobilization into the bloodstream occurs through an increase in bone marrow eNOS activation as a result of hyperoxia. The increased eNOS stimulates NO production, which in turn helps to produce EPCs from the bone marrow. Though hyperoxia can increase the levels of circulating EPCs in the bloodstream, the cells are not effectively mobilized to the wound site, creating another roadblock in the path to healing.

The term “homing” relates to the signals that attract and stimulate the cells involved in healing to migrate to sites of injury and aid in repair. EPC recruitment to the wound site depends on ischemia-induced upregulation of stromal cell–derived factor–1α (SDF-1α). Gallagher et al. (18) also report a decrease in SDF-1α expression particularly by epithelial cells and myofibroblasts derived from wounds of streptozocin-induced diabetic mice. The decrease in SDF-1α was found to be responsible for decreased EPC homing. This defect was largely corrected by the simultaneous administration of HBO and SDF-1α at the wound site. These results imply ischemic conditions, as may be the case in diabetic wounds complicated by decreased peripheral blood flow. In this issue of the JCI, Gallagher and colleagues (18) show that EPCs in the bone marrow respond to ischemia by following chemokine gradients, which results in the homing of these cells to sites of hypoxia, where they then participate in the formation of new blood vessels. Bone marrow–derived EPCs are mobilized by eNOS activation in the bone marrow, a process that the authors hypothesized is impaired in diabetics, thus preventing these cells from reaching the wound site in significant numbers. Hyperoxia has been shown to stimulate eNOS activation in some tissues (19). Therefore, to test the effect of hyperoxia on eNOS activation and EPC recruitment, the authors wounded and later exposed diabetic and nondiabetic mice to hyperbaric oxygen therapy (HBO). Results showed that although the total numbers of active EPCs were much lower in diabetic mice than in controls, hyperoxia does indeed spur the mobilization of EPCs from the bone marrow to the bloodstream (Figure 1). EPC mobilization into the bloodstream occurs through an increase in bone marrow eNOS activation as a result of hyperoxia. The increased eNOS stimulates NO production, which in turn helps to produce EPCs from the bone marrow. Though hyperoxia can increase the levels of circulating EPCs in the bloodstream, the cells are not effectively mobilized to the wound site, creating another roadblock in the path to healing.

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that the decreased expression of SDF-1α by epithelial cells and myofibroblasts may be responsible for the lack of EPC homing to the periphery of diabetic wounds. This work is extremely important because it underscores the complexity of regulatory responses in diabetic wounds and explains the inconsistent response to currently approved hyperoxia protocols (i.e., HBO) in patients with diabetes (20).

From bench to bedside
A combination of therapeutic approaches is likely to lead to a successful treatment outcome for diabetic wounds. Although EPCs seem to be ideal candidates for in vivo cell-based therapies for ischemia, Gallagher and colleagues show here that singularly targeting one aspect of EPC function is not completely efficient in correcting diabetic vascular complications due to the deficits affecting both the cellular and cytokine-mediated responses to ischemia (18). Therefore, future therapies for ischemic complications will have to be based on correcting multiple deficits simultaneously. Therapeutic interventions, including correcting both EPC activation via HBO and EPC homing via administration of SDF-1α, may significantly accelerate diabetic wound healing by correcting the EPC deficit inherent to diabetic wounds.

Clinically, approved evidence-based protocols based on adequate off-loading (21) coupled with the use of FDA-approved biological therapies that have undergone rigorous controlled randomized trials (5, 22) must be utilized. These FDA-approved treatments are clinically efficacious and include PDGF-BB (23), fibroblasts delivered in an absorbable mesh (24), and fibroblasts and keratinocytes delivered in type 1 collagen (25). If simultaneously combined with current therapies, potential treatments targeting eNOS activation and EPC recruitment might further stimulate healing. Wound progression should be monitored with a Wound Electronic Medical Record. This allows nonhealing wounds (those that have not healed after 2 weeks of treatment) to be objectively measured and treatment to be tailored accordingly.

From the laboratory perspective, there is a compelling need to develop technologies to rapidly convert current discoveries into effective therapy for people with diabetes as well as to provide understanding of the molecular and cellular etiologies of patients with foot ulcers. Identification of the DFU genome and correlation of transcriptional profiles with clinical outcomes will determine specific genes that prevent a wound from healing and should further be utilized to develop therapies to prevent wound progression and promote rapid healing.

The promise of the future
Technologies for various molecular analyses (such as genomics, proteomics, transgenic mice), systems for sustained topical delivery (such as polymers and adenovirus vectors), major advances in tissue engineering (such as human skin engineering, cellular matrices, and bone marrow–derived cell therapy), novel discoveries of disease molecular pathogenesis from studies of patient biopsies and animal models, and developments in molecular targeting (in areas such as antisense oligonucleotides, siRNA, antibodies, and small molecules), coupled with breakthroughs in stem cell research, hold the promise of a bright future. All of these powerful technological tools could potentially be applied to people with diabetic wounds in the near future. One of the major remaining steps is the integration of these resources into a coordinated effort to make the technology developed at the bench available to patients at the bedside. It is these synergistic therapies, starting at the molecular and cellular level, that will help to eliminate amputations in patients with diabetes.

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