ORIGINAL ARTICLE

Macrophage Dynamics in Diabetic Wound Healing

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Abstract Wound healing in diabetes is a complex process, characterised by a chronic inflammation phase. The exact mechanism by which this occurs is not fully understood, and whilst several treatments for healing diabetic wounds exist, very little research has been conducted towards the causes of the extended inflammation phase. We describe a mathematical model which offers a possible explanation for diabetic wound healing in terms of the distribution of macrophage phenotypes being altered in the diabetic patient compared to normal wound repair. As a consequence of this, we put forward a suggestion for treatment based on rectifying the macrophage phenotype imbalance.

Keywords Macrophages · Wound healing · Diabetes

1. Introduction

It has been known for many years that wounds in diabetic patients can take longer to heal than similar wounds in non-diabetics (Mulder et al., 1998). Typically healing takes several months, and many wounds do not heal for 12–18 months or more. The normal wound healing mechanism is obviously disrupted in some way, although despite intensive research a comprehensive understanding of this disruption, or its extent, has not yet been realised. There are, however, pieces of this complex jigsaw which have been identified and by combining these pieces together it becomes possible to present an initial mathematical model of the wound healing process as affected by diabetes mellitus.

The first stage of the wound healing process is the inflammation stage (Robson et al., 2001), and macrophages are among the first cells to arrive at the wound site in response to chemical signals (growth factors) released by platelets (Singer and Clark, 1999). Growth factors stimulate the chemotaxis and mitosis of both endothelial cells and fibroblasts, and are thus vital for the second stage of wound

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repair, the proliferative stage. In diabetic patients, macrophages are known to persist past the inflammatory stage in chronic non-healing wounds, and data from Loots et al. (1998) shows that significant numbers of these cells have been measured on day 28 of healing, long after macrophages are no longer seen in similar wounds in the control (non-diabetic) subjects.

Macrophages themselves are differentiated monocytes, and result from monocytes responding to certain chemical stimuli. There are known to be three types of macrophage important to the wound environment—cytocidal, inflammatory and repair macrophages (Riches, 1996). Monocytes become inflammatory macrophages in the presence of $1,3-\beta$ -glucan, and differentiate into repair macrophages in the presence of hyaluronan. Cytocidal macrophages are formed in response to polyinosinate-polycytidylate and are 'killer' cells that remove bacteria and other debris from the wound site. Since we are not considering the effects of a bacterial load on the wound site or debris removal in the model, this macrophage phenotype will not be discussed further in this paper.

Each type of macrophage produces different growth factors and cytokines. Inflammatory macrophages produce platelet-derived growth factor (PDGF), transforming growth factor- β (TGF- β) and basic fibroblast growth factor (Singer and Clark, 1999; Robson et al., 2001), which attract fibroblasts and endothelial cells to the wound site and encourage them to proliferate. Repair macrophages, which help remodel the extracellular matrix of the wound, produce insulin-like growth factor 1 and also PDGF (Singer and Clark, 1999; Robson et al., 2001).

It is the balance between the inflammatory and repair macrophage populations (Zykova et al., 2000) that appears to be crucial for successful wound healing. Since monocytes become repair macrophages in the presence of hyaluronan, and hyaluronan is produced by fibroblasts, it follows that if the balance between the macrophage populations is disturbed, as suspected in diabetic wounds, then the hypothesis is that there could be an insufficient amount of hyaluronan being produced by fibroblasts, resulting in the repair macrophage population being too low for healing to be completed.

The effects of diabetes on the wound healing process are the impairment of cellular proliferation for most cell types (Sank et al., 1994; Hehenberger et al., 1998; Lerman et al., 2003), increased apoptosis of endothelial cells (Lorenzi et al., 1985; Baumgartner-Parzer et al., 1995; Darby et al., 1997), increased average blood glucose level (Williams and Pickup, 2001), impairment of blood vessel regrowth (Loots et al., 1998; Singer and Clark, 1999), inadequate flow through blood vessels (Singer and Clark, 1999; Greenhalgh, 2003) and decreased collagen deposition at the wound site (Black et al., 2003). Furthermore, it is likely that growth factor expression is altered (Shukla et al., 1998; Wetzler et al., 2000; Robson et al., 2001; Greenhalgh, 2003), and nitric oxide secretion (Bulgrin et al., 1995; Schaeffer et al., 1996; Schaeffer et al., 1997; Zykova et al., 2000) and macrophage removal to the lymph nodes may also be impaired (Bellingan et al., 1996).

Mathematical models of wound repair and healing have thus far been directed towards the proliferation and repair stages of the wound healing process, but it is evident that for diabetic wounds, the inflammatory phase should be modelled in the first instance, as this is when macrophages are most involved. The aim of this paper is to investigate the inflammatory stage of the wound healing process, in both diabetic and normal patients, in terms of the macrophage population behaviour. In the next section, we will derive a basic mathematical model capturing the main features of the inflammatory stage, concentrating on the behaviour of macrophages. In Section 3 we proceed to numerically simulate our model, which leads to a fuller analysis of the steady states in Section 4. This enables us to identify the number of steady-states and their nature, and whether these states can be correlated with the diabetic and normal wound healing states seen in patients. Finally, in Section 5, we discuss our results with reference to available biological data and suggest possible suitable treatment strategies.

2. A basic mathematical model

Very little work has been done in this context from a mathematical viewpoint. Our mathematical model is deliberately simple and concentrates on the involvement of inflammatory and repair macrophages in the inflammatory stage of the wound healing process. The model consists of three variables, considered on a timescale starting approximately 1 day post-wounding. In comparison to other models (Pettet et al., 1996; Cobbold and Sherratt, 2000) we examine the inflammatory stage, and how events at this stage may have a knock-on effect on the latter stages of the wound healing process. By looking only at this stage we are able to ignore the effects of fibroblasts and endothelial cells (Singer and Clark, 1999), and concentrate on the macrophage dynamics. Also, as we are not investigating bacterial cell populations we can ignore the behaviour of cyctocidal (killer) macrophages. In our model, we choose the inflammatory macrophage density $\phi_{\rm I}(t)$ (cells mm⁻³) and the repair macrophage density $\phi_{\rm R}(t)$ (cells mm⁻³) as variables, as they are the key cell populations during the inflammatory stage (Singer and Clark, 1999); t denotes the time in days. Furthermore, we include the TGF- β concentration T(t) (pg mm⁻³) as this growth factor is released by platelets upon wounding, and macrophage precursors (monocytes) migrate to the wound in response to the TGF- β concentration (Riches, 1996; Robson et al., 2001).

2.1. Inflammatory macrophages

In modelling the inflammatory macrophage population, we represent the migration of monocytes to the wound site in response to TGF- β concentration by a function *K*, with α being the proportion of the monocytes that become inflammatory macrophages in response to chemical stimuli at the wound site. Mitotic division of the cells is represented by a logistic growth term, although the macrophage population mainly increases due to migration: there is still a significant amount of mitosis (Miyasaki, 2002). There is also a crowding term representing the effect of the repair macrophage population. Finally, we have a linear decay term representing the removal of the macrophages from the wound site. Note that macrophages do not actually die at the wound site, but travel to the lymph node system, the spleen or the liver (Bellingan et al., 1996). Combining these terms gives the following ordinary differential equation representing the rate of change of the inflammatory macrophage cell density:

$$\frac{d\phi_{\rm I}}{dt} = \alpha K(T) + k_1 k_2 \phi_I (1 - k_3(\phi_{\rm I} + \phi_R)) - d_1 \phi_{\rm I}$$
(1)

In this equation, k_1 represents the proportion of macrophages undergoing mitotic division at any one time and k_2 is the macrophage growth rate. In the model k_1 and k_2 only appear as a product but we include them both explicitly to facilitate comparison with other mathematical models, in which $k_1 = 1$, and so is not included as a separate parameter. In our context, a typical value of k_1 is 0.05 because the number of macrophages undergoing mitotic division is low (Miyasaki, 2002).

2.2. Repair macrophages

We assume that repair macrophages exhibit the same characteristics as inflammatory macrophages regarding cell migration, mitosis and removal; there is very little data available about this. As α is the fraction of monocytes that become inflammatory macrophages then it follows that $(1 - \alpha)$ is the monocyte fraction that becomes repair macrophages, as both macrophage phenotypes originate from the same pool of available cells. Thus, the ordinary differential equation representing the rate of change of the repair macrophage cell density is:

$$\frac{d\phi_{\rm R}}{dt} = (1 - \alpha)K(T) + k_1k_2\phi_{\rm R}(1 - k_3(\phi_{\rm I} + \phi_{\rm R})) - d_1\phi_{\rm R}$$
(2)

2.3. Transforming growth factor- β , TGF- β

TGF- β is only produced by inflammatory macrophages (Singer and Clark, 1999), and again the production of TGF- β by these cells may be considered as constant. The decay rate may also be considered as linear, and therefore the ordinary differential equation representing the rate of change of TGF- β concentration is:

$$\frac{\mathrm{d}T}{\mathrm{d}t} = k_4 \phi_\mathrm{I} - d_2 T \tag{3}$$

3. Numerical solution of equations

Numerical solutions of our model (Eqs. (1)–(3)) were carried out using ode15s, a stiff ODE solver in Matlab. While investigating the model we were able to obtain simulations of both diabetic and normal wound healing. To simulate normal wound healing, we assume that $\alpha = 0.5$ (i.e. that there are equal numbers of inflammatory and repair macrophages formed), and we take $\alpha = 0.8$ in a diabetic wound (i.e. that there are more inflammatory macrophages formed than repair



Fig. 1 A qualitative form of K(T) representing monocyte migration, based on the data available in Wahl et al. (1987). Note that when T = 0, the macrophage migration is small but positive. The curve plotted in the figure is the cubic $K(T) = \tau_1 T^3 + \tau_2 T^2 + \tau_3 T + \tau_4$ with $\tau_1 = -2.47$, $\tau_2 =$ 21.94, $\tau_3 = 6.41$ and $\tau_4 = 1.75$. Note that a cubic is the simplest mathematical form for K(T) giving the appropriate behaviour. In particular, a quadratic form for K(T) does not give the two distinct behaviours shown in Fig. 2.

macrophages). Unfortunately, there is currently no quantitative data on which the value of α for diabetic wounds can be based.

There is also little actual data available to construct a suitable function of K(T) but Wahl et al. (1987) presents data on the migration of monocytes in vivo in response to TGF- β . Figure 1 shows a qualitative form of K(T) constructed using this data. When the TGF- β concentration T is zero, there is a low background level of macrophage migration. As T increases, the migration increases initially, reaching a maximum at about $T = 6.0 \text{ pg mm}^{-3}$, and then decreasing again.

Figure 2 shows typical macrophage and TGF- β profiles obtained for both normal and diabetic wounds by numerical solution of the model (Eqs. (1)–(3)). The numerical results suggest that there are two stable steady states for our system of equations—one that is near zero, and one that is away from zero. In a normal wound the macrophage population decreases to the near zero steady state, but in a diabetic wound the macrophage population continues to increase as the macrophages persist in the wound environment, and the other steady state is reached. This suggests a detailed study of the nature of these steady states to see if these correlate with behaviour observed for normal and diabetic wounds.

4. Steady state and stability analysis

When solving for steady states, it is convenient to keep the form of K(T) general and to express solutions in terms of T. This shows that the steady states satisfy



Fig. 2 Numerical simulation of our model (Eqs. (1)–(3)) illustrating normal and diabetic wound healing. The diabetic wound simulation shows that macrophages persist in the wound environment far longer than they do in the normal wound simulation, and that the level of TGF- β is far higher than in the normal wound. The initial conditions are $\phi_{\rm I}(0) = \phi_{\rm R}(0) = 200$ cells mm⁻³, T(0) = 6 pg mm⁻³. The parameters are $k_1 = 0.05$ (5%), $k_2 = 0.693$ day⁻¹, $k_3 = 0.002$ (cells mm⁻³)⁻¹, $k_4 = 0.07$ pg cells⁻¹ day⁻¹, $d_1 = 0.2$ day⁻¹, $d_2 = 9.1$ day⁻¹. The function K(T) used in the simulation is the same as illustrated in Fig. 1.

either:

(i)

$$\phi_{\rm I} = \frac{d_2 T}{k_4}, \phi_{\rm R} = \frac{(k_1 k_2 - d_1)}{k_1 k_2 k_3} - \frac{d_2 T}{k_4}$$
$$K(T) = 0$$



Fig. 3 Plots of q and r as α and β are varied. The *plain solid line* is a plot of (10) while the line with crosses shows (7). This figure shows that a small imbalance in the inflammatory and repair macrophage populations can result in wound healing being severely disrupted. All other parameter values are as in Figs. 1 and 2.

(ii)

$$\phi_{\rm I} = \frac{d_2 T}{k_4}, \ \phi_{\rm R} = \frac{d_2 T (1 - \alpha)}{k_4 \alpha}$$
$$K(T) = \frac{k_1 k_2 k_3 d_2^2 T^2}{k_4^2 \alpha^2} + \frac{(d_1 - k_1 k_2) d_2 T}{k_4 \alpha}$$
(4)

Solution set (i) implies that there is no migration of monocytes to the wound site, and since we know this not to be true even in diabetic wounds we disregard this solution set and instead concentrate on solution set (ii).

As α increases from 0 to 1, the number of solutions of (4) varies from one to three. Numerical stability analysis (not included) of the steady states shows that when there is only one steady state, it is stable, and when there are three intersections, the sequence is stable-unstable-stable. This is commonly seen in biological problems, and we use a method adapted from the study of spruce budworm population dynamics (Murray, 2002) to investigate the changes in steady state number.

4.1. Bifurcation location

The simplest mathematical form for K(T) having the appropriate qualitative form (see Fig. 1) is a cubic

$$K(T) = \tau_1 T^3 + \tau_2 T^2 + \tau_3 T + \tau_4 \tag{5}$$

with $\tau_1 < 0$ and $\tau_2, \tau_3, \tau_4 > 0$. (Note that a quadratic form for K(T) does not give the required asymmetry and would not allow for the possibility of three solutions to (4)). At the steady states we know that this cubic form of K(T) will satisfy (4), so that

$$\tau_1 T^3 + \tau_2 T^2 + \tau_3 T + \tau_4 = r T^2 + q T$$
(6)

where for our model:

$$r = \frac{k_1 k_2 k_3 d_2^2}{k_4^2 \alpha^2} \qquad q = \frac{(d_1 - k_1 k_2) d_2}{k_4 \alpha} \,. \tag{7}$$

Rearranging Eq. (6) gives:

$$T^{3} + \left(\frac{\tau_{2} - r}{\tau_{1}}\right)T^{2} + \left(\frac{\tau_{3} - q}{\tau_{1}}\right)T + \frac{\tau_{4}}{\tau_{1}} = 0.$$
(8)

At the transition between one and three real roots, Eq. (8) will have a double root. A relation between q and r at this transition can thus be obtained by comparing (8) with the general form of a cubic equation with a double root, namely

$$(T - \beta)^{2}(T - \gamma) \equiv T^{3} - (2\beta + \gamma)T^{2} + (\beta^{2} + 2\gamma\beta)T - \beta^{2}\gamma = 0$$
(9)

where β and γ are steady state values of *T*. If we then use this equation to determine *r* and *q* as functions of β alone we obtain:

$$r = \tau_2 + 2\tau_1\beta - \frac{\tau_4}{\beta^2}; \qquad q = \tau_3 + 2\frac{\tau_4}{\beta} - \tau_1\beta^2.$$
 (10)

Figure 3 shows the relationship between r and q given by (10), treating β as a parameter linking r and q. At the cusp,

$$\frac{\mathrm{d}r}{\mathrm{d}\beta} = \frac{\mathrm{d}q}{\mathrm{d}\beta} = 0$$

which implies that

$$\beta = \sqrt[3]{\frac{-\tau_4}{\tau_1}}.$$

For our parameters, this gives $\beta = 0.89$, i.e. $T = 0.89 \text{ pg mm}^{-3}$. In Fig. 3 we also plot the relationship between r and q implied by (7). Here α has been allowed to vary with the other model parameters fixed. This shows that as α is increased, there is a bifurcation at $\alpha = 0.54$, at which the number of steady states changes from one

to three. Crucially, the value of $\alpha = 0.5$ corresponding to normal wound healing is below this threshold, while the value of $\alpha = 0.8$ corresponding to diabetic healing is above it. Our model predicts that this is the key factor underlying the difference in repair of normal and diabetic wounds.

5. Discussion

Although the model we utilised is a simple one, both the numerical solution and the steady state analysis indicate that this model can provide insights into wound healing in normal and diabetic patients. However, this model has only investigated the behaviour of macrophages and TGF- β and there are many more cell populations and growth factors present during the wound healing process. Therefore, our model cannot claim to give a definitive answer to the problems encountered in the healing of diabetic wounds.

5.1. Macrophage dynamics

Figure 1 shows that the model captures the key qualitative features of wound repair in the two cases of normal and diabetic wound healing; in particular, that the macrophage population decreases to almost zero in normal wound healing, whilst the macrophages persist in the diabetic case. Moreover, our analysis shows the two different wound healing responses seen clinically, and demonstrates that there is a sudden shift between the two healing states, as the balance between inflammatory and repair macrophages changes. The critical value of α for which the system changes from having one to three steady states is 0.54 which occurs when T = 3.55pg mm⁻³. This is less than values typically reported in diabetic wounds, and provides a possible explanation for why diabetic wounds do not heal. There is an excess of TGF- β in the wound environment, which in turn attracts more monocytes to the wound site, leading to a higher level of inflammatory macrophages. These cells produce more TGF- β , perpetuating this vicious circle of events.

5.2. Treatment suggestions

Our model predicts that there is an imbalance between the inflammatory and repair macrophage populations in the diabetic wound environment. Therefore, one course of action to aid healing would be to address this imbalance, and return it to that which is found in normal patients. This requires a reduction of α . In principle, this could be done in one of two ways, either by decreasing the amount of 1,3- β -glucan in the wound environment or by increasing the amount of hyaluronan available.

Figure 4 demonstrates a hypothetical treatment applied to a diabetic wound 10 days after formation. This treatment reduces α to 0.2 and returns the diabetic wound to a healed state, compared to a diabetic wound which was left untreated. Of course, this treatment may not work in practice because the resultant effect on other cell populations, such as fibroblasts, has not been considered in our model, and any alterations at the inflammation stage may well alter the latter stages of wound healing.



Fig. 4 Comparison of a diabetic wound left untreated and one to which a therapy reducing α to 0.2 is applied at Day 10. In the diabetic wound the macrophage population persists, but in the treated wound the macrophage population decreases in accordance with behaviour seen in healing wounds. All other parameters as before.

6. Conclusion

The results of our modelling are consistent with the hypothesis that the balance of macrophage phenotypes is disrupted in diabetic wound healing. Possible extensions of the model could be to include equations for other cell populations, such as fibroblasts and keratinocytes, to enable the modelling of current treatments for diabetic wound management. In conclusion, our work highlights the importance of macrophages in diabetic wound healing, and supports the hypothesis that diabetic wounds appear to be 'stuck' in the inflammation stage and can be 'jump-started' into healing by the application of an appropriate treatment.

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