Biomaterials - Tissue Interactions

Homework #8

New hypothetical mechanism for tissue regeneration by a collagen-GAG scaffold It has recently been observed that TGF β 1 binds extensively on the large surface of a highly porous collagen-GAG scaffold. It will be assumed that bound TGFb1 is unavailable for regulation of cell function and that only unbound, or "free", TGFb1 is involved in cell regulation. You are a researcher exploring the consequences of this new finding with the objective of using this fact in the design of new biomaterials-based approaches to regeneration.

A. The researcher is familiar with data ("contraction-blocking data") showing that collagen-GAG scaffolds block wound contraction most effectively when the average pore size lies between approximately 20 and 140 μ m. See Fig. 1. Having conducted preliminary studies you have found out that the mass of bound TGF β 1, **m**, is directly proportional to the specific surface, **o**, of a series of scaffolds.

$m = C\sigma$

where C is a constant of proportionality. The specific surface, σ , of scaffolds is known to depend on the pore size, **d**, by the following relation (Gibson and Ashby, 1997):

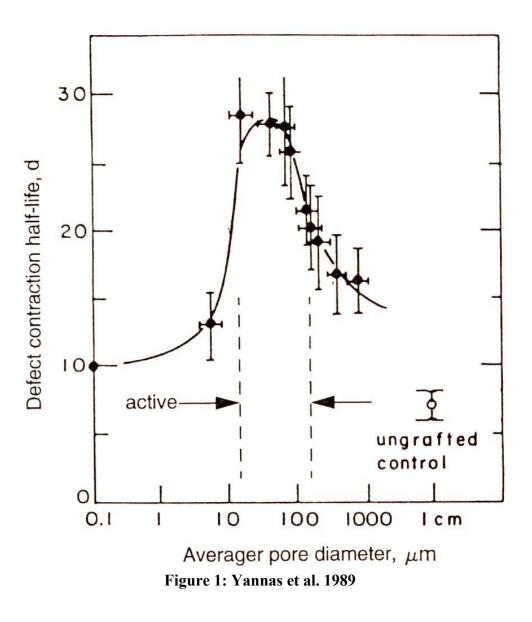
 $\sigma = K/d$

where **K** is a constant of proportionality. Write the relation between the mass of bound TGF β 1, **m**, and the average pore size, **d**, of a series of scaffolds.

B. Collagen-GAG scaffolds degrade during implantation. Set up and solve a linear differential equation with one or more constant coefficients which can be used as a model to describe the decay of the mass of bound TGFb1, **m**, with time of implantation. Assume that the mass of bound TGFb1, **m**, is directly proportional to **M**, the total scaffold mass.

C. Describe briefly the hypothetical role that extensive binding of TGF β 1 on the scaffold surface plays in scaffold- induced organ regeneration. Describe one or a small number of measurements, conducted in vivo, that could be used to test the validity of such a mechanism.

D. Study the "contraction-blocking data" in Fig. 1. Can the entire set of these data be explained by the relation you have derived above between **m** and **d**? Explain briefly.



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