

Lecture 4: Degradable Materials with Biological Recognition (part II)

Last time:	Biological recognition <i>in vivo</i> Engineering biological recognition of biomaterials: adhesion/migration peptides
Today:	Engineering biological recognition of biomaterials: enzymatic recognition and cytokine signaling
Reading:	S.E. Sakiyama-Elbert and J.A. Hubbell, 'Functional Biomaterials: Design of Novel biomaterials,' <i>Annu. Rev. Mater. Sci.</i> 31 , 183-201 (2001) J.C. Schense et al., 'Enzymatic incorporation of bioactive peptides into fibrin matrices enhances neurite extension,' <i>Nat. Biotech.</i> 18 , 415-419 (2000)
Supplementary Reading:	'The Extracellular Matrix,' pp. 1124-1150, <i>Molecular Biology of the Cell</i> , Lodish et al.

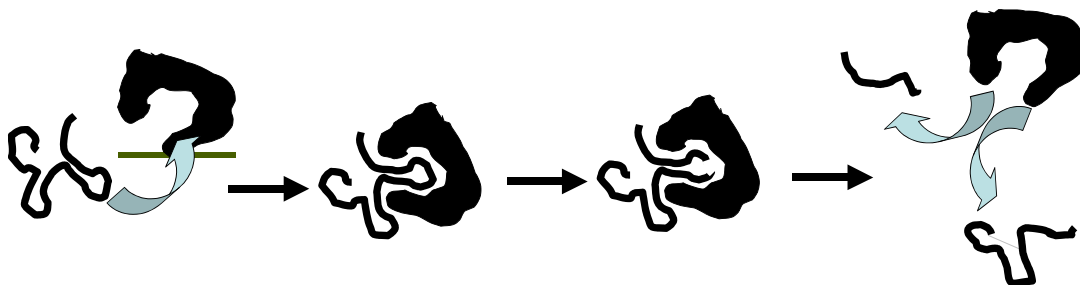
Recognition of Biomaterials by Proteases: Engineering Enzyme-mediated degradation of polymers

Basic concept: include peptide sequences in the polymer chain which are cleaved by enzymatic activity of serum proteins/cellular secreted products (active breakdown) [hydrolysis active but slow...]

- Amide bond $-(NH)-(CO)-$ provides natural hydrolytic mechanism for degradation, but breaks down very slowly in physiological conditions

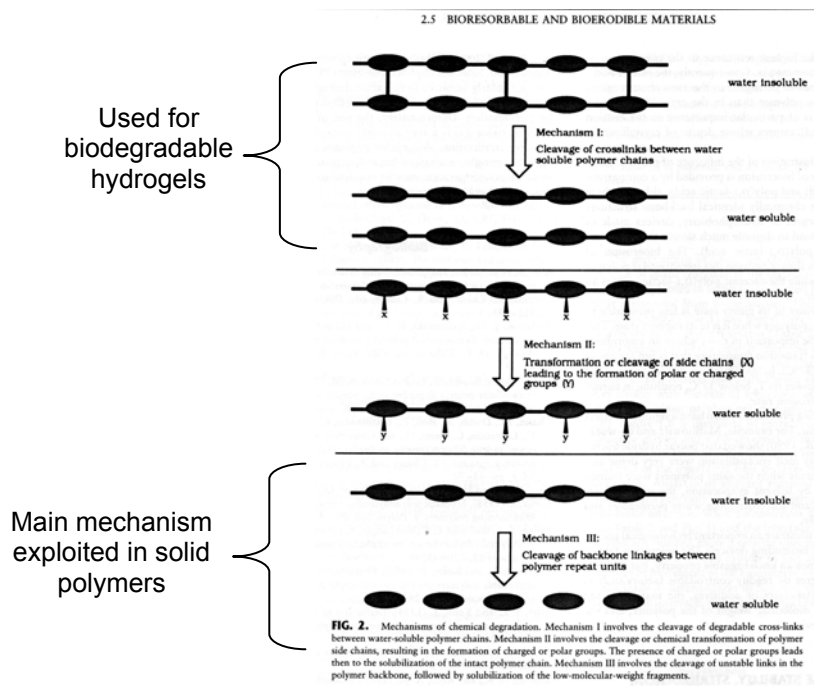
Remodeling enzymes *in vivo*:

- 1) binding of target by enzyme
- 2) specific site cleavage



- What is enzymatic cleavage used for *in vivo*?¹ (Reading Ch.1 4 Voet and Voet "Enzymatic catalysis")
 - Remodling ECM, migration of cells through matrix
 - Removing functional groups from signaling molecules (phosphorylation/dephosphorylation)
- Utility in the design of biomaterials:
 - Enzymatic cleavage can breakdown polymers more quickly than hydrolysis
 - Polymers can be designed to be surface-eroding easily (enzyme can't access interior)
 - Degradation could be localized to tissue where enzyme is produced

Reminder of breakdown mechanisms:



(Ratner, Biomaterials Science)

(SLIDE)

Cleavage of synthetic polymers by enzymes

Cell source	Enzyme	Native function	Acts on	Degradation Mechanism	Result
Various bacteria	lipases	protease	Polyesters, polyesteramides	III	Monomers or dimers
<i>Tritirachium album</i> (mold)	Proteinase K	Protease	Poly(lactide)	III	Monomers or dimers
Mammalian cells	esterases	protease	Poly(alkyl cyanoacrylates)	II	Water-soluble polymers
Mammalian cells	Papain, pepsin	proteases	polyesteramides ²	III	Untested
Mammalian cells	α -chymotrypsin	Serine protease	Aromatic peptides in polyesteramides ³ (e.g. Ala, Val, Leu)	III	Untested
Mammalian cells	elastase	protease	Polyesteramides	III	untested

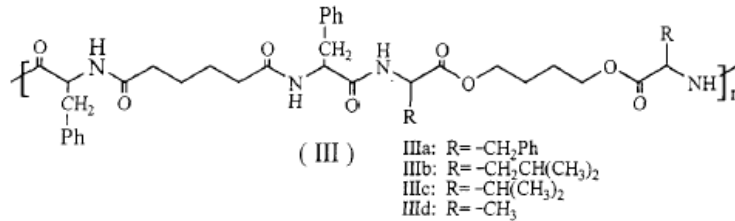
- Pepsin: protease from papaya
- Papain: main protease in gastric juice of stomach
- Chymotrypsin: digestive enzyme

- Mold and bacterial proteases not relevant *in vivo*, but make these polymers also of significant interest for environmentally-friendly packaging
- Of interest in the use of biodegradable materials in environmentally-friendly packaging, but not a concern for *in vivo* applications

- Data comparing *in vitro* and *in vivo* degradation rates indicates that enzymatic cleavage of most synthetic polymers is negligible
- Polyesteramides have been synthesized with enzymatic recognition:

Enzymatic attack on polyesteramides

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- Breakdown by mechanism III
- Enzymatic breakdown of polymers is fast relative to simple hydrolysis:

Polyesteramide breakdown by papain:

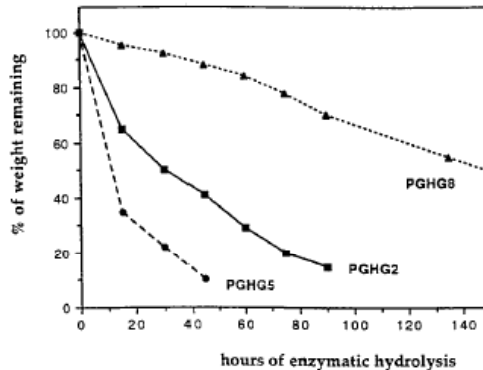
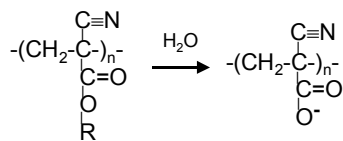


Figure 10. Weight changes of three representative polymers during *in vitro* enzymatic degradation with papain in a 0.05M phosphate buffer. PGHG5 corresponds to the polymer of the studied poly(ester amide) series that has a faster degradation rate.

- What does papain do *in vivo*?

Esterase action on poly(alkyl cyanoacrylates)⁴:



Poly(alkyl cyanoacrylates):

formation of poly(2-cyanoacrylic acid)

- Breakdown by mechanism II

Mechanism:

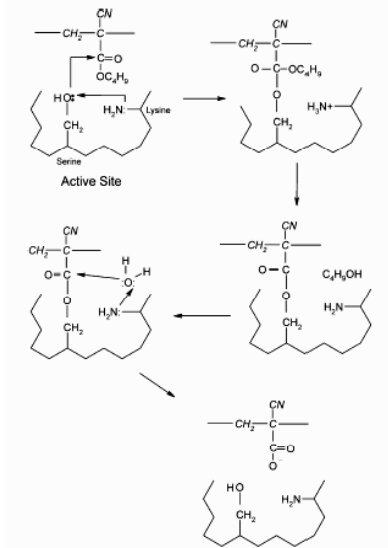


Fig. 11. Enzyme catalysis of esterase involving the ester hydrolysis of poly (butyl cyanoacrylate).

data on degradation of 250nm-diam. porous particles:
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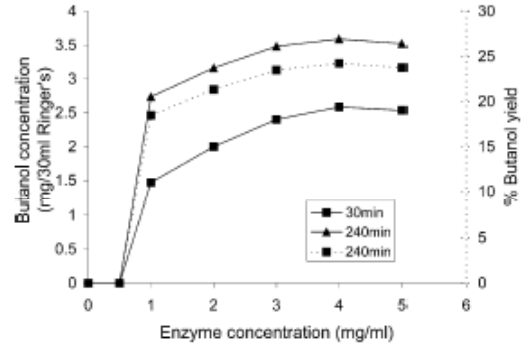


Fig. 2. Butanol production with variation in enzyme concentration (pH 7, 37 °C).

- What does esterase do in vivo?

Enzymatic activity in vivo on peptide sequences: ^{5,6}

Cleavage Enzyme	Functions <i>in vivo</i>	Target amino acid sequences
Plasminogen activator (urokinase or tissue-type plasminogen activator) / plasminogen → plasmin	Degradation of fibrin matrices, angiogenesis, tumor progression; urokinase can bind to cell surface receptor	on fibrinogen: Arg ₁₀₄ -Asp ₁₀₅ , Arg ₁₁₀ -Val ₁₁₁ , Lys ₂₀₆ -Met ₂₀₇ , Arg ₄₂ -Ala ₄₃ , Lys ₁₃₀ -Glu ₁₃₁ , Lys ₈₄ -Ser ₈₅ , Lys ₈₇ -Met ₈₈
Matrix metalloproteinases (soluble and cell-surface): e.g. Fibroblast Collagenase (MMP I)	Facilitate cell migration	Type I collagen: Gly ₇₇₅ -Ile ₇₇₆ In smaller peptides: Gly-Leu or Gly Ile bonds
Elastase	Elastin remodeling	Poly(Ala) sequences

REFS for MMP: (REF J. Biol. Chem. 256, 9511 (1981); J. Biol. Chem. 264, 393 (1989); J. Biol. Chem. 266, 6747 (1991))

Note that proteases often have complementary protease inhibitors:
e.g. plasminogen activators inhibited by plasminogen activator inhibitor type-1 (a serine protease inhibitor)

Examples of peptide sequences used to allow enzymatic cleavage in biomaterials

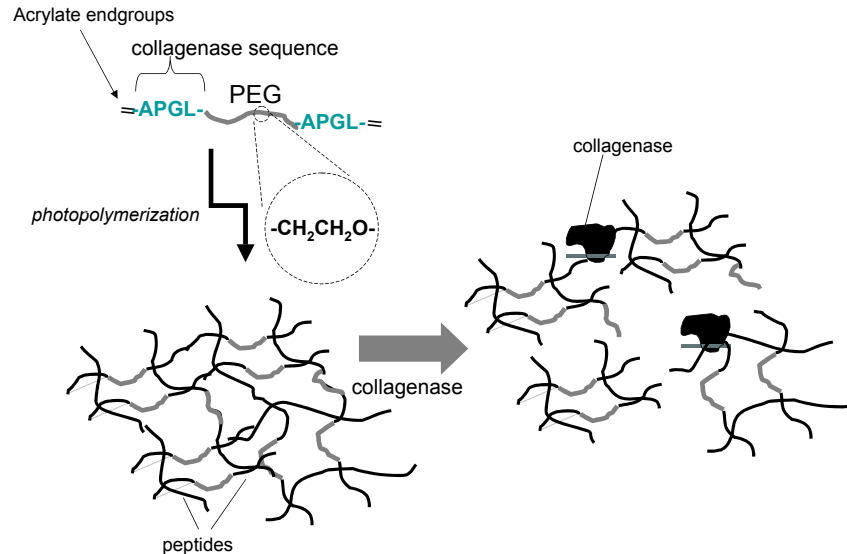
(Work led by J. West and J.A. Hubbell⁷⁻⁹)

recognition by collagenase: -Ala-Pro-Gly-*I*-Leu-

recognition by plasmin: -Val-Ala-*I*-Asn-

recognition by elastase: -Ala-Ala-Ala-Ala-Ala- (polyalanine sequence)

Example: poly(ethylene glycol) networks: acrylate-**APGL**-PEG-**LGPA**-acrylate (**ON BOARD**)



network structure formed by stitching together short strings of acrylate endgroups

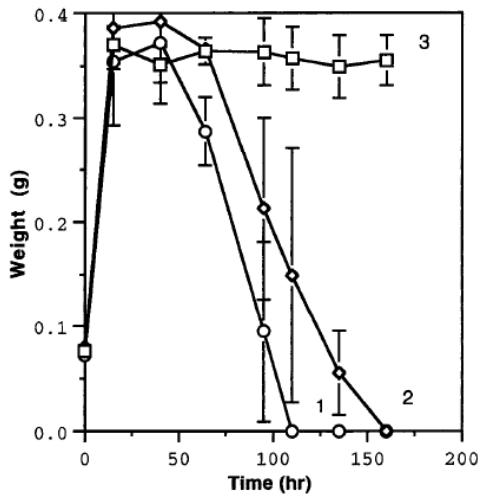


Figure 1. Enzymatic degradation of a hydrogel material formed by photopolymerization of Acr-(APGL)-PEG-(APGL)-Acr in the presence of 2 mg/mL collagenase (curve 1) or 0.2 mg/mL collagenase (curve 2), but not in the presence of 0.2 U/mL plasmin (curve 3). Each point represents the mean of four samples, and the standard error of the mean is shown.

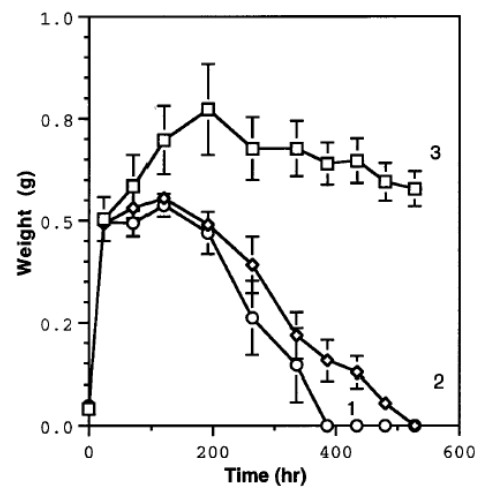


Figure 2. Enzymatic degradation of a hydrogel material formed by photopolymerization of Acr-(VRN)-PEG-(VRN)-Acr in the presence of 2 U/mL plasmin (curve 1) or 0.2 U/mL plasmin (curve 2), but not in the presence of 0.2 mg/mL collagenase (curve 3). Each point represents the mean of four samples, and the standard error of the mean is shown.

- Initially, swelling of network increases and wet weight goes up as first cross-links are broken, then as chains are freed and begin to diffuse out, weight goes down

degradation rate of network depends on collagenase concentration (from Mann et al.):

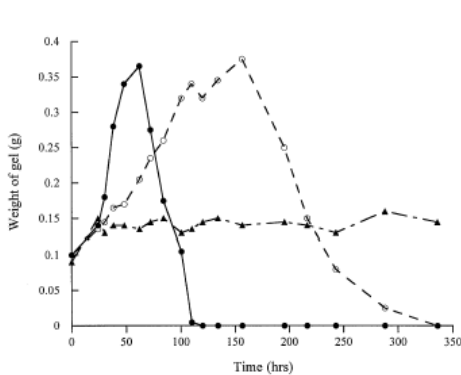


Fig. 1. Degradation of GGLGPAGGK-derivatized PEG hydrogels in solutions containing collagenase. (●) 2 mg/ml collagenase; (○) 0.2 mg/ml collagenase; (▲) no collagenase.

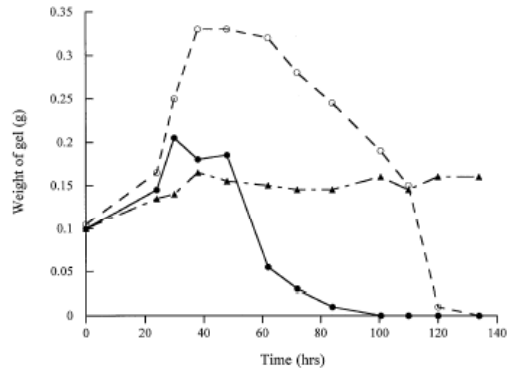


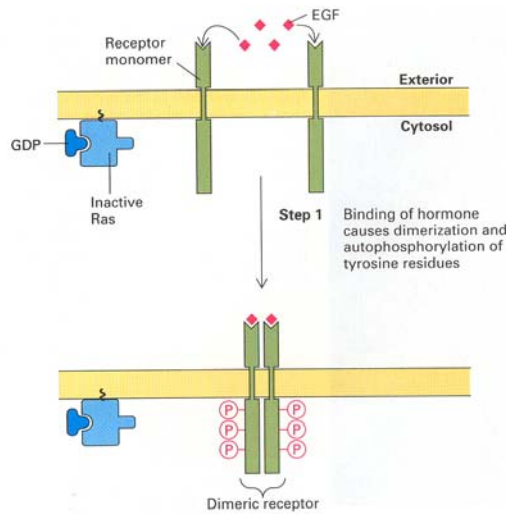
Fig. 2. Degradation of AAAAAAAAAAK-derivatized PEG hydrogels in solutions containing elastase. (●) 2 mg/ml elastase; (○) 0.2 mg/ml elastase; (▲) no elastase.

- note that these are hydrogels which has a major impact on the degradation rate...
- degradation rate is controlled by enzyme concentration and is selective for the enzyme targeted

Any examples tested in vivo?

Recognition of Biomaterials by Cytokine Receptors: Engineering growth and differentiation of cells on biomaterials via cytokine peptides¹⁰⁻¹²

- Growth factors small proteins (ca. 50 aa)
- Can be immobilized to polymer chains present at surfaces of biomaterials much like adhesion peptides and presented to receptors of cells
- Many growth factors signal by DIMERIZATION and autophosphorylation
 - Again, spatial distribution may be key in controlling signaling
 - Cytokines immobilized in close proximity may favor signaling



(Lodish Fig. 20-32)

- INTERNALIZATION: used to down-regulate signals
 - Immobilized cytokines may show more potent signaling due to lack of this down-regulation pathway
- One of the first examples- Griffith lab work with epidermal growth factor immobilized on star polymers:

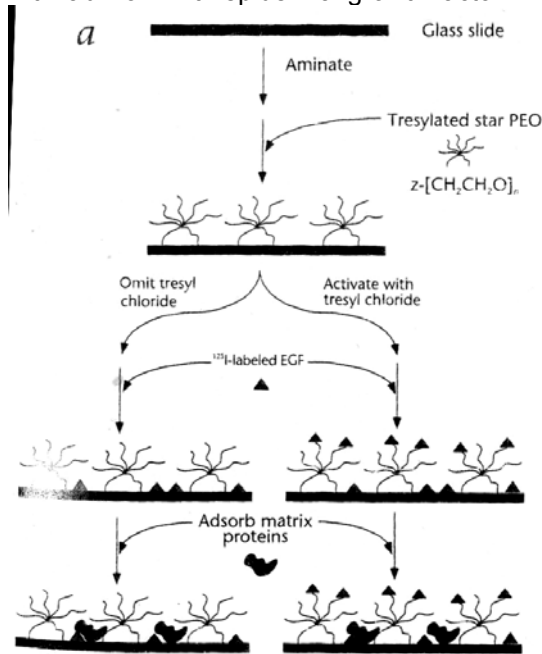
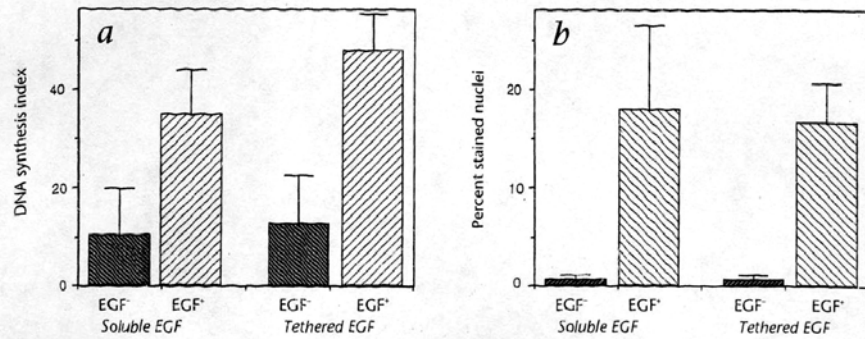


Fig. 2 Tethered EGF produces the same increase in DNA synthesis as soluble EGF for primary rat hepatocytes. *a*, Soluble EGF case contained 10 ng/ml EGF (EGF⁺) or 0 ng/ml EGF (EGF⁻). Tethered EGF was prepared with star PEO (*f* = 180 and *M_n* = 10,000) as shown in Fig. 1*a*; surfaces present either 3.2 ng/cm² covalently tethered EGF and 1.9 ng/cm² nonspecifically adsorbed EGF (EGF⁺) or 0 ng/cm² covalently tethered EGF and 1.9 ng/cm² nonspecifically adsorbed EGF (EGF⁻). Staining index represents the number of nuclei stained per area covered by cells, and represent the mean ± s.d. for at least 100 nuclei. *b*, Soluble case is the same as (*a*). Tethered EGF was prepared by the solution-first method with star PEO (*f* = 70, *M_n* = 5200) as shown in Fig. 1*b*; surfaces present either 0.4 ng/cm² covalently tethered EGF and <0.006 ng/cm² nonspecifically adsorbed EGF (EGF⁺) or 0 ng/cm² covalently tethered EGF and <0.006 ng/cm² nonspecifically adsorbed EGF (EGF⁻). DNA synthesis was measured as a percentage of the total nuclei that had synthesized DNA and each point represents the mean ± s.d. for three separate surfaces with at least 100 nuclei counted per surface.



References

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