



ANKARA ÜNİVERSİTESİ
BIYOTEKNOLOJİ
E N S T I T Ü S Ü

sentetik biyoloji çağında terapötik peptidler

doruk engin

daha özgül hedefleme

daha iyi farmakodinamik ve
farmakokinetik özellikler

kimyasal maddeler ile elde edilmesi
mümkin olmayan etkiler

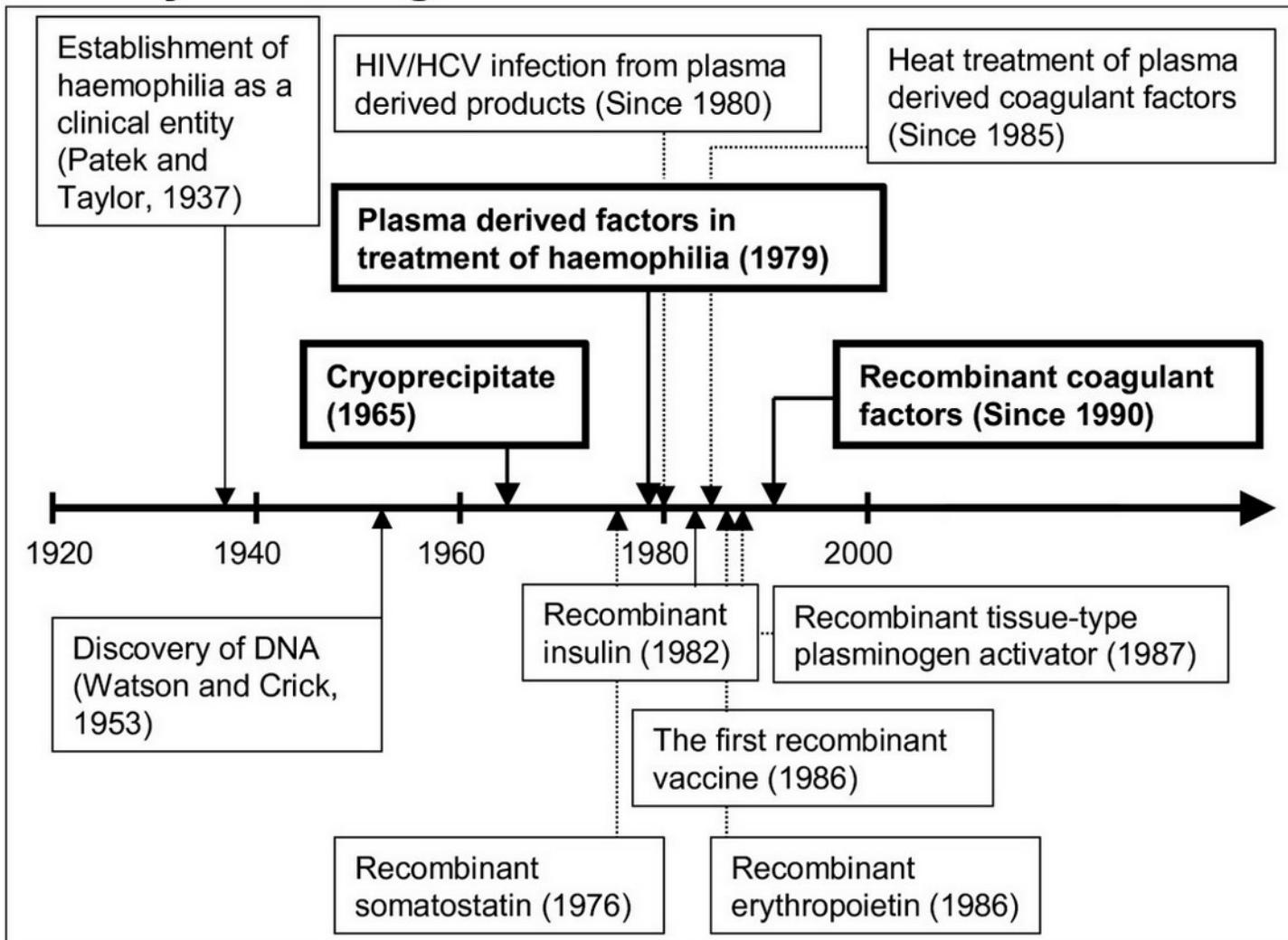
immün yanıt

klinik çalışmalar daha kısa sürede

patent koruması

rekombinant DNA teknolojisi

Fikir yeni değil...

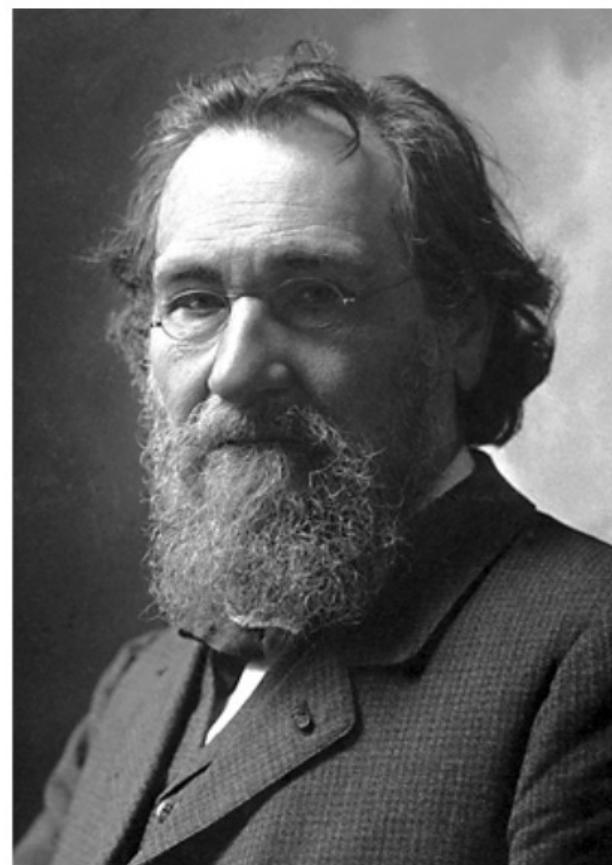


...ancak "hümanize" formların oluşturulması için sentetik biyoloji!



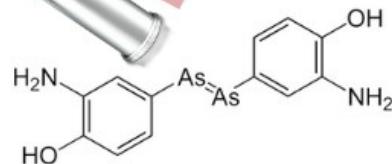
Paul Ehrlich

Paul Ehrlich



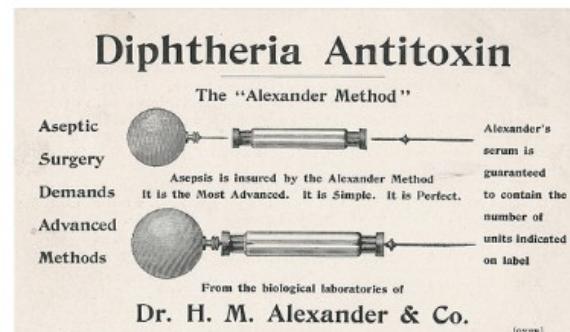
Élie Metchnikoff



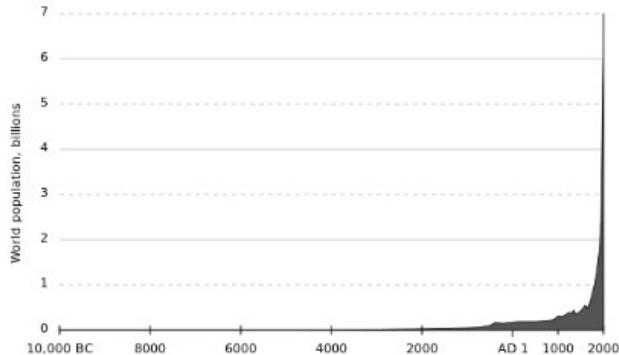


Arsenik bileşikleri mikropları öldürüyor

Antikorlar mikropları öldürüyor



Sihirli mermi - vuracağı hedefi biliyor!



World population milestones in billions (USCB estimates)									
Population	1	2	3	4	5	6	7	8	9
Year	1804	1927	1960	1974	1987	1999	2011	2024	2042
Years elapsed	—	123	33	14	13	12	12	13	18



Antikorlar, mikroorganizmaları her zaman öldürmüyorum!

Üretici - 1



Son kullanıcı Üretici - 1



Üretici - 2

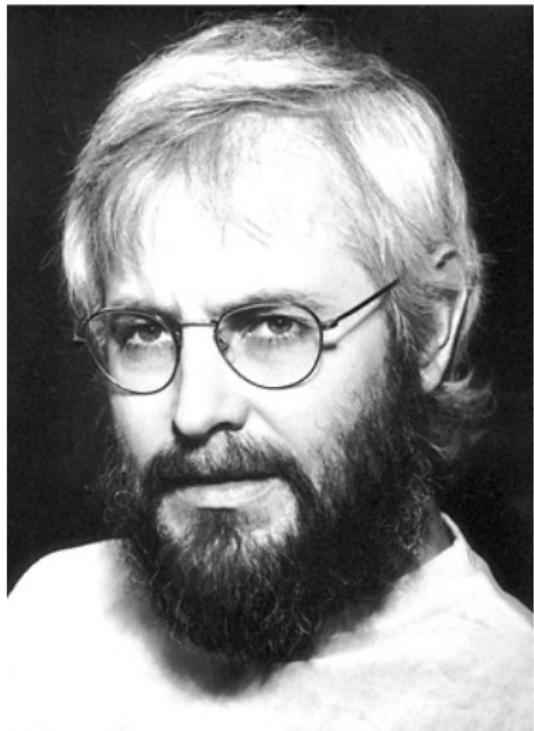


Aracı



Poliklonal antikorlar/antiserum

- tanı
- tedavi



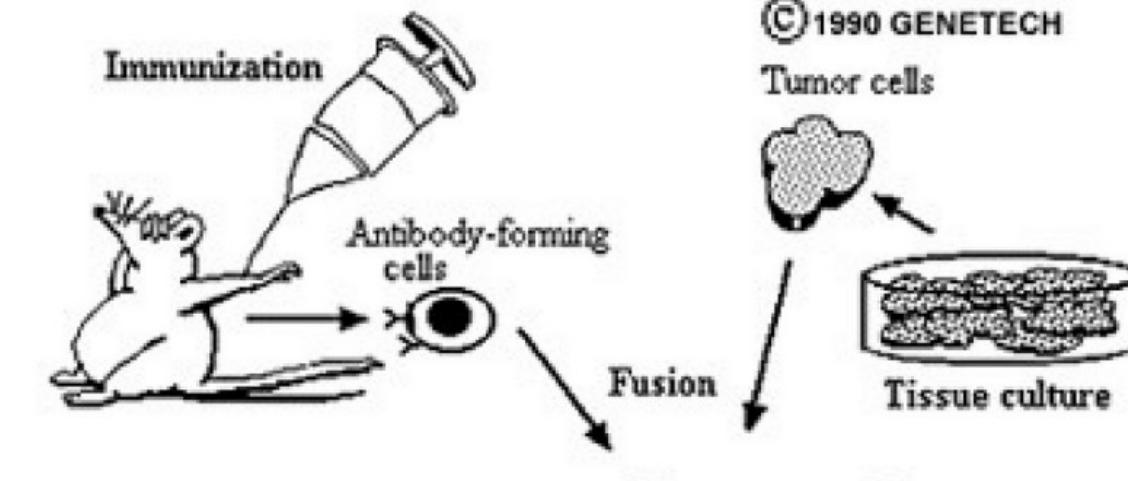
Georges Köhler



César Milstein

'75

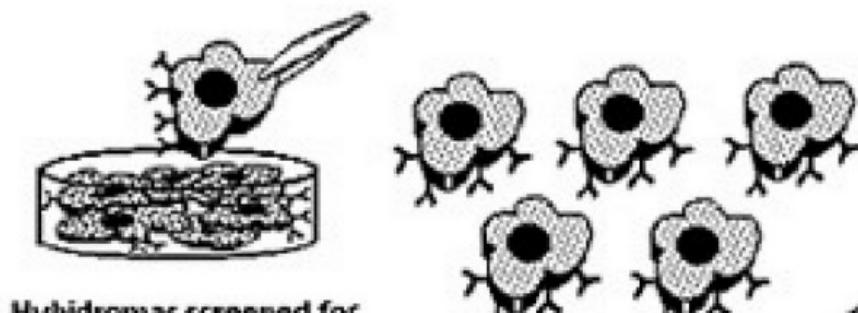
©1990 GENETECH



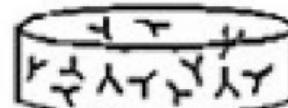
Hybridomas

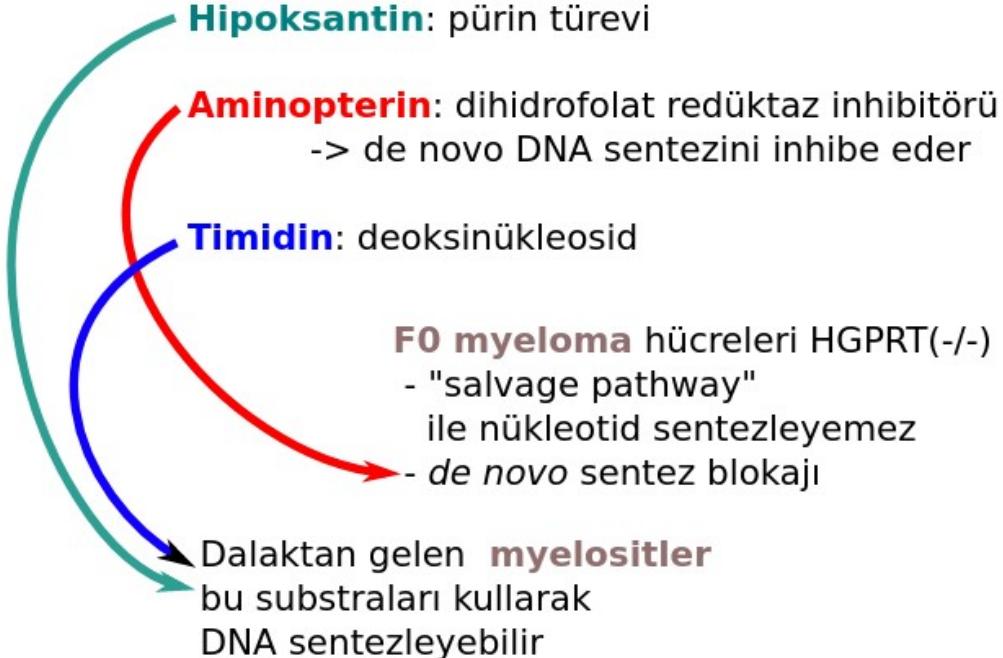
=

hipoksantin-aminopterin-timidin
besiyeri



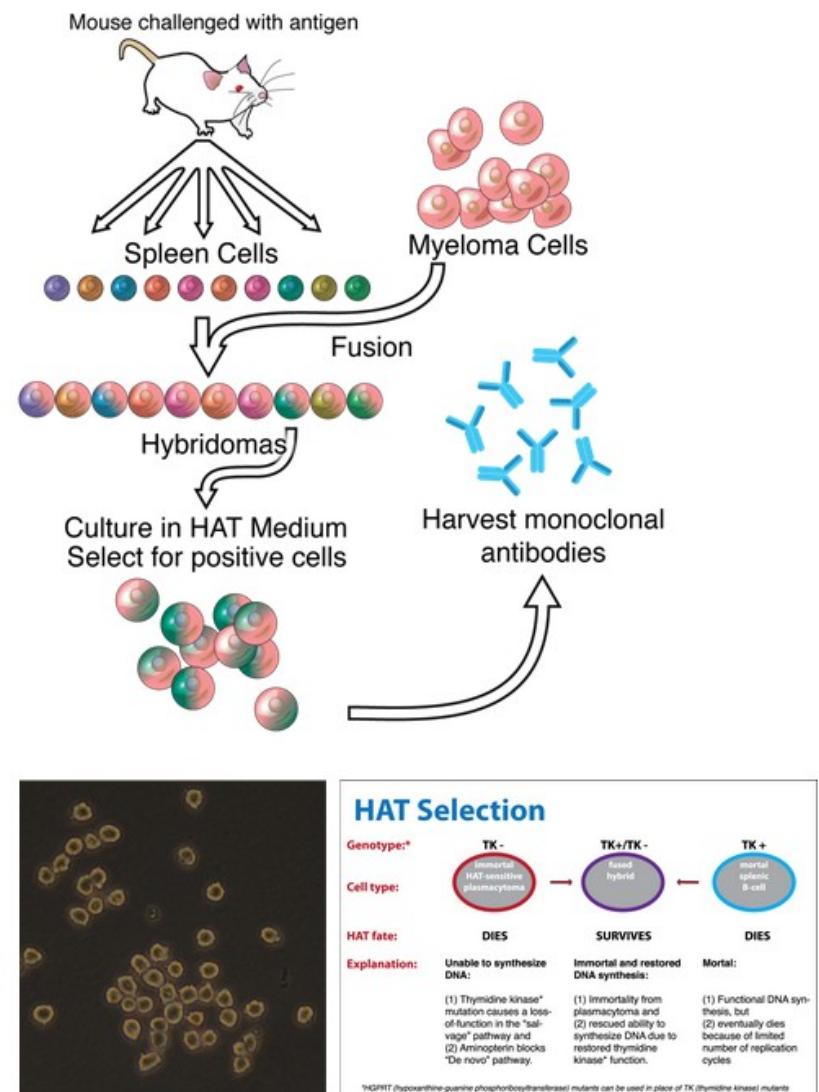
Antibody-producing hybridomas cloned

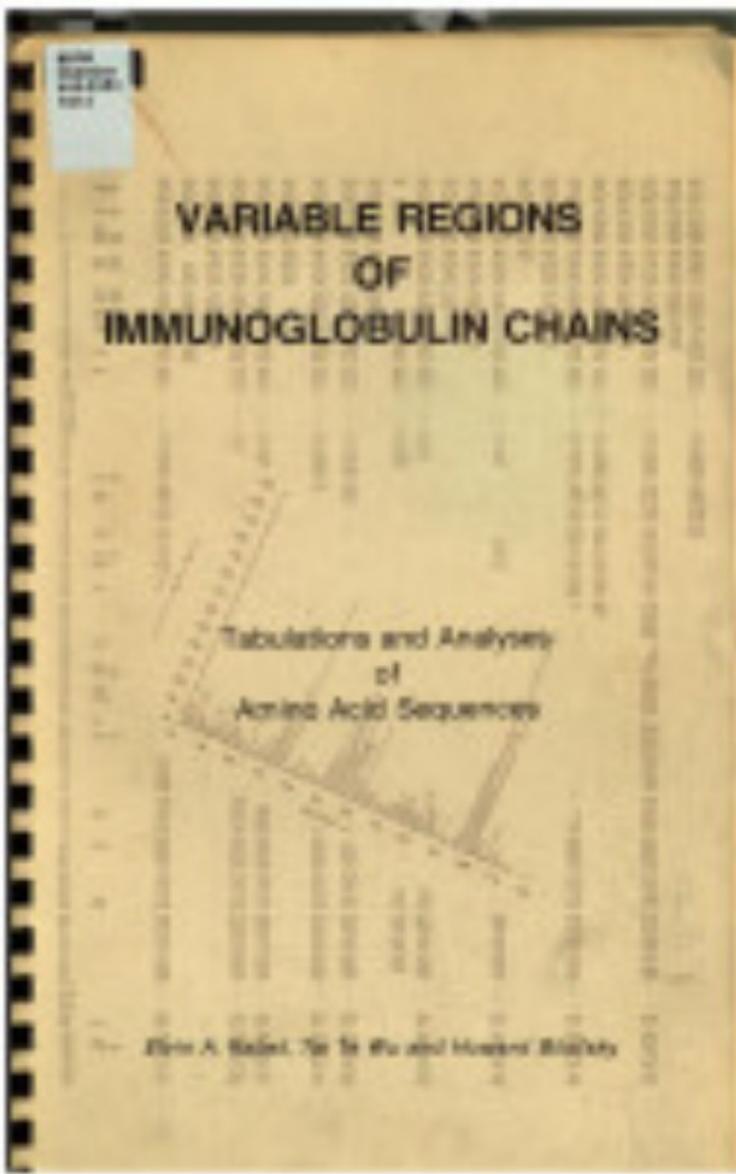




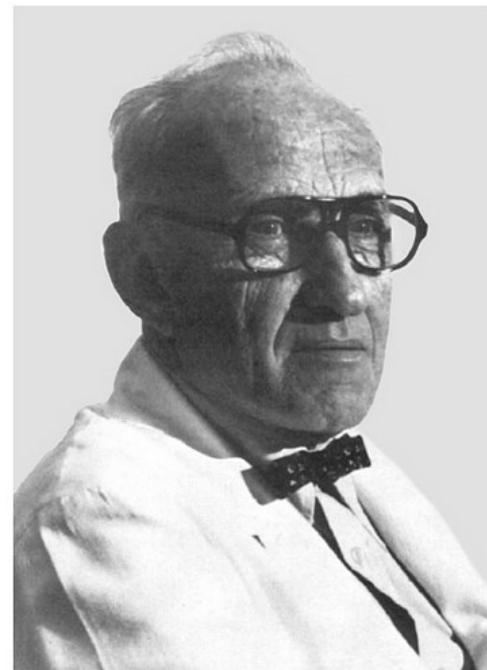
hipoksantin-guanin fosforiboziltransferaz (HGPRT)

HAT besiyeri ile 14 gün seçim...



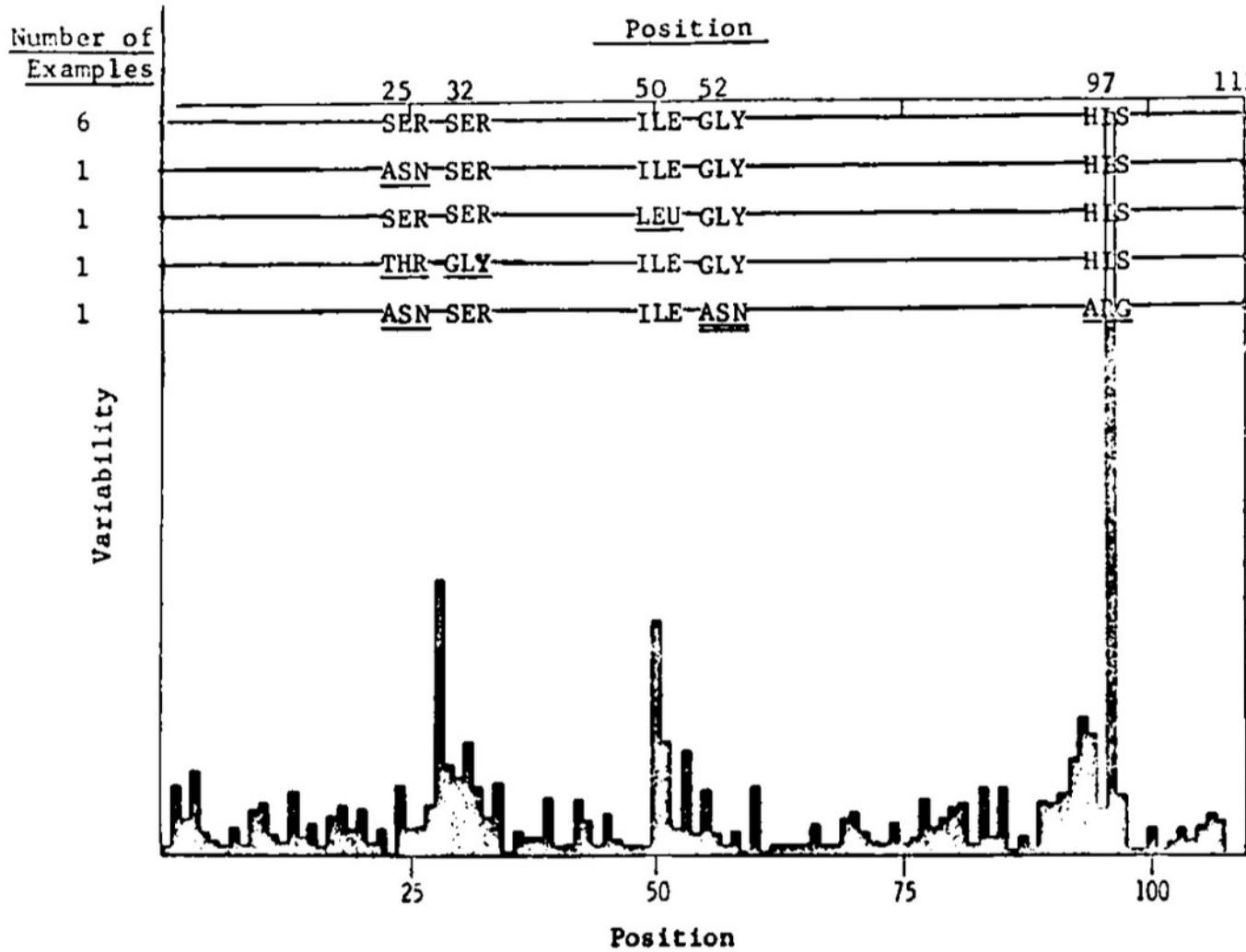


Variable Regions of Immunoglobulin Chains: Tabulations and Analyses of Amino Acid Sequences

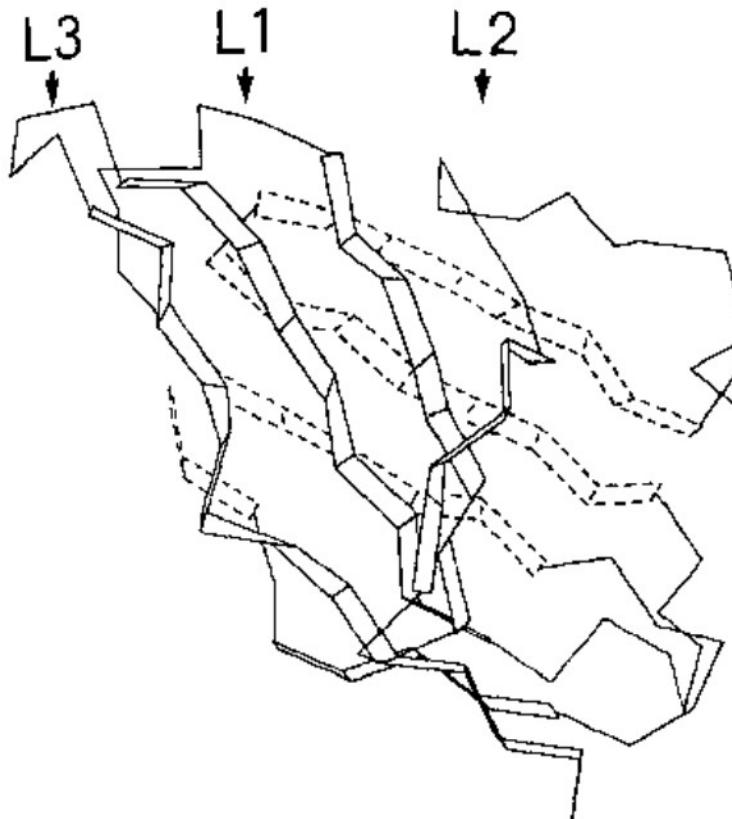
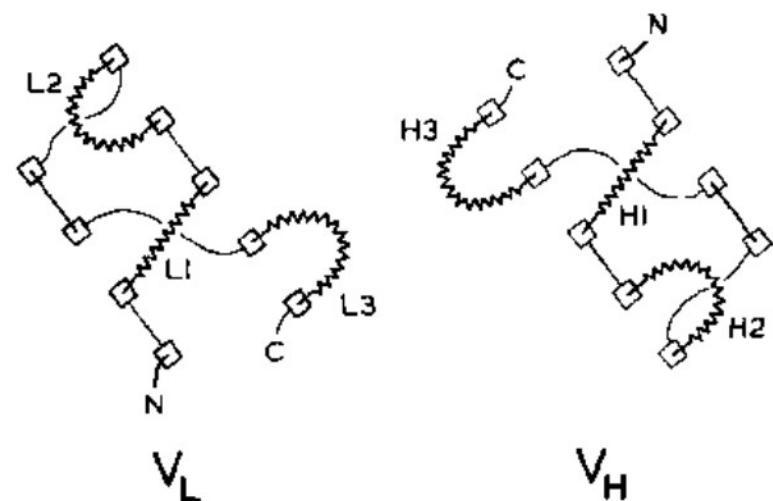
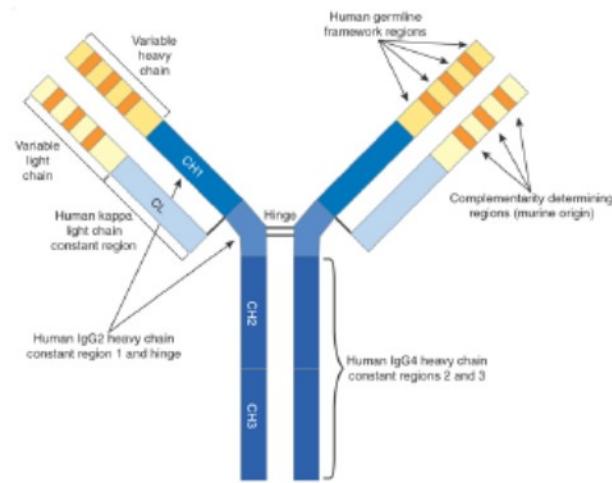


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Elmer A. Kabat

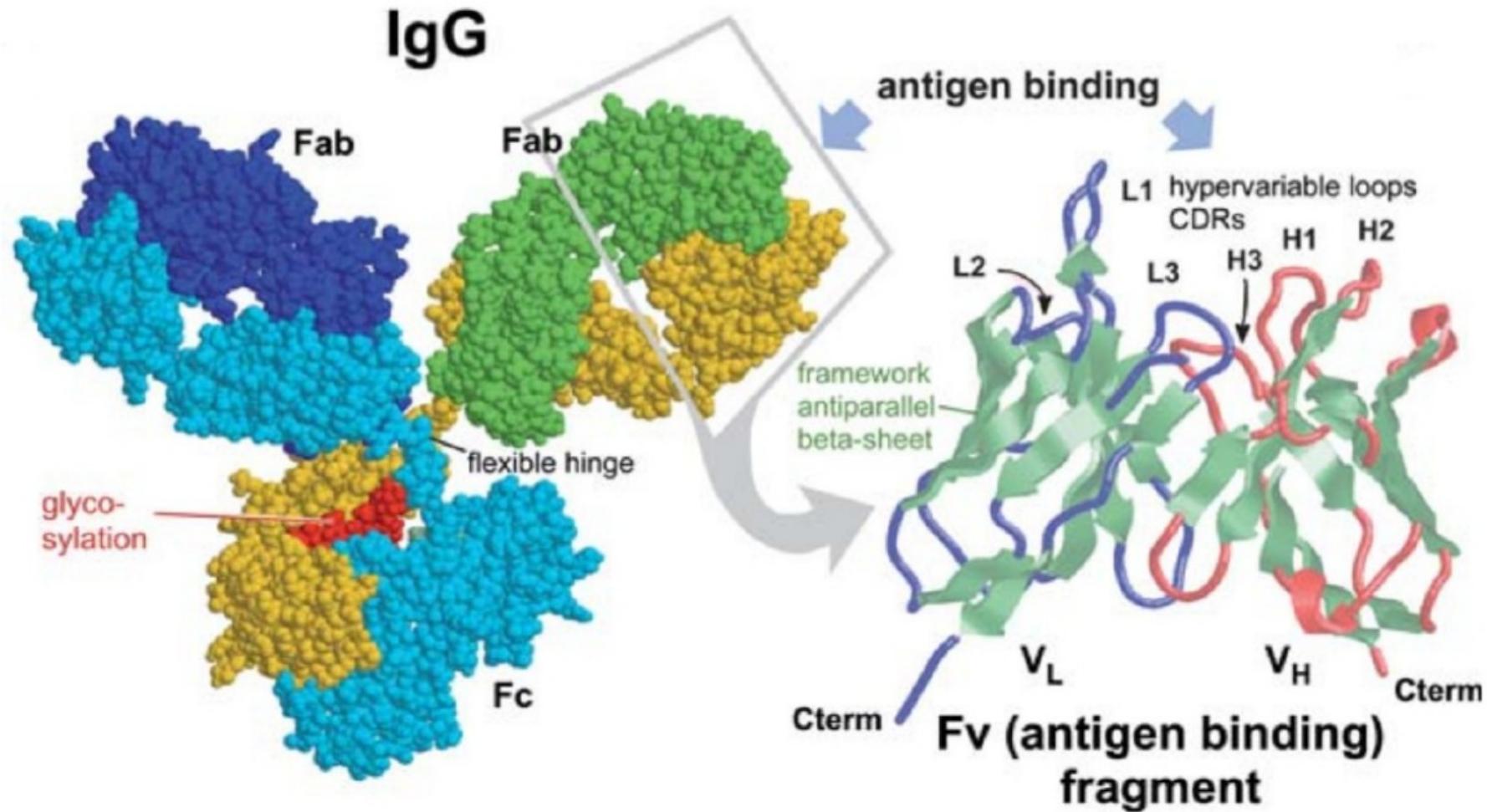


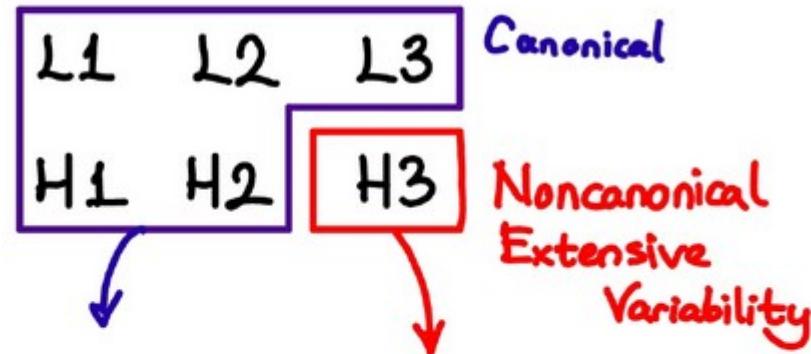
Kabat EA, Wu TT. Attempts to Locate Complementarity-Determining Residues in the Variable Positions of Light and Heavy Chains *. Annals of the New York Academy of Sciences 1971;190(1):382-393.



Chothia C, Lesk AM. Canonical structures for the hypervariable regions of immunoglobulins. J. Mol. Biol. 1987 Aug;196(4):901-917.

Bağlanmanın anatomisi...





Canonical sınıf üyeleri hemen hemen aynı H3 tahmini

omurga konformasyonuna sahip

— loop uzunluğu

— CDR'lardaki anahtar amino
asitler

— Framework'deki anahtar aminoasitler

— dB taraması

— Ab initio modeller

— Conformasyonel arama

— Hepsi :)

CDR-L1
 5 belirgin <
 4 belirgin olmayan >
 1 belirgin olmayan K
 Toplam 10 sınıf

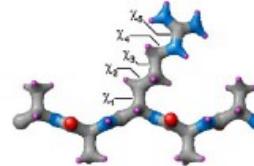
CDR-L2
 Fügülük bir yapıya uyuyor

CDR-L3
 Sınıf 1+2 → 9 üye
 Sınıf 3 → 8 üye
 Benzer hairpin loop'larından oluşuyor
 Farklılıklar Prolinin pozisyonundan
 kaynaklanıyor
 Sınıf 4 → 7 üye
 Sınıf 5 → 8 üye

e:
CDR-H1
 Sınıf 1 → 10 üye
 Sınıf 2 → ?
 Sınıf 3 → 11 üye
 Sınıf 4 → 12 üye

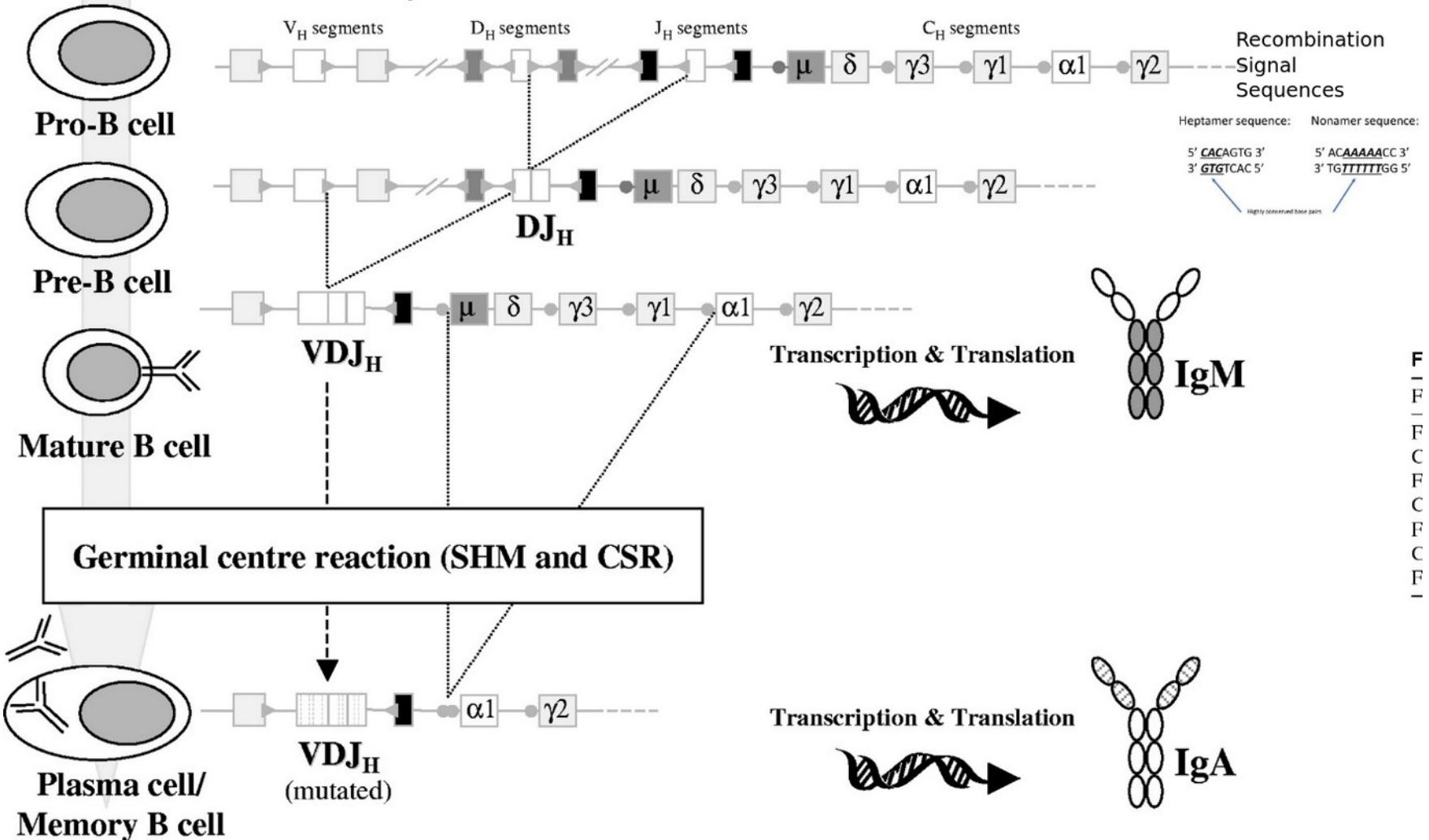
CDR-H2
 Sınıf 1 → 9 üye
 Sınıf 2 → 10 üye
 Sınıf 3 → 10 üye
 Sınıf 4 → 12 üye

CDR-H3
 Canonical sınıflar yerine korunmuş
 yan zincirler bulunuyor

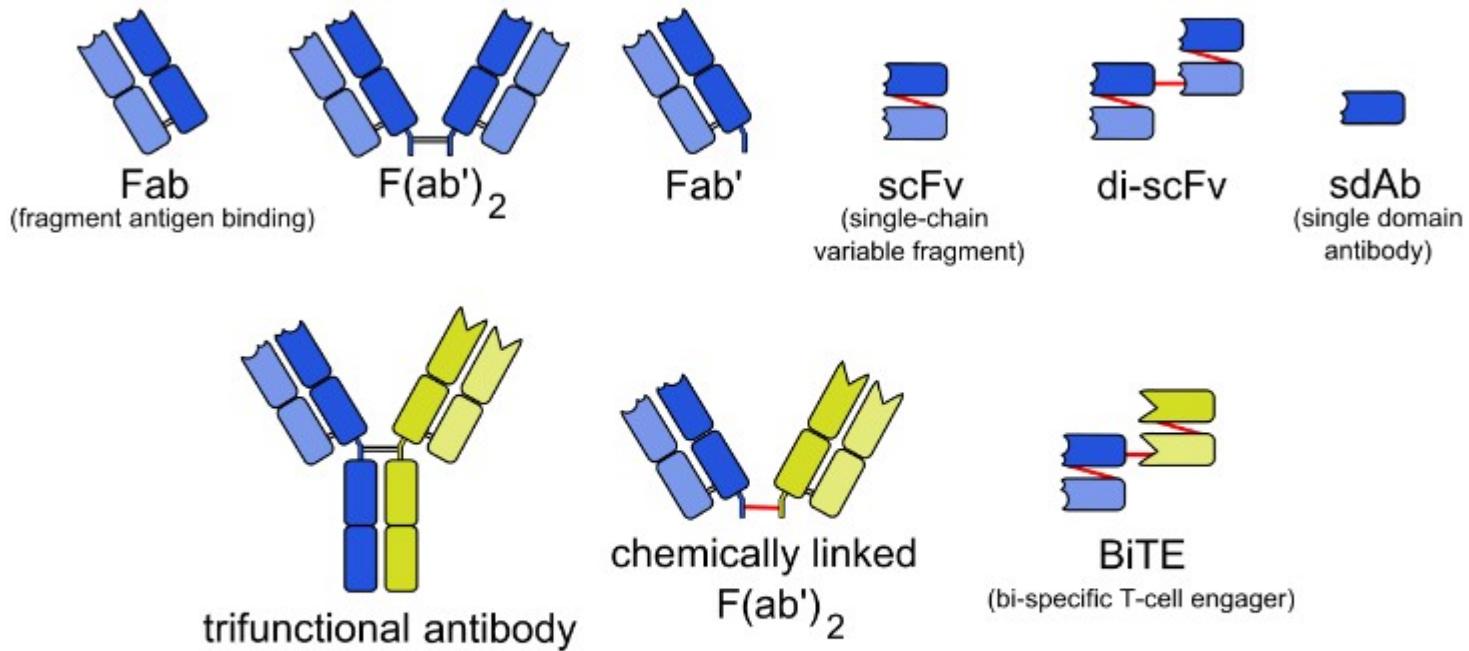


FRs and CDRs of Antibody and TCR Variable Regions

FR or CDR	V _L	V _H	V _α	V _β	V _γ	V _δ
FR1	1–23	1–22	1–22	1–23	1–21	1–22
CDR1	24–34	31–35B	23–33	24–33	22–34	23–34A
FR2	35–49	36–49	34–47	34–49	35–49	35–49
CDR2	50–56	50–65	48–56	50–56	50–59	50–57
FR3	57–88	66–91	57–92	57–94	60–95	58–89
CDR3	89–97	95–102	93–105	95–107	96–107	90–105
FR4	98–107	103–113	106–116	108–116A	108–116C	106–116



Immunoglobulin gene rearrangements and the pathogenesis of multiple myeloma
 David González, Mirjam van der Burg, Ramón García-Sanz, James A. Fenton, Anton W. Langerak, Marcos González, Jacques J. M. van Dongen, Jesus F. San Miguel and Gareth J. Morgan. Blood 2007 110:3112-3121



daha küçük... daha kompakt... hücre içi hedefler?
 çoklu hedefleme...

Daha yaratıcı kombinasyonlar...

