

Background I

Celiac Disease (CD) is an intestinal disorder with multifactorial etiology. This chronic inflammatory disease is triggered by the ingestion of wheat **gluten** or related proteins from rye and barely in genetically-predisposed individuals. The disease is associated with specific **HLA alleles** indicating that HLA genes contribute to CD genetic susceptibility. Given the indisputable role of gluten in causing inflammation and immune-mediated tissue damage, CD represents a unique model of autoimmunity in which, in contrast to other autoimmune diseases, a close association with HLA genes (**DQ2** and/or **DQ8**), a highly specific humoral autoimmune response (autoantibodies to tissue transglutaminase), and, most importantly, the triggering environmental factor (gluten), are known. The disease shows a strong human HLA association predominantly to HLA-DQ2 (**A1*0501, B1*0201**) and/or DQ8 (**B1*0302**) heterodimer (Fig. 1).

Linkage disequilibrium

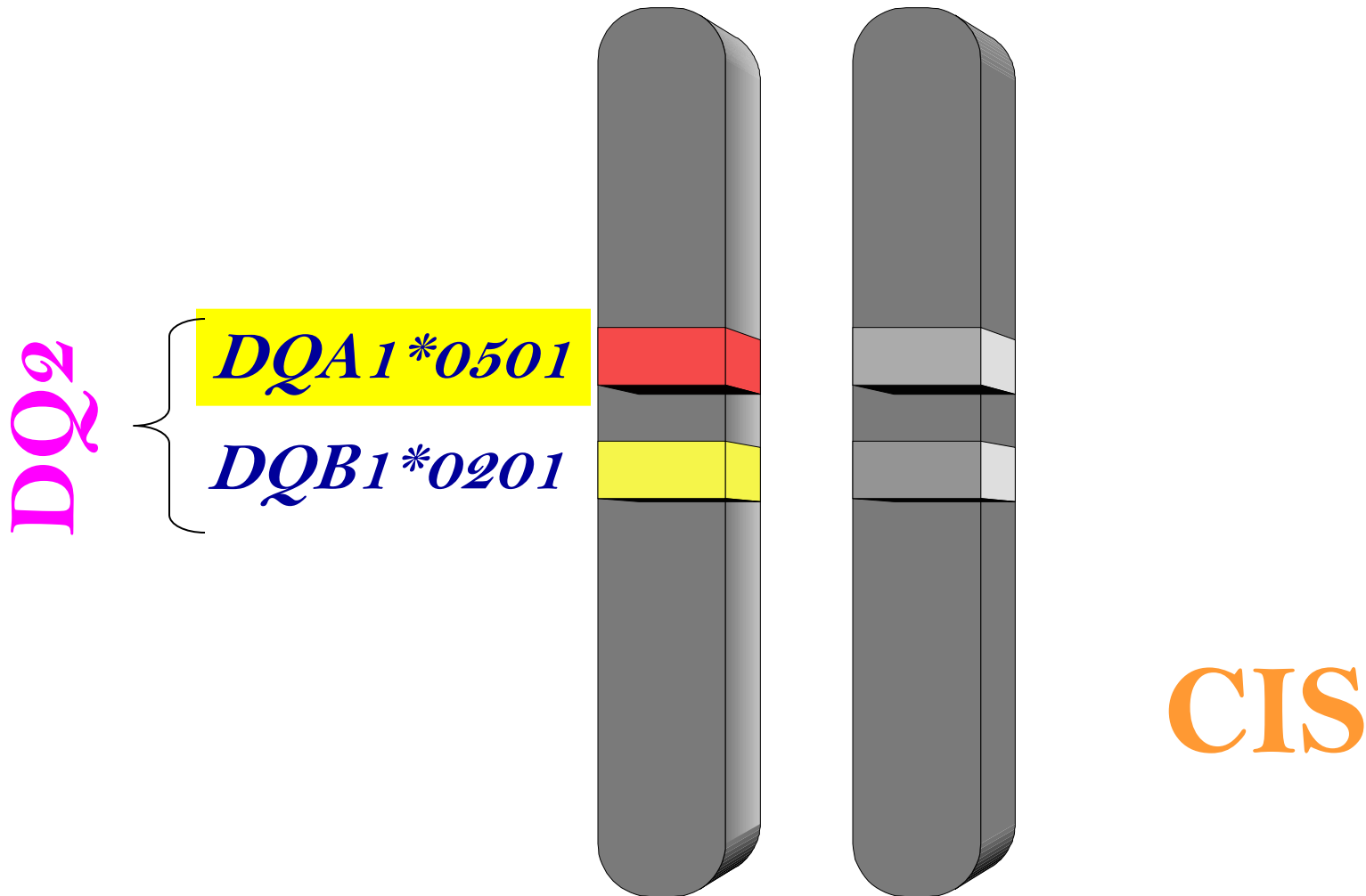


Figure 1

Linkage disequilibrium

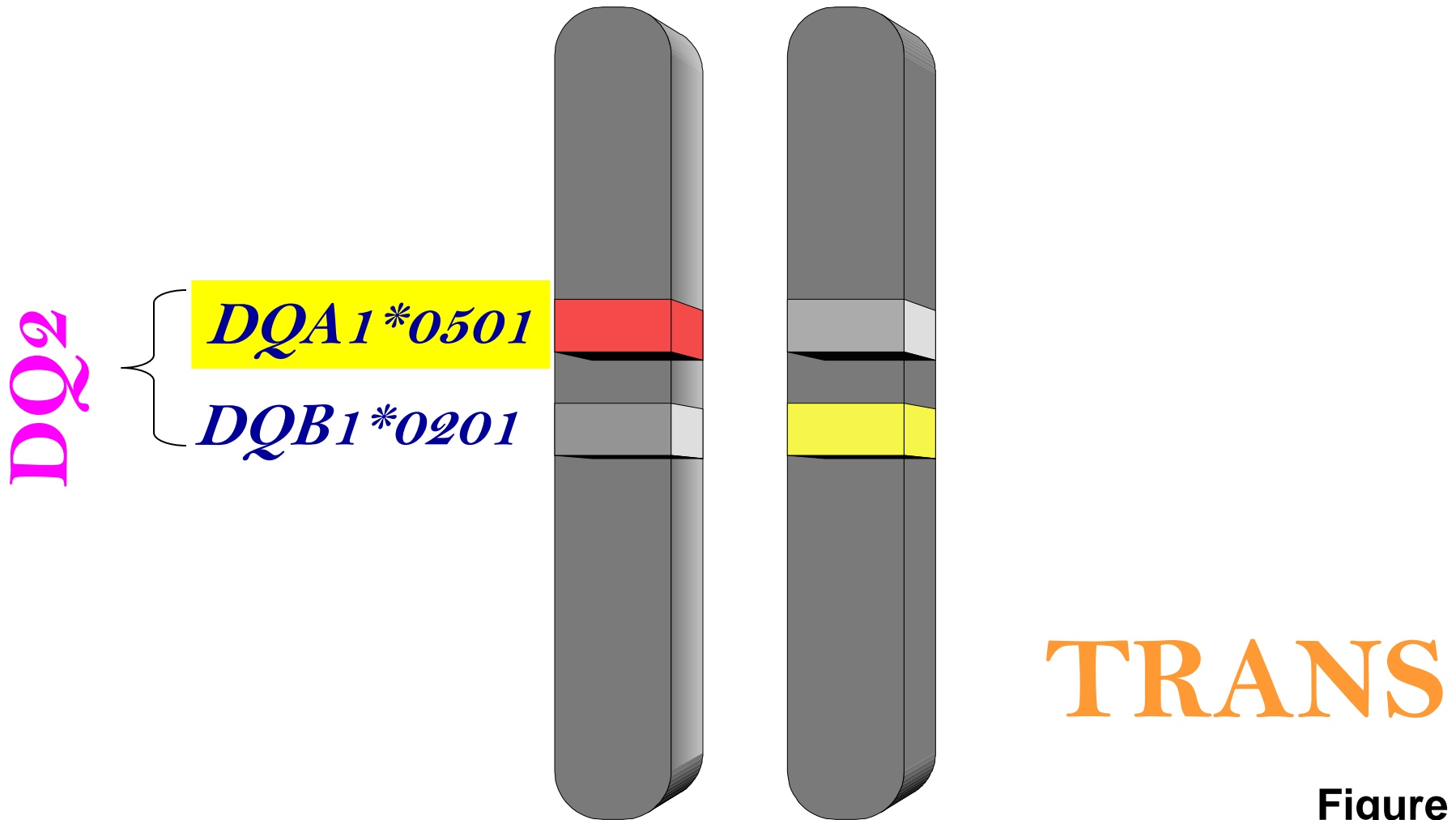
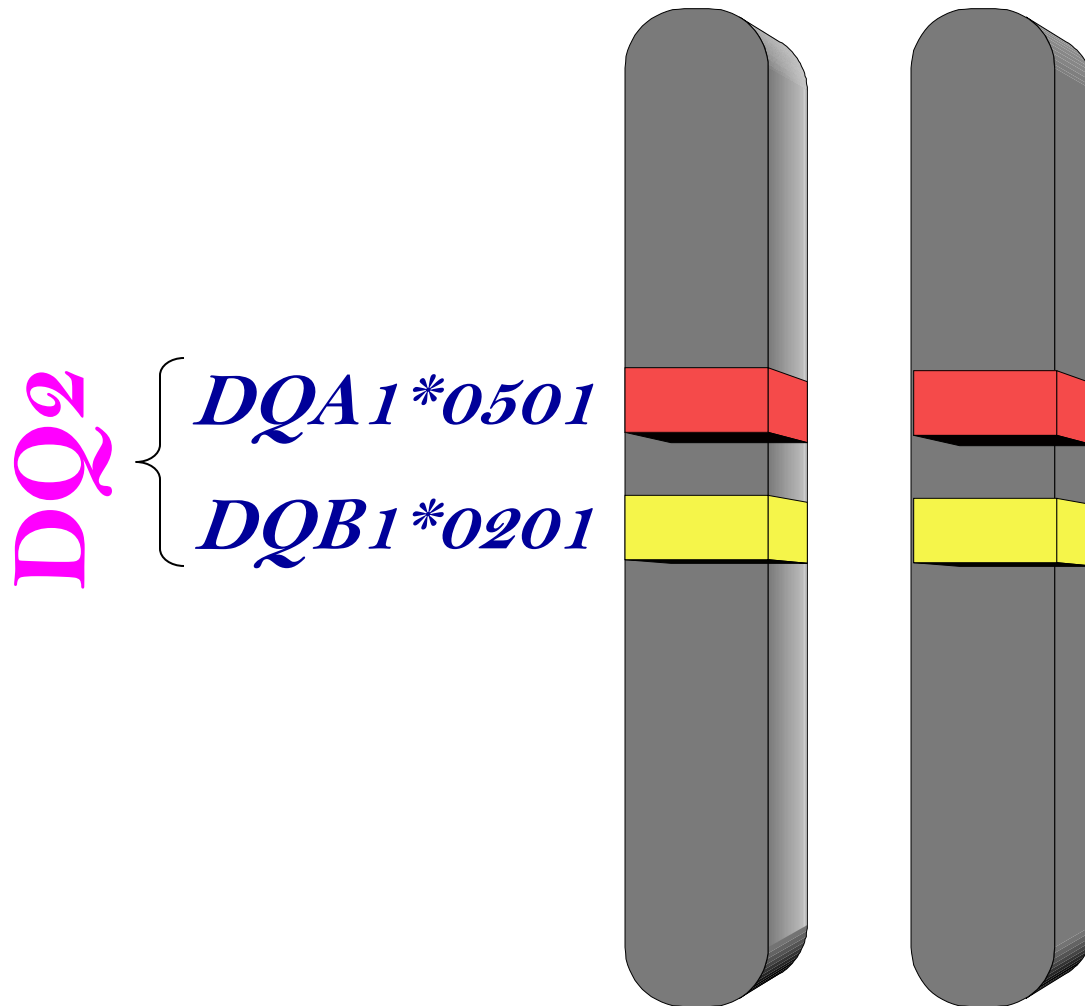


Figure 1

Linkage disequilibrium



CIS

Figure 1

Linkage disequilibrium

DQ₈ { *DQA: Any*
*DQB1*0302*

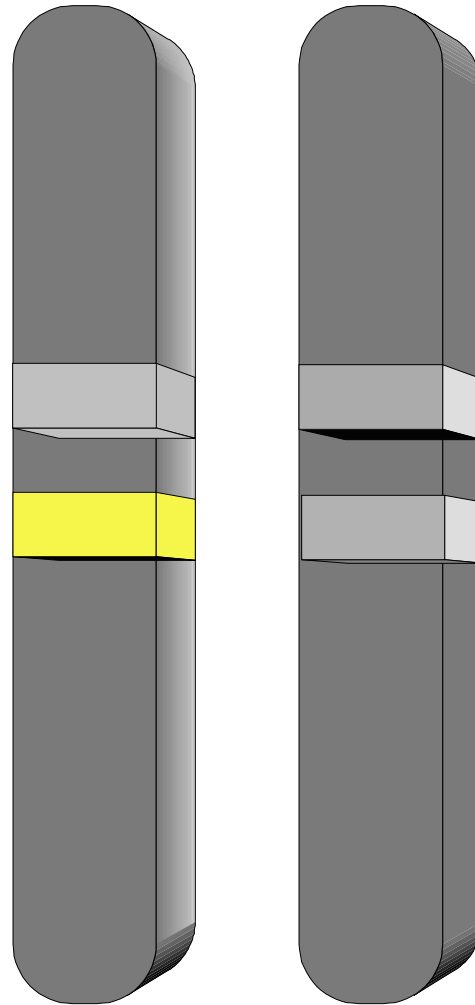


Figure 1

DQ2

A1*0501

B1*0201

DQ8

—

B1*0302



95% CD patients

5% DQ2/DQ8 negative????

Background II

The **HLA class II** genes play a key role in CD pathogenesis. HLA DQ2 and, less efficiently, DQ8 are the antigen presenting cells surface receptors for **deamidated toxic gliadin fragments** (Fig 2). Despite that their presence is considered necessary for CD pathogenesis, there is a low percentage of CD patients that are DQ2/DQ8 negative. However it has recently been reported that HLA DQ2 haplotype can be coded by the previously not described **A1*0201** allele.

Antigen Presenting Cell and its Role in CD

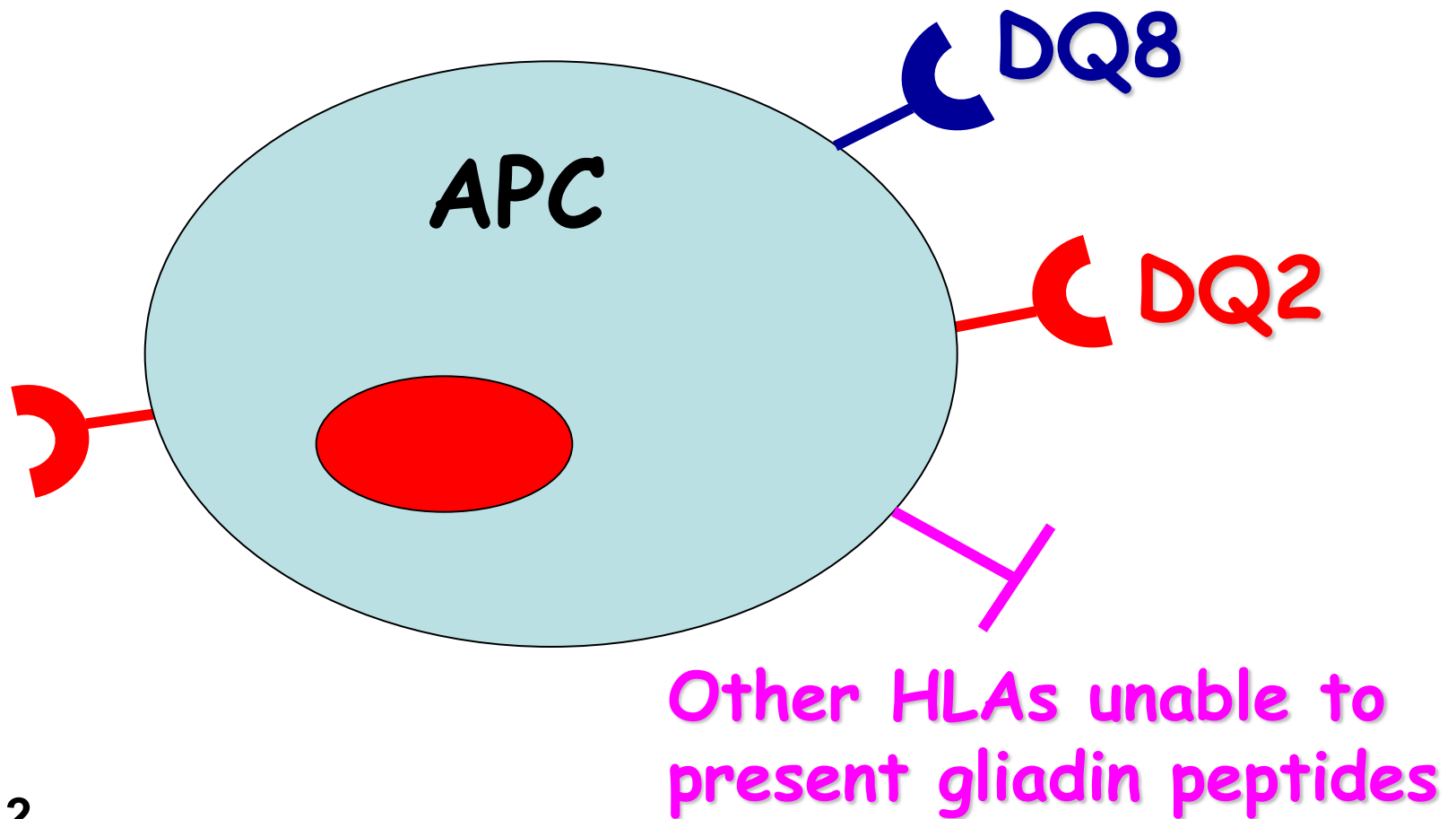


Figure 2

Aim

To verify whether patients classified as DQ2/DQ8 negative harbor the **A1*0201** allele both in European and North American populations.

Methods

The **HLA typing** was performed in biopsy-proven CD patients.

Genomic DNA was extracted from whole blood samples; the DNA obtained was subjected to **PCR** using the Eu-DQ® Kit (Eurospital, Trieste - Italy). This Kit contains multiplex PCR reactions for **DQA1*0501**, **DQA1*0201**, **DQ B1*0302**, and **DQ B1*0201** primers. Beta globin primers were used as internal control. The amplicons were resolved on 3% agarose gel and stained using ethidium bromide.

Results I

	U.S.A.				Europe			
	CD Patients		Controls		CD Patients		Controls	
Total (N)	78		40		100		60	
	N	%	N	%	N	%	N	%
DQ2 (A1*0501/B1*0201)	45	57.69	9	22.50	58	58.00	14	23.33
DQ2 (A1*0501+ A1*0201/B1*0201)	14	17.95	2	5.00	16	16.00	2	3.33
DQ2 (A1*0201/B1*0201)	6	7.69	2	5.00	5	5.00	3	5.00
DQ2/DQ8 (A1*0501/B1*0201 & B1*0302)	6	7.69	4	10.00	8	8.00	3	5.00
DQ2/DQ8 (A1*0201/B1*0201 & B1*0302)	0	0.00	2	5.00	1	1.00	1	1.67
DQ8 (B1*0302)	7	8.97	6	15.00	12	12.00	3	5.00
DQ2/DQ8 negative	0	0.00	15	37.50	0	0.00	34	56.67

Table 1

Results II

As reported in literature 95% of European CD patients tested positive for DQ2 and/or DQ8 using conventionally genotyping. The remaining 5% harbored the **A1*0201/B1*0201 haplotype** (see Table 1). Similarly in American CD patients 7.7% of DQ2/DQ8 negative subjects as established by the classical HLA genotyping also tested positive for A1*0201/B1*0201 allele. (see Table 1).

Therefore the A1*0201 allele accounted for the previously described DQ2/DQ8 negative CD patients, irrespective of geographical differences.

DQ2

DQ8

A1*0501

B1*0201



95% CD patients

B1*0302



A1*0201

B1*0201



5% CD patients
(previously negative)

Conclusions

Our data showed that **A1*0201 allele** accounts for all DQ2/DQ8 negative, biopsy-proven CD patients.

These results suggest that this allele should be included for the appropriate **genomic screening** of CD patients

The presence of **DQ2** and/or **DQ8** is confirmed to be crucial for deamidated gliadin presentation by antigen presenting cells as a key step in the pathogenesis of the intestinal damage typical of celiac disease (figure 3).

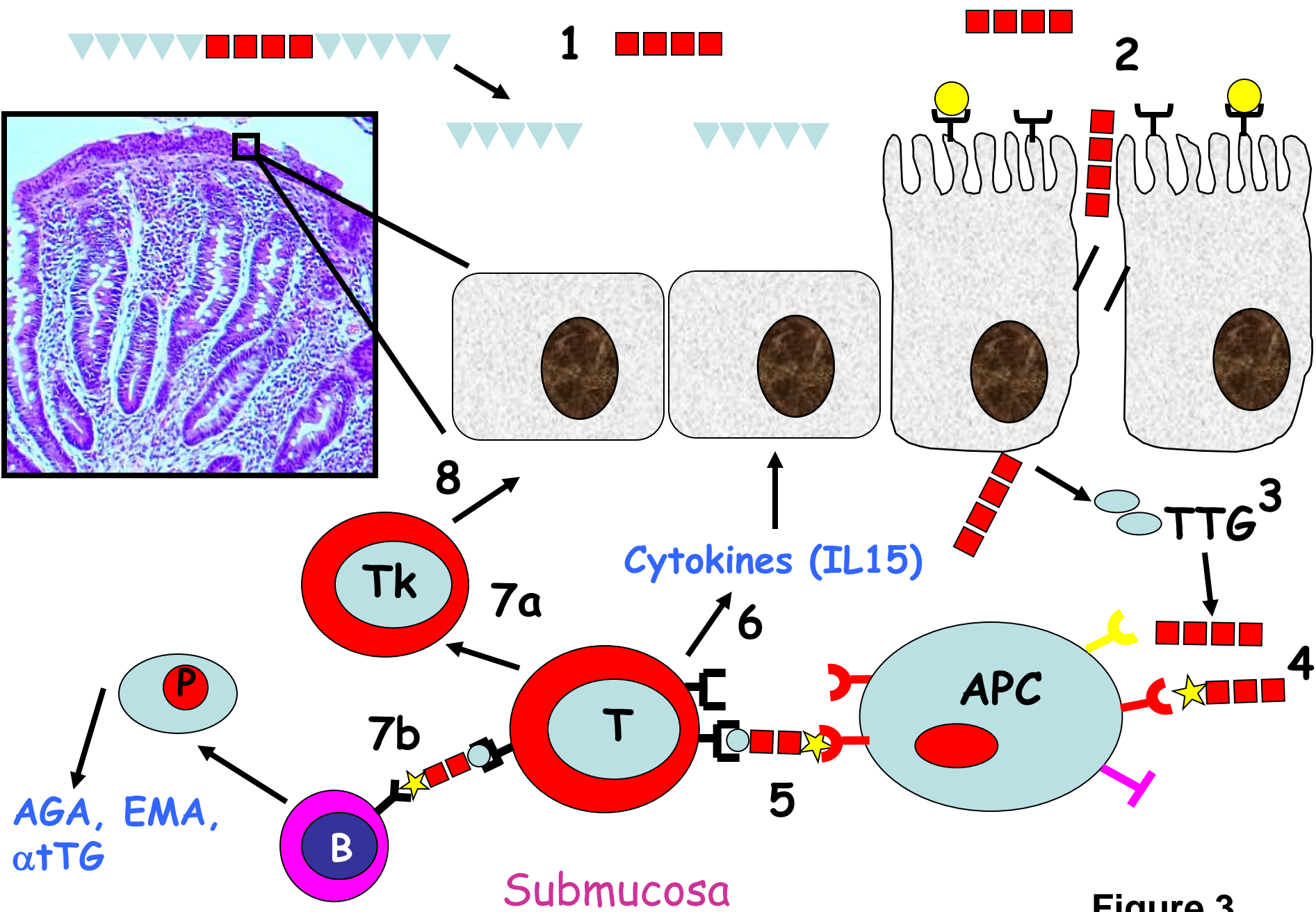


Figure 3