

Eksozomlar

Prof. Dr. Tlay Kılıçaslan Ayna

**İzmir Katip Çelebi Üniversitesi Tıp Fakltesi Tıbbi Biyoloji AD.
TEAH Doku Tipleme Lab.**

REVIEW

Exosomes and microvesicles in kidney transplantation:
the long road from trash to gold

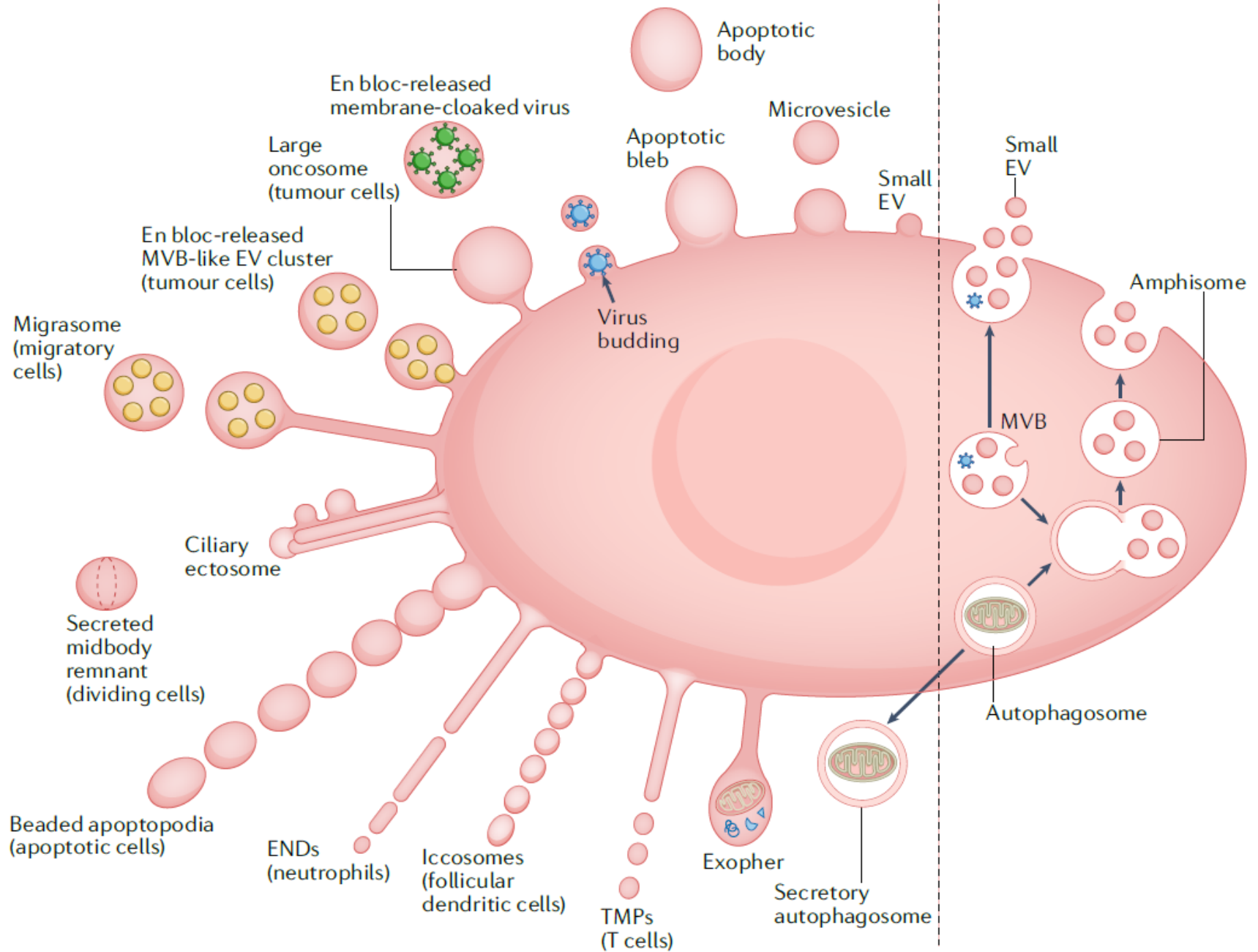
LUIS RAMALHETE^{1,2,3}, RUBEN ARAÚJO², ANÍBAL FERREIRA^{2,4},
CECÍLIA R. C. CALADO^{5,6}

Yıllar	Yayın sayısı
1991-2014	43
EV-2024	4455
Exo-2024	3213

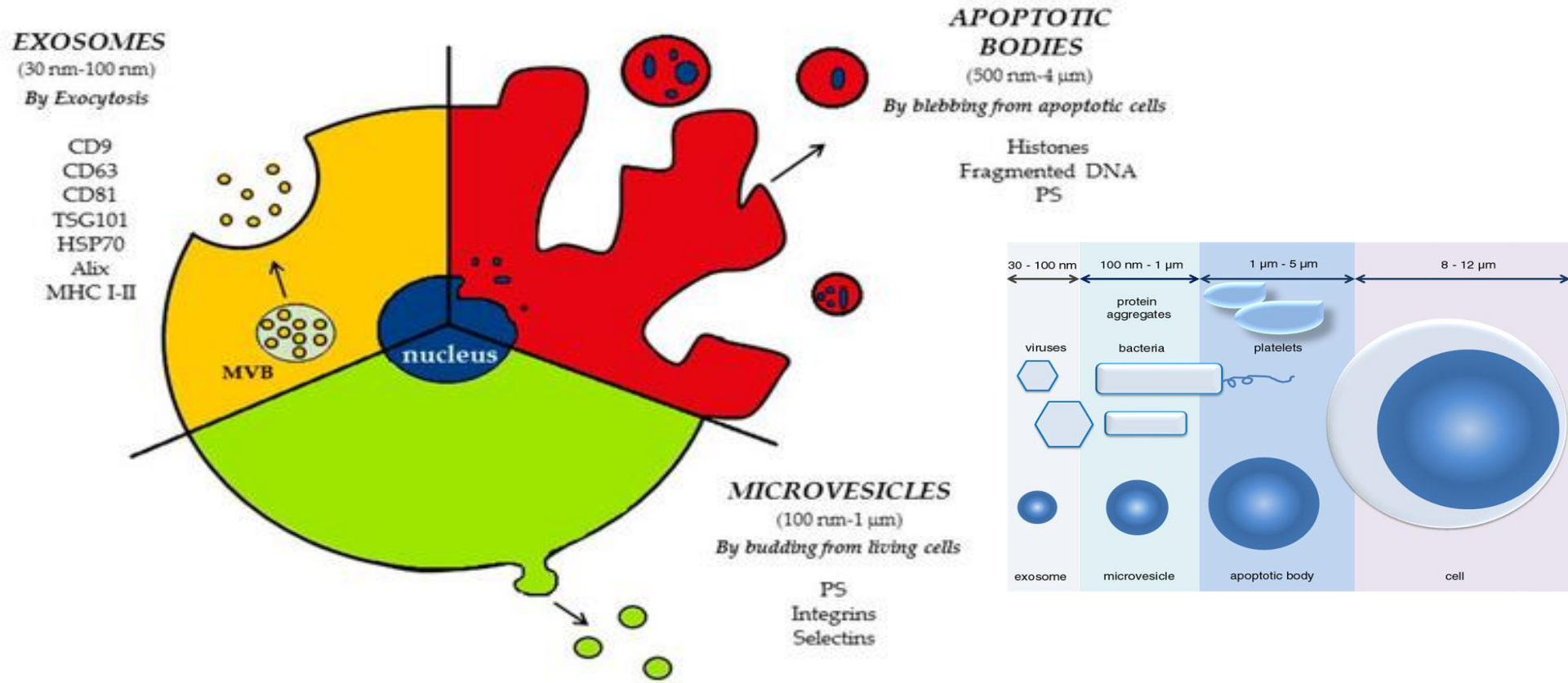
Protein	41860
RNA	7540
miRNA	764
Lipit	1116

Ectosomes

Exosomes



Uluslararası Extrasellüler veziküller Derneğinin (ISEV) yönergesi;



EV'lerin biyokimyasal bileşimi farklıdır; ancak ISEV, EV'lerin izolasyonu, karakterizasyonu ve onayı için yönergeler tanımlamıştır.

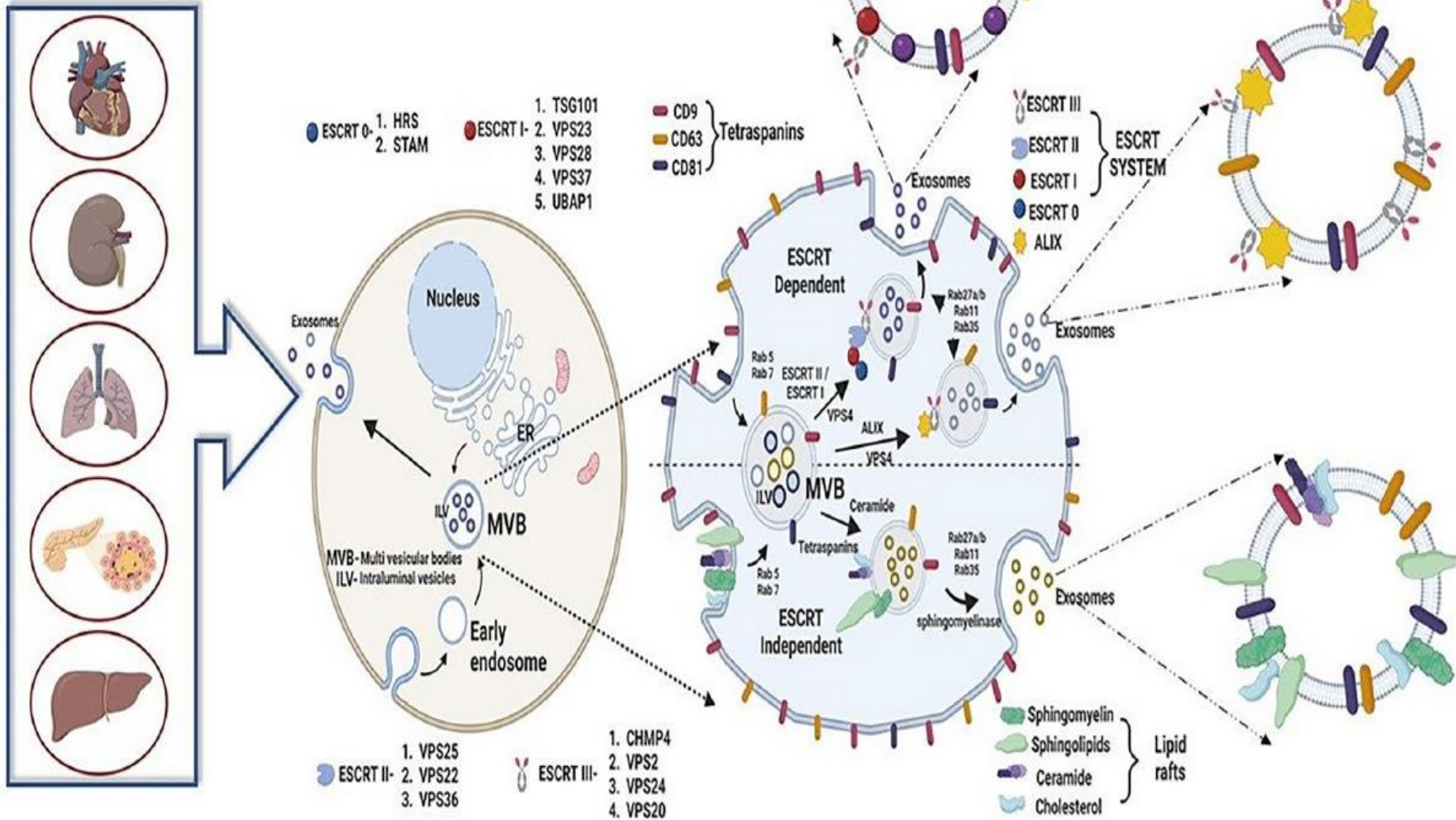
Table 1 Main characteristics of exosomes, microvesicles, and apoptotic bodies

	Exosomes	Microvesicles	Apoptotic bodies	References
Size	30–100 nm	100–1,000 nm	1–5 μ m	[3]
Origin	Intraluminal vesicles within multivesicular bodies	Plasma membrane and cellular content	Plasma membrane, cellular fragments	[4]
Mechanism of formation	Fusion of multivesicular bodies with the plasma membrane	Outward blebbing of the plasma membrane	Cell shrinkage and programmed cell death	[5, 6]
Release	Constitutive and/or cellular activation	Constitutive and/or cellular activation	Apoptosis	[4]
Time of release	Ten minutes or more	Few seconds	–	[7, 8]
Pathways	ESCRT-dependent Tetraspanin-dependent Ceramide-dependent Stimuli-dependent	Ca ²⁺ -dependent Stimuli- and cell-dependent	Apoptosis-related	[3]
Lipid membrane composition	Enriched in cholesterol and ceramide, expose phosphatidylserine, contain lipid rafts	Expose phosphatidylserine, enriched in cholesterol and diacylglycerol, contain lipid rafts	–	[3, 9]
Content	Proteins, mRNA, miRNA, lipids	Proteins, mRNA, miRNA, lipids	Cell organelles, proteins, nuclear fractions, DNA, coding and noncoding RNA, lipids	[3]

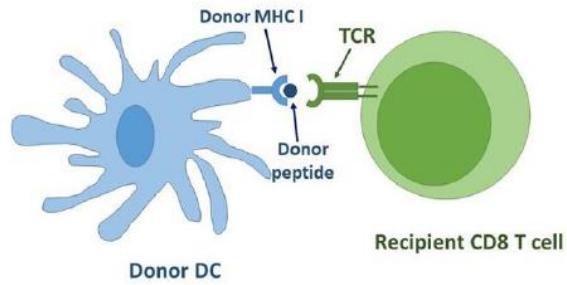
ESCRT endosomal sorting complex required for transport

endosomal-sorting complex required for transport (ESCRT)

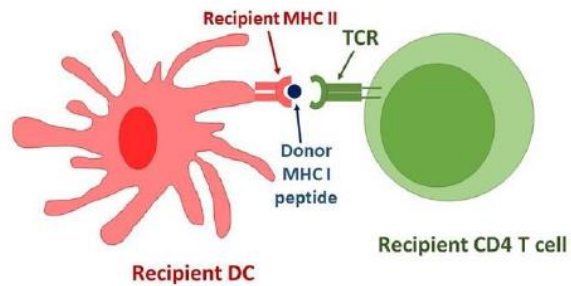
Exosomes secreted from the transplanted organs



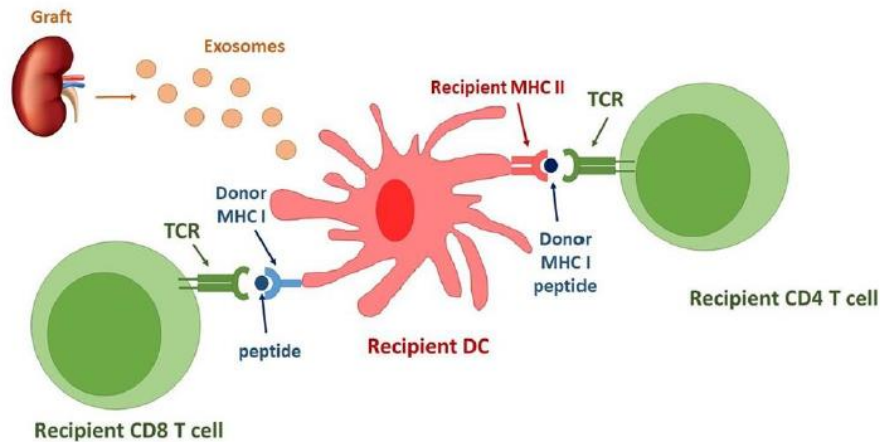
A. DIRECT PATHWAY OF ALLORECOGNITION



B. INDIRECT PATHWAY OF ALLORECOGNITION

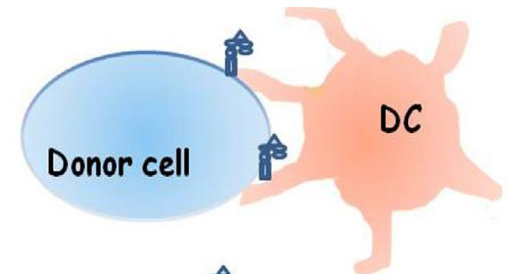


C. SEMI-DIRECT PATHWAY OF ALLORECOGNITION

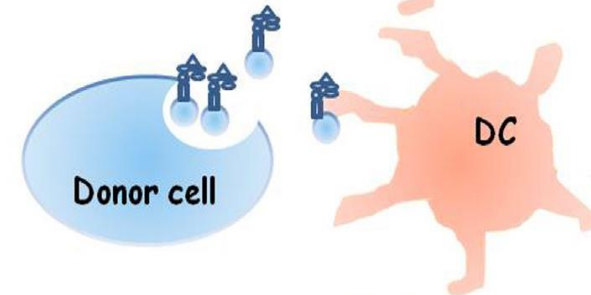


Cross dressing

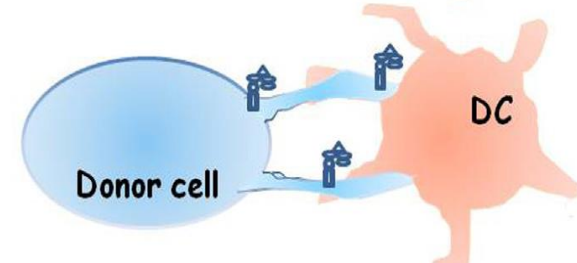
Trogocytosis



Exosomes



Tunneling Nanotubes





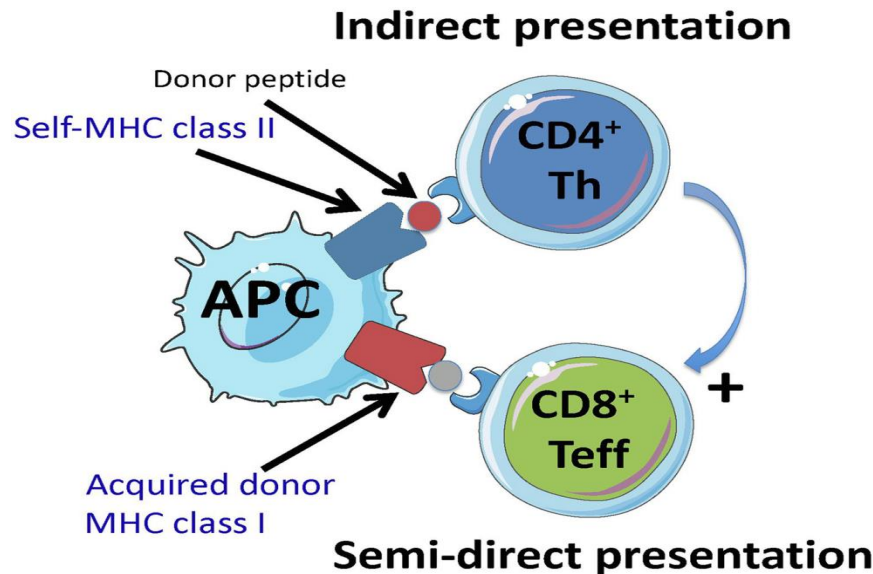
Frelinger Jeffrey-1974

MHC'lerin lökositler
arasında transferi



Brian Dolan-2006

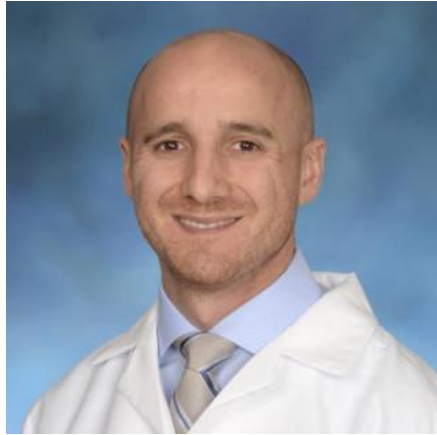
Transferden sonra MHC-
peptit kompleksleri iki gün
boyunca APC de exp.



Lesley Smythe 2013

Alıcı DC'leri trf
allogeneik MHC-sınıf I
kazanımının,
transplantasyonu
takiben en az 1 ay
boyunca
gerçekleştiğini ve CD8
T hücre
aktivasyonunun temel
sebebi olduğunu
göstermişlerdir.

2016

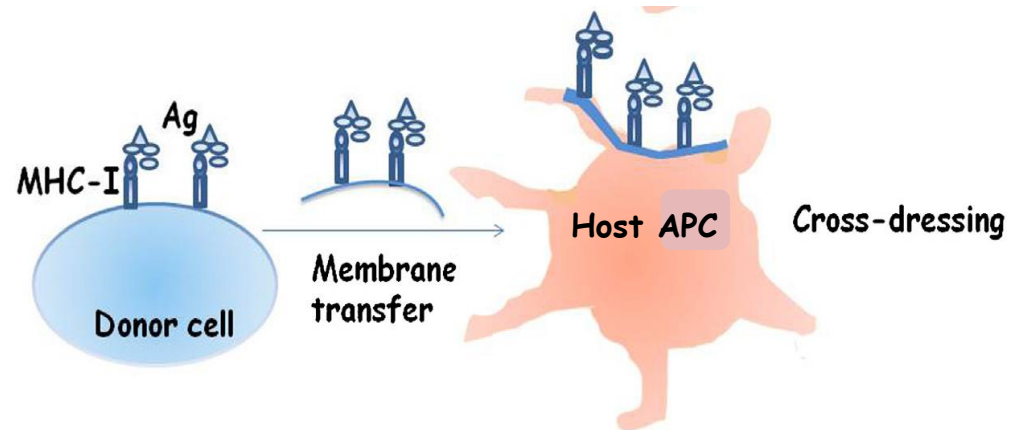
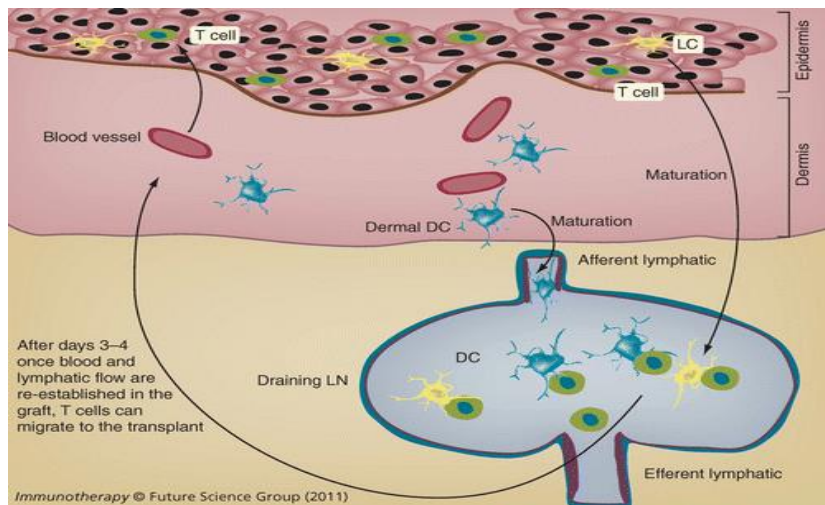


Jose Marino

100/1000000 donör DC
90000/1000000 Cross dressing alıcı APC



Mohamed H. Babiker-Mohamed

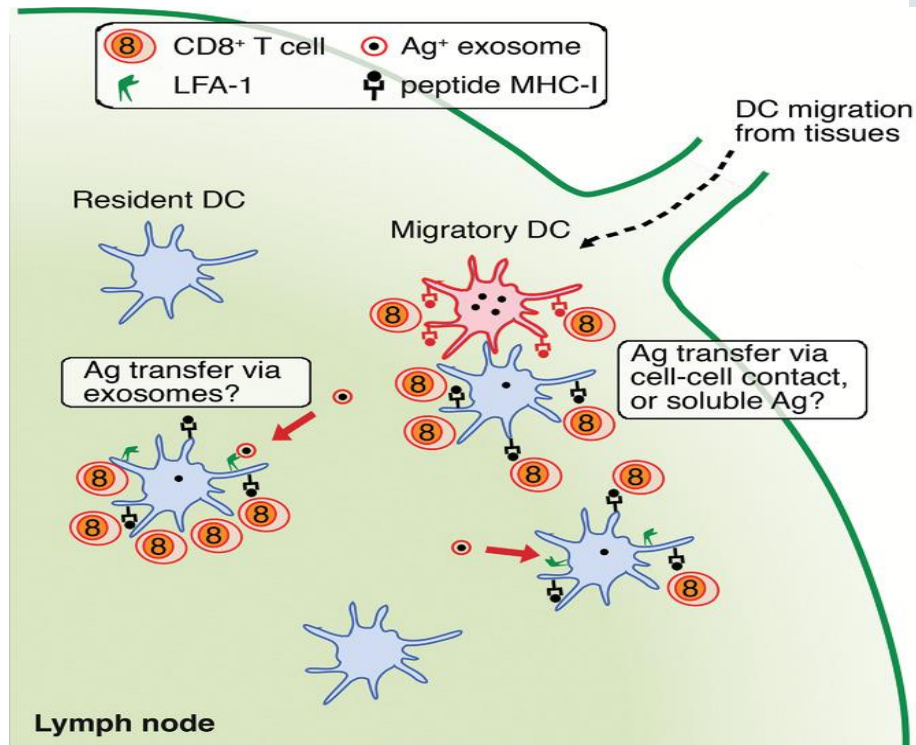


Marino J. Donor exosomes rather than passenger leukocytes initiate alloreactive T cell responses after transplantation, Science Immunology, 2016.

2016



Adrian Morelli

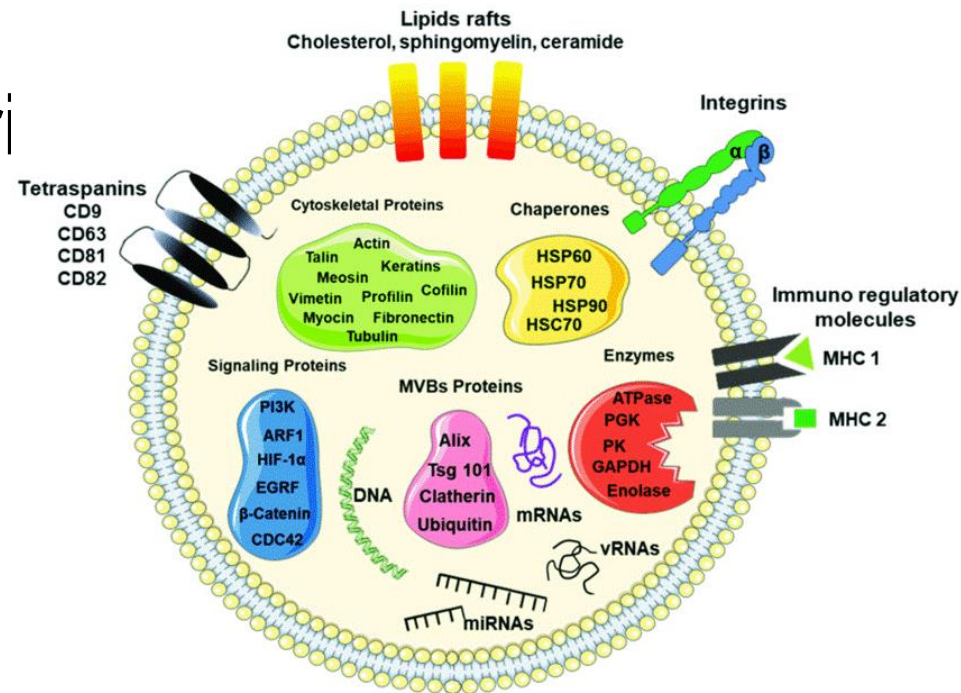


Quan Liu

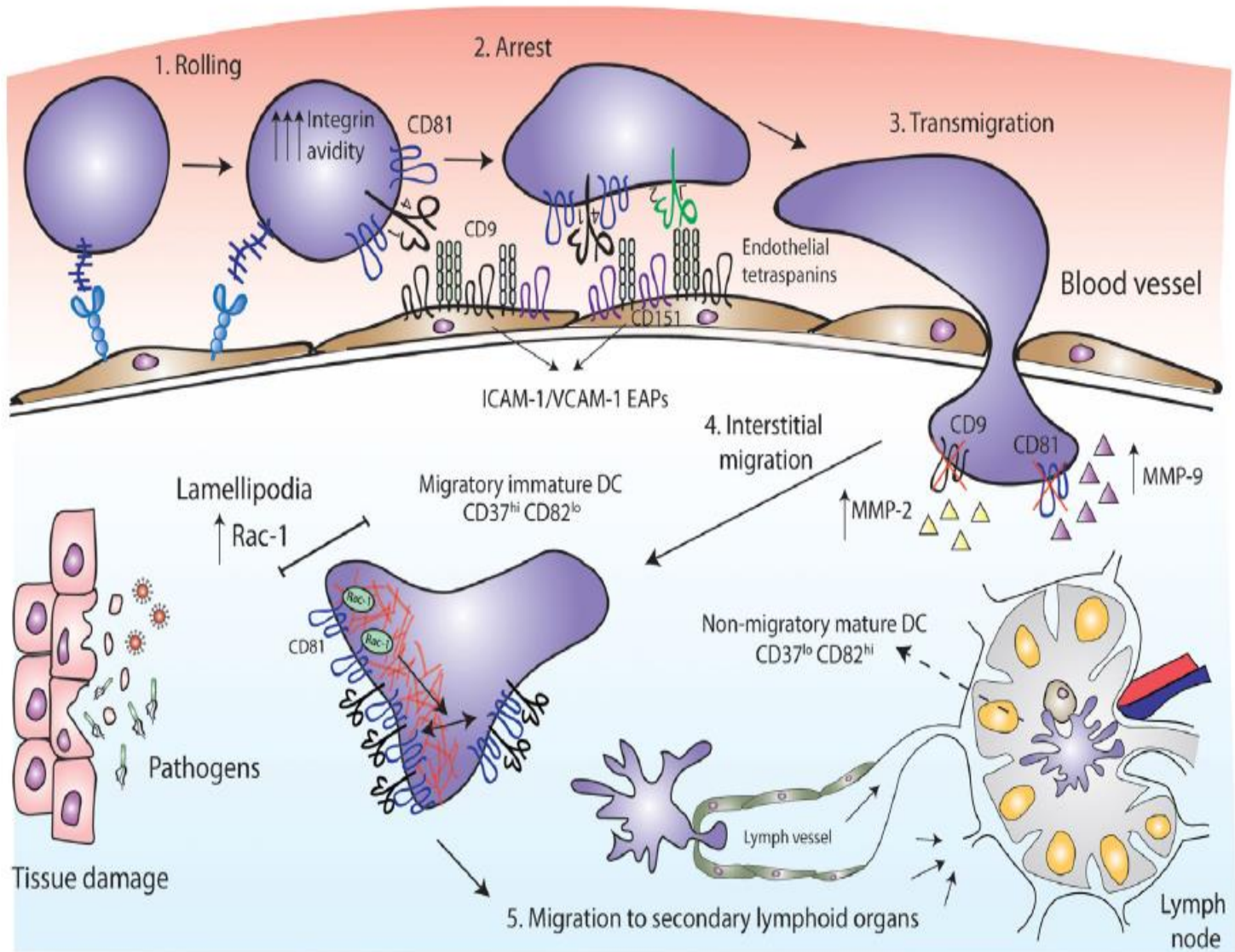
Liu Q, Morelli AE, Donor dendritic cell-derived exosomes promote allograft-targeting immune response, J Clin Invest, 2016

Tetraspaninler

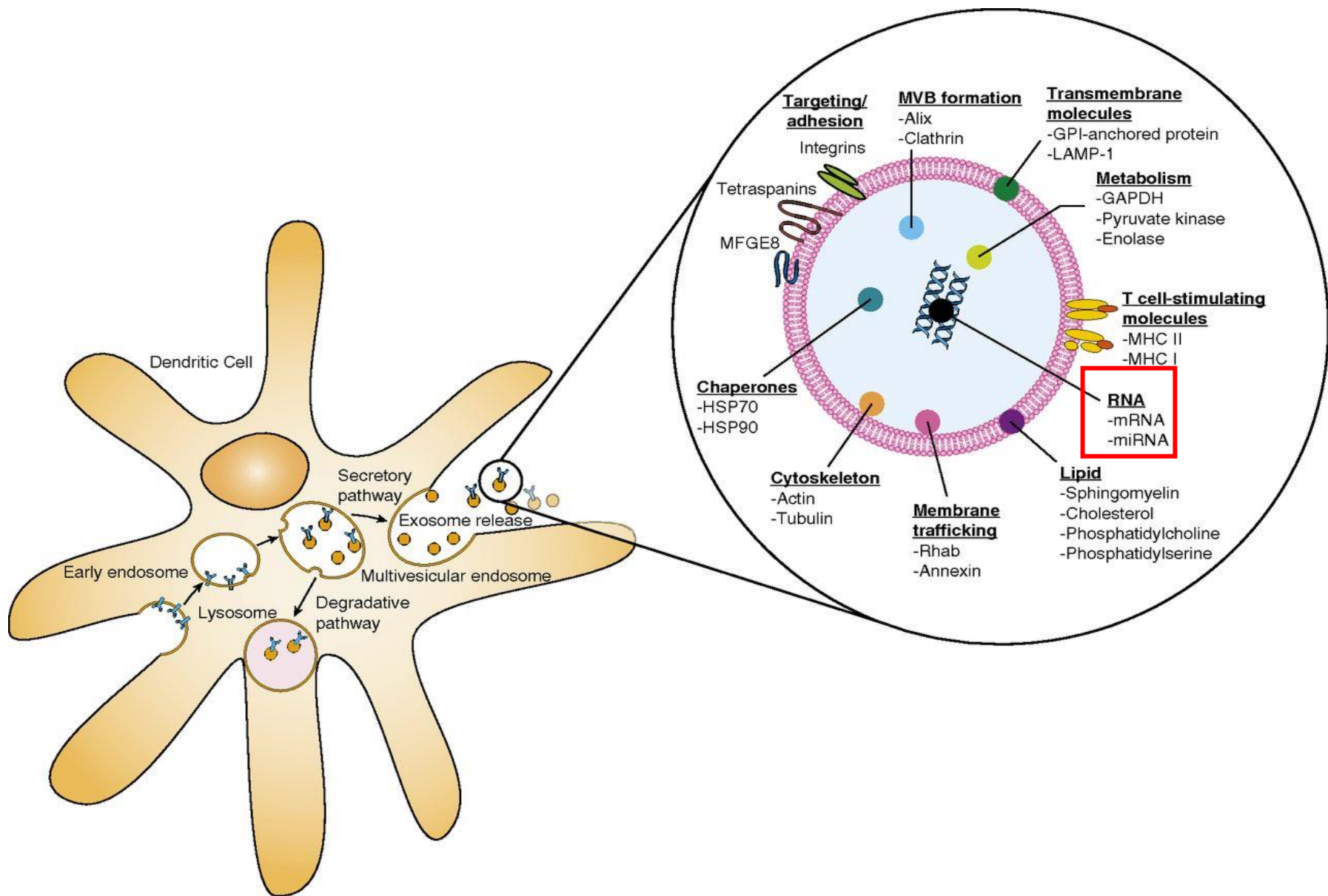
- ✓ PRR
- ✓ MHC II
- ✓ Adezyon molekülleri
- ✓ Sinyal molekülleri



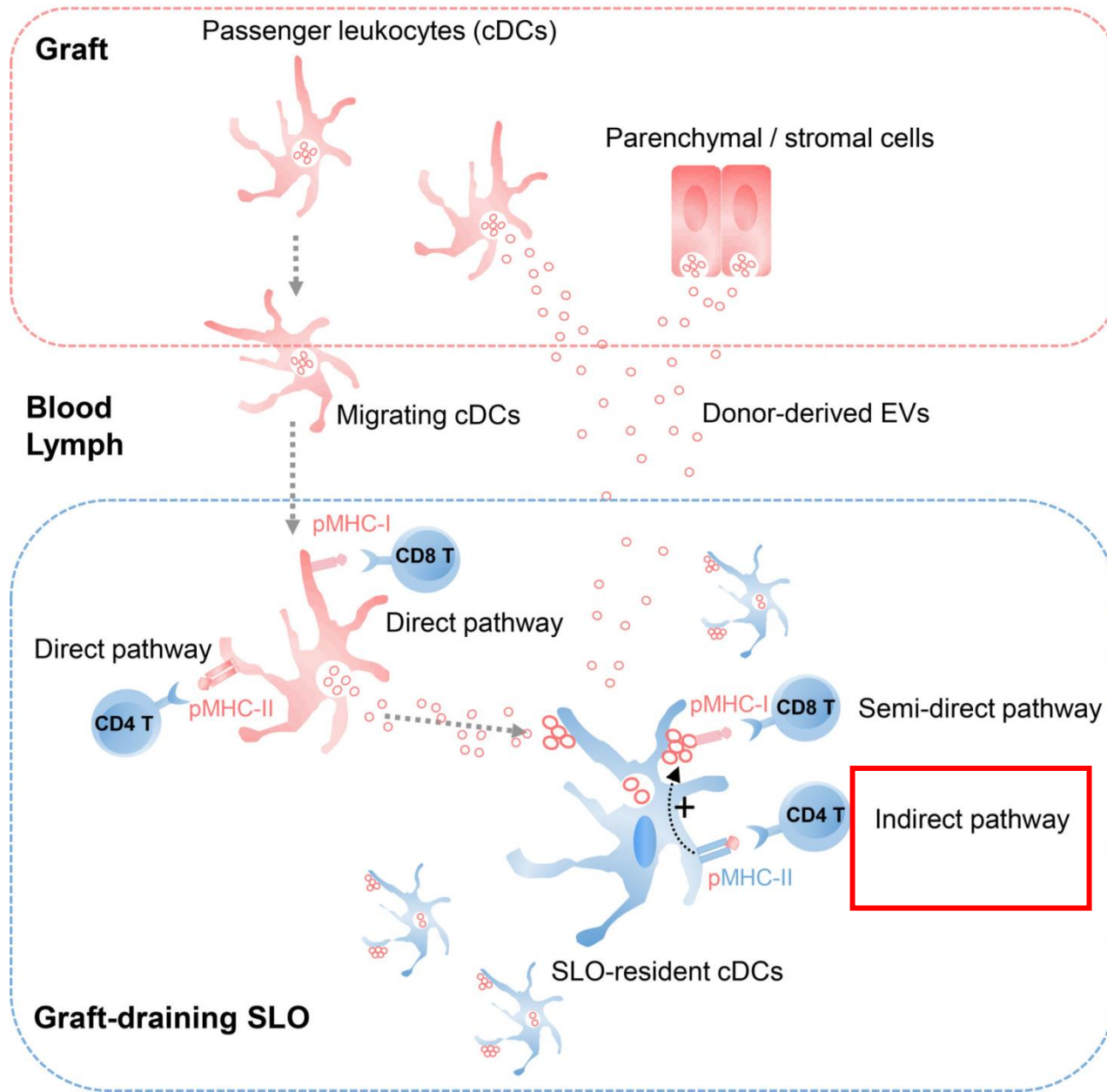
Tetraspanins act as key players in APC migration

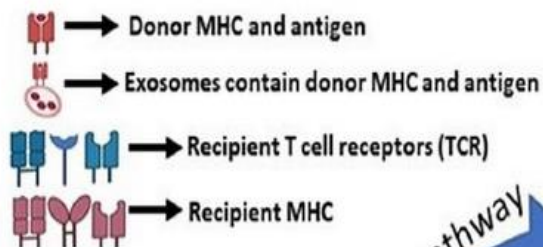
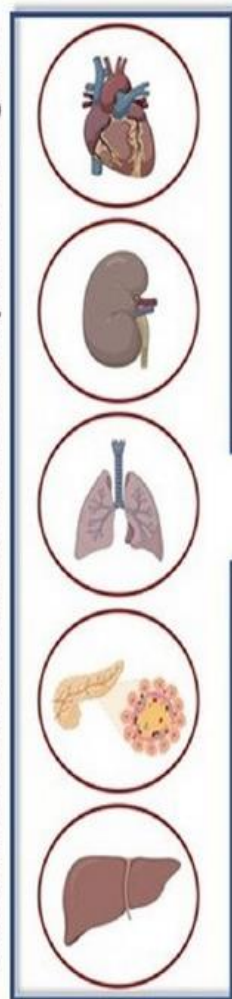


EV'yi salgılayan Hücre!!!!!! farklı T hücre yanıtları
EV'lerin transfer ettiği kargo !!!!!!!!! Sayı!!!!!!



Donor-derived
Recipient-derived

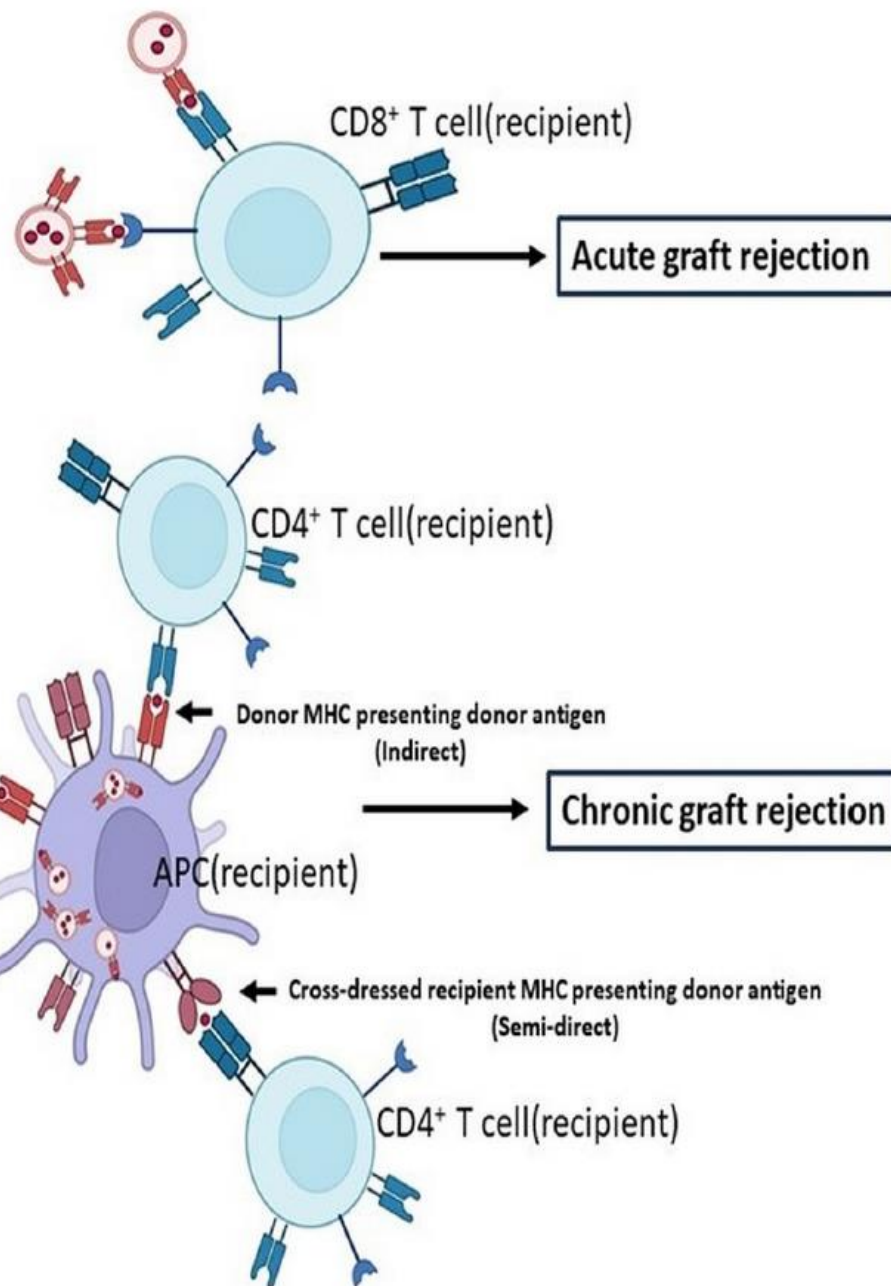




Antigen and exosomes from allograft

Direct pathway

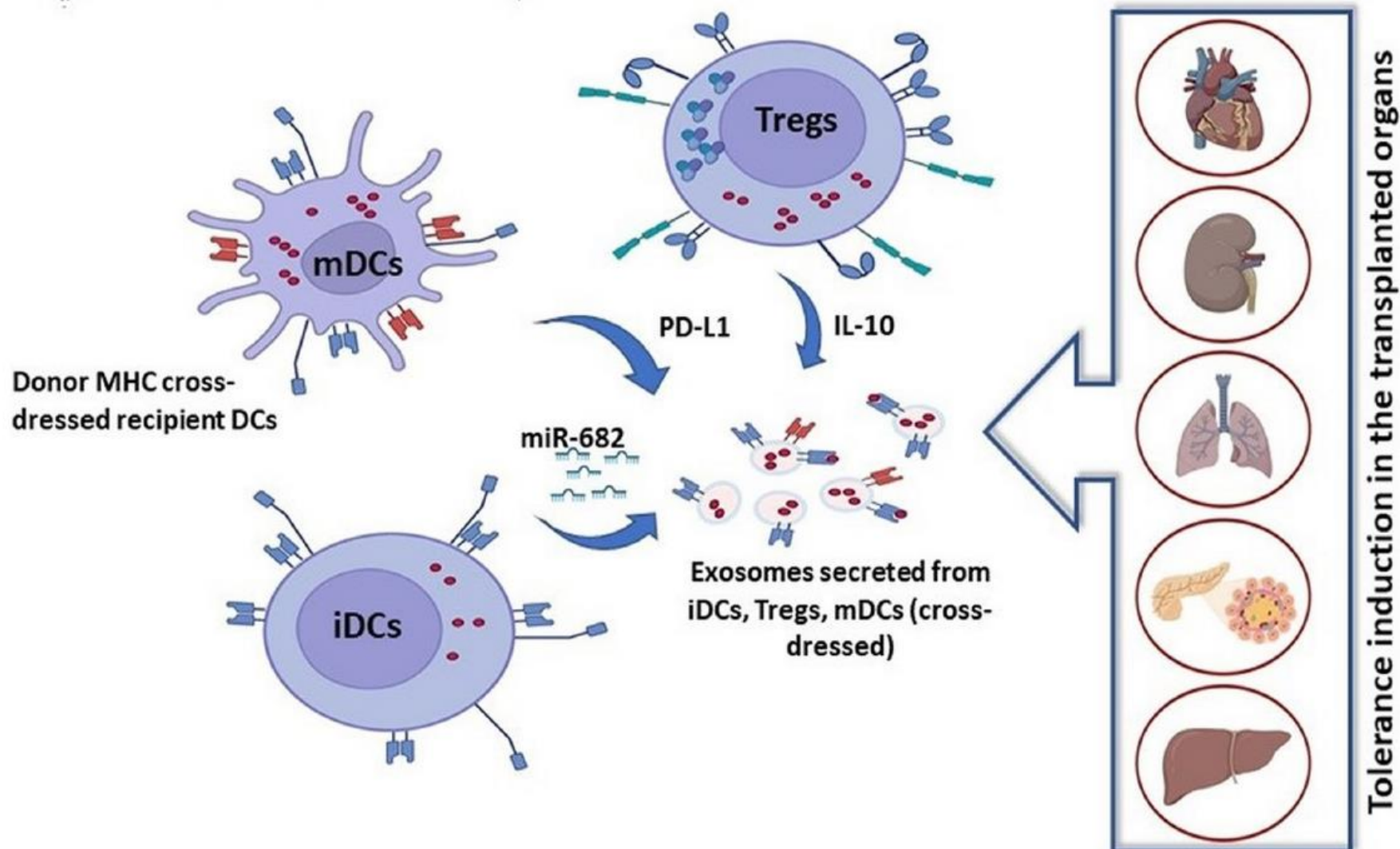
Semi-direct or Indirect pathway

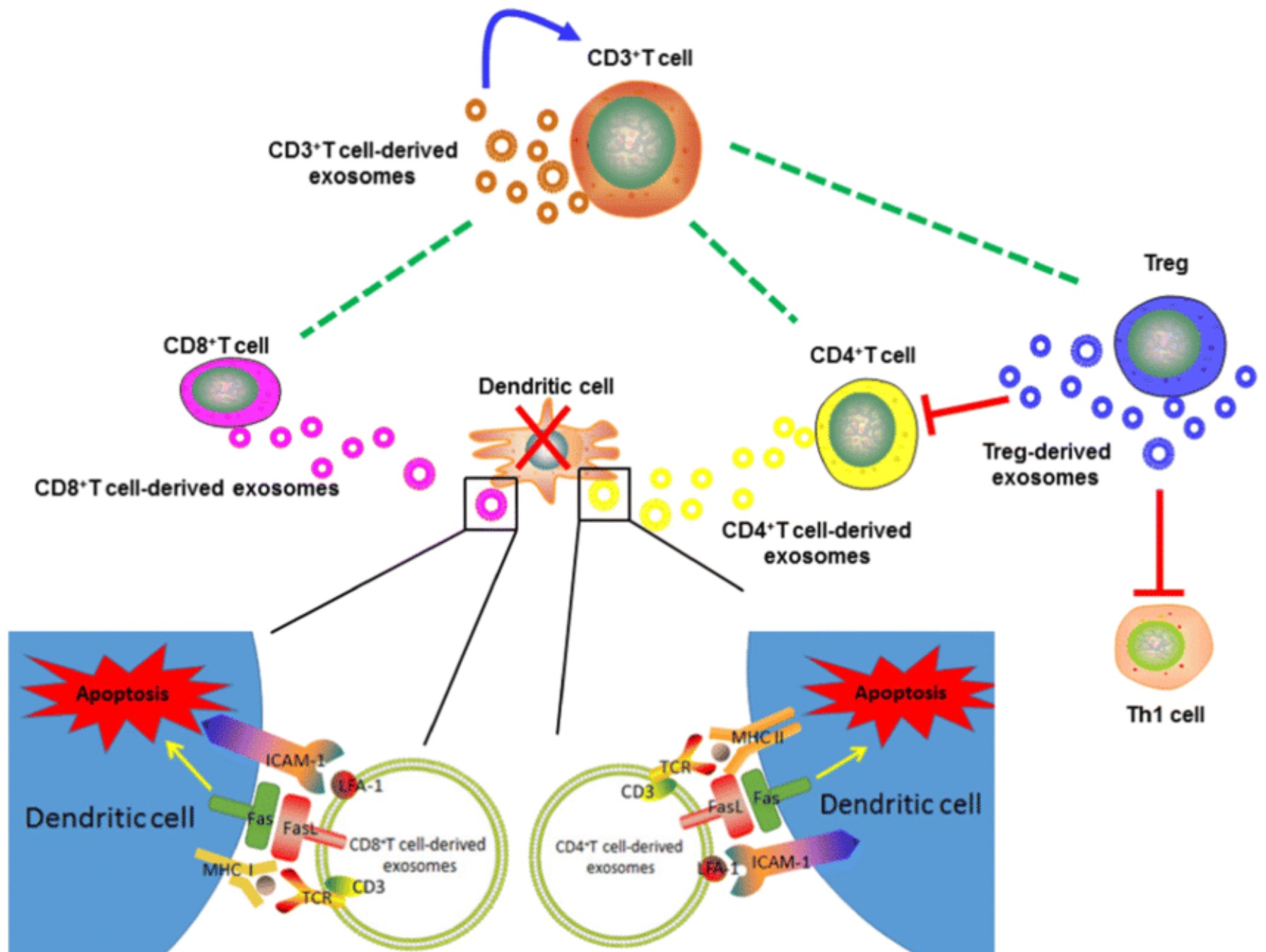


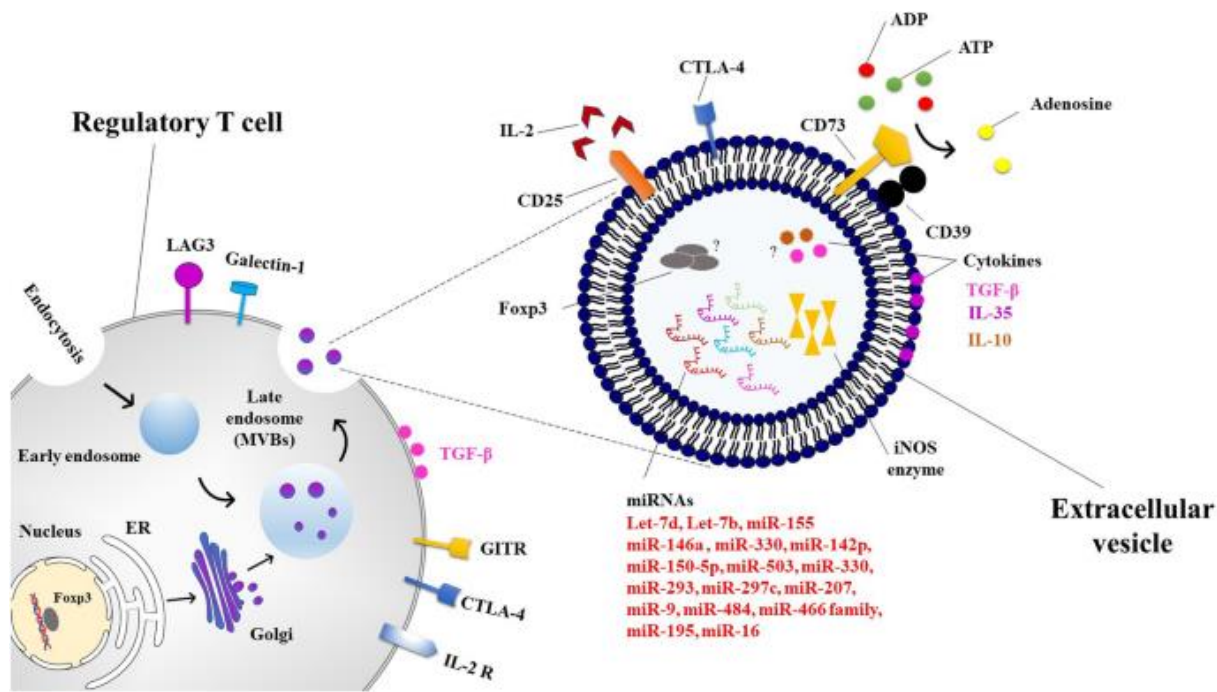
- Recipient's MHC
- Donor MHC and antigen
- Foxp3
- T cell receptor (TCR)

- CD11c
- Interleukin 10 (IL-10)
- CD4

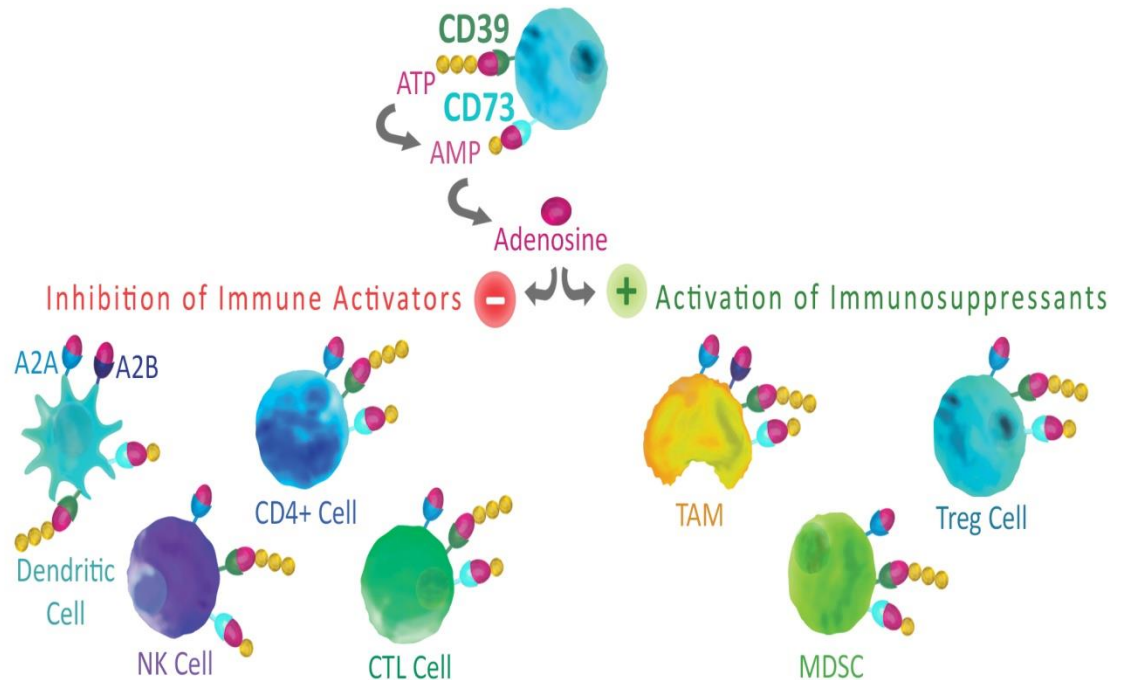
- *mDCs – Mature dendritic cells
- *iDCs – Immature dendritic cells
- *Tregs – T regulatory cells







Komplexe pro
Kostimulatoren



Non invasif tanı aracı olarak Exosomlar

Kan,

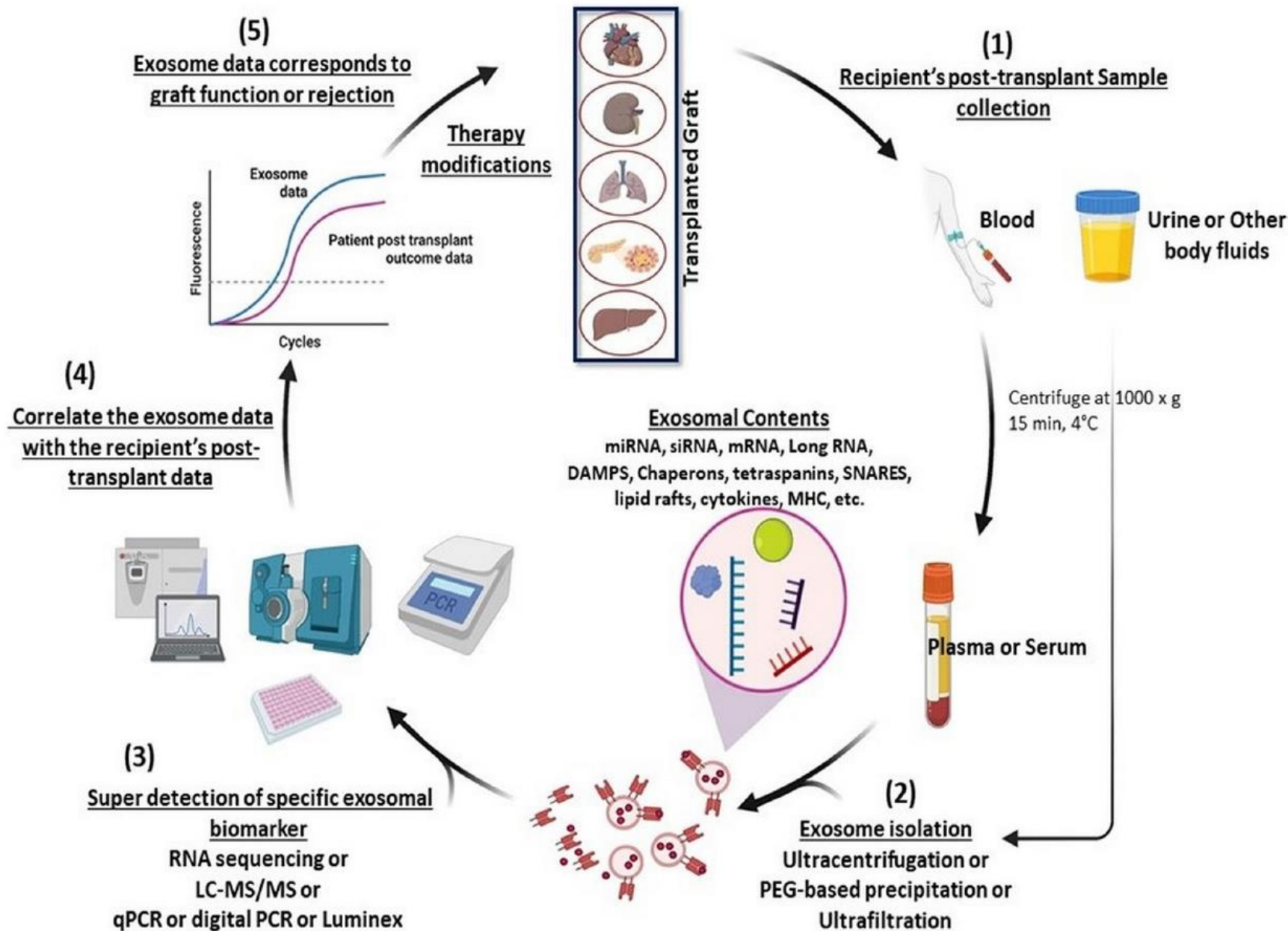
İdrar

Bronkoalveolar lavaj (BAL),

Tükürük,

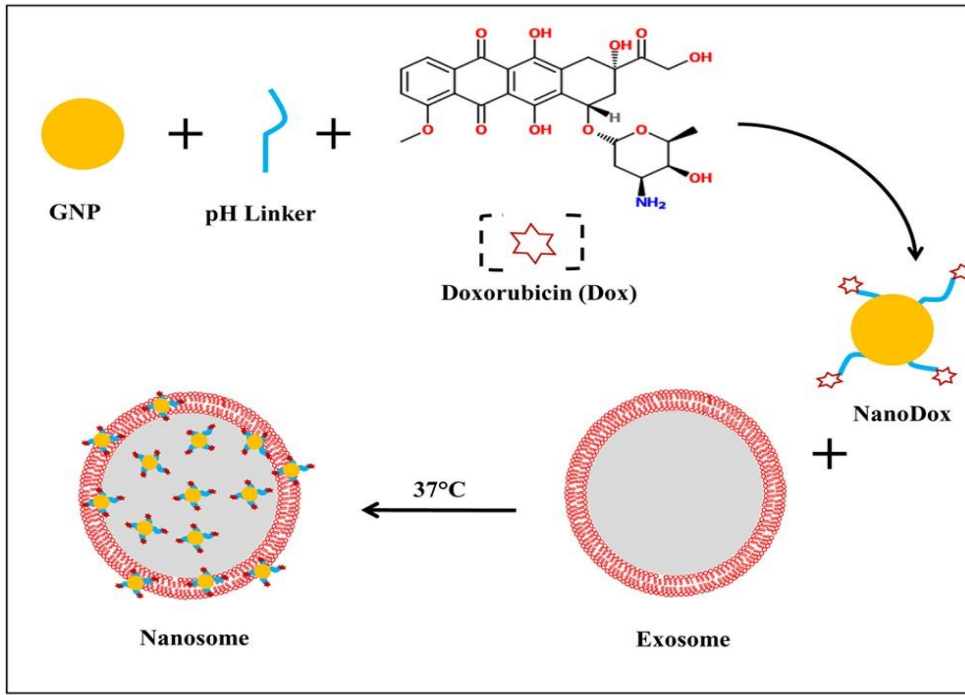
Süt,

Amniyotik sıvı



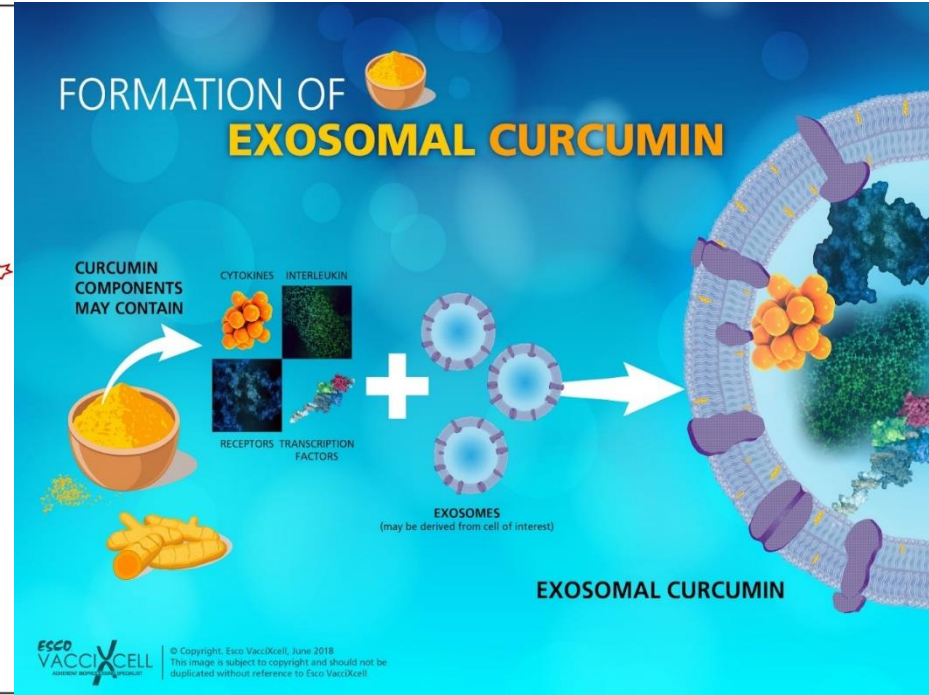
Exosome temelli tedaviler

- Lipit bariyer ilacın korunması açısından önemli
- Alıcı hücrelere endositozu kolaylaştırır.
- Exosomların düşük immunojenitesi yan etkiyi azaltır.
- Kan beyin bariyerini geçer



Anti kanser etki

Yan etki



Stabilite

Kan konsantrasyonunda

**Lipopolisakkaritlerle
indüklenen septik şoka
karşı koruyucu**

Table 1

Recipient's exosomes were used as a potential biomarker in clinical transplantation.

	Transplant Type	Methods used	The outcome of the study
1	Kidney	LC-MS/MS	<ul style="list-style-type: none"> > The global protein contents were profiled from the exosomes isolated from the urine samples of transplant recipients > About 17 proteins were seen to increase in the exosomes of the transplant recipients diagnosed with T cell-mediated rejection (TCMR) > Tetraspanin-1 and hemopexin in the exosomes were significantly higher in TCMR recipients [33].
2	Kidney	qPCR	<ul style="list-style-type: none"> > Total RNA was extracted from the plasma samples collected from the transplant recipients > About 21 candidate mRNA levels are high in the exosomes of recipients diagnosed with antibody-mediated rejection (AMR) > Among them, gp130, SH2D1B, TNFα, and CCL4 mRNAs were significantly higher in AMR cohort exosomes than in other groups [32].
3	Kidney	iKEA (Biochip)	<ul style="list-style-type: none"> > Exosomes were directly captured from the urine samples collected from the recipients > The ikea captured high levels of CD3 (T- cell marker) positive exosomes in allograft-rejecting recipient's urine samples with 91.1 % accuracy. [72]
4	Kidney	qPCR (TaqMan)	<ul style="list-style-type: none"> > Exosomes were isolated from the recipient urine samples and the miRNAs were quantified using TaqMan qPCR > The elevated amount of BK viral miRNAs B1-5p and 3p in the urine exosomes of renal transplant recipients with BK virus nephropathy (BKVN). > These exosomal viral miRNAs could be used as a surrogate marker for the diagnosis of BKVN [116]

Table 1 (*continued*)

	Transplant Type	Methods used	The outcome of the study
5	Pancreatic Islet	HLA-specific NTA and Immunoblotting	<ul style="list-style-type: none">➤ Quantitation and characterization of allo-islet exosomes in the recipient's plasma➤ The donor islet-specific exosome numbers decreased in the recipient plasma before islet graft dysfunction➤ The donor-specific islet exosomes contain the exosome markers (CD63, flotillin), and transmembrane protein (FXYD2) along with donor-specific HLA molecules. [71]
6	Pancreatic Islet	NTA	<ul style="list-style-type: none">➤ Quantitation and characterization of allo-islet exosomes in the recipient's plasma➤ There is time specific increase of auto antigen GAD65 in the plasma exosomes of these recipients indicating the recurrence of autoimmunity in T1D recipients after islet transplantation [117]

(continued on next page)

Table 1 (continued)

	Transplant Type	Methods used	The outcome of the study
7	Pancreatic Islet	RNA sequencing and qPCR	<ul style="list-style-type: none"> ➤ Total RNAs were isolated from the Islet transplant recipient's plasma and miRNAs present were profiled using RNA sequencing and qPCR ➤ Islets release a significant amount of ER stress-induced miRNAs (miR-29b and miR-216a) and damage associated miRNAs (miR-375 and miR-148a) to the circulation during peritransplant time through exosomes ➤ The high amount of these exosomes release during the peritransplant period correlated with poor transplant outcomes in Islet transplants. [7,118]
8	Pancreatic Islet	LC-MS/MS and ELISA	<ul style="list-style-type: none"> ➤ Proteomic profile of the exosomes isolated in samples collected during clinical islet isolations ➤ Analysis of the global protein content in the exosomes isolated during isolation reveals a high amount of DAMPs like histone (H2B, H3, and H4) and Chaperons (HSP 70, HSPA1) ➤ The high amount of DAMPs/Islet equivalence (IEQ) released during isolation positively correlated with the recipient's post-transplant insulin requirement [80,81]
9	Heart	LC-MS/MS	<ul style="list-style-type: none"> ➤ Proteomic profiling of serum

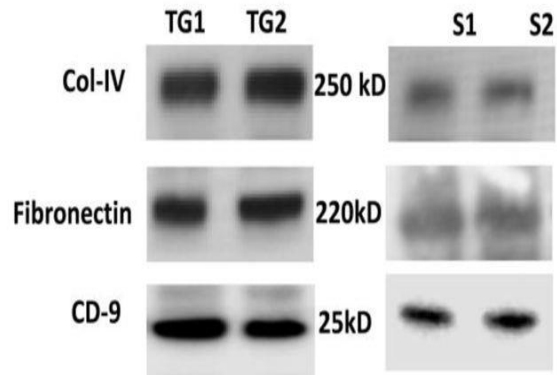
Table 1 (continued)

	Transplant Type	Methods used	The outcome of the study
			<ul style="list-style-type: none"> exosomes of the allograft-rejecting recipients. Further comparative pathway analysis revealed the possible mechanisms regulated by these miRNAs such as Pi3K-AKT, Wnt, endocytosis, focal adhesion, ubiquitin-mediated proteolysis, MAPK kinase, and TGF-β. [18]
12	Lung	Luminex, ELISPOT, and Immunoblotting	<ul style="list-style-type: none"> ➤ The exosomes were isolated from the transplant recipient's serum and BALF ➤ The exosomes isolated from the serum of lung transplant recipients with BOS showed a significantly high amount of SAGs, Col-V, Kα1T, 20s proteasome α3 subunit, along with costimulatory molecules (CD80, CD86, CD40, MHC-II), and transcription factors (NF-kB, HIF-1A, MHC CIITA, IRAK1, and MyD88) [17,119,120]
13	Lung	RNA sequencing	<ul style="list-style-type: none"> ➤ The exosomes were isolated from the transplant recipient's BALF. The Global RNA content was profiled. ➤ About 29 inflammatory and immune-related RNAs were seen significantly high in the exosomes isolated from the BALF of recipients with acute rejection. ➤ These exosomal RNAs could be a possible biomarker for acute rejection [121]

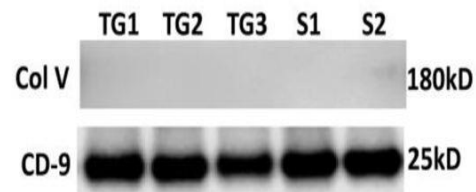
Table 1 (*continued*)

	Transplant Type	Methods used	The outcome of the study
9	Heart	LC-MS/MS	<ul style="list-style-type: none"> ➤ Proteomic profiling of serum exosomes isolated from transplant recipients ➤ About 15 proteins were differentially expressed in the exosomes of the allograft rejected recipients compared to non-rejected recipients. ➤ Within these 15 proteins, 8 proteins were known for their role in immune regulation, complement activation, adaptive immunity, and coagulation. [75]
10	Heart	qPCR array	<ul style="list-style-type: none"> ➤ Serum exosomes were isolated from the transplant recipients and the miRNA present were profiled using qPCR array ➤ The exosomes isolated from the allograft acute rejecting recipients showed enrichment of 4 miRNAs (miR-142, miR-92a, miR-339, and miR-21). ➤ The study has identified a novel mechanism of miR-142, which is transferred to endothelial cells from acute rejecting allograft exosomes. MiR-142 regulates the recipient endothelial function by targeting RAB11FIP2. [27]
11	Lung	qPCR array and immune blotting	<ul style="list-style-type: none"> ➤ The exosomes were isolated from the transplant recipient's serum and bronchoalveolar lavage fluid (BALF) ➤ Donor HLA and lung-associated self-antigens (SAGs), Collagen V (Col-V), and Kα1 tubulin were detected in the exosomes of allograft-rejecting recipients only. ➤ About 123 miRNAs were differentially expressed in the

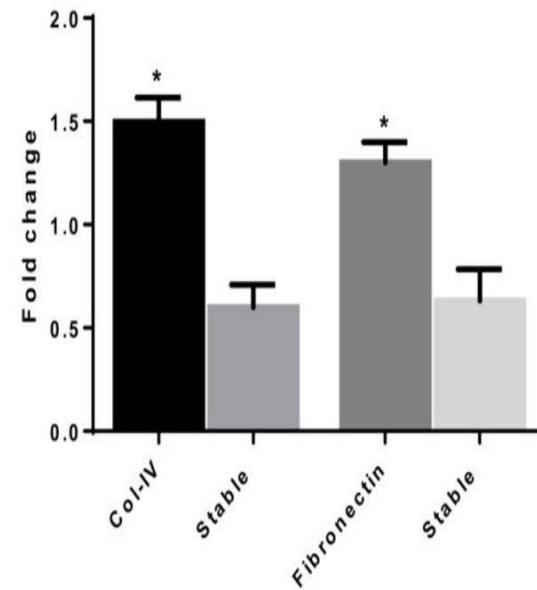
(A)



(C)



(B)



Teşekkürler