

EV-METRIC

EV-TRACK: transparent reporting and centralizing knowledge in extracellular vesicle research

EV-TRACK Consortium*

We argue that the field of extracellular vesicle (EV) biology needs more transparent reporting to facilitate interpretation and replication of experiments. To achieve this, we describe EV-TRACK, a crowdsourcing knowledgebase (<http://evtrack.org>) that centralizes EV biology and methodology with the goal of stimulating authors, reviewers, editors and funders to put experimental guidelines into practice.

2017. 03 .06

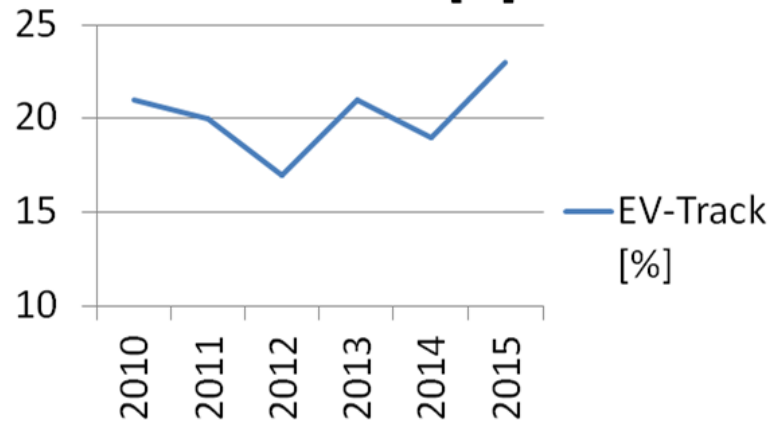




EV-METRIC

- Reported
- Not reported
- Not applicable

EV-Track [%]



All experiments on the same sample type

EV-enriched proteins

Non EV-enriched protein

Qualitative and quantitative analysis

Electron microscopy images

Density gradient

EV density

Ultracentrifugation specifics

Antibody specifics

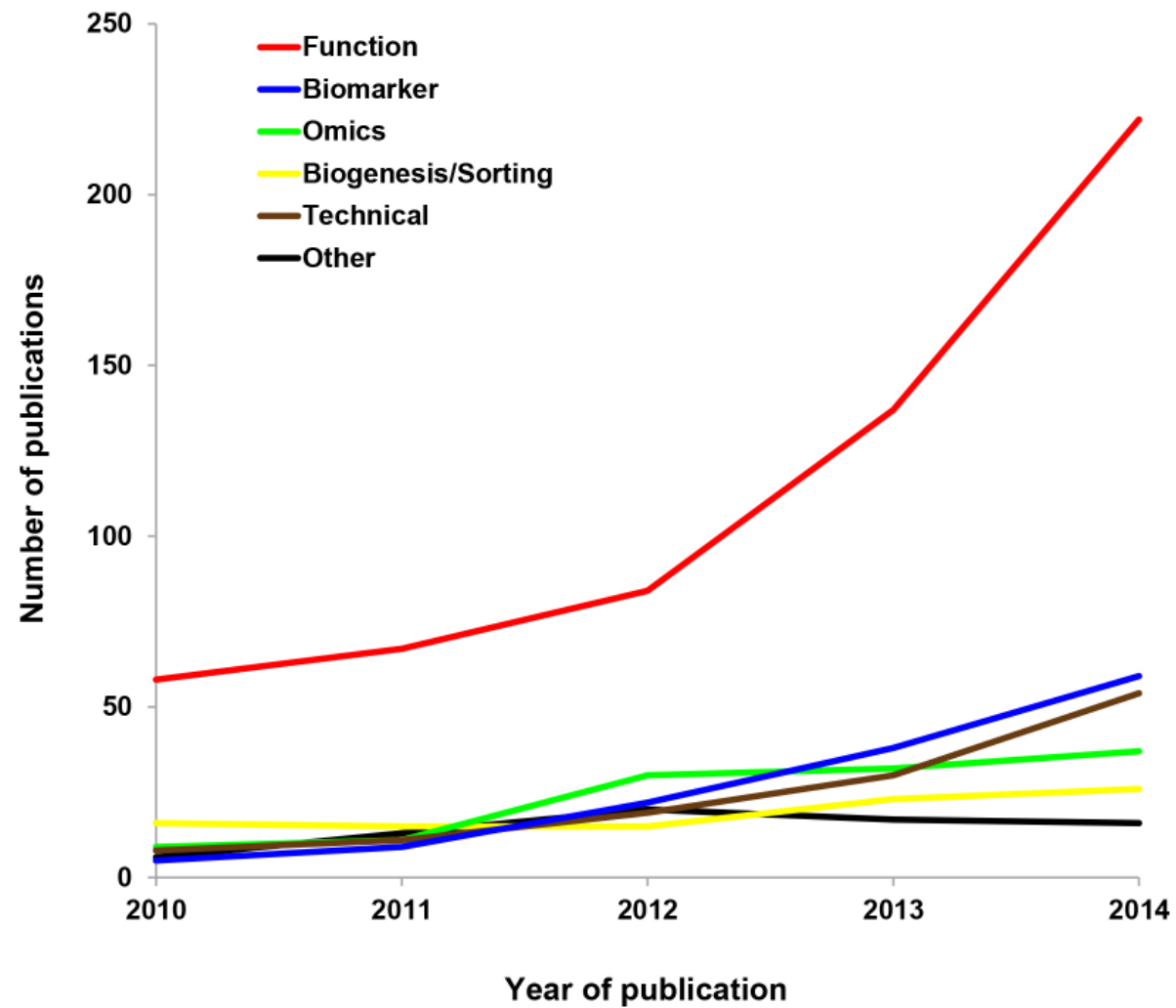
Lysate preparation

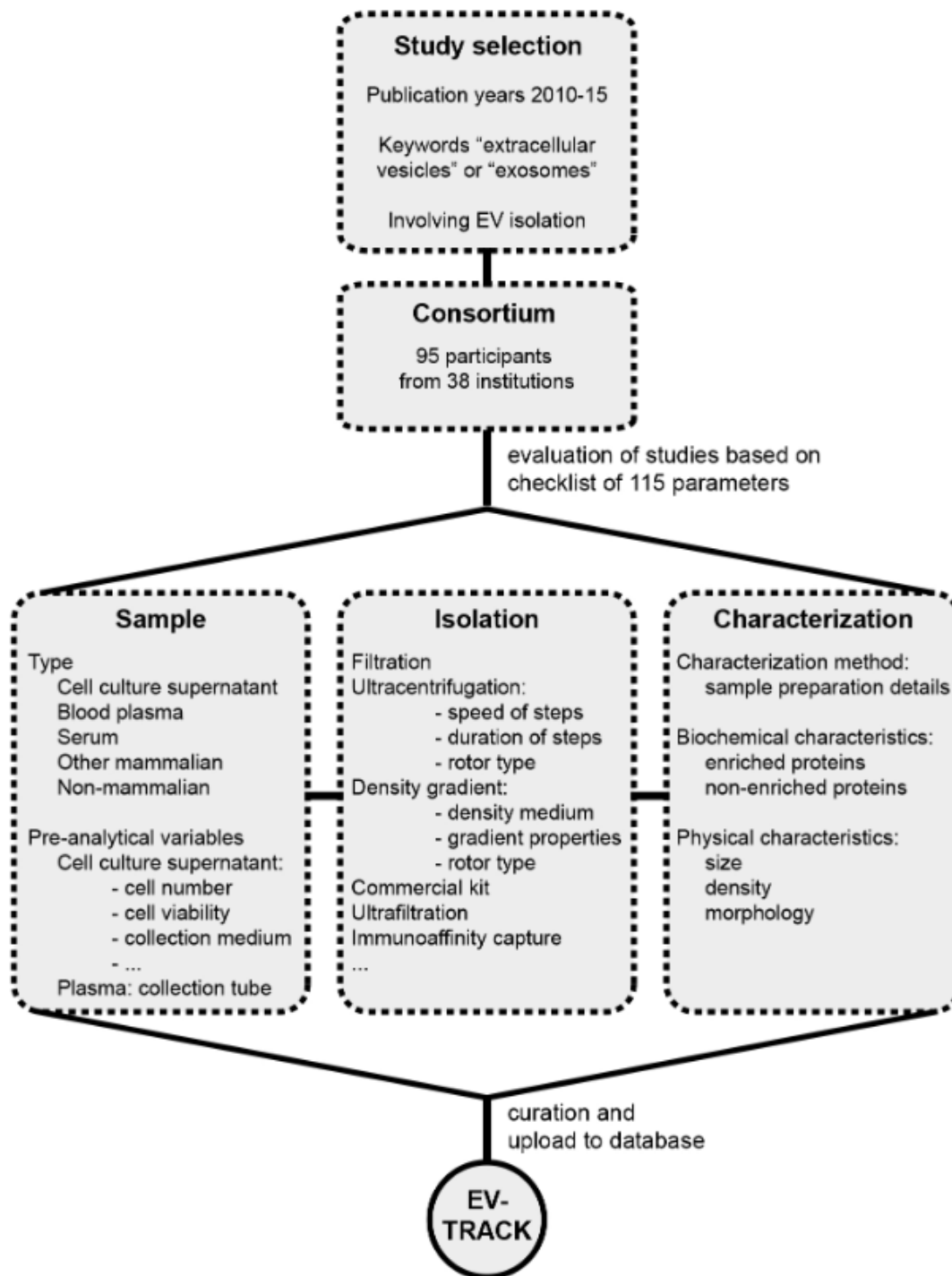


Legnagyobb EV track pontszámú cikk:

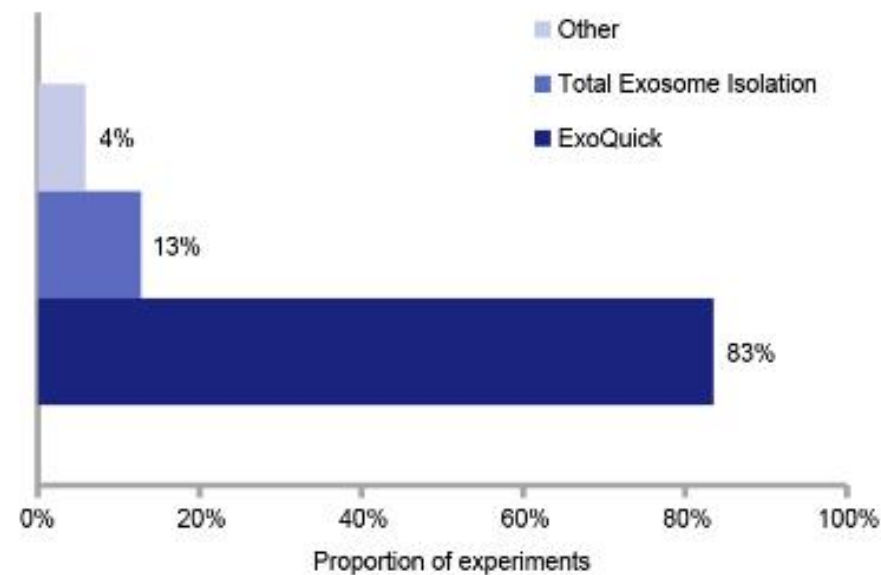
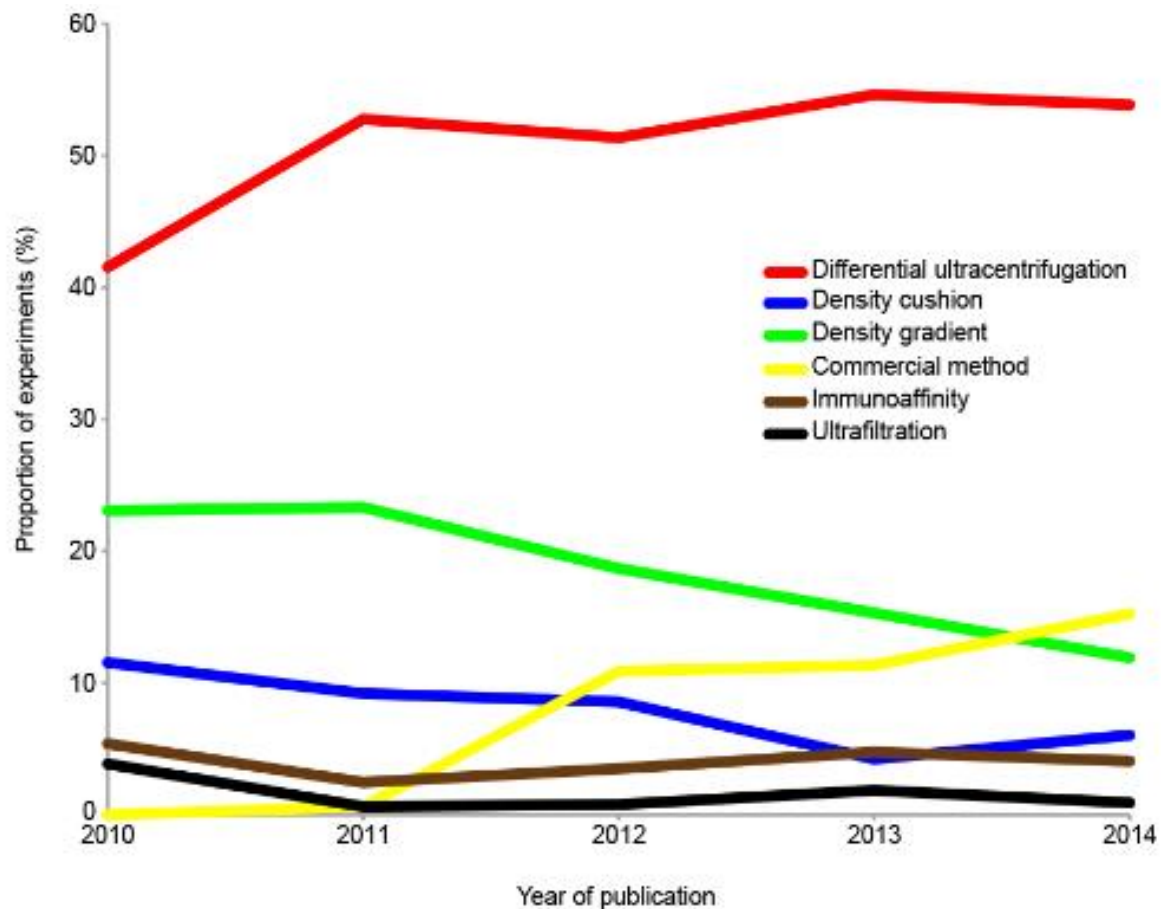
Details	EV-TRACK ID	Experiment nr. ?	Species ?	Sample type	Isolation protocol ?	First author	Year	EV-METRIC ?
+	EV140001	2/4	Homo sapiens	Cell culture supernatant	UF Iodixanol-DG 0.45 µm 0.2 µm filter	Van Deun J	2014	88%

EV tanulmányok célkitűzései

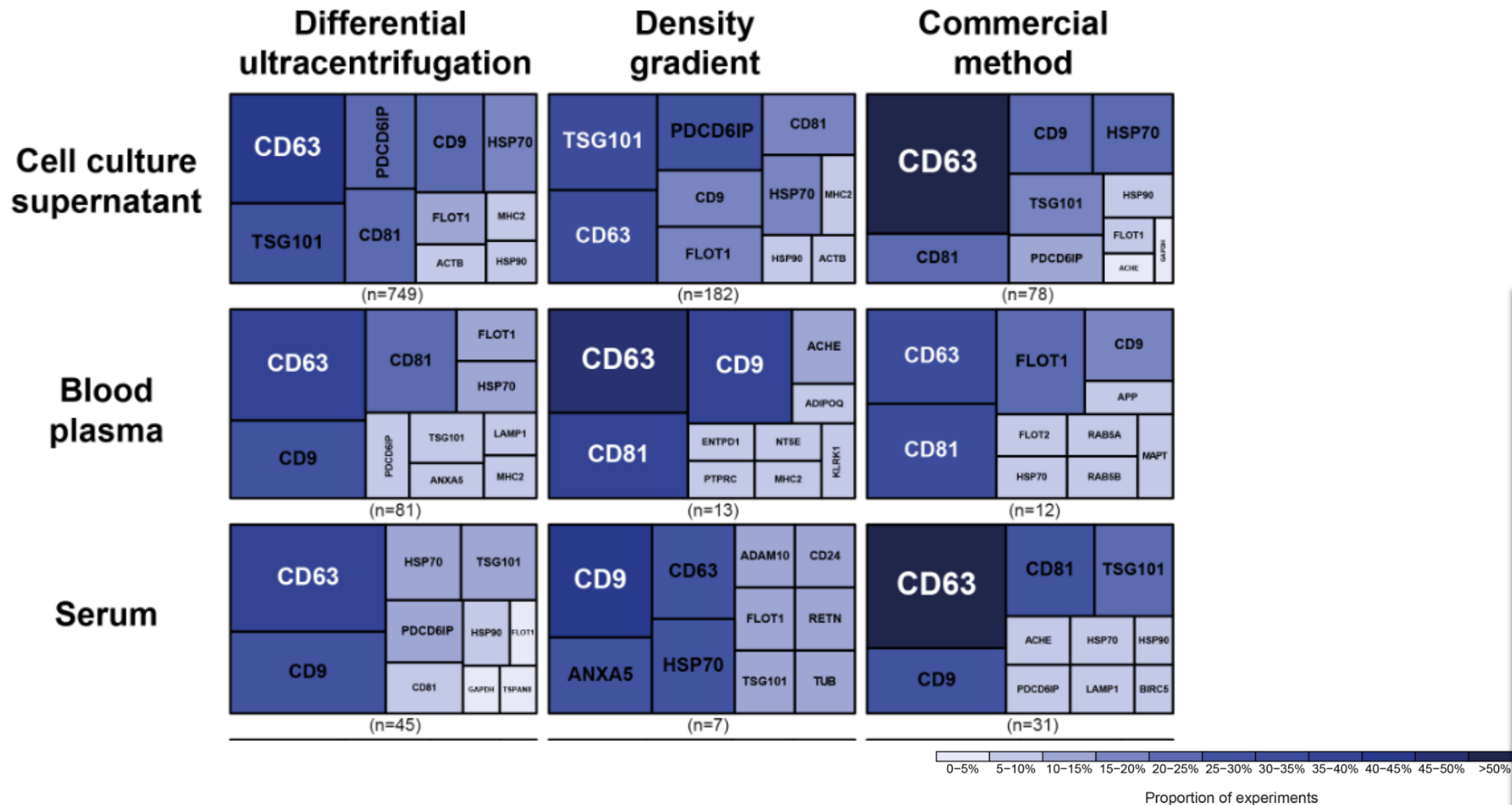




EV izolálási módszerek

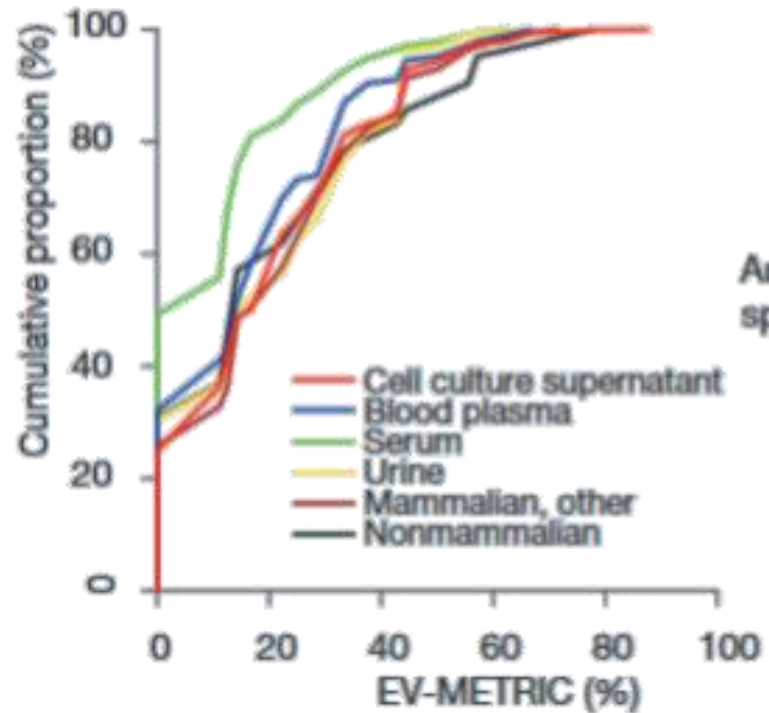


EV izolálási módszerek és jellegzetes EV markerek

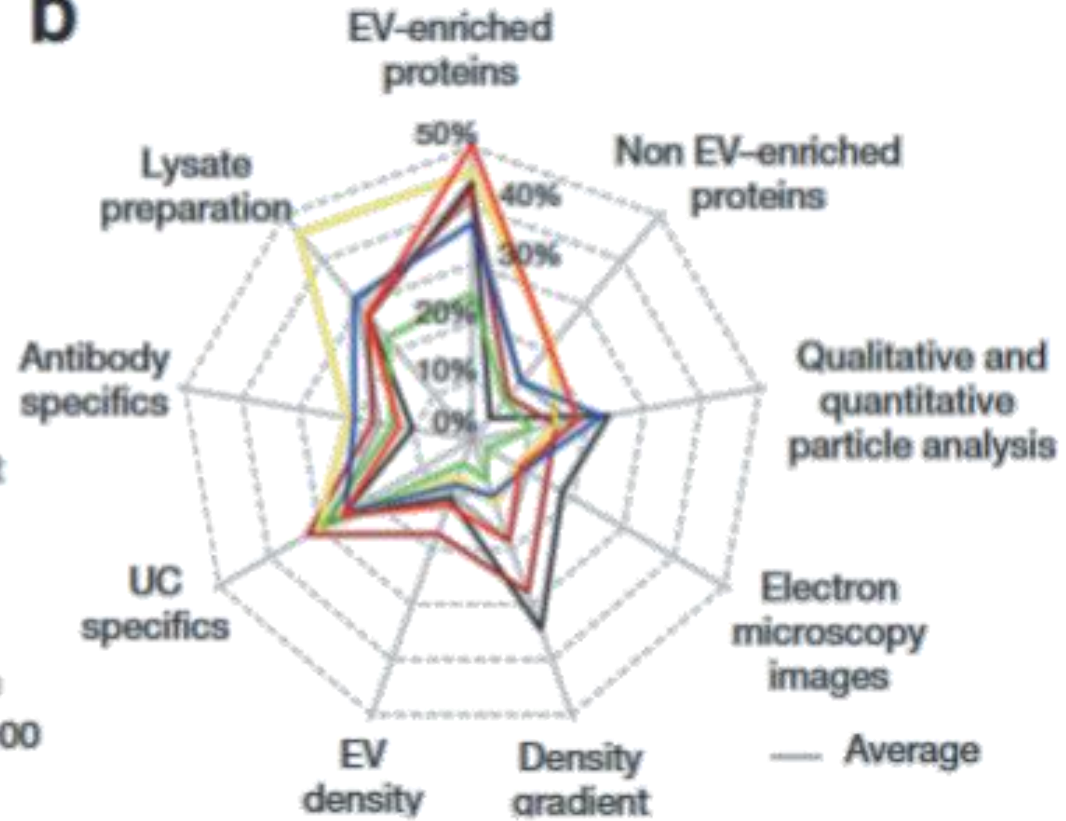


EV jellemzési módszerek = EV-METRIC

a



b





The EV-TRACK platform

Features

- ✓ Prior to submitting a manuscript for peer-review, you are encouraged to upload the methodological specifications that allow inclusion of the assigned EV-TRACK ID in the material and methods section of your manuscript (see [My EV-TRACK](#))
- ✓ A unique identifier (EV-TRACK ID) is assigned to each publication, along with an experiment number
- ✓ EV-TRACK IDs and metrics will be offered when you submit the required specifications along with the associated publications to EV-TRACK (see [My EV-TRACK](#))
- ✓ EV-related experiments can be queried based on different search criteria to discover EV research (see [Search](#))
- ✓ For each study, a summary is included that provides a short overview of the implemented isolation protocol and characteristics of the vesicles that were isolated (terminology, (non-)enriched proteins, size, density)

Most recent

- [Zhu Y](#)
Year: 2015
EV-METRIC: 33%
- [Zhang L](#)
Year: 2015
EV-METRIC: 11%
- [Zhang J](#)
Year: 2015
EV-METRIC: 11%
- [DeClercq V](#)
Year: 2015
EV-METRIC: 33%
- [Yu DD](#)
Year: 2015
EV-METRIC: 0%

SEARCH FOR MORE

Statistics

1226 publications
1741 experiments

NA
0 experiments
Avg. EV-METRIC: 0%

2015
150 experiments
Avg. EV-METRIC: 21%

2014
663 experiments
Avg. EV-METRIC: 20%

2013
379 experiments
Avg. EV-METRIC: 17%

2012
256 experiments
Avg. EV-METRIC: 21%

1,742 experiments that are recorded in EV-TRACK report 190 unique isolation methods and 1,038 unique protocols to retrieve EVs from biofluids

1	EV-TRACK ID	protein analysis: EV enriched	protein analysis: non EV-enriched	complementary particle analysis	electron microscopy images	density gradient	vesicle density	UC specific	antibody specific	lysis buffer	Score
2	EV120025	0	0	0	0	0	0	1	1	0	22%
3	EV120025	1	0	1	0	1	1	NA	1	0	63%
4	EV120025	0	0	0	0	1	1	NA	1	0	38%
5	EV140292	1	0	0	0	0	0	1	0	0	22%
6	EV140292	1	0	0	0	0	0	1	0	0	22%
7	EV130156	0	0	0	0	0	0	0	NA	NA	0%
8	EV140055	1	0	0	0	0	0	0	0	1	22%
9	EV140019	0	0	0	1	0	0	0	1	1	33%
10	EV130077	0	0	0	0	0	0	0	0	0	0%
11	EV130077	1	0	0	0	1	0	NA	0	0	25%
12	EV110030	1	0	0	0	0	0	1	0	0	22%
13	EV140293	1	0	0	0	0	0	0	0	0	11%
14	EV140020	0	0	0	1	0	0	1	1	1	44%
15	EV110082	0	0	0	0	0	0	0	0	0	0%
16	EV130006	0	0	0	1	0	0	1	1	1	44%
17	EV130157	0	0	0	0	0	0	NA	0	0	0%
18	EV130157	0	0	0	0	0	0	0	NA	NA	0%
19	EV140294	0	0	0	0	0	0	NA	0	0	0%
20	EV140295	0	0	0	0	0	0	NA	NA	NA	0%
21	EV100071	1	0	0	0	0	0	0	NA	NA	14%
22	EV120113	0	0	0	0	0	0	0	NA	NA	0%
23	EV110031	0	0	0	0	0	0	0	0	1	11%
24	EV120010	1	0	0	0	0	0	NA	1	1	38%
25	EV120010	1	0	0	0	0	0	0	1	1	33%
26	EV120010	1	0	0	0	0	0	1	1	1	44%
27	EV120010	1	0	0	0	0	0	NA	1	1	38%
28	EV120010	1	0	0	0	0	0	NA	1	1	38%
29	EV120010	1	0	0	0	0	0	1	1	1	44%
30	EV130158	0	0	0	0	0	0	0	0	0	0%
31	EV110012	1	0	1	0	0	0	0	0	0	22%
32	EV110032	0	0	1	0	0	0	0	0	0	11%

Blad1

Blad2

Blad3

EV Track pontozási rendszer (1)

1. Study	
Study number	Specify
First author	Specify
Year of publication	Specify
Number of citations	Specify
2. Journal	
Name	Specify
Impact factor from year of publication	Specify
3. Title	
Exosomes	0 or 1
Microvesicles	0 or 1
Microparticles	0 or 1
Extracellular vesicles	0 or 1
Other vesicle-related term, e.g. Protasome, ...	NA
4. Keywords	
Are keywords available?	0 or 1
Exosomes	0 or 1
Microvesicles	0 or 1
Microparticles	0 or 1
Extracellular vesicles	0 or 1
Other vesicle-related term, e.g. Protasome, ...	NA
5. Study aim	
Choose from drop-down list	-Select-
If "other" was selected: specify	NA
6. Sample	
Cell culture supernatant	
Cell culture supernatant	0 or 1
Serum-free medium	0 or 1
EV-depleted serum containing medium	0 or 1
How was the serum depleted of EVs?	NA
Serum EV depletion protocol	0 or 1
Cell viability	0 or 1
Specification of cell number?	0 or 1

Blood plasma	0 or 1
Collection Tube: EDTA	0 or 1
Collection Tube: Citrate	0 or 1
Collection tube: other	NA
If necessary: select other biofluid from drop-down list	
If "other" was selected: specify	NA
7. Isolation method	
(Differential) (ultra)centrifugation	0 or 1
Density gradient	0 or 1
Commercial	0 or 1
If commercial: specify	Specify
Microfluidics	0 or 1
Other	0 or 1
If other: specify	Specify
Company	NA
Academic: own group or third party paper	NA
If academic reference: specify	NA
Number of papers to get protocol	Specify
Pre-processing of sample by filtration	
0.8 µm	0 or 1
0.45 µm	0 or 1
0.2 or 0.22 µm	0 or 1
0.1 µm	0 or 1
Differential (ultra)centrifugation	
<= 500 g	0 or 1
> 500 g AND < 10,000 g	0 or 1
>= 10,000 g AND < 50,000 g	0 or 1
>= 50,000 g AND < 100,000 g	0 or 1
>= 100,000 g AND < 150,000 g	0 or 1
>= 150,000 g	0 or 1
Duration of final pelleting (before washing step) (min)	Specify
Rotor type	Specify
Swinging bucket or fixed angle rotor?	NA

EV Track pontozási rendszer (2)

Adjusted K-factor	Specify
Wash step included after first-time pelleting?	0 or 1
Wash step volume	Specify
Rotor type for wash step centrifugation	Specify
Adjusted K-factor for wash step centrifugation	Specify
Density gradient centrifugation	
Only as validation/ selected experiments?	0 or 1
Type: choose from drop-down list	NA
Lowest percentage/molarity (upper layer)	Specify
Highest percentage/molarity (lower layer)	Specify
Top-down or bottom-up gradient	NA
Speed of fractions pelleting (in g)	Specify
Wash of fractions included	0 or 1
Volume of wash step	Specify
Rotor type for first-time fractions pelleting	Specify
Swinging bucket or fixed angle rotor?	NA
Adjusted k-factor	Specify
Vesicle density (g/mL)	Specify
Sucrose cushion	
Only as validation/ selected experiments?	0 or 1
Ultrafiltration	
Only as validation/ selected experiments?	0 or 1
Immuno-affinity capture (beads)	
Only as validation/ selected experiments?	0 or 1
For which surface proteins was selected?	Specify
8. Characterisation	
Protein analysis	
Western blot	0 or 1
ELISA	0 or 1
(antibody-coated) beads for flow cytometry	0 or 1
Proteomics	0 or 1
EV enriched proteins	
Alix	0 or 1
CD63	0 or 1
CD81	0 or 1
CD9	0 or 1
Flotillin-1	0 or 1
HSP90	0 or 1

HSP70	0 or 1
Syntenin	0 or 1
TSG101	0 or 1
Other	NA
Total number of EV-enriched proteins checked	Specify
Non-EV enriched proteins	
Cell organelle protein	0 or 1
Albumin	0 or 1
Uromodulin	0 or 1
Other	NA
Antibodies	
Catalogue number/clone specified?	0 or 1
Dilution factor of antibodies specified?	0 or 1
Sample lysate conditions specified?	0 or 1
EV database used for characterization?	0 or 1
Data submitted to database?	0, 1 or NS
Particle analysis (Size and number)	
Dynamic light scattering	0 or 1
High resolution flow cytometry	0 or 1
Nanoparticle tracking analysis	0 or 1
Tunable resistive pulse sensing	0 or 1
Other	NA
Microscopy	
Performed?	0 or 1
Transmission electron microscopy (no immuno)	0 or 1
Immuno-electron microscopy	0 or 1
If immuno: protein marker(s) used	Specify
Cryo-electron microscopy	0 or 1
Scanning electron microscopy	0 or 1
Atomic force microscopy	0 or 1
Other	NA
Large field with multiple vesicles (>=5)	0 or 1
Close-up of single vesicles	0 or 1
Mean vesicle size (+-stdev)	Specify
Remarks (optional)	Specify

k-faktor

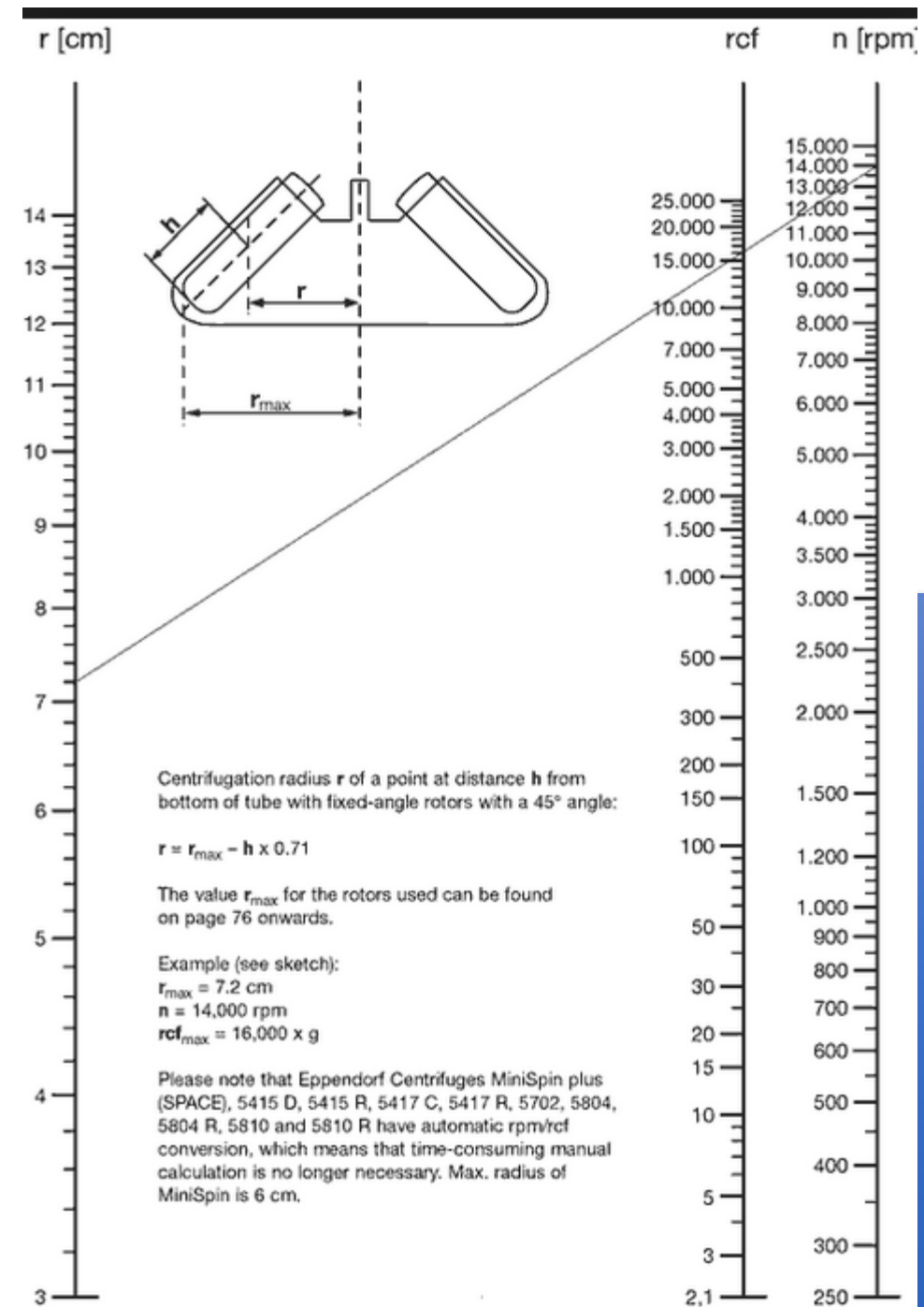
The k-factor or clearance factor of a rotor is a measure of the pelleting efficiency of a rotor that runs at its maximum velocity and is calculated by the equation:

$$k = (2.53 \cdot 10^5 \cdot \ln(r_{\max}/r_{\min})) / (\text{RPM}/100)^2$$

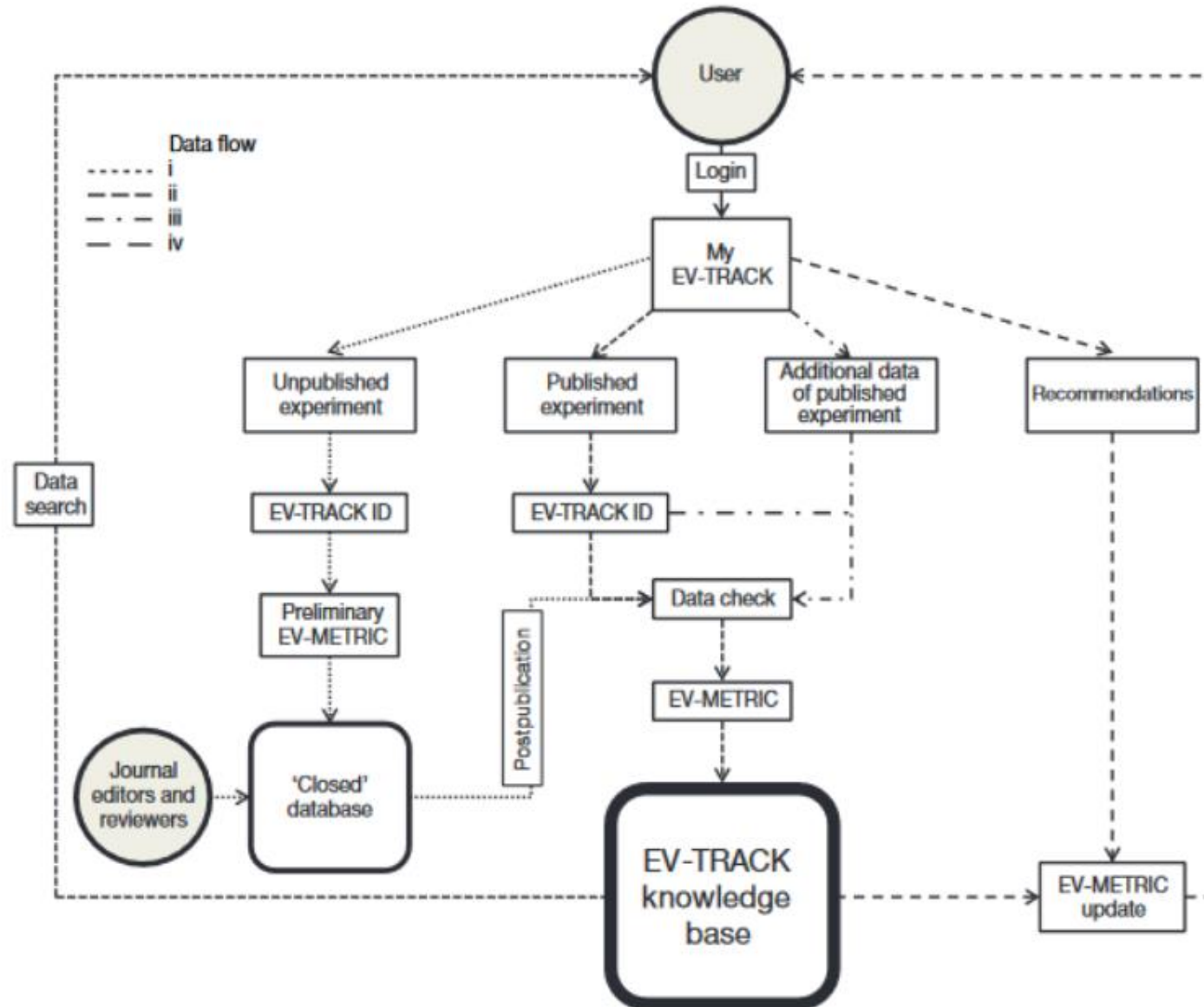
If a rotor is run below its maximum velocity, the following equation should be used to calculate the adjusted k-factor:

$$k_{\text{adj}} = k \cdot (\text{max RPM}/\text{actual RPM})^2$$

(RPM: revolutions per minute; r_{max}/r_{min}: maximum/minimum distance from the rotational axis).



EV TRACK működési algoritmus



Köszönöm szépen a figyelmet!

