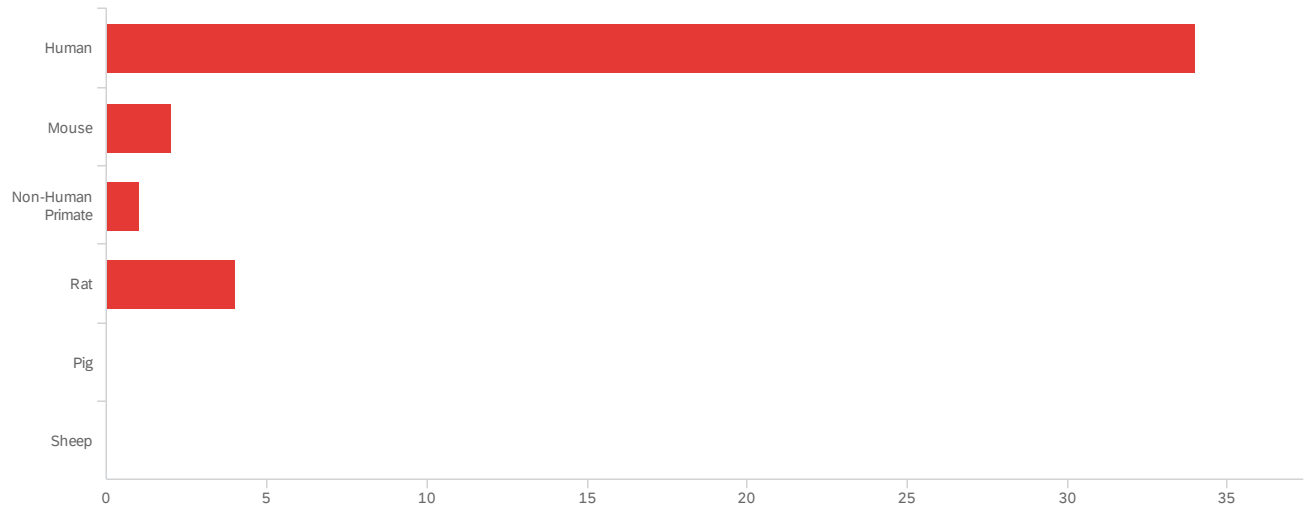


Q1 - Q1. What is the source of your CSF? Please select all that apply.



#	Field	Choice Count
8	Human	82.93% 34
9	Mouse	4.88% 2
10	Non-Human Primate	2.44% 1
11	Rat	9.76% 4
12	Pig	0.00% 0
13	Sheep	0.00% 0

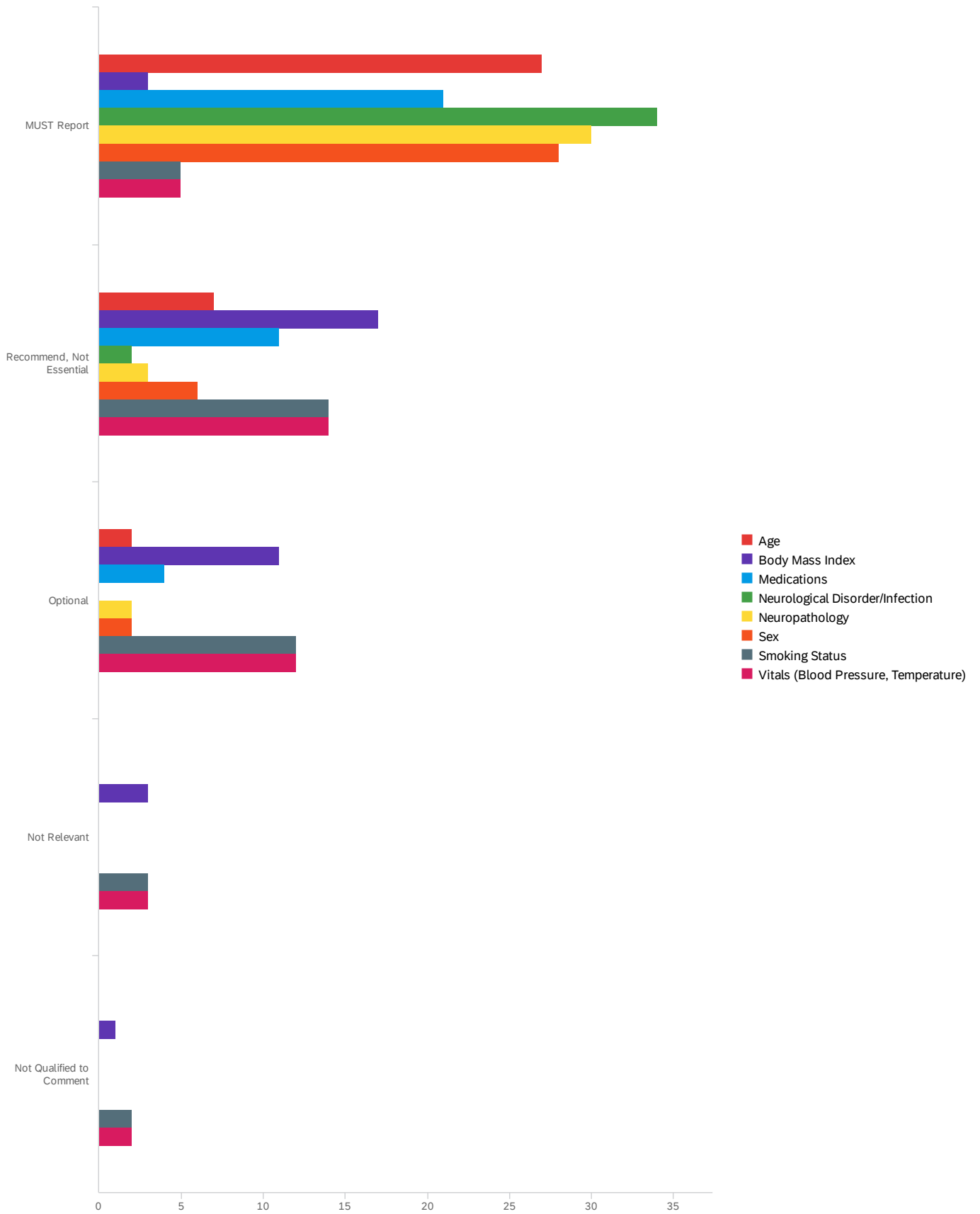
Q1.1 - Q1.1 Other:

Q1.1 Other:

---

Q2#1 - Q2. What parameters might form reporting guidelines for CSF donor information?

Please rate the im... - Click to write Column 1



#	Field	Minimum	Maximum	Mean	Std Deviation	Variance	Count
1	Age	1.00	3.00	1.31	0.57	0.32	36

#	Field	Minimum	Maximum	Mean	Std Deviation	Variance	Count
2	Body Mass Index	1.00	5.00	2.49	0.87	0.76	35
3	Medications	1.00	3.00	1.53	0.69	0.47	36
4	Neurological Disorder/Infection	1.00	2.00	1.06	0.23	0.05	36
5	Neuropathology	1.00	3.00	1.20	0.52	0.27	35
6	Sex	1.00	3.00	1.28	0.56	0.31	36
7	Smoking Status	1.00	5.00	2.53	1.01	1.03	36
8	Vitals (Blood Pressure, Temperature)	1.00	5.00	2.53	1.01	1.03	36

#	Field	MUST Report	Recommend, Not Essential	Optional	Not Relevant	Not Qualified to Comment	Total
1	Age	75.00% 27	19.44% 7	5.56% 2	0.00% 0	0.00% 0	36
2	Body Mass Index	8.57% 3	48.57% 17	31.43% 11	8.57% 3	2.86% 1	35
3	Medications	58.33% 21	30.56% 11	11.11% 4	0.00% 0	0.00% 0	36
4	Neurological Disorder/Infection	94.44% 34	5.56% 2	0.00% 0	0.00% 0	0.00% 0	36
5	Neuropathology	85.71% 30	8.57% 3	5.71% 2	0.00% 0	0.00% 0	35
6	Sex	77.78% 28	16.67% 6	5.56% 2	0.00% 0	0.00% 0	36
7	Smoking Status	13.89% 5	38.89% 14	33.33% 12	8.33% 3	5.56% 2	36
8	Vitals (Blood Pressure, Temperature)	13.89% 5	38.89% 14	33.33% 12	8.33% 3	5.56% 2	36

Showing rows 1 - 8 of 8

## Q2.1 - Q2.1 Other:

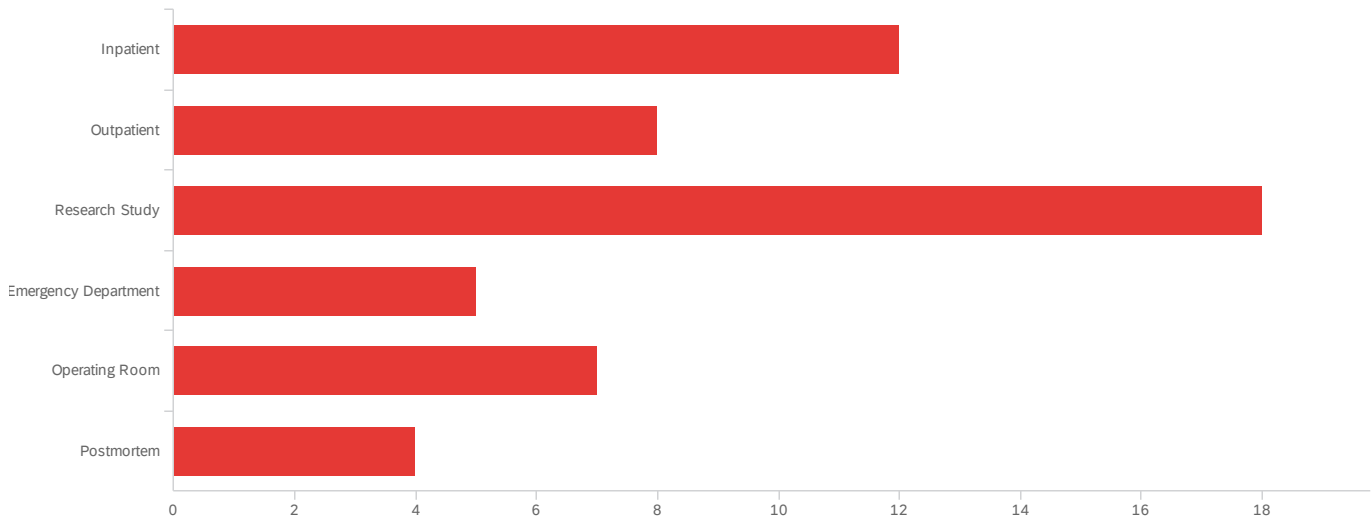
Q2.1 Other:

---

Unfortunately, rankings on your list require context. For many of them, ranking will depend on the targets and the pathological state of interest.

Cardiovascular disease, Inflammatory disorder

Q3 - Q3. What is the setting for your CSF collection? Please select all that apply.



#	Field	Choice Count
1	Inpatient	22.22% 12
2	Outpatient	14.81% 8
3	Research Study	33.33% 18
4	Emergency Department	9.26% 5
6	Operating Room	12.96% 7
7	Postmortem	7.41% 4

54

Showing rows 1 - 7 of 7

## Q3.1 - Q3.1 Other:

Q3.1 Other:

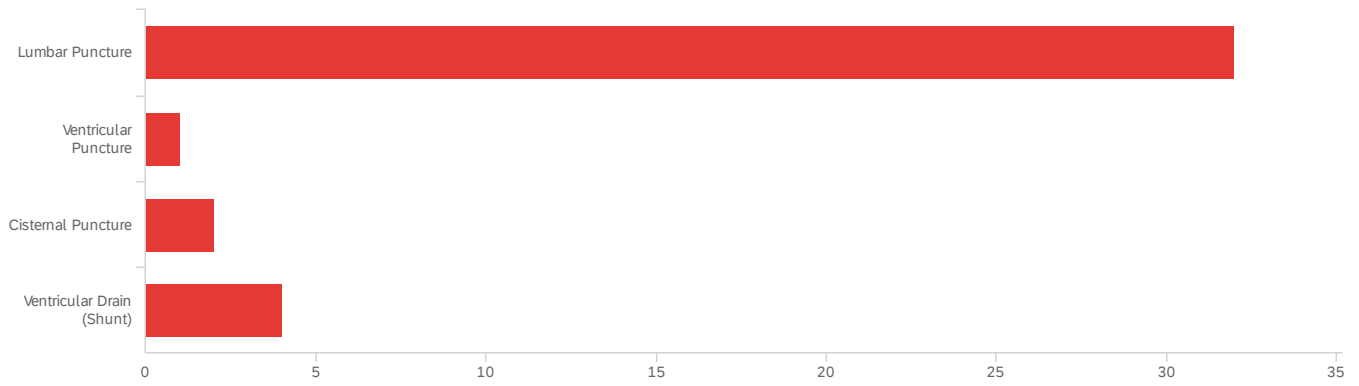
---

in vivo

Postmortem CSF is a strange fluid and should not be used in my opinion.



Q4 - Q4. What is the CSF sampling site? Please select all that apply.



#	Field	Choice Count
1	Lumbar Puncture	82.05% 32
2	Ventricular Puncture	2.56% 1
3	Cisternal Puncture	5.13% 2
6	Ventricular Drain (Shunt)	10.26% 4

39

Showing rows 1 - 5 of 5

## Q4.1 - Q4.1 Other:

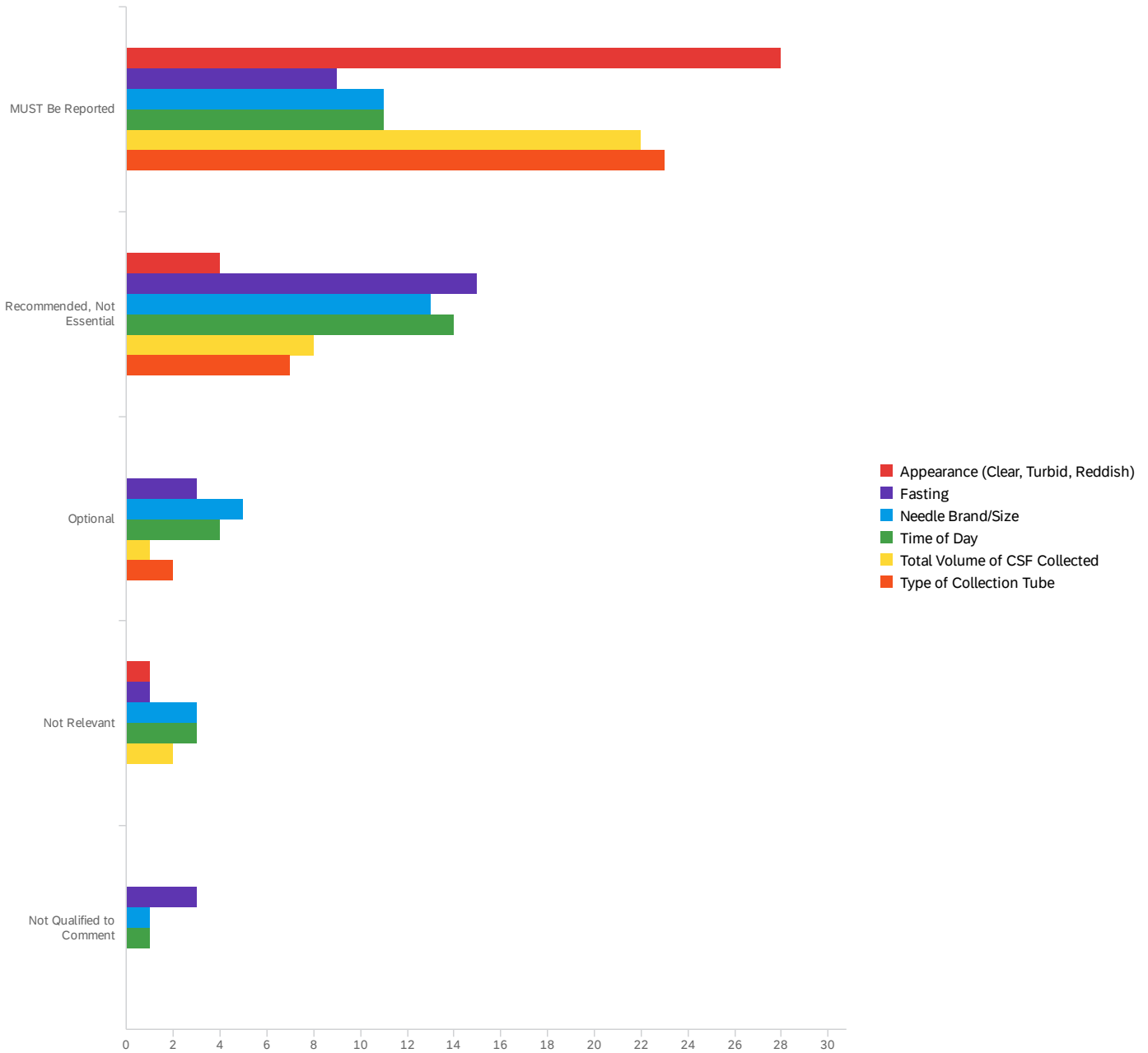
Q4.1 Other:

---

brain -pre-operative sampling after dura opened

Q5#1 - Q5. What parameters might form reporting guidelines for CSF collection? Please

rate the importanc... - Click to write Column 1



#	Field	Minimum	Maximum	Mean	Std Deviation	Variance	Count
1	Appearance (Clear, Turbid, Reddish)	1.00	4.00	1.21	0.59	0.35	33
2	Fasting	1.00	5.00	2.16	1.17	1.36	31
3	Needle Brand/Size	1.00	5.00	2.09	1.05	1.11	33

#	Field	Minimum	Maximum	Mean	Std Deviation	Variance	Count
4	Time of Day	1.00	5.00	2.06	1.04	1.09	33
5	Total Volume of CSF Collected	1.00	4.00	1.48	0.82	0.67	33
6	Type of Collection Tube	1.00	3.00	1.34	0.59	0.35	32

#	Field	MUST Be Reported	Recommended, Not Essential	Optional	Not Relevant	Not Qualified to Comment	Total
1	Appearance (Clear, Turbid, Reddish)	84.85% 28	12.12% 4	0.00% 0	3.03% 1	0.00% 0	33
2	Fasting	29.03% 9	48.39% 15	9.68% 3	3.23% 1	9.68% 3	31
3	Needle Brand/Size	33.33% 11	39.39% 13	15.15% 5	9.09% 3	3.03% 1	33
4	Time of Day	33.33% 11	42.42% 14	12.12% 4	9.09% 3	3.03% 1	33
5	Total Volume of CSF Collected	66.67% 22	24.24% 8	3.03% 1	6.06% 2	0.00% 0	33
6	Type of Collection Tube	71.88% 23	21.88% 7	6.25% 2	0.00% 0	0.00% 0	32

Showing rows 1 - 6 of 6

## Q5.1 - Q5.1 Other:

Q5.1 Other:

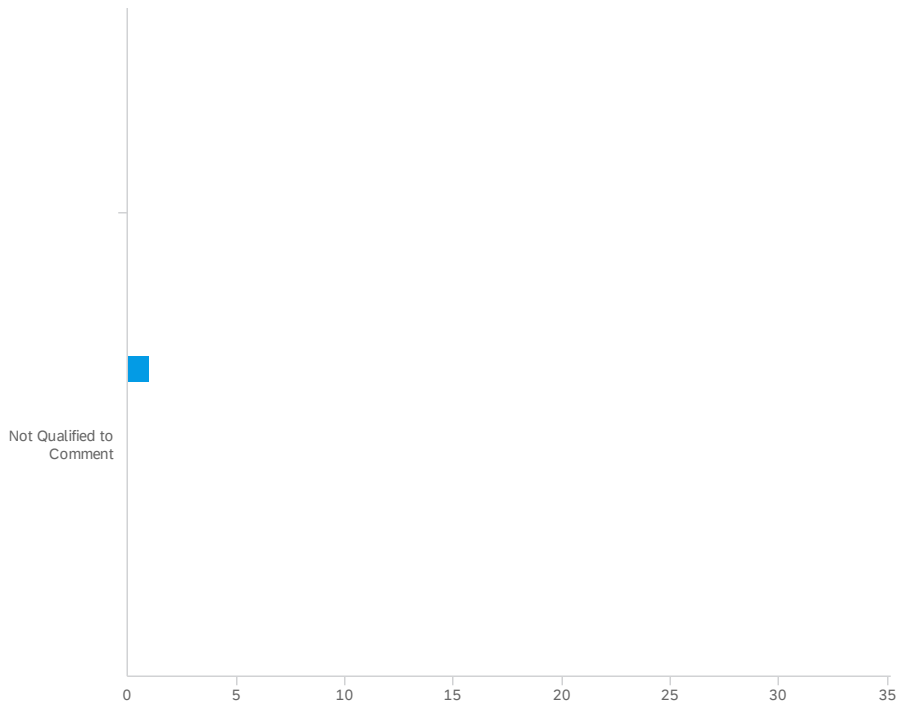
---

It would be ideal to assess parameters by some means beyond visual examination. Like spectroscopy for presence of hemoglobin or measuring specific gravity or protein content.

Anesthesia

When was the CSF sample collected (i.e. first sample after puncture, last sample after puncture) when sample is taken via Lumbar puncture





#	Field	Minimum	Maximum	Mean	Std Deviation	Variance	Count
1	Centrifugation Speed & Time	1.00	2.00	1.03	0.17	0.03	33
2	CSF Additives (Preservatives, Protease Inhibitors)	1.00	2.00	1.06	0.24	0.06	33
3	CSF Pooled Before Aliquoting (Gradient Effect)	1.00	5.00	1.30	0.80	0.64	33
4	Freezing Method	1.00	3.00	1.21	0.48	0.23	33
5	Freeze/Thaw Cycles	1.00	3.00	1.18	0.52	0.27	33
6	Storage Tube Type	1.00	3.00	1.72	0.67	0.45	32
7	Temperature Prior to Freezing	1.00	3.00	1.42	0.61	0.37	31
8	Temperature for Storage	1.00	2.00	1.12	0.33	0.11	33
9	Thawing Methodology	1.00	3.00	1.36	0.54	0.29	33
10	Time Between Draw and Storage	1.00	3.00	1.55	0.70	0.49	33
11	Years in Storage	1.00	3.00	1.52	0.56	0.31	33

#	Field	MUST Be Reported	Recommended, Not Essential	Optional	Not Relevant	Not Qualified to Comment	Total
1	Centrifugation Speed & Time	96.97% 32	3.03% 1	0.00% 0	0.00% 0	0.00% 0	33

#	Field	MUST Be Reported	Recommended, Not Essential	Optional	Not Relevant	Not Qualified to Comment	Total
2	CSF Additives (Preservatives, Protease Inhibitors)	93.94% 31	6.06% 2	0.00% 0	0.00% 0	0.00% 0	33
3	CSF Pooled Before Aliquoting (Gradient Effect)	81.82% 27	12.12% 4	3.03% 1	0.00% 0	3.03% 1	33
4	Freezing Method	81.82% 27	15.15% 5	3.03% 1	0.00% 0	0.00% 0	33
5	Freeze/Thaw Cycles	87.88% 29	6.06% 2	6.06% 2	0.00% 0	0.00% 0	33
6	Storage Tube Type	40.63% 13	46.88% 15	12.50% 4	0.00% 0	0.00% 0	32
7	Temperature Prior to Freezing	64.52% 20	29.03% 9	6.45% 2	0.00% 0	0.00% 0	31
8	Temperature for Storage	87.88% 29	12.12% 4	0.00% 0	0.00% 0	0.00% 0	33
9	Thawing Methodology	66.67% 22	30.30% 10	3.03% 1	0.00% 0	0.00% 0	33
10	Time Between Draw and Storage	57.58% 19	30.30% 10	12.12% 4	0.00% 0	0.00% 0	33
11	Years in Storage	51.52% 17	45.45% 15	3.03% 1	0.00% 0	0.00% 0	33

Showing rows 1 - 11 of 11



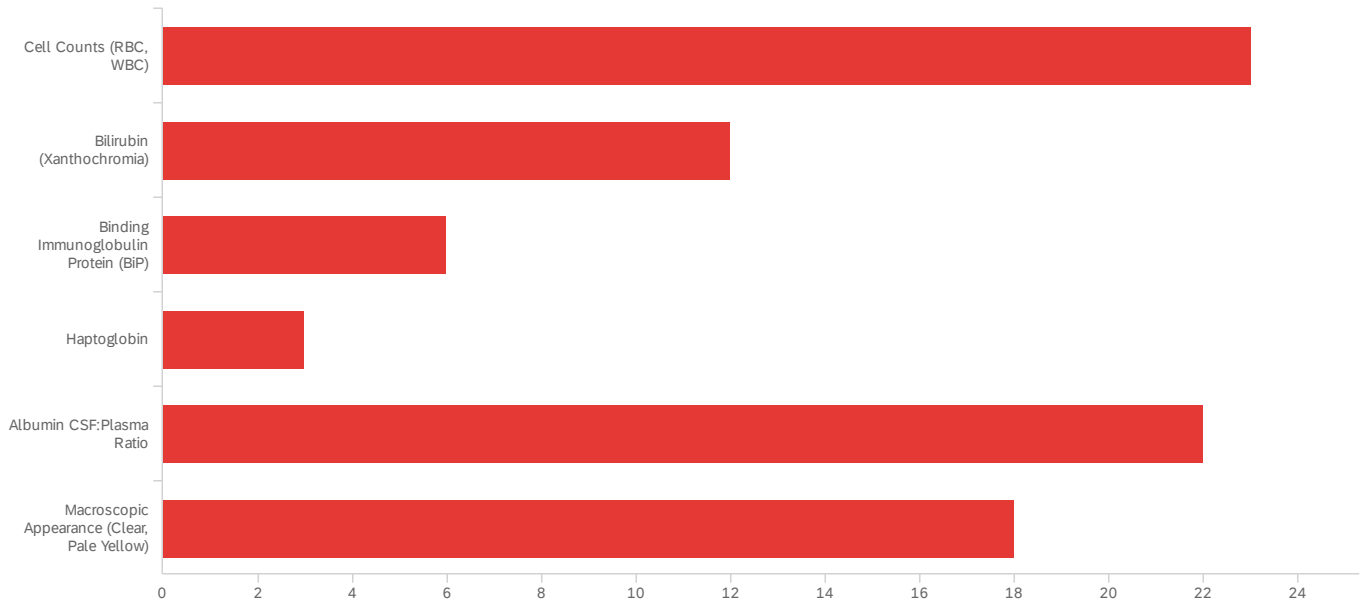
## Q6.1 - Q6.1 Other:

Q6.1 Other:

---

more information is always better

Q7 - Q7. What markers should be considered as CSF contaminants? Please select all that apply.



#	Field	Choice Count
1	Cell Counts (RBC, WBC)	27.38% 23
2	Bilirubin (Xanthochromia)	14.29% 12
3	Binding Immunoglobulin Protein (BiP)	7.14% 6
4	Haptoglobin	3.57% 3
5	Albumin CSF:Plasma Ratio	26.19% 22
7	Macroscopic Appearance (Clear, Pale Yellow)	21.43% 18

84

Showing rows 1 - 7 of 7

## Q7.1 - Q7.1 Other:

Q7.1 Other:

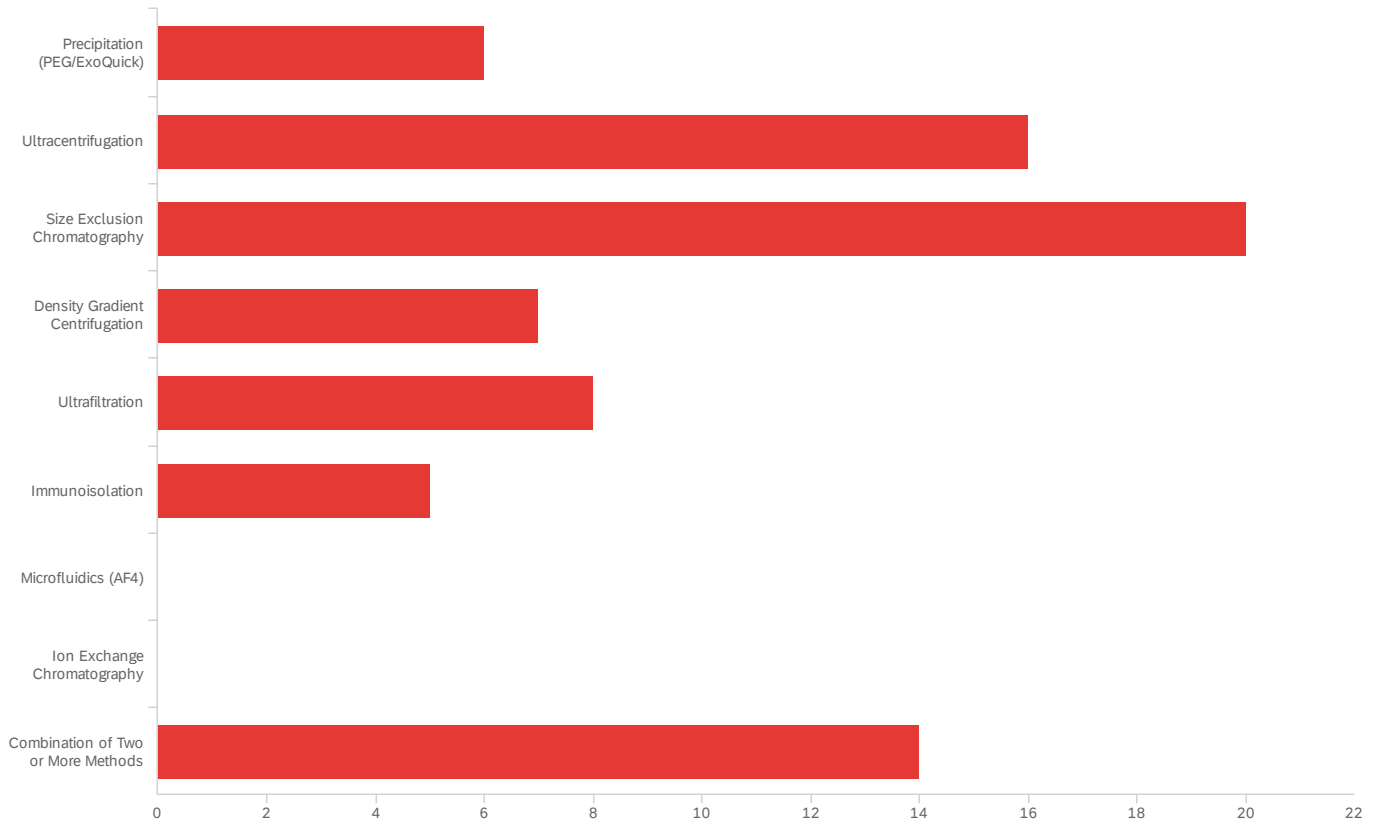
---

I am not sure how to answer this question, since I was unsure if this meant contaminants of CSF (stuff that isn't supposed to be there) or contaminants of EV preps.

Hemoglobin

Q8 - Q8. What method(s) do you use to fractionate/concentrate your CSF EVs? Please

select all that apply.



#	Field	Choice Count
1	Precipitation (PEG/ExoQuick)	7.89% 6
2	Ultracentrifugation	21.05% 16
3	Size Exclusion Chromatography	26.32% 20
4	Density Gradient Centrifugation	9.21% 7
5	Ultrafiltration	10.53% 8
6	Immunoisolation	6.58% 5
7	Microfluidics (AF4)	0.00% 0
8	Ion Exchange Chromatography	0.00% 0
9	Combination of Two or More Methods	18.42% 14
		76

## Q8.1 - Q8.1 Other:

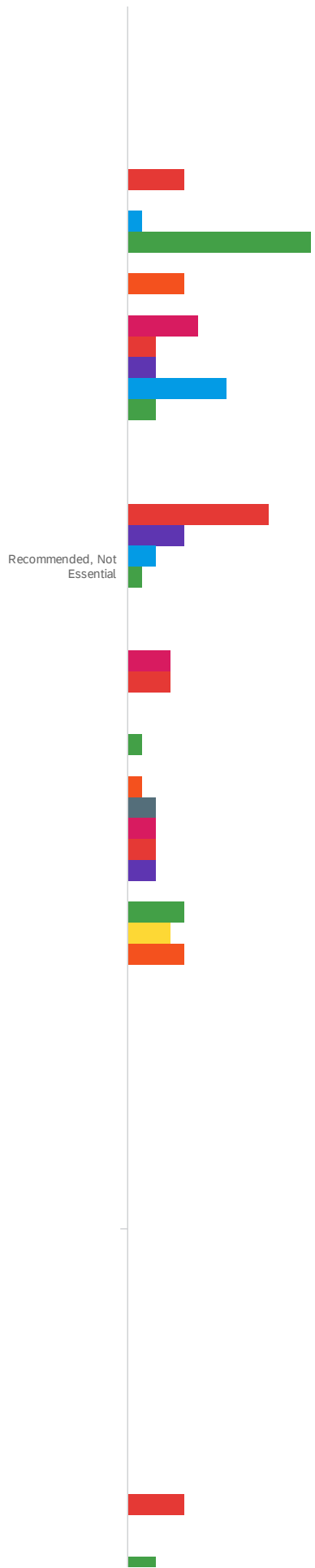
Q8.1 Other:

---

size exclusion chromatography followed by ultrafiltration to concentrate EV fractions

rotovap





- Centrifuge Model & Rotor
- Speed (x g)
- Temperature
- Tube Type



- Time
- Centrifuge Model & Rotor
- Densities/Number of Layers
- Fraction Collection (Visible Bands, Sequential Fraction Removal)
- Fractions Containing EVs
- Medium (Iodixanol, Sucrose)
- Pooling Strategy
- Post-Gradient Processing (Dialysis, Ultracentrifugation, Ultrafiltration, Pr...
- Sample Loading (Top, Bottom)
- Speed (x g)
- Temperature
- Time
- Tube Type
- Concentration (40x, 100x)
- Membrane Material
- Molecular Weight Cut-Off
- Capture Molecule Information (Source, Resin Type, Pore Size)
- Elution Buffer Components & Volume
- Elution Steps
- Sample Volume & Dilution
- Wash Buffer & Number of Washes
- Column Information (Source, Pre-Pack, Resin Type, Pore Size)
- Elution Buffer Components & Volume
- Number & Volume of Collected Fractions
- Reporting Which Fractions Contain EVs
- Pooling Strategy
- Sample Dilution & Volume
- Column Information (Source, Pre-Pack, Resin Type, Pore Size)
- Elution Buffer Components & Volume
- Number & Volume of Collected Fractions
- Reporting Which Fractions Contain EVs
- Pooling Strategy
- Sample Dilution & Volume
- Concentration (40x, 100x)

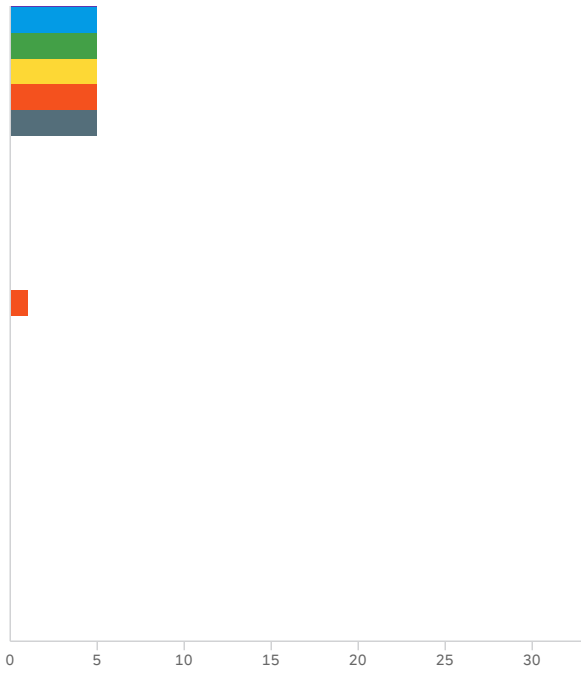


Not Relevent



Not Qualified to  
Comment





#	Field	Minimum	Maximum	Mean	Std Deviation	Variance	Count
1	Centrifuge Model & Rotor	3.00	7.00	3.53	0.96	0.92	30
2	Speed (x g)	3.00	7.00	3.13	0.71	0.50	31
3	Temperature	3.00	7.00	3.16	0.72	0.52	31
4	Tube Type	3.00	7.00	3.68	0.86	0.73	31
5	Time	3.00	7.00	3.13	0.71	0.50	31
6	Centrifuge Model & Rotor	3.00	7.00	3.85	1.46	2.13	27
7	Densities/Number of Layers	3.00	7.00	3.59	1.42	2.02	27
8	Fraction Collection (Visible Bands, Sequential Fraction Removal)	3.00	7.00	3.81	1.41	2.00	26
9	Fractions Containing EVs	3.00	7.00	3.69	1.43	2.06	26
10	Medium (Iodixanol, Sucrose)	3.00	7.00	3.69	1.43	2.06	26
11	Pooling Strategy	3.00	7.00	4.00	1.41	2.00	25
12	Post-Gradient Processing (Dialysis, Ultracentrifugation, Ultrafiltration, Precipitation)	3.00	7.00	3.74	1.43	2.04	27
13	Sample Loading (Top, Bottom)	3.00	7.00	3.81	1.47	2.15	27
14	Speed (x g)	3.00	7.00	3.59	1.42	2.02	27

#	Field	Minimum	Maximum	Mean	Std Deviation	Variance	Count
15	Temperature	3.00	7.00	3.59	1.42	2.02	27
16	Time	3.00	7.00	3.59	1.42	2.02	27
17	Tube Type	3.00	7.00	4.15	1.38	1.90	27
38	Concentration (40x, 100x)	3.00	7.00	3.34	0.84	0.71	29
19	Membrane Material	3.00	7.00	3.46	1.12	1.25	26
20	Molecular Weight Cut-Off	3.00	7.00	3.35	1.07	1.15	26
21	Capture Molecule Information (Source, Resin Type, Pore Size)	3.00	7.00	3.46	1.28	1.63	26
22	Elution Buffer Components & Volume	3.00	7.00	3.48	1.30	1.69	25
23	Elution Steps	3.00	7.00	3.56	1.33	1.77	25
24	Sample Volume & Dilution	3.00	7.00	3.68	1.32	1.74	25
25	Wash Buffer & Number of Washes	3.00	7.00	3.68	1.32	1.74	25
26	Column Information (Source, Pre-Pack, Resin Type, Pore Size)	3.00	7.00	3.83	1.62	2.64	24
27	Elution Buffer Components & Volume	3.00	7.00	3.83	1.62	2.64	24
28	Number & Volume of Collected Fractions	3.00	7.00	3.96	1.62	2.62	24
29	Reporting Which Fractions Contain EVs	3.00	7.00	3.92	1.63	2.66	24
30	Pooling Strategy	3.00	7.00	3.96	1.62	2.62	24
31	Sample Dilution & Volume	3.00	7.00	4.00	1.61	2.58	24
32	Column Information (Source, Pre-Pack, Resin Type, Pore Size)	3.00	4.00	3.07	0.26	0.07	27
33	Elution Buffer Components & Volume	3.00	4.00	3.07	0.26	0.07	27
34	Number & Volume of Collected Fractions	3.00	4.00	3.07	0.26	0.07	27
35	Reporting Which Fractions Contain EVs	3.00	3.00	3.00	0.00	0.00	27
36	Pooling Strategy	3.00	5.00	3.21	0.49	0.24	28
37	Sample Dilution & Volume	3.00	5.00	3.18	0.47	0.22	28
38	Concentration (40x, 100x)	3.00	7.00	3.34	0.84	0.71	29

#	Field	MUST be Reported		Recommended, Not Essential		Optional		Not Relevant		Not Qualified to Comment		Total
1	Centrifuge Model & Rotor	70.00%	21	13.33%	4	13.33%	4	0.00%	0	3.33%	1	30
2	Speed (x g)	96.77%	30	0.00%	0	0.00%	0	0.00%	0	3.23%	1	31
3	Temperature	93.55%	29	3.23%	1	0.00%	0	0.00%	0	3.23%	1	31
4	Tube Type	48.39%	15	41.94%	13	6.45%	2	0.00%	0	3.23%	1	31
5	Time	96.77%	30	0.00%	0	0.00%	0	0.00%	0	3.23%	1	31
6	Centrifuge Model & Rotor	66.67%	18	14.81%	4	0.00%	0	3.70%	1	14.81%	4	27
7	Densities/Number of Layers	85.19%	23	0.00%	0	0.00%	0	0.00%	0	14.81%	4	27
8	Fraction Collection (Visible Bands, Sequential Fraction Removal)	65.38%	17	19.23%	5	0.00%	0	0.00%	0	15.38%	4	26
9	Fractions Containing EVs	76.92%	20	7.69%	2	0.00%	0	0.00%	0	15.38%	4	26
10	Medium (Iodixanol, Sucrose)	76.92%	20	7.69%	2	0.00%	0	0.00%	0	15.38%	4	26
11	Pooling Strategy	52.00%	13	28.00%	7	4.00%	1	0.00%	0	16.00%	4	25
12	Post-Gradient Processing (Dialysis, Ultracentrifugation, Ultrafiltration, Precipitation)	74.07%	20	7.41%	2	3.70%	1	0.00%	0	14.81%	4	27
13	Sample Loading (Top, Bottom)	74.07%	20	0.00%	0	11.11%	3	0.00%	0	14.81%	4	27
14	Speed (x g)	85.19%	23	0.00%	0	0.00%	0	0.00%	0	14.81%	4	27
15	Temperature	85.19%	23	0.00%	0	0.00%	0	0.00%	0	14.81%	4	27
16	Time	85.19%	23	0.00%	0	0.00%	0	0.00%	0	14.81%	4	27
17	Tube Type	40.74%	11	37.04%	10	3.70%	1	3.70%	1	14.81%	4	27
18	Concentration (40x, 100x)	79.31%	23	13.79%	4	3.45%	1	0.00%	0	3.45%	1	29
19	Membrane Material	80.77%	21	7.69%	2	3.85%	1	0.00%	0	7.69%	2	26
20	Molecular Weight Cut-Off	88.46%	23	3.85%	1	0.00%	0	0.00%	0	7.69%	2	26
21	Capture Molecule Information (Source, Resin Type, Pore Size)	88.46%	23	0.00%	0	0.00%	0	0.00%	0	11.54%	3	26
22	Elution Buffer Components & Volume	88.00%	22	0.00%	0	0.00%	0	0.00%	0	12.00%	3	25
23	Elution Steps	84.00%	21	0.00%	0	4.00%	1	0.00%	0	12.00%	3	25
24	Sample Volume & Dilution	72.00%	18	12.00%	3	4.00%	1	0.00%	0	12.00%	3	25
25	Wash Buffer & Number of Washes	72.00%	18	12.00%	3	4.00%	1	0.00%	0	12.00%	3	25

#	Field	MUST be Reported		Recommended, Not Essential		Optional		Not Relevent		Not Qualified to Comment		Total
26	Column Information (Source, Pre-Pack, Resin Type, Pore Size)	79.17%	19	0.00%	0	0.00%	0	0.00%	0	20.83%	5	24
27	Elution Buffer Components & Volume	79.17%	19	0.00%	0	0.00%	0	0.00%	0	20.83%	5	24
28	Number & Volume of Collected Fractions	70.83%	17	4.17%	1	4.17%	1	0.00%	0	20.83%	5	24
29	Reporting Which Fractions Contain EVs	75.00%	18	0.00%	0	4.17%	1	0.00%	0	20.83%	5	24
30	Pooling Strategy	70.83%	17	4.17%	1	4.17%	1	0.00%	0	20.83%	5	24
31	Sample Dilution & Volume	66.67%	16	8.33%	2	4.17%	1	0.00%	0	20.83%	5	24
32	Column Information (Source, Pre-Pack, Resin Type, Pore Size)	92.59%	25	7.41%	2	0.00%	0	0.00%	0	0.00%	0	27
33	Elution Buffer Components & Volume	92.59%	25	7.41%	2	0.00%	0	0.00%	0	0.00%	0	27
34	Number & Volume of Collected Fractions	92.59%	25	7.41%	2	0.00%	0	0.00%	0	0.00%	0	27
35	Reporting Which Fractions Contain EVs	100.00%	27	0.00%	0	0.00%	0	0.00%	0	0.00%	0	27
36	Pooling Strategy	82.14%	23	14.29%	4	3.57%	1	0.00%	0	0.00%	0	28
37	Sample Dilution & Volume	85.71%	24	10.71%	3	3.57%	1	0.00%	0	0.00%	0	28
38	Concentration (40x, 100x)	79.31%	23	13.79%	4	3.45%	1	0.00%	0	3.45%	1	29

Showing rows 1 - 38 of 38

## Q9.1 - Q9.1 Other:

Q9.1 Other:

---

All

Q10 - Q10. What is the minimal starting volume of CSF needed for your fractionation method(s)?

Q10. What is the minimal starting volume of CSF needed for your fractionati...

500 ul

100-500uL

100ul

need to be found out

200 -500 micro liter

There is no single answer. For a high-abundance RNA target, we might need only 50 or 100 ul. For some downstream applications, several mL are needed or even pooling of multiple samples.

1 mL

100ul is the absolute minimum, we prefer 250ul

1 ml

0,5 ml

500µl

500µL

1 mL

3 ml

500ul

200 µL

0.2 ml

i recall we had useable material from 2-3 ml csf

1ml

50 ul

Q10. What is the minimal starting volume of CSF needed for your fractionati...

---

3 ml

0.5 mL

500 microliters

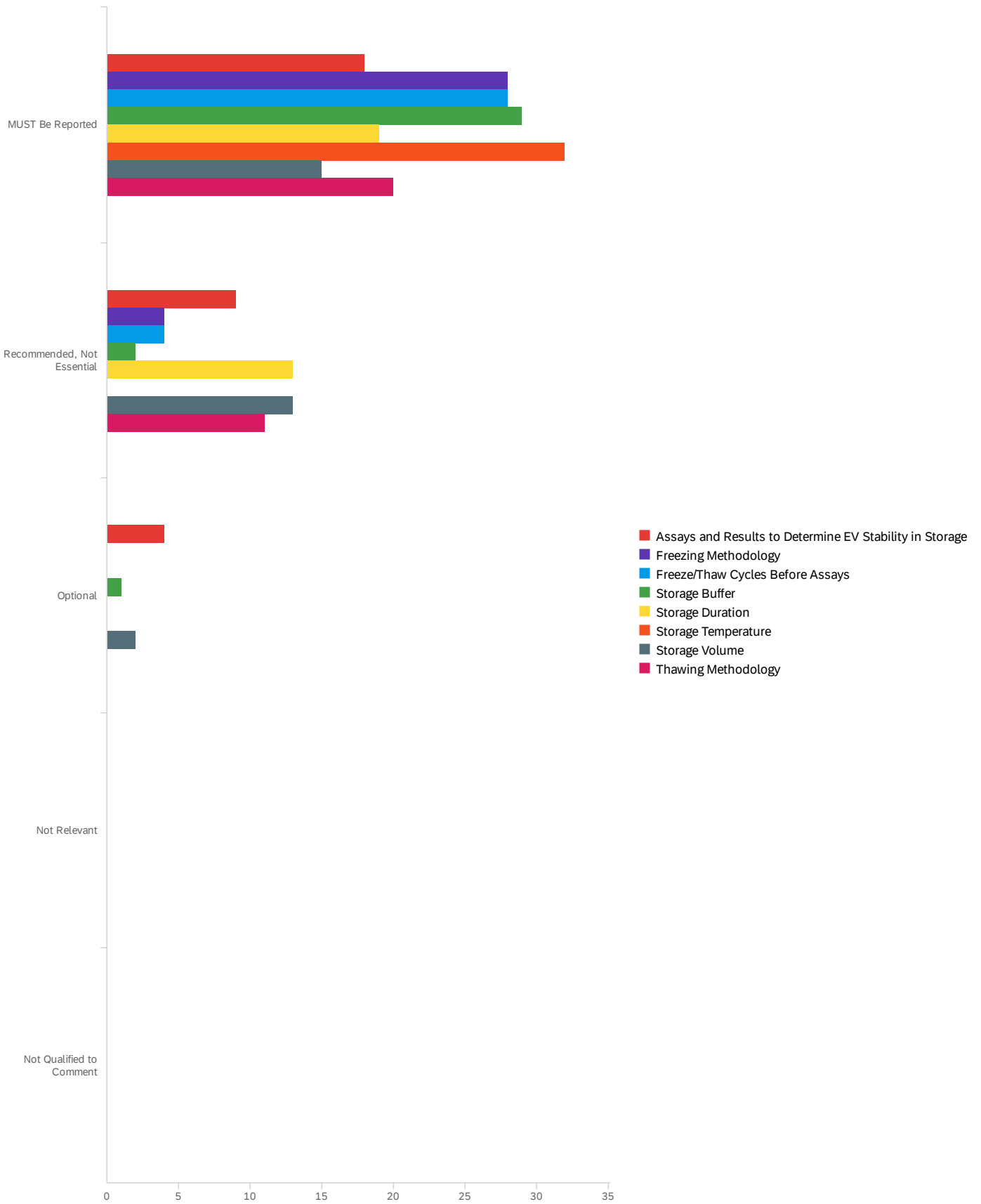
500 uL

500 ul for SEC if downstream application is qPCR. 5mLs CSF for SEC for WB, TEM.



Q11#1 - Q11. What parameters might form reporting guidelines for post-fractionation

storage of CSF EV pre... - Click to write Column 1



#	Field	Minimum	Maximum	Mean	Std Deviation	Variance	Count
1	Assays and Results to Determine EV Stability in Storage	1.00	3.00	1.55	0.71	0.51	31

#	Field	Minimum	Maximum	Mean	Std Deviation	Variance	Count
2	Freezing Methodology	1.00	2.00	1.13	0.33	0.11	32
3	Freeze/Thaw Cycles Before Assays	1.00	2.00	1.13	0.33	0.11	32
4	Storage Buffer	1.00	3.00	1.13	0.41	0.17	32
5	Storage Duration	1.00	2.00	1.41	0.49	0.24	32
6	Storage Temperature	1.00	1.00	1.00	0.00	0.00	32
7	Storage Volume	1.00	3.00	1.57	0.62	0.38	30
8	Thawing Methodology	1.00	2.00	1.35	0.48	0.23	31

#	Field	MUST Be Reported	Recommended, Not Essential	Optional	Not Relevant	Not Qualified to Comment	Total
1	Assays and Results to Determine EV Stability in Storage	58.06% 18	29.03% 9	12.90% 4	0.00% 0	0.00% 0	31
2	Freezing Methodology	87.50% 28	12.50% 4	0.00% 0	0.00% 0	0.00% 0	32
3	Freeze/Thaw Cycles Before Assays	87.50% 28	12.50% 4	0.00% 0	0.00% 0	0.00% 0	32
4	Storage Buffer	90.63% 29	6.25% 2	3.13% 1	0.00% 0	0.00% 0	32
5	Storage Duration	59.38% 19	40.63% 13	0.00% 0	0.00% 0	0.00% 0	32
6	Storage Temperature	100.00% 32	0.00% 0	0.00% 0	0.00% 0	0.00% 0	32
7	Storage Volume	50.00% 15	43.33% 13	6.67% 2	0.00% 0	0.00% 0	30
8	Thawing Methodology	64.52% 20	35.48% 11	0.00% 0	0.00% 0	0.00% 0	31

Showing rows 1 - 8 of 8

## Q11.1 - Q11.1 Other:

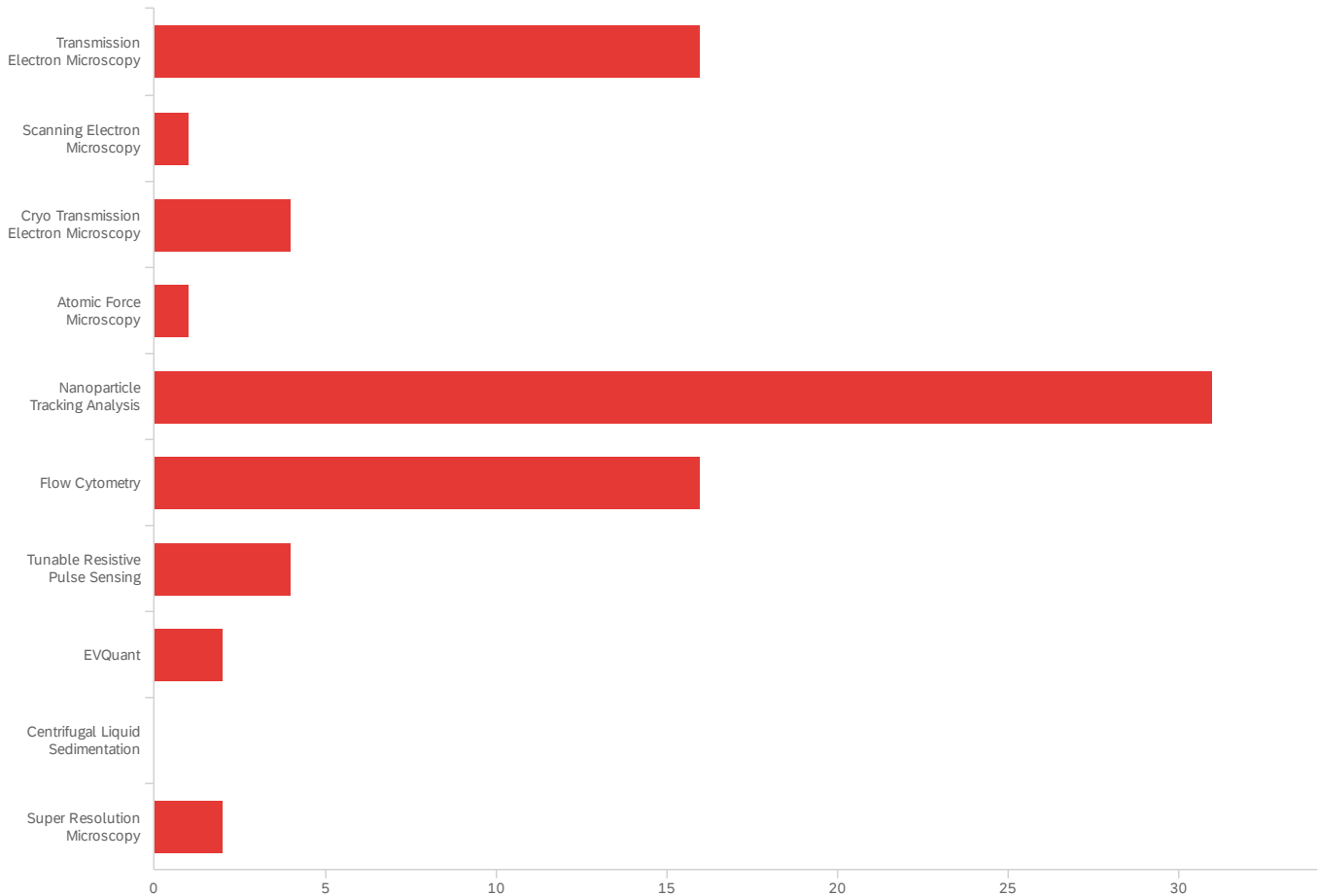
Q11.1 Other:

---

I think that EVs are fairly stable during storage, although many would disagree with that.

## Q12 - Q12. What methods do you use for analyzing CSF EV size and/or concentration?

Please select all that apply.



#	Field	Choice Count
1	Transmission Electron Microscopy	20.78% 16
2	Scanning Electron Microscopy	1.30% 1
3	Cryo Transmission Electron Microscopy	5.19% 4
4	Atomic Force Microscopy	1.30% 1
6	Nanoparticle Tracking Analysis	40.26% 31
9	Flow Cytometry	20.78% 16
10	Tunable Resistive Pulse Sensing	5.19% 4
12	EVQuant	2.60% 2
17	Centrifugal Liquid Sedimentation	0.00% 0

#	Field	Choice Count
18	Super Resolution Microscopy	2.60% 2

77

Showing rows 1 - 11 of 11

## Q12.1 - Q12.1 Other:

Q12.1 Other:

---

Zetaview particle counts

AF4,

DLS

immuno-phenotyping assays

MRPS, multiplex

Q13 - Q13. What is the minimal starting volume of CSF needed for your EV size and/or concentration methods?

Q13. What is the minimal starting volume of CSF needed for your EV size and...

0.8 ml

5-10uL

5-10ul

10ul

not yet determined

200-500

Up to several mL

200ul

Depending on the concentration, volume of EV fraction needed: NTA: 10-100  $\mu$ L Flow cytometry: 25-100  $\mu$ L

5-8 ML

10 microliters

500 $\mu$ l

500 $\mu$ L

500UL

1 mL

3 ml

500ul

if measuring direct on CSF 200ul is plenty

1ml

50 ul

3 ml



Q13. What is the minimal starting volume of CSF needed for your EV size and...

5 mL

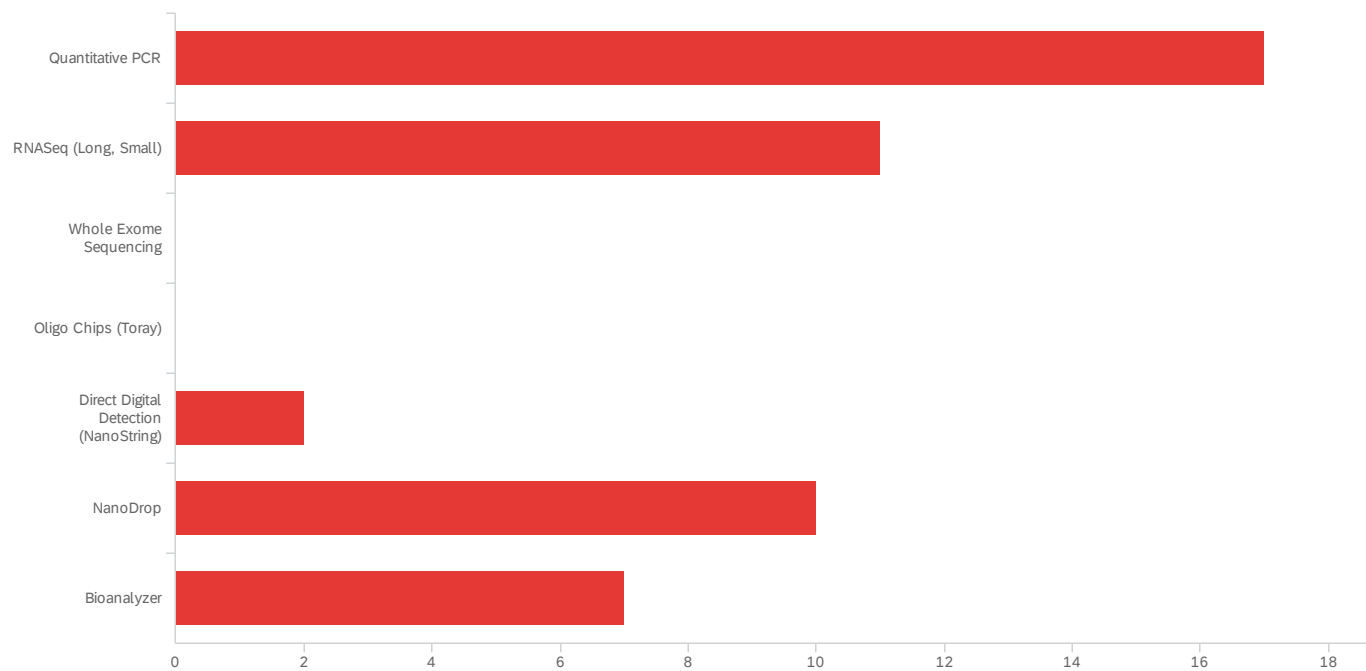
0.5 mL

500

5 mL

For fNTA and TEM 5mL CSF prior to SEC, 500 ul for flow cytometry.

Q14 - Q14. What method do you use for EV RNA cargo analysis? Please select all that apply.



#	Field	Choice Count
1	Quantitative PCR	36.17% 17
2	RNASeq (Long, Small)	23.40% 11
4	Whole Exome Sequencing	0.00% 0
5	Oligo Chips (Toray)	0.00% 0
7	Direct Digital Detection (NanoString)	4.26% 2
10	NanoDrop	21.28% 10
11	Bioanalyzer	14.89% 7
		47

Showing rows 1 - 8 of 8

## Q14.1 - Q14.1 Other:

Q14.1 Other:

---

surface cargo - flow cytometry

to be determined

Qubit

I do proteomics

## Q15 - Q15. What is the minimal starting volume for your EV RNA analysis method?

Q15. What is the minimal starting volume for your EV RNA analysis method?

---

0.5 ml

5-10uL

10-20ul

250ul

to be determined

200-500 micro liter

Ranges from tens of microliters (single qPCR) to many mL (sequencing).

200ul

2-5 µL

5ml

200 microliters

500µL

500ul

we have not yet managed this with CSF

1ml

3 ml

5mL

Subjective

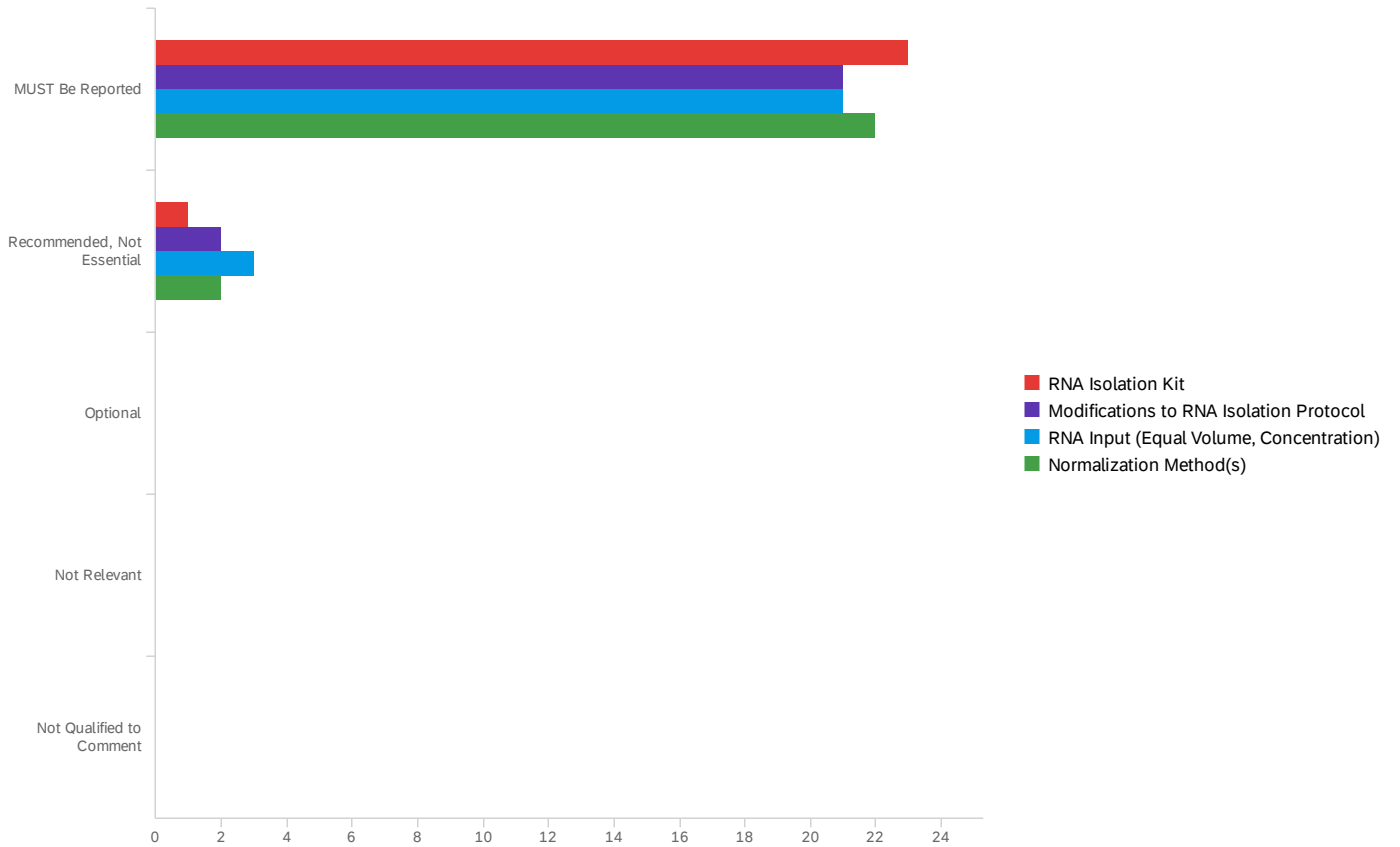
50

500 uL

500ul prior to qPCR.

# Q16#1 - Q16. What parameters might form reporting guidelines for CSF EV RNA Cargo

analysis? Please rate t... - Click to write Column 1



#	Field	Minimum	Maximum	Mean	Std Deviation	Variance	Count
1	RNA Isolation Kit	1.00	2.00	1.04	0.20	0.04	24
2	Modifications to RNA Isolation Protocol	1.00	2.00	1.09	0.28	0.08	23
3	RNA Input (Equal Volume, Concentration)	1.00	2.00	1.13	0.33	0.11	24
4	Normalization Method(s)	1.00	2.00	1.08	0.28	0.08	24

#	Field	MUST Be Reported	Recommended, Not Essential	Optional	Not Relevant	Not Qualified to Comment	Total
1	RNA Isolation Kit	95.83% 23	4.17% 1	0.00% 0	0.00% 0	0.00% 0	24
2	Modifications to RNA Isolation Protocol	91.30% 21	8.70% 2	0.00% 0	0.00% 0	0.00% 0	23

#	Field	MUST Be Reported	Recommended, Not Essential	Optional	Not Relevant	Not Qualified to Comment	Total
3	RNA Input (Equal Volume, Concentration)	87.50% 21	12.50% 3	0.00% 0	0.00% 0	0.00% 0	24
4	Normalization Method(s)	91.67% 22	8.33% 2	0.00% 0	0.00% 0	0.00% 0	24

Showing rows 1 - 4 of 4

## Q16.1 - Q16.1 Other:

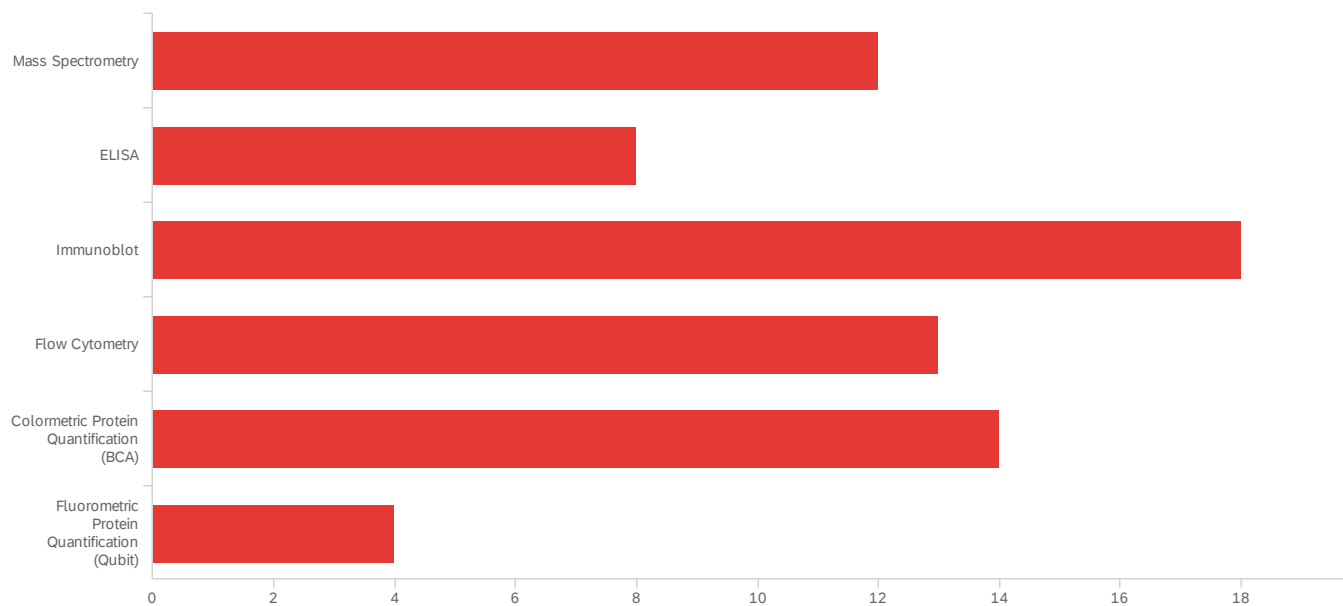
Q16.1 Other:

---

If the goal is to measure EV RNA, it must be shown that the RNA is in or attached to the EVs. In many studies, it cannot be excluded that most RNA is not associated with EVs and is simply a co-isolate. I often worry that even a tiny influence of cells (including those coming from the puncture) might contribute more RNA to CSF "EV" measurements than the actual CSF EVs.

lysis procedure, buffers used, detection dye/kit, cDNA synthesis step, amplification method, primer/probe sequences used, etc

Q17 - Q17. What method do you use for CSF EV protein analysis? Please check all that apply.



#	Field	Choice Count
1	Mass Spectrometry	17.39% 12
3	ELISA	11.59% 8
4	Immunoblot	26.09% 18
6	Flow Cytometry	18.84% 13
8	Colormetric Protein Quantification (BCA)	20.29% 14
9	Fluorometric Protein Quantification (Qubit)	5.80% 4

69

Showing rows 1 - 7 of 7



# Q17.1 - Q17.1 Other:

Q17.1 Other:

---

to be determined

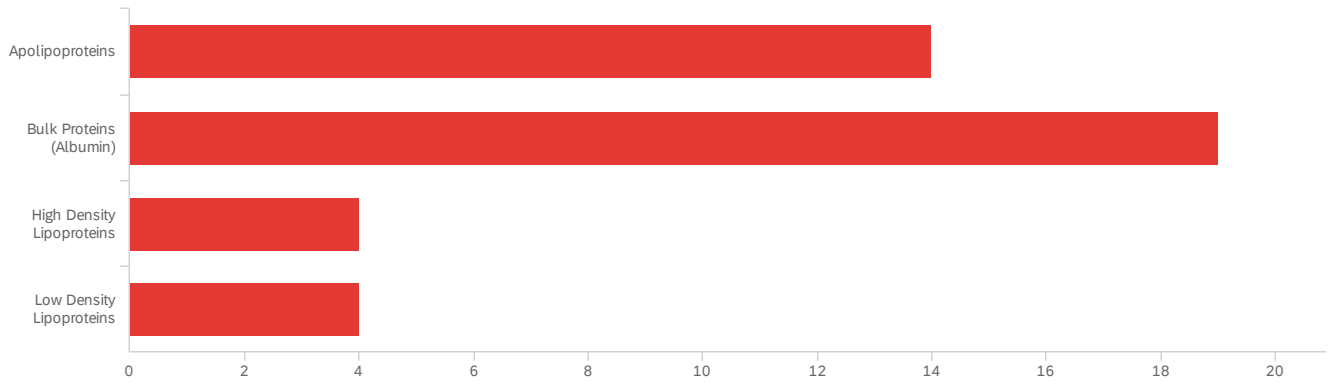
fluoNTA, Nanoview

Nanoview ExoView

Multiplex

# Q18 - Q18. What non-EV markers do you use for CSF EV protein analysis and/or purity?

Please select all that apply.



#	Field	Choice Count
1	Apolipoproteins	34.15% 14
2	Bulk Proteins (Albumin)	46.34% 19
3	High Density Lipoproteins	9.76% 4
4	Low Density Lipoproteins	9.76% 4

41

Showing rows 1 - 5 of 5

## Q18.1 - Q18.1 Other:

Q18.1 Other:

---

histone 4

IgG

## Q19 - Q19. What is the minimal starting volume of CSF for your EV protein analysis?

Q19. What is the minimal starting volume of CSF for your EV protein analysi...

---

10-20ul

250ul

TBD

200-500 micro liter

1 mL

0.5 ml CSF

2ML

400 microliters

500µl

500µL

1 mL

3 ml

500ul

200µL

0.5 ml

as for our purificaion 1-2mls

1ml

100

3 ml

5mL

0.5 mL

500

Q19. What is the minimal starting volume of CSF for your EV protein analysi...

---

5 mL

5mL for Immunoblot, 500ul for flow cytometry.

## Q20 - Q20. What neuronal and/or glial EV markers do you use in CSF studies?

Q20. What neuronal and/or glial EV markers do you use in CSF studies?

Anti Mog

under investigation

syn

MMP2 and EMMPRIN but these are specific to our disease of interest

L1CAM, NCAM, CD11b

Neurofilament light protein and GFAP

L1CAM, NCAM

NCAM, L1CAM, GLAST

ICAM

L1CAM CD63 CD9

L1CAM (neurons)

NCAM, GFAP, Vimentin

proprietary astrocytoma antigen markers and astrocytoma-associated miRs

L1CAM, TMEM119

L1CAM

NSE, Synaptophysin

Yet to find a neuro-specific marker

Glast, NCAM

Glast, NCAM

## Q21 - Q21. What antibodies (and source) do you recommend for CSF EV positive & negative protein markers?

Q21. What antibodies (and source) do you recommend for CSF EV positive & ne...

---

TBD

cell signalling

positive: CD81, CD9, CD63 - Santa Cruz (western blot), Biolegend (flow cytometry) negative: ApoA-I - Santa Cruz

ongoing research

Positive: CD9, CD63, CD81 Negative: Albumin

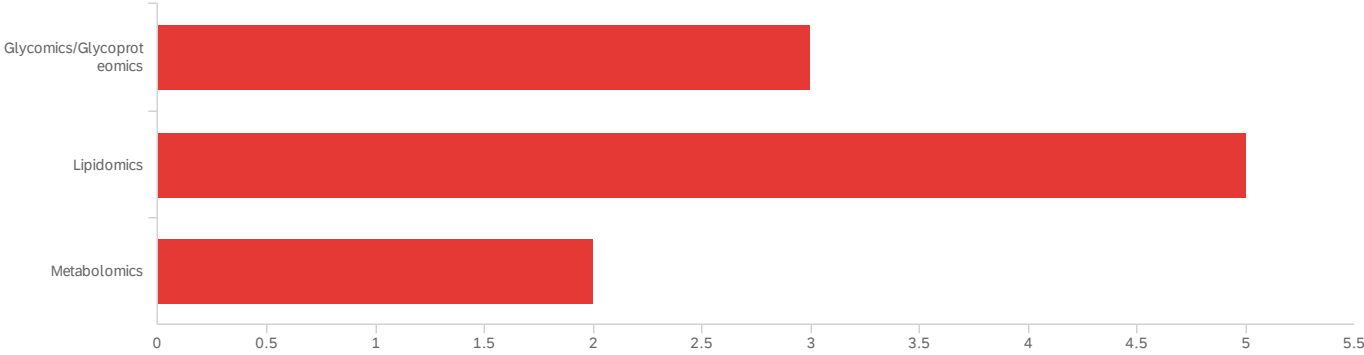
positive: tetraspanins (abcam), NCAM (abcam) , negative: calnexin (abcam)

ITGB1, IGF-1R, IGF-2R for positive, source variable

CD9, CD63, CD81, TSG101, Flotillin, APOA1, Albumin Novus antibodies

ApoA1, Santa Cruz, (12C8) 80551; Albumin, Cell Signaling, #4929; Flotillin, Abcam, AB133497; CD81, Santa Cruz, (B-11) sc-166029; GLAST, Novus, NB100-1869

Q22 - Q22. What other methods do you use to analyze EV components? Please select all that apply.



#	Field	Choice Count
1	Glycomics/Glycoproteomics	30.00% 3
2	Lipidomics	50.00% 5
3	Metabolomics	20.00% 2
		10

Showing rows 1 - 4 of 4



## Q22.1 - Q22.1 Other:

Q22.1 Other:

---

we aim for lipidomics and metabolomics

Immunomodulatory assays

proteomics !

proteomics MS/MS

## Q23 - Q23. What is the minimal starting volume of CSF for other EV analysis methods?

Q23. What is the minimal starting volume of CSF for other EV analysis metho...

---

10-20ul

TBD

200-500 micro liter

500µL

1 mL

3 ml

500ul

0.5 ml

1ml

3 ml

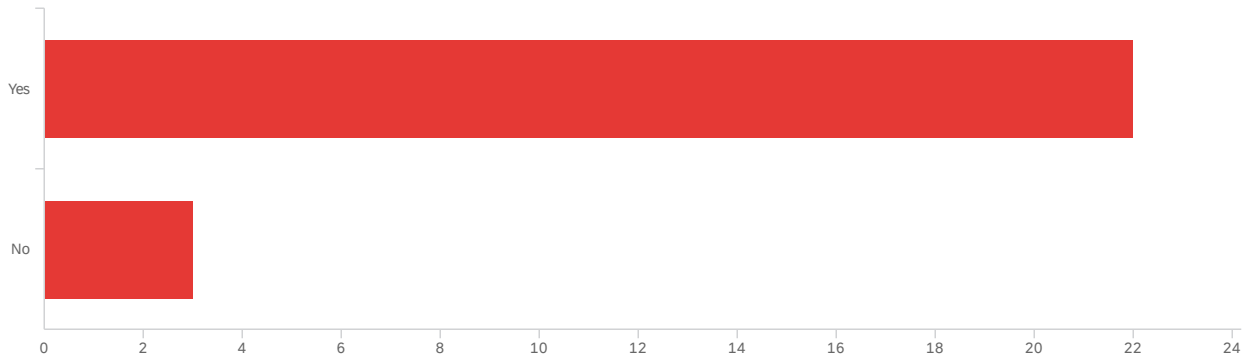
5mL

subjective

500

5 mL

Q24 - Q24. Should MISEV2018 guidelines be followed for CSF EV measures in publications (e.g. quantification of protein, lipid, nucleic acids, surface markers), with compliance stated?



#	Field	Minimum	Maximum	Mean	Std Deviation	Variance	Count
1	Q24. Should MISEV2018 guidelines be followed for CSF EV measures in publications (e.g. quantification of protein, lipid, nucleic acids, surface markers), with compliance stated?	1.00	2.00	1.12	0.32	0.11	25

#	Field	Choice Count
1	Yes	88.00% 22
2	No	12.00% 3
		25

Showing rows 1 - 3 of 3

Q25 - Please cite key papers on CSF EVs (PMID or DOI), or add additional comments

here.

Please cite key papers on CSF EVs (PMID or DOI), or add additional comments...

---

NA

doi: 10.7150/thno.31502; doi: 10.1002/pmic.201800257; doi: 10.1080/20013078.2017.1369805; doi: 10.1080/20013078.2017.1317577; doi: 10.1093/brain/awv346; doi: 10.1080/20013078.2017.1348885; doi: 10.1016/j.celrep.2019.08.036; 10.1186/s12014-020-09294-7

<https://doi.org/10.1186/s12974-019-1617-y>

doi: 10.1080/20013078.2017.1369805. this is our only experience/concerted effort with CSF and we havent been able to progress it in MS

**End of Report**