



Recommendations from JBF discussion group, DG2013-01 on the preparation of calibration standards and QC samples for bioanalysis of small molecular drugs

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Overview

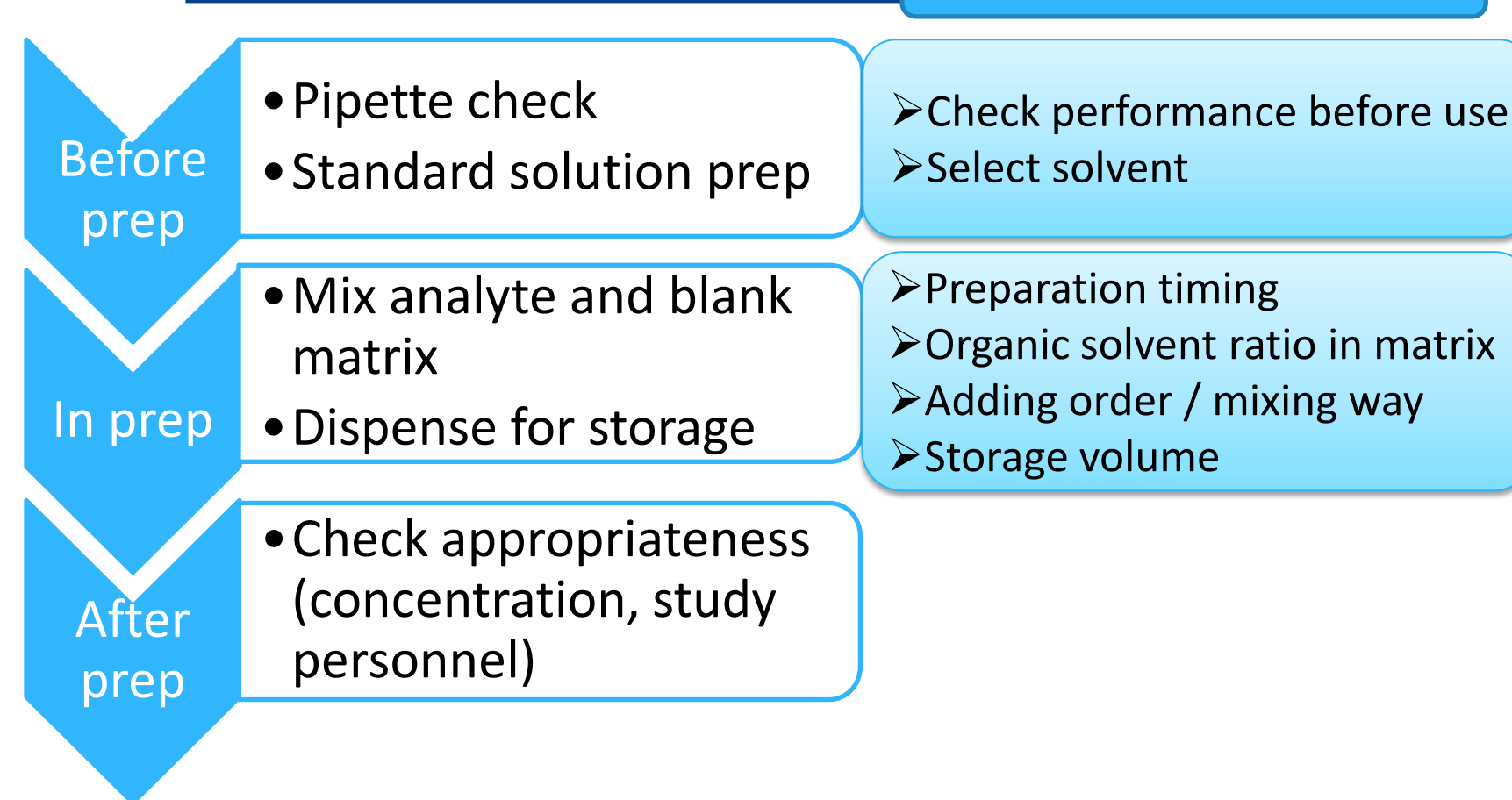
- Practical preparation methods of calibration standards (CS) and QC samples are not covered in FDA/EU/MHLW BMV guidelines.
- Broad range of opinions on the preparation methods of CS and QC samples in the questionnaires survey from 148 bioanalysts in Japan.
- Present our recommendable procedures on the preparation of CS and QC samples with pros and cons.

Focusing on small molecules

Points to consider

- Sample preparation processes:
Accurate and reproducible as possible even in different occasions, analysts or labs
- Analytes dispersion into matrix solution:
Uniform as possible
- QC samples:
Similar to actual samples as possible

Process flow



Recommendable procedures

Pipette check

- A mechanical pipette used for diluting standard solutions should be checked just before use.
- Systematic error : should be within $\pm 2\%$
 - To add the analyte accurately and reproducibly as possible in different occasions.
 - In practically, the systematic error to be kept within $\pm 1\%$ in DG 2013-01 member's experience.
- Random error : To set appropriate criteria based on the systematic error

Solvent choice

- Consider based on the **analyte solubility** and **adsorption**, etc.
- Many kinds of solvents can be used, but consider to **minimize the protein denaturation** as possible:
 - Mixture of organic solvent and water
 - Organic solvent (Alcohol > Acetonitrile)

Pros :

- Improve a difficulty of handling organic solvents
- Avoid bacteria growth in aqueous solutions

Preparation timing

Sample	Recommendable timing	Reason
CS	Prep in use	Multiple prep is effective to avoid the specific bias depending on a single preparation
QC A&P Stability Sample analysis	Batch prep	To check if QC samples prepared in a batch are measured accurately and reproducibly throughout the study

A&P: accuracy and precision

Solvent ratio

Sample	Recommendable ratio*	Reason
CS	Not limited	Need QC samples quantified appropriately
QC A&P ¹⁾ Stability ²⁾ Sample analysis ¹⁾	Within 2% (max 5%)	To be same as actual samples as possible and avoid protein denaturation

*Solvent ratio to blank matrix

- 1) Not limited when prepare in use (e.g. rare matrix)
2) Stability is not evaluated when prepare in use (e.g. rare matrix)

Adding order, mix, storage volume

Process	Recommendable process	Reason
Adding order	Dispense blank matrix → add SS* to the matrix	To avoid an adsorption of analyte and a contamination to blank matrix
Mix	Vortex mix and then inversion	To avoid insufficient dispersion
Storage volume	Aliquot more than required volume of a sample analysis	Enable to take an aliquot as same as study samples on the day of analysis

Check after prep

Check point	Sample	Recommended timing to check	Reason
Conc.	QC A&P Stability Sample analysis	Before use Just after prep Before use	To check accurately prepared and to avoid a reject of analytical run
Study personnel*	CS, A&P, etc	Before sample analysis	To verify appropriateness of study personnel

*Note: A partial validation is required if study personnel, who does not perform the validation study, is participated to the sample analysis.

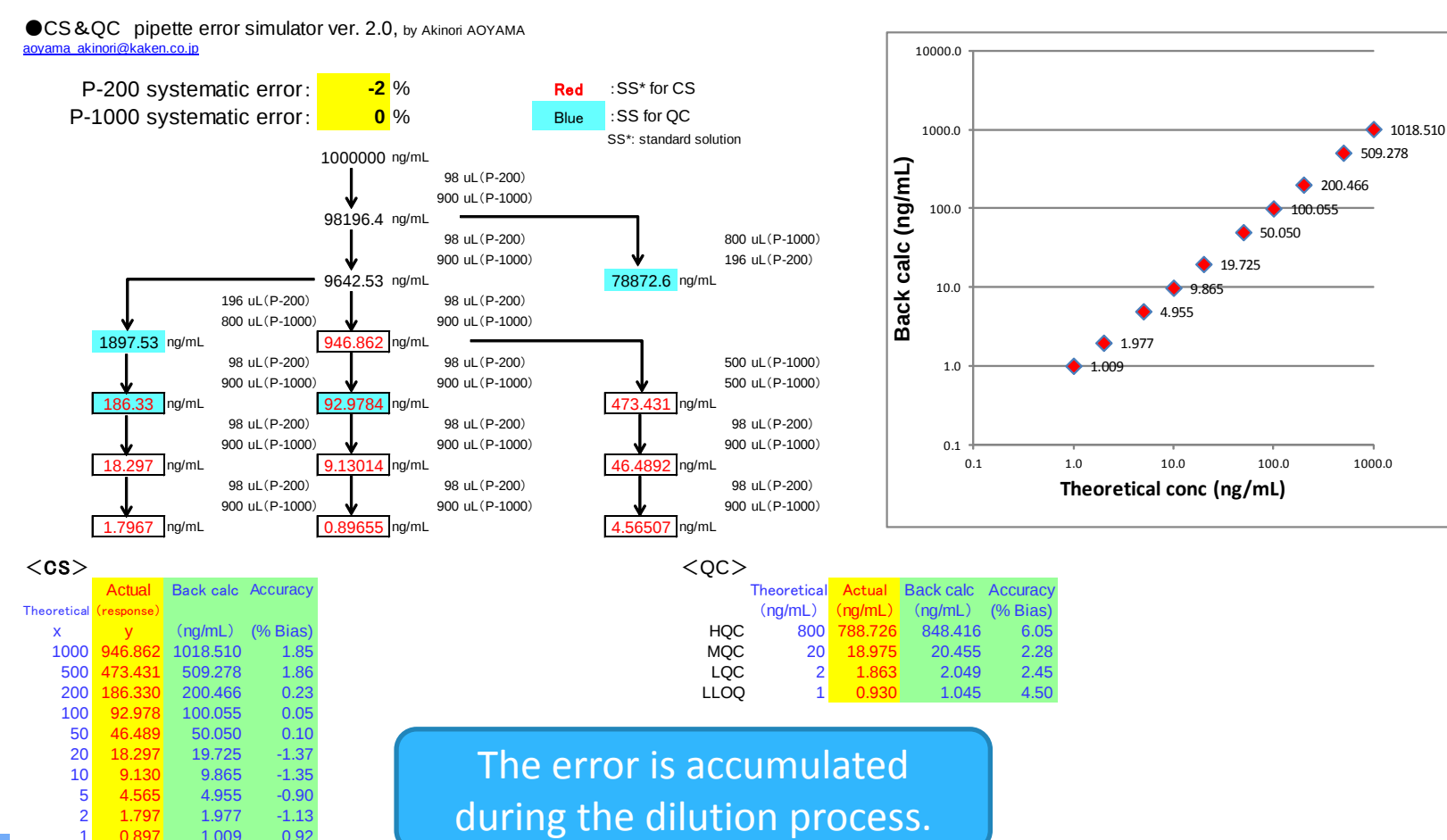
Summary

Recommendable process

- Before prep
- Check pipette just before use, systematic error: $\leq \pm 2\%$
 - SS* prep: consider analyte solubility, adsorption and protein denaturation
- In prep
- Prep timing: CS: prep in use, QC: prep in batch
 - Solvent ratio: CS: depends on QC, QC: $\leq \pm 2\%$ (max 5%)
 - Dispense blank matrix → add ss → mix vortex and inversion
 - Aliquot more than analytical volume, then store
- After prep
- Check conc; QC (stability): just after prep, QC others: before use
 - Check study personnel: before sample analysis
- SS*: standard solution

Questionnaires and discussion

Simulation: accumulating error caused by pipette error



Simulation : $\pm 2\%$ systematic error

Original: -2% systematic error							ISR: 2% systematic error						
Normal conc.		Actual conc.	Concentration	Back calc.	Conc. error	Judge	Normal conc.		Actual conc.	Concentration	Back calc.	Conc. error	Judge
			(ng/mL)	(ng/mL)	(%)					(ng/mL)	(ng/mL)	(%)	
1000	948.862	5.3	1018.510	1.8	OK	1000	1054.866	5.5	992.841	-1.7	OK		
500	474.431	6.3	509.270	1.8	OK	500	527.433	5.5	490.397	-1.7	OK		
200	186.330	6.8	200.466	0.2	OK	200	214.335	7.2	199.664	-0.2	OK		
100	92.978	7.0	100.054	0.1	OK	100	107.382	7.4	100.008	0.0	OK		
50	46.489	7.0	50.050	0.1	OK	50	53.691	7.4	49.981	0.0	OK		
20	18.297	8.5	19.725	-1.4	OK	20	21.819	9.1	20.263	-1.4	OK		
10	9.130	8.7	9.889	-1.3	OK	10	10.931	9.3	10.139	-1.4	OK		
5	4.565	8.7	4.900	-0.8	OK	5	5.466	9.3	5.046	-0.8	OK		
2	1.797	-10.2	1.977	-1.1	OK	2	2.221	11.1	2.023	-1.1	OK		
1	0.897	-10.3	1.009	-0.9	OK	1	1.113	11.3	0.990	-1.0	OK		
HQC	800	789.726	1.4	804.410	0.1	OK	HQC	800	811.127	1.4	795.734	-0.9	OK
MQC	20	18.976	-6.1	20.455	2.3	OK	MQC	20	21.055	5.3	19.572	-2.1	OK
LQC	2	1.863	-6.8	2.049	2.5	OK	LQC	2	2.143	7.2	1.951	-2.5	OK
LLOQ	1	0.930	-7.0	1.045	0.6	OK	LLOQ	1	1.074	7.4	0.954	-0.6	OK
UK sample (High)	950	950.000	None	1021.880	7.6	ISR	950	950.000	None	885.131	-14.3%	ISR	
UK sample (Low)	2.000	2.000	None	2.139	6.9	ISR	2.000	2.000	None	1.831	-10.9%	ISR	

ISR (ISR criteria: Original conc. (average of High and Low) + 10%)

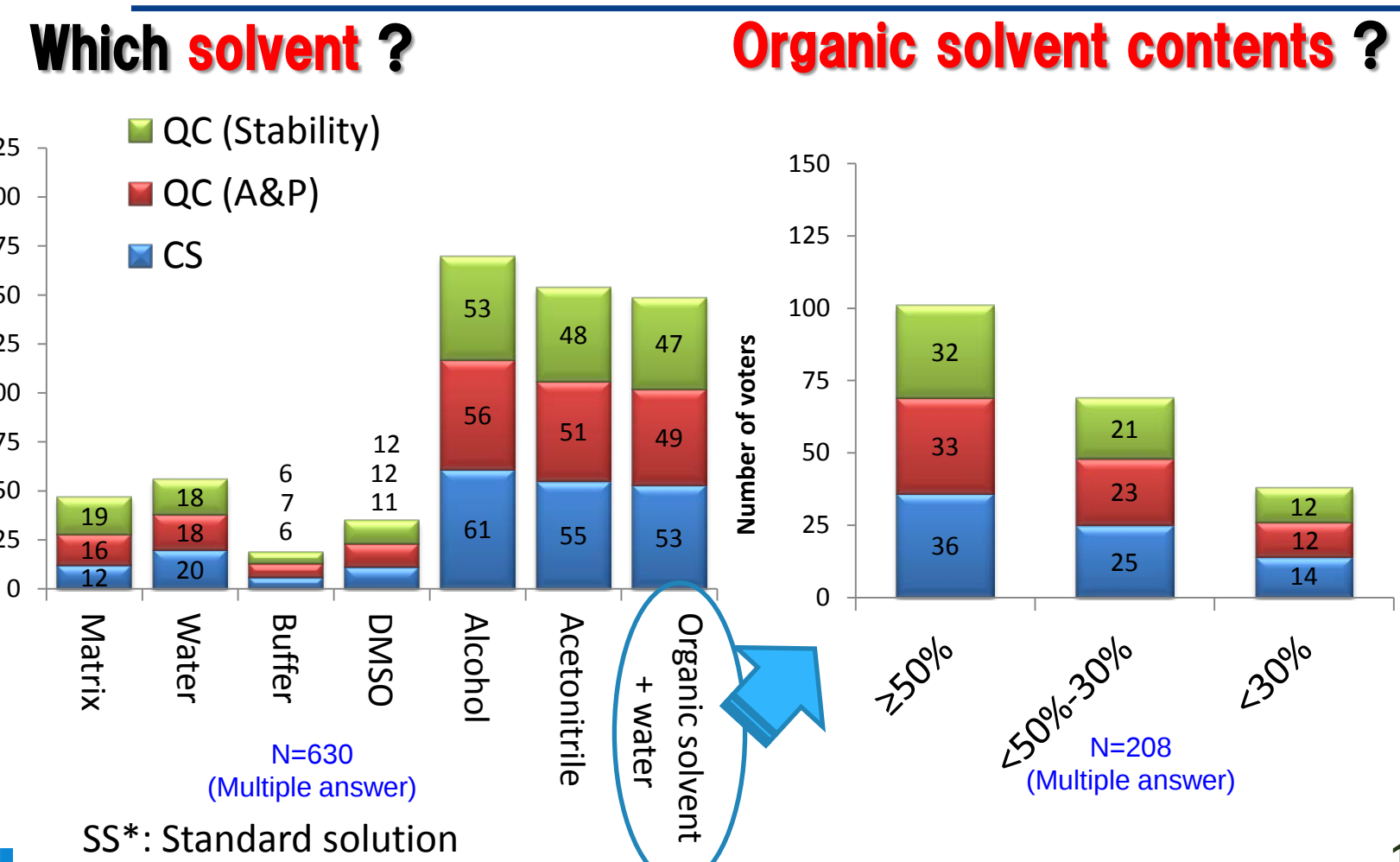
% Bias of Unknown samples:
within 10%

**Meet ISR criteria, within ±20%,
but barely**

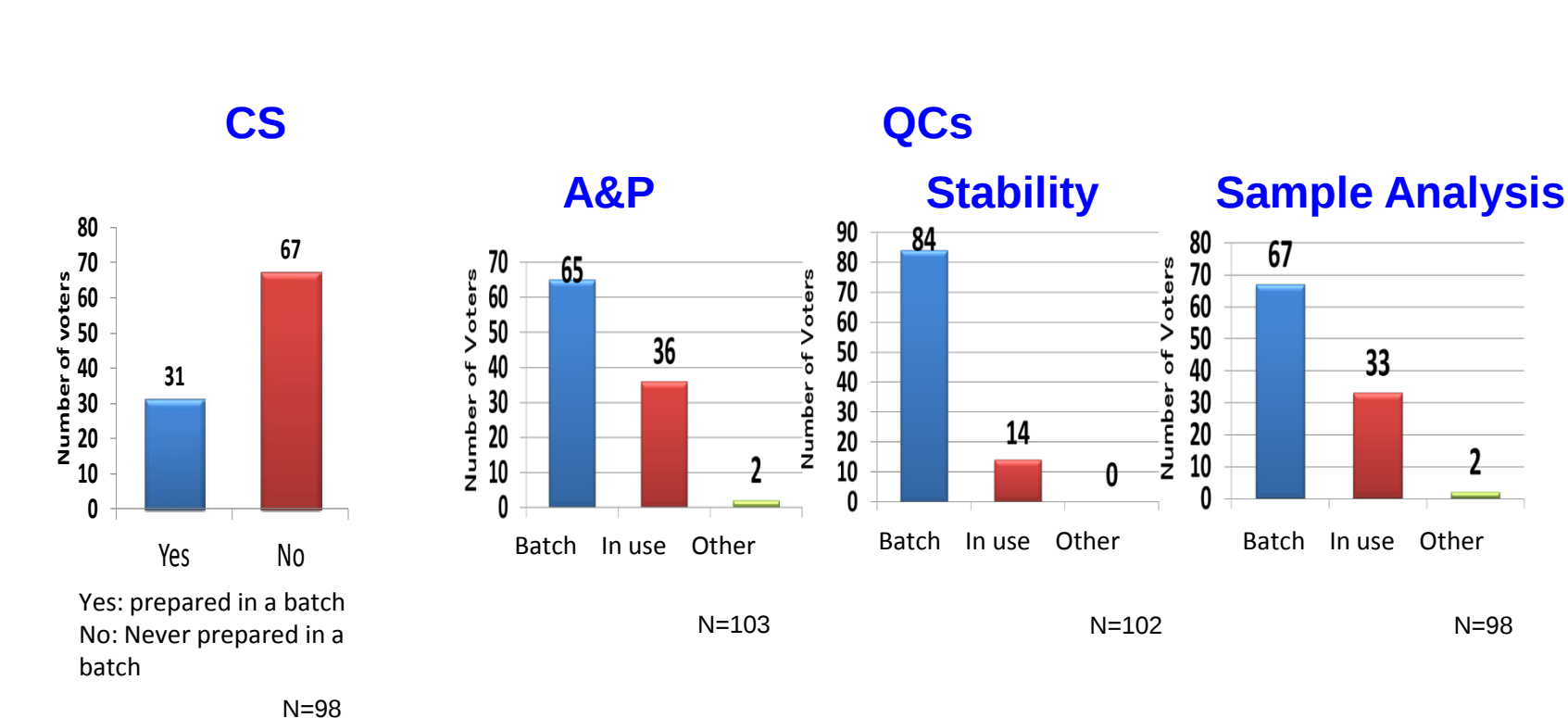
% Bias of Unknown samples: within 10%

Meet ISR criteria, within $\pm 20\%$, but barely

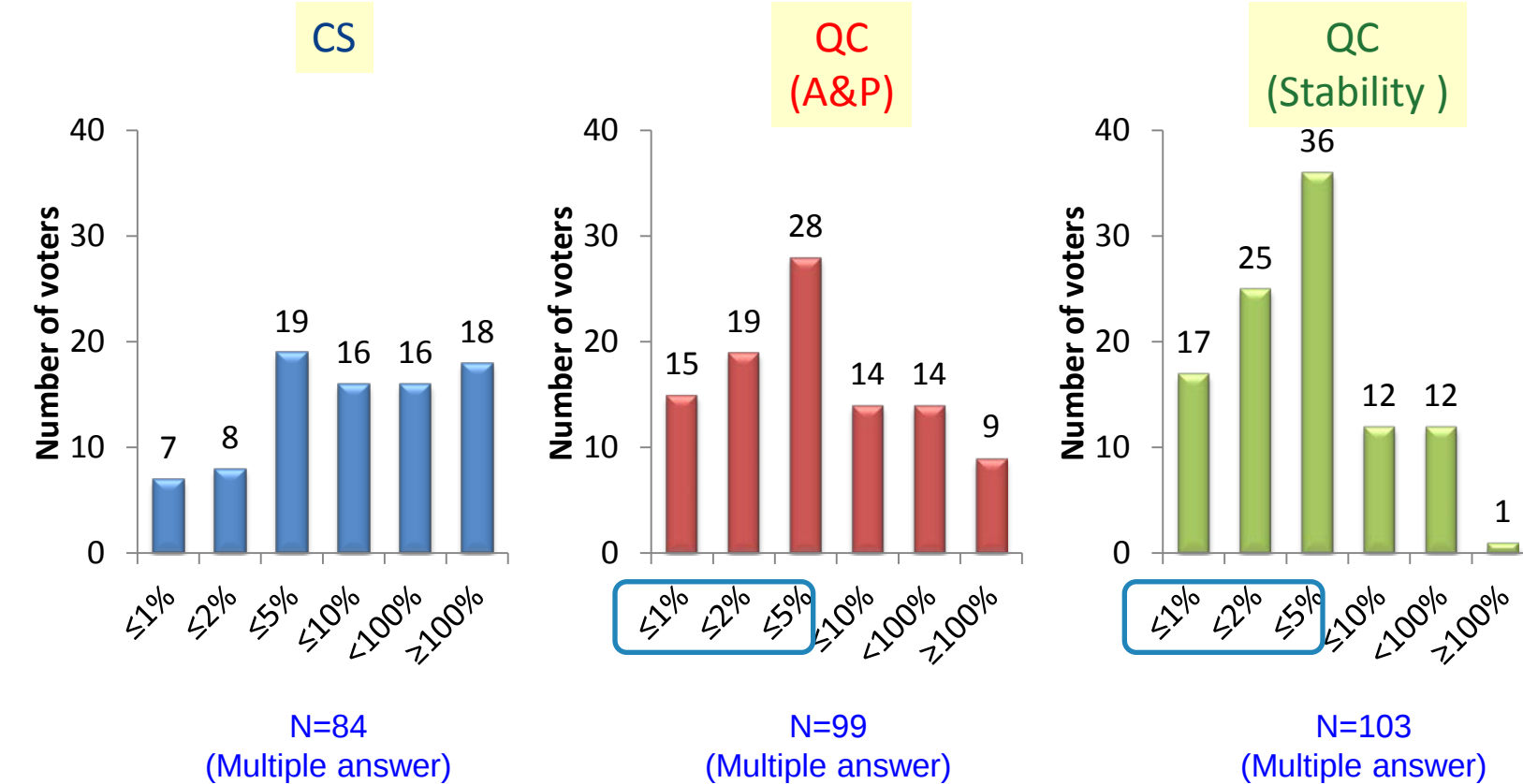
Q : SS* solvent choice



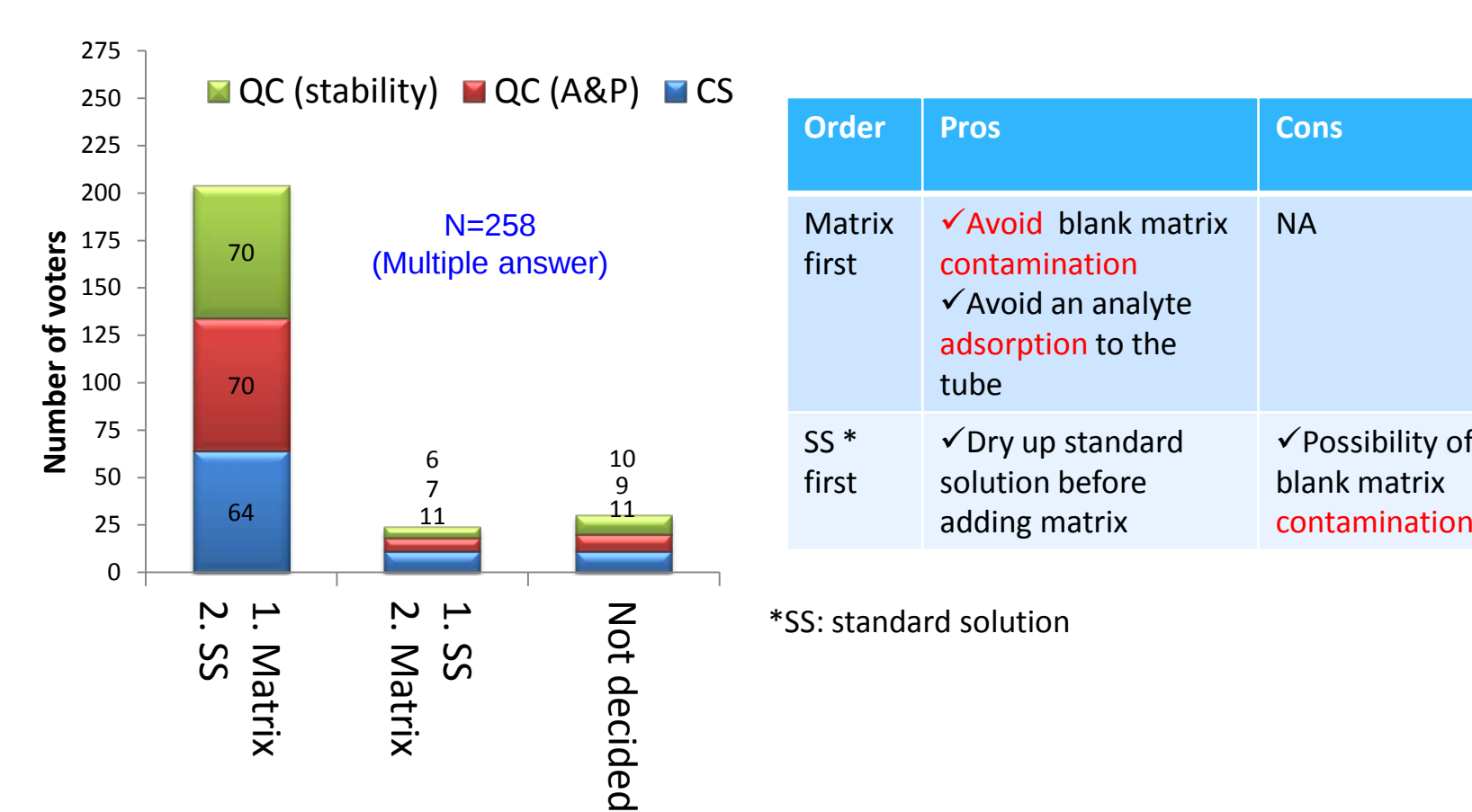
Q : Prepare Batch or In Use?



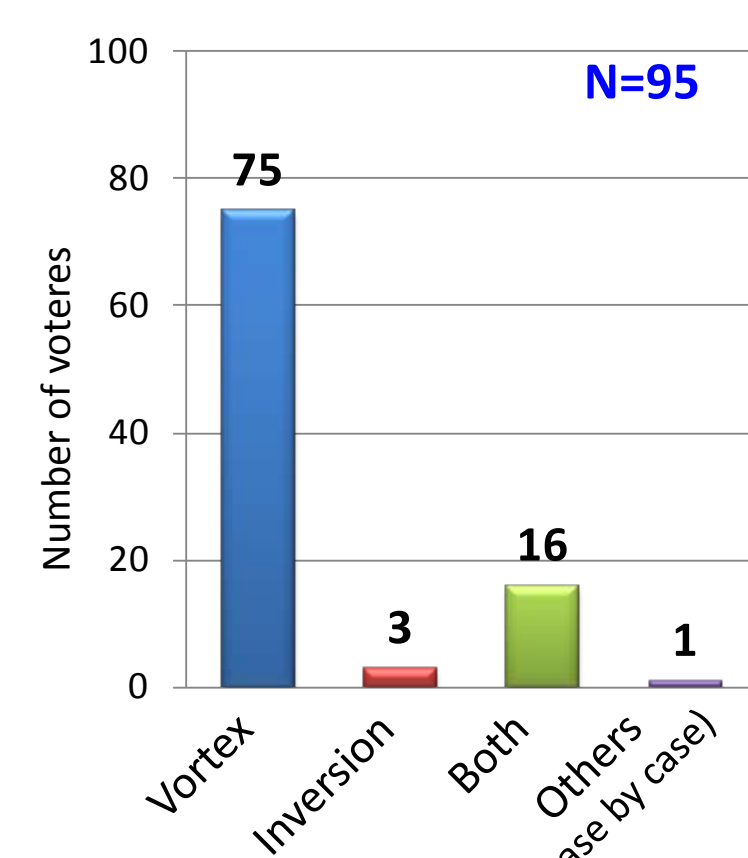
Q : SS* Ratio to blank matrix?



Q : Which first?



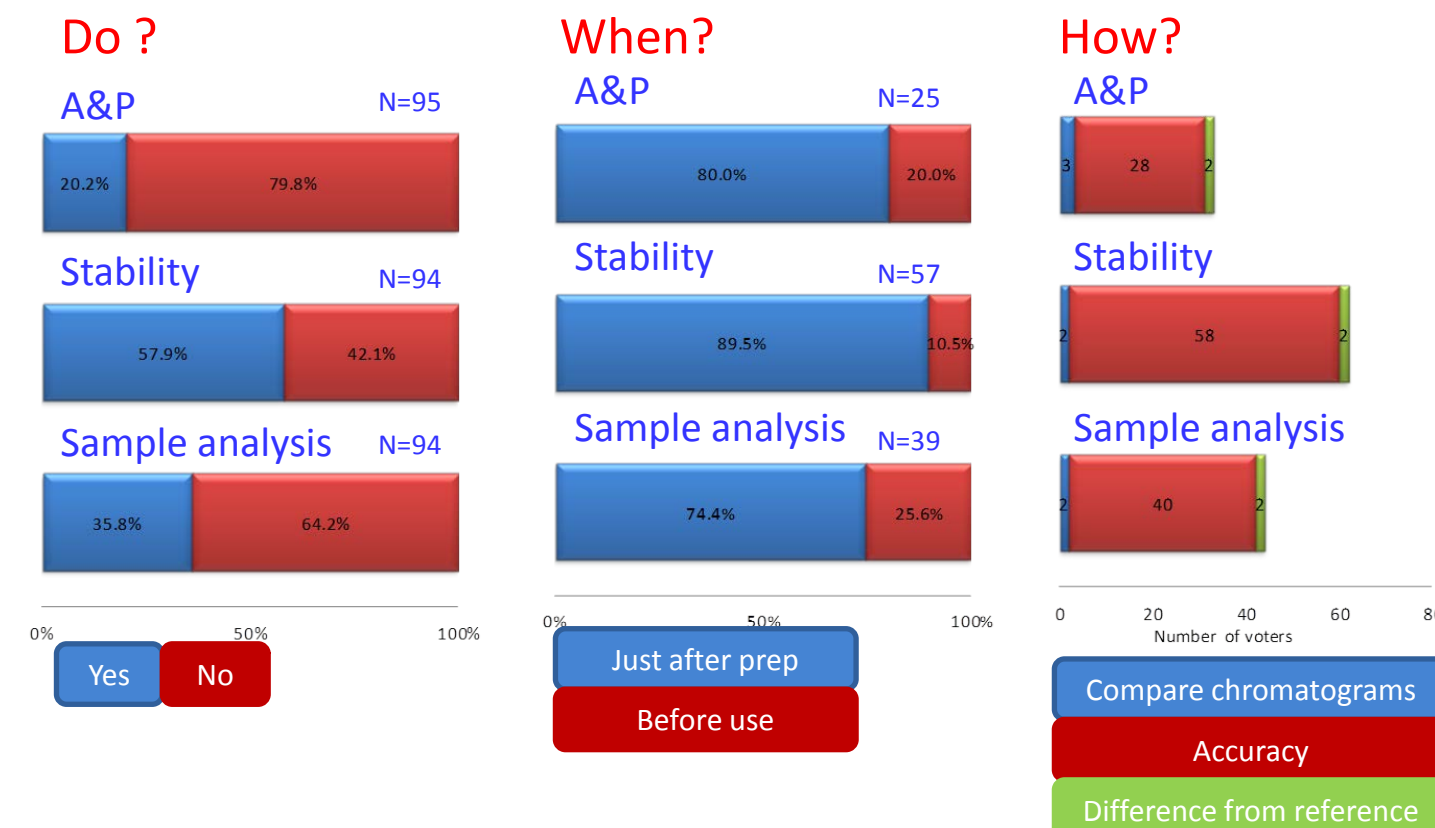
Q: How do you mix?



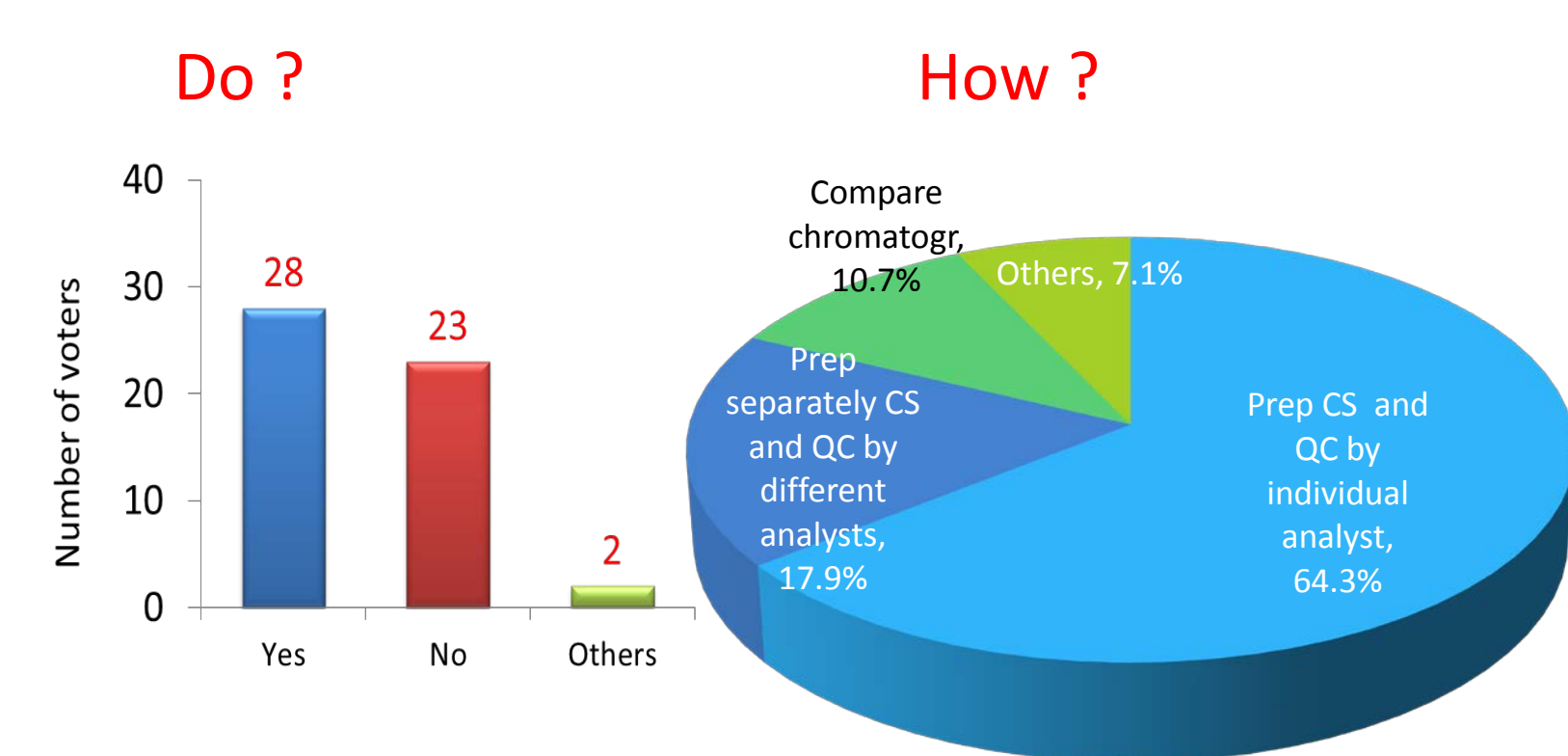
➤ Vortex mix isn't adequate to fully disperse analytes in matrix solutions.

➤ The insufficient dispersion causes a large variability in measured concentrations

Q: Check after prep



Q: Check personnel



Note: The views and opinions expressed herein represent those of the DG2013-01 and do not necessarily represent the views, opinions or practices of GlaxoSmithKline K.K. or Japan Bioanalysis Forum.