

RoM = robust average	AV = assigned value	Dmax = acceptable percent difference
SD = standard deviation	CRV = certified reference value	LL = lower limit
CV = coefficient of variation	RV = reference value	UL = upper limit
Ntot = total number of participants	CVE = consensus value from experts	Neva = number of evaluated participants
Nout = number of results excluded before calculation	CVP = consensus value from all participants	Nsuc = number of successful participants
	CVPG = consensus value from participants groups	Srel = success (relative)
	U_{AV} = expanded uncertainty of the assigned value ($k = 2$)	

Test	[unit]	Comparability												
		RoM	SD	CV [%]	N _{tot}	N _{out}	AV	U_{AV}	D _{max}	LL	UL	N _{eva}	N _{suc}	S _{rel}
(247) Gammopathies (quan.)					14							14	14	100%
Samples and groups	[g/L]													
Sample A		14,5	3,0	21	14	CVP	15,5	0,80	33%	10,3	20,7	14	14	100%
All results		14,5	3,0	21	14	0						14		
(279) Total protein (plasma)					14							14	14	100%
Samples and groups	[g/L]													
Sample A		62,5	1,5	2,4	14	CVP	62,9	0,46	8%	57,8	68	14	14	100%
All results		62,5	1,5	2,4	14	0						14		
(314) FLC kappa					11							11	9	82%
Samples and groups	[mg/L]													
Sample A		155	69	44	11	CVP	168	23	50%	84	252	11	9	82%
All results		155	69	44	11	0						11		
(315) FLC lambda					11							11	11	100%
Samples and groups	[mg/L]													
Sample A		2,53	0,85	34	11	CVP	3,21	0,56	100%	0	6,42	11	11	100%
All results		2,53	0,85	34	11	0						11		
(545) Gammopathies (quan.)					10							10	10	100%
Samples and groups	[g/L]													
Sample B		0,579	0,24	41	10	CVP	0,545	0,070	80%	0,109	0,981	10	10	100%
All results		0,579	0,24	41	10	0						10		
(550) Total protein (urine)					12							12	12	100%
Samples and groups	[g/L]													
Sample B		0,963	0,095	9,8	12	CVP	0,824	0,046	30%	0,576	1,08	12	12	100%
All results		0,963	0,095	9,8	12	0						12		