

Sannova achieves single-digit CV's for most analyses:

Small Molecules: Summary of all calibration curves for the validation of Dextroamphetamine

Batch ID	Date of batch Injected	STD A	STD B	STD C	STD D	STD E	STD F	STD G	STD H
		0.366	0.732	1.464	2.440	4.880	9.761	19.522	39.043
		ng/mL	ng/mL	ng/mL	ng/mL	ng/mL	ng/mL	ng/mL	ng/mL
003-005	16-Jan-12	0.368	0.720	1.498	2.411	4.773	10.098	19.304	38.994
003-008	20-Jan-12	0.379	0.678	1.485	2.370	4.945	9.932	20.272	38.386
003-007	17-Jan-12	0.369	0.712	1.495	2.439	4.755	10.194	19.261	38.723
003-009	17-Jan-12	0.360	0.733	1.529	2.567	4.734	10.081	18.417	38.012
003-012	20-Jan-12	0.377	0.715	1.336	2.459	4.988	9.874	19.312	41.005
003-013	20-Jan-12	0.366	0.721	1.482	2.537	4.801	10.292	18.383	38.434
Mean		0.370	0.713	1.471	2.464	4.833	10.079	19.158	38.926
SD		0.007	0.019	0.068	0.075	0.107	0.157	0.699	1.071
%CV		1.9	2.7	4.6	3.0	2.2	1.6	3.6	2.8
% ACCURACY		101.1	97.4	100.5	101.0	99.0	103.3	98.1	99.7
n		6	6	6	6	6	6	6	6

Large Molecules: Summary of all calibration standard concentrations in Rat Plasma for Protein by digestion

Batch ID	Date of batch Injected	STD A	STD A'	STD B	STD C	STD D	STD E	STD F	STD G	STD H	STD H'
		2.904	2.904	5.186	8.644	14.406	20.580	29.400	42.000	60.000	60.000
		µg/mL	µg/mL	µg/mL	µg/mL	µg/mL	µg/mL	µg/mL	µg/mL	µg/mL	µg/mL
012-001	18-Sep-15	2.895	2.902	5.192	8.859	13.986	20.726	29.327	42.008	61.445	58.704
013-001	22-Sep-15	2.956	2.877	5.180	8.378	14.033	21.342	30.087	42.700	56.912	61.383
008-001	24-Sep-15	2.871	2.858	5.493	8.636	13.887	21.665	27.258	42.315	61.409	59.650
006-001	25-Sep-15	2.849	2.897	5.444	8.571	14.158	20.967	28.418	41.951	62.467	58.327
014-001	29-Sep-15	2.726	3.018	5.398	8.736	14.326	20.373	27.108	40.605	64.752	57.025
013-002	29-Sep-15	2.829	2.932	5.315	8.944	13.882	19.898	30.175	41.827	57.512	62.724
006-003	01-Oct-15	2.818	2.961	5.400	8.304	14.211	20.669	30.255	41.826	58.506	61.034
014-006	07-Oct-15	2.898	2.870	5.300	8.875	13.473	20.833	29.459	41.326	60.166	60.588
Mean		2.855	2.914	5.340	8.663	13.995	20.809	29.011	41.820	60.396	59.929
SD		0.068	0.054	0.114	0.235	0.262	0.546	1.277	0.632	2.658	1.858
%CV		2.4	1.9	2.1	2.7	1.9	2.6	4.4	1.5	4.4	3.1
% ACCURACY		98.3	100.3	103.0	100.2	97.1	101.1	98.7	99.6	100.7	99.9
n		8	8	8	8	8	8	8	8	8	8

In CMC: Summary of all Aminocaproic Acid Residual Solvents

Solvent	System Precision		Limit of Quantitation				Linearity		Stability of sample at room temperature	
	Spec	Result (%)	Spec	Result (%)	Spec	Result	Spec	Result	Spec (%)	Result (%)
Methanol	% RSD must be NMT 15.0	0.7	% RSD must be NMT 25.0	2.0	S/N must be NLT 10	28	The correlation coefficient value must be ≥ 0.98	1.00	The recovery of stability sample must be 80.0 to 140.0	97.1
Acetone		0.5		1.6		210		1.00		99.2
Isopropyl Alcohol		0.3		1.7		82		1.00		102
Dichloromethane		0.9		2.0		26		1.00		90.2
n-Hexane		1.0		2.1		69		0.99		96.7
Cyclohexanone		0.6		2.9		48		1.00		91.5

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Solvent	Accuracy			Method Precision		Intermediate Precision		Ruggedness	
	Spec (%)	Result (%)	Result (%) (LOQ)	Spec	Result	Spec	Result	Spec	Result
Methanol		94.7 - 100	96.0 - 103.6		3.1%		2.0%		3.63%
Acetone		96.3 - 98	98.4 - 107.3		2.6%		3.3%		2.1%
Isopropyl Alcohol	Recovery 80.0 - 140.0	103.1 - 105.4	98.0 - 105.3	% RSD must be NMT 15.0	1.9%	% RSD must be NMT 15.0	2.0%	% RSD must be NMT 20.0	2.5%
Dichloromethane	Recovery 70.0 - 140.0 at LOQ	93.2 - 96.6	92.4 - 100.9		2.2%		4.0%		3.5%
n-Hexane		110.1 - 115.0	111.4 - 122.4		2.1%		4.4%		15.0%
Cyclohexanone		89.5 - 92.8	93.8 - 101.9		2.6%		2.6%		3.2%