

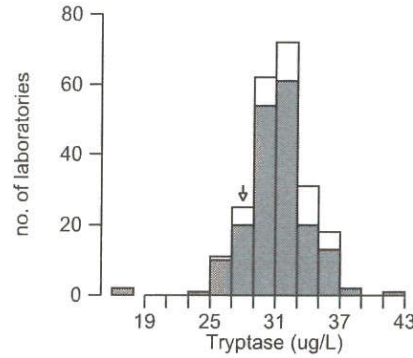
225 / 247 (91%) laboratories returned results for this distribution.
Samples 183-1 and 183-2 were identical samples made from pooled normal human serum with added tryptase.
Laboratories should analyse their data to ensure within assay imprecision between identical samples is acceptable.

☐ All methods
■ Phadia 250

Your MRVIS is 51 ●
Your MRBIS is +10 ●
Your SDBIS is 56 ●

Specimen : 183-1

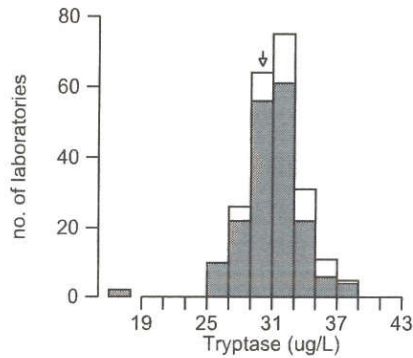
	n	Mean	SD	CV(%)
All methods [ALTM]	225	31.45	2.56	8.1
Phadia 100	11	34.78	1.65	4.7
Phadia 1000	29	31.68	2.03	6.4
Phadia 250	184	31.28	2.54	8.1
Phadia 2500	1	31.40		



Your result 28.9
Target value (ALTM) 31.45
Your BIS -101
Standard Uncertainty 0.21
CCV 8

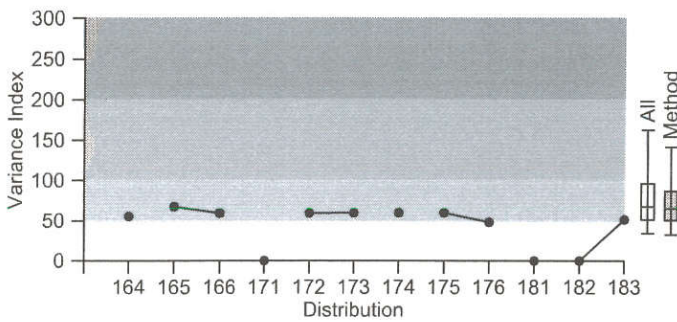
Specimen : 183-2

	n	Mean	SD	CV(%)
All methods [ALTM]	224	31.29	2.35	7.5
Phadia 100	11	35.08	1.72	4.9
Phadia 1000	29	31.32	1.66	5.3
Phadia 250	183	31.08	2.32	7.5
Phadia 2500	1	32.60		

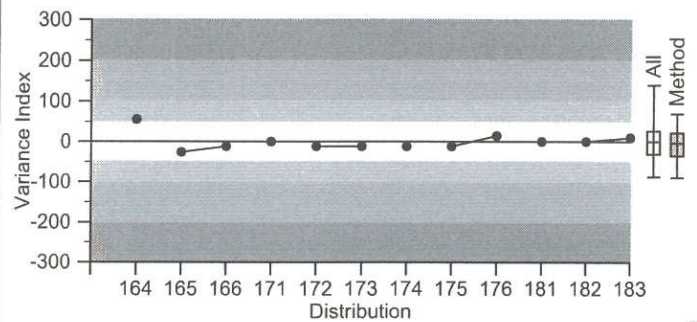


Your result 29.3
Target value (ALTM) 31.29
Your BIS -79
Standard Uncertainty 0.20
CCV 8

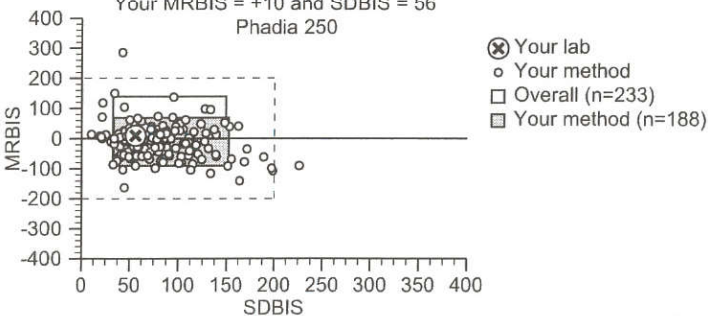
Your MRVIS = +51



Your MRBIS = +10



Scatter plot of MRBIS and SDBIS
Your MRBIS = +10 and SDBIS = 56
Phadia 250



Performance Criteria

- Ideal MRVIS < 50
- Good MRVIS 50-100
- Adequate MRVIS 101-200
- Poor MRVIS > 200 or SDBIS > 200

For further performance criteria information please see our website at www.immqas.org.uk.
If laboratories require further assistance please contact the centre.