

Variables	GE - VSS	GE - VIS	Philips - VSS	Philips - VIS	P-value
<i>Apical Posterior Strain</i>					
<i>N (N missing)</i>	32 (3)	31 (4)	26 (9)	32 (3)	
<i>Mean (SD)</i>	-16.8 (9.2)	-17.4 (10.7)	-19.8 (7.0)	-14.8 (10.7)	0.04
<i>Basal Anteroseptal Strain</i>					
<i>N (N missing)</i>	35 (0)	35 (0)	25 (10)	34 (1)	
<i>Mean (SD)</i>	-16.2 (5.2)	-18.0 (6.8)	-15.1 (4.8)	-15.8 (6.2)	0.20
<i>Basal Posterior Strain</i>					
<i>N (N missing)</i>	29 (6)	28 (7)	25 (10)	26 (9)	
<i>Mean (SD)</i>	-22.1 (7.0)	-20.9 (7.8)	-14.9 (6.3)	-23.7 (7.3)	0.001
<i>Mid Anteroseptal Strain</i>					
<i>N (N missing)</i>	35 (0)	35 (0)	28 (7)	34 (1)	
<i>Mean (SD)</i>	-17.8 (4.8)	-19.0 (6.2)	-15.4 (6.1)	-19.1 (4.8)	0.05
<i>Mid Posterior Strain</i>					
<i>N (N missing)</i>	30 (5)	25 (10)	26 (9)	27 (8)	
<i>Mean (SD)</i>	-21.1 (6.5)	-17.2 (8.9)	-20.2 (8.8)	-18.8 (8.8)	0.07
<u>4 Chamber</u>					
<i>Apical Lateral Strain</i>					
<i>N (N missing)</i>	33 (2)	34 (1)	32 (3)	35 (0)	
<i>Mean (SD)</i>	-21.5 (8.2)	-21.4 (10.7)	-19.7 (5.1)	-22.3 (9.0)	0.18
<i>Apical Septal Strain</i>					
<i>N (N missing)</i>	35 (0)	35 (0)	31 (4)	35 (0)	
<i>Mean (SD)</i>	-15.6 (7.7)	-16.4 (7.2)	-24.8 (6.3)	-18.2 (6.7)	<.0001
<i>Basal Lateral Strain</i>					
<i>N (N missing)</i>	34 (1)	33 (2)	32 (3)	34 (1)	
<i>Mean (SD)</i>	-18.4 (7.5)	-20.7 (8.2)	-14.7 (5.2)	-20.0 (7.9)	<.0001
<i>Basal Septal Strain</i>					
<i>N (N missing)</i>	35 (0)	35 (0)	32 (3)	35 (0)	
<i>Mean (SD)</i>	-15.2 (3.2)	-16.8 (5.8)	-14.9 (3.8)	-16.7 (6.0)	0.15
<i>Mid Lateral Strain</i>					
<i>N (N missing)</i>	34 (1)	34 (1)	32 (3)	33 (2)	
<i>Mean (SD)</i>	-22.1 (6.7)	-19.7 (7.5)	-21.9 (5.2)	-19.1 (9.0)	0.10
<i>Mid Septal Strain</i>					
<i>N (N missing)</i>	35 (0)	35 (0)	32 (3)	35 (0)	
<i>Mean (SD)</i>	-16.1 (4.0)	-18.1 (4.1)	-15.3 (4.0)	-19.6 (5.5)	0.001

Supplementary Table 1. Assessment of effects of scanner and software (and their interaction) by repeated Measures ANOVA.

<i>Variable</i>	<i>Scanner Effect</i>		<i>Software Effect</i>		<i>Interaction</i>	
	<i>F</i>	<i>P-value</i>	<i>F</i>	<i>P-value</i>	<i>F</i>	<i>P-value</i>
<u>Global Strain</u>						
<i>2 Chamber Strain</i>	1.99	0.17	32.48	<0.01	3.08	0.06
<i>3 Chamber Strain</i>	0.02	0.90	1.79	0.18	0.29	0.75
<i>4 Chamber Strain</i>	0.92	0.34	47.82	<0.01	0.15	0.86
<i>Average Strain</i>	1.21	0.28	39.57	<0.01	0.93	0.40
<u>Regional Strain</u>						
2 Chamber						
<i>Apical Anterior Strain</i>	15.98	<0.01	6.15	0.02	6.76	0.01
<i>Apical Inferior Strain</i>	1.00	0.33	0.21	0.65	12.75	<0.01
<i>Basal Anterior Strain</i>	7.98	<0.01	9.46	<0.01	0.83	0.37
<i>Basal Inferior Strain</i>	1.10	0.30	2.70	0.11	3.90	0.06
<i>Mid Anterior Strain</i>	0.15	0.70	31.45	<0.01	0.12	0.73
<i>Mid Inferior Strain</i>	0.03	0.87	1.24	0.27	0.06	0.80
3 Chamber						
<i>Apical Anterior Strain</i>	9.44	<0.01	0.01	0.94	19.42	<0.01
<i>Apical Posterior Strain</i>	0.37	0.55	5.25	0.03	4.08	0.05
<i>Basal Anteroseptal Strain</i>	3.55	0.07	1.83	0.19	0.37	0.55
<i>Basal Posterior Strain</i>	3.46	0.07	6.78	0.02	17.55	<0.01
<i>Mid Anteroseptal Strain</i>	1.16	0.29	7.78	<0.01	2.04	0.16
<i>Mid Posterior Strain</i>	0.10	0.75	4.59	0.04	1.18	0.29
4 Chamber						
<i>Apical Lateral Strain</i>	0.65	0.43	1.12	0.30	4.03	0.05
<i>Apical Septal Strain</i>	40.40	<0.01	7.71	<0.01	14.73	<0.01
<i>Basal Lateral Strain</i>	4.61	0.04	18.33	<0.01	2.20	0.15
<i>Basal Septal Strain</i>	0.07	0.79	5.68	0.02	0.03	0.87
<i>Mid Lateral Strain</i>	0.03	0.86	7.09	0.01	0.05	0.83
<i>Mid Septal Strain</i>	0.35	0.56	21.02	<0.01	2.64	0.11

Supplemental Table 2. Agreement of TomTec output between GE and Philips images. Strains are in %. Positive mean bias indicates Philips was higher (more negative), negative mean bias indicates GE was higher (more negative).

<i>Variable</i>	<i>Mean (SD)</i>	<i>Mean difference (SD_b)</i>	<i>P-value</i>	<i>95% LOA</i>	<i>COV</i>
<u>Global Strain</u>					
<i>2 Chamber Average Strain</i>	-18.5 (4.8)	0.9 (3.7)	0.16	-6.4, 8.3	14.0
<i>2 Chamber LoL Strain</i>	-20.1 (4.9)	-0.0 (3.5)	0.87	-7.0, 6.9	12.2

<i>Variable</i>	<i>Mean (SD)</i>	<i>Mean difference (SD_b)</i>	<i>P-value</i>	<i>95% LOA</i>	<i>COV</i>
3 Chamber Average Strain	-18.1 (4.3)	-0.2 (3.4)	0.72	-7.1, 6.7	13.4
3 Chamber LoL Strain	-19.1 (4.8)	0.0 (3.9)	0.79	-7.7, 7.8	14.4
4 Chamber Average Strain	-18.6 (4.0)	0.4 (3.1)	0.42	-5.8, 6.7	11.9
4 Chamber LoL Strain	-20.1 (4.2)	0.3 (3.0)	0.62	-5.7, 6.3	10.6
Average Strain	-18.4 (4.1)	0.4 (2.3)	0.35	-4.2, 5.0	8.8
LoL Average Strain	-19.8 (4.3)	0.1 (2.3)	0.73	-4.5, 4.8	8.3
<u>Regional Strain</u>					
<u>2 Chamber</u>					
Apical Anterior Strain	-18.0 (8.0)	2.2 (9.0)	0.26	-15.8, 20.1	35.4
Apical Inferior Strain	-22.4 (6.2)	3.3 (9.1)	0.02	-14.8, 21.5	28.6
Basal Anterior Strain	-17.2 (5.5)	1.7 (8.0)	0.16	-14.2, 17.6	32.8
Basal Inferior Strain	-17.8 (7.8)	0.5 (8.0)	0.76	-15.5, 16.5	31.9
Mid Anterior Strain	-18.3 (6.0)	0.7 (8.1)	0.63	-15.5, 16.9	31.4
Mid Inferior Strain	-20.1 (7.1)	0.0 (7.1)	0.96	-14.2, 14.2	24.9
<u>3 Chamber</u>					
Apical Anterior Strain	-20.7 (7.7)	-0.7 (7.2)	0.66	-15.1, 13.7	24.6
Apical Posterior Strain	-16.7 (9.6)	-1.8 (10.3)	0.43	-22.4, 18.7	43.5
Basal Anteroseptal Strain	-16.9 (5.4)	-2.2 (7.6)	0.09	-17.4, 13.0	31.8
Basal Posterior Strain	-22.9 (6.0)	2.6 (8.1)	0.10	-13.6, 18.9	25.2
Mid Anteroseptal Strain	-19.0 (4.1)	0.1 (7.6)	0.92	-15.0, 15.2	28.1
Mid Posterior Strain	-17.4 (8.0)	1.3 (8.7)	0.36	-16.1, 18.7	35.2
<u>4 Chamber</u>					
Apical Lateral Strain	-21.9 (9.1)	0.9 (7.7)	0.52	-14.5, 16.4	25.0
Apical Septal Strain	-17.3 (6.3)	1.8 (5.8)	0.08	-9.8, 13.4	23.7
Basal Lateral Strain	-20.4 (6.4)	-0.2 (9.0)	0.86	-18.3, 17.8	31.3
Basal Septal Strain	-16.7 (4.6)	-0.1 (7.4)	0.95	-14.8, 14.6	31.1
Mid Lateral Strain	-19.7 (7.3)	-0.1 (6.6)	0.81	-13.3, 13.1	23.7
Mid Septal Strain	-18.9 (3.7)	1.5 (6.2)	0.17	-10.9, 13.8	23.2

VIS = vendor-independent software; LOA = limits of agreement; COV = coefficient of variation

Supplemental Table 3. Agreement of global and regional strain values between GE and Philips VSS. Strains are in %. Positive mean bias indicates Philips was higher (more negative), negative mean bias indicates GE was higher (more negative).

<i>Variable</i>	<i>Mean (SD)</i>	<i>Mean difference (SD_b)</i>	<i>P-value</i>	<i>95% LOA</i>	<i>COV</i>
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<i>Variable</i>	<i>Mean (SD)</i>	<i>Mean difference (SD_b)</i>	<i>P-value</i>	<i>95% LOA</i>	<i>COV</i>
<u>Global Strain</u>					
<i>2 Chamber Strain</i>	-18.2 (4.7)	1.6 (3.4)	0.05	-5.2, 8.3	13.1
<i>3 Chamber Strain</i>	-18.5 (4.4)	0.1 (3.1)	0.80	-6.0, 6.3	11.8
<i>4 Chamber Strain</i>	-18.7 (4.1)	0.3 (2.6)	0.36	-5.0, 5.5	9.9
<i>Average Strain</i>	-18.4 (4.1)	0.5 (2.2)	0.15	-3.8, 4.9	8.4
<u>Regional Strain</u>					
<u>2 Chamber</u>					
<i>Apical Anterior Strain</i>	-20.8 (6.4)	8.8 (10.0)	0.00	-11.1, 28.8	33.9
<i>Apical Inferior Strain</i>	-21.8 (5.9)	-1.5 (5.6)	0.08	-12.7, 9.8	18.3
<i>Basal Anterior Strain</i>	-14.3 (5.1)	3.7 (6.0)	0.00	-8.3, 15.8	29.9
<i>Basal Inferior Strain</i>	-16.4 (5.0)	-1.9 (5.1)	0.02	-12.1, 8.4	22.0
<i>Mid Anterior Strain</i>	-14.0 (4.1)	1.2 (6.2)	0.88	-11.2, 13.7	31.6
<i>Mid Inferior Strain</i>	-21.0 (5.4)	0.4 (5.5)	0.74	-10.6, 11.5	18.6
<u>3 Chamber</u>					
<i>Apical Anterior Strain</i>	-20.4 (6.6)	8.3 (9.6)	0.00	-10.9, 27.6	33.4
<i>Apical Posterior Strain</i>	-19.2 (6.6)	3.7 (9.9)	0.08	-16.1, 23.5	36.5
<i>Basal Anteroseptal Strain</i>	-15.7 (3.5)	-1.2 (5.7)	0.34	-12.6, 10.3	25.9
<i>Basal Posterior Strain</i>	-18.7 (5.1)	-7.9 (9.1)	0.00	-26.1, 10.3	34.4
<i>Mid Anteroseptal Strain</i>	-16.4 (4.4)	-2.1 (6.9)	0.09	-15.9, 11.8	29.9
<i>Mid Posterior Strain</i>	-21.0 (6.9)	-0.5 (8.0)	0.59	-16.6, 15.6	27.0
<u>4 Chamber</u>					
<i>Apical Lateral Strain</i>	-20.7 (6.2)	-2.3 (6.4)	0.04	-15.1, 10.5	21.8
<i>Apical Septal Strain</i>	-20.1 (5.5)	9.4 (9.1)	0.00	-8.8, 27.6	32.0
<i>Basal Lateral Strain</i>	-16.7 (5.9)	-3.6 (6.0)	0.00	-15.5, 8.4	25.4
<i>Basal Septal Strain</i>	-15.1 (2.8)	-0.4 (4.4)	0.68	-9.1, 8.4	20.5
<i>Mid Lateral Strain</i>	-22.4 (5.4)	-0.1 (4.3)	0.98	-8.7, 8.4	13.5
<i>Mid Septal Strain</i>	-15.8 (3.3)	-0.8 (4.7)	0.42	-10.2, 8.6	21.1

VSS = vendor-specific software; LOA = limits of agreement; COV = coefficient of variation

Supplemental Table 4. Agreement of global and regional strain values for GE (VSS) with Tomtec (VIS). Strains are in %. Positive mean bias indicates VIS (Tomtec) was higher (more negative), negative mean bias indicates that VSS was higher (more negative).

<i>Variable</i>	<i>Mean (SD)</i>	<i>Mean difference (SD_b)</i>	<i>P-value</i>	<i>95% LOA</i>	<i>COV</i>
<u>Global Strain</u>					

<i>Variable</i>	<i>Mean (SD)</i>	<i>Mean difference (SD_b)</i>	<i>P-value</i>	<i>95% LOA</i>	<i>COV</i>
<i>2 Chamber - VSS Global Strain vs VIS Average Global Strain</i>	<i>-18.1 (5.3)</i>	<i>0.1 (3.1)</i>	<i>0.91</i>	<i>-6.2, 6.3</i>	<i>12.3</i>
<i>2 Chamber - VSS Global Strain vs VIS LoL Global Strain</i>	<i>-19.1 (5.4)</i>	<i>2.2 (3.1)</i>	<i><.01</i>	<i>-4.1, 8.4</i>	<i>11.6</i>
<i>3 Chamber - VSS Global Strain vs VIS Average Global Strain</i>	<i>-18.3 (4.4)</i>	<i>-0.1 (3.3)</i>	<i>0.92</i>	<i>-6.6, 6.4</i>	<i>12.6</i>
<i>3 Chamber - VSS Global Strain vs VIS LoL Global Strain</i>	<i>-18.6 (4.6)</i>	<i>0.5 (3.9)</i>	<i>0.41</i>	<i>-7.2, 8.3</i>	<i>14.7</i>
<i>4 Chamber - VSS Global Strain vs VIS Average Global Strain</i>	<i>-18.4 (4.2)</i>	<i>0.0 (2.9)</i>	<i>0.95</i>	<i>-5.7, 5.8</i>	<i>11.0</i>
<i>4 Chamber - VSS Global Strain vs VIS LoL Global Strain</i>	<i>-19.2 (4.2)</i>	<i>1.6 (2.9)</i>	<i><.01</i>	<i>-4.2, 7.4</i>	<i>10.8</i>
<i>VSS Global Strain vs VIS LoL Global Strain</i>	<i>-19.0 (4.5)</i>	<i>1.4 (2.0)</i>	<i><.01</i>	<i>-2.5, 5.4</i>	<i>7.4</i>
<i>VSS Global Strain vs VIS Average Global Strain</i>	<i>-18.3 (4.4)</i>	<i>0.0 (2.0)</i>	<i>0.98</i>	<i>-3.9, 3.9</i>	<i>7.6</i>
<u>Regional Strain</u>					
<u>2 Chamber</u>					
<i>Apical Anterior Strain</i>	<i>-17.4 (7.3)</i>	<i>0.0 (10.2)</i>	<i>0.86</i>	<i>-20.4, 20.4</i>	<i>41.4</i>
<i>Apical Inferior Strain</i>	<i>-22.2 (7.2)</i>	<i>-2.8 (8.1)</i>	<i>0.09</i>	<i>-18.9, 13.4</i>	<i>25.7</i>
<i>Basal Anterior Strain</i>	<i>-14.7 (6.4)</i>	<i>3.3 (6.4)</i>	<i>0.01</i>	<i>-9.6, 16.1</i>	<i>31.0</i>
<i>Basal Inferior Strain</i>	<i>-17.9 (6.5)</i>	<i>0.3 (8.3)</i>	<i>0.75</i>	<i>-16.4, 17.0</i>	<i>33.0</i>
<i>Mid Anterior Strain</i>	<i>-16.2 (6.6)</i>	<i>3.7 (6.1)</i>	<i><.01</i>	<i>-8.5, 16.0</i>	<i>26.7</i>
<i>Mid Inferior Strain</i>	<i>-20.8 (6.3)</i>	<i>-0.6 (6.2)</i>	<i>0.50</i>	<i>-13.1, 11.9</i>	<i>21.2</i>
<u>3 Chamber</u>					
<i>Apical Anterior Strain</i>	<i>-19.3 (7.3)</i>	<i>4.3 (8.0)</i>	<i>0.01</i>	<i>-11.8, 20.4</i>	<i>29.5</i>
<i>Apical Posterior Strain</i>	<i>-17.9 (8.7)</i>	<i>0.0 (8.7)</i>	<i>0.94</i>	<i>-17.4, 17.4</i>	<i>34.4</i>
<i>Basal Anteroseptal Strain 3</i>	<i>-17.1 (4.9)</i>	<i>1.8 (7.1)</i>	<i>0.14</i>	<i>-12.4, 16.0</i>	<i>29.3</i>
<i>Basal Posterior Strain</i>	<i>-21.8 (5.6)</i>	<i>-1.7 (9.4)</i>	<i>0.42</i>	<i>-20.4, 17.1</i>	<i>30.4</i>
<i>Mid Anteroseptal Strain</i>	<i>-18.4 (4.4)</i>	<i>1.2 (6.9)</i>	<i>0.32</i>	<i>-12.6, 14.9</i>	<i>26.4</i>
<i>Mid Posterior Strain</i>	<i>-19.4 (7.2)</i>	<i>-4.1 (6.6)</i>	<i>0.01</i>	<i>-17.4, 9.2</i>	<i>24.2</i>
<u>4 Chamber</u>					
<i>Apical Lateral Strain</i>	<i>-21.2 (8.8)</i>	<i>-0.5 (7.4)</i>	<i>0.78</i>	<i>-15.4, 14.4</i>	<i>24.9</i>
<i>Apical Septal</i>	<i>-16.0 (6.5)</i>	<i>0.8 (7.4)</i>	<i>0.51</i>	<i>-13.9, 15.6</i>	<i>32.7</i>
<i>Basal Lateral Strain</i>	<i>-19.7 (6.6)</i>	<i>2.0 (8.4)</i>	<i>0.12</i>	<i>-14.7, 18.8</i>	<i>30.0</i>
<i>Basal Septal Strain</i>	<i>-16.0 (3.9)</i>	<i>1.5 (5.2)</i>	<i>0.09</i>	<i>-8.8, 11.9</i>	<i>22.8</i>
<i>Mid Lateral Strain</i>	<i>-20.9 (6.4)</i>	<i>-2.4 (6.2)</i>	<i>0.03</i>	<i>-14.7, 10.0</i>	<i>20.9</i>
<i>Mid Septal Strain</i>	<i>-17.1 (3.1)</i>	<i>2.1 (5.1)</i>	<i>0.02</i>	<i>-8.2, 12.3</i>	<i>21.2</i>

VSS = vendor-specific software; VIS = vendor-independent software; LOA = limits of agreement; COV = coefficient of variation

Supplemental Table 5. Agreement of global and regional strain values for Philips (VSS) with Tomtec (VIS). Strains are in %. Positive mean bias indicates VIS (Tomtec)

was higher (more negative), negative mean bias indicates that VSS was higher (more negative).

<i>Variable</i>	<i>Mean (SD)</i>	<i>Mean difference (SD_b)</i>	<i>P-value</i>	<i>95% LOA</i>	<i>COV</i>
<u>Global Strain</u>					
<i>2 Chamber - VSS Global Strain vs VIS Average Global Strain</i>	<i>-18.4 (4.9)</i>	<i>-0.1 (3.0)</i>	<i>0.48</i>	<i>-6.2, 5.9</i>	<i>11.6</i>
<i>2 Chamber - VSS Global Strain vs VIS LoL Global Strain</i>	<i>-19.3 (4.9)</i>	<i>1.6 (3.2)</i>	<i>0.14</i>	<i>-4.7, 7.9</i>	<i>11.6</i>
<i>3 Chamber - VSS Global Strain vs VIS Average Global Strain</i>	<i>-18.3 (4.2)</i>	<i>-0.3 (3.4)</i>	<i>0.51</i>	<i>-7.0, 6.5</i>	<i>13.1</i>
<i>3 Chamber - VSS Global Strain vs VIS LoL Global Strain</i>	<i>-18.7 (4.5)</i>	<i>0.6 (3.9)</i>	<i>0.41</i>	<i>-7.1, 8.3</i>	<i>14.6</i>
<i>4 Chamber - VSS Global Strain vs VIS Average Global Strain</i>	<i>-18.6 (4.0)</i>	<i>-0.1 (2.9)</i>	<i>0.96</i>	<i>-5.8, 5.6</i>	<i>10.9</i>
<i>4 Chamber - VSS Global Strain vs VIS LoL Global Strain</i>	<i>-19.3 (4.1)</i>	<i>1.4 (2.9)</i>	<i>0.01</i>	<i>-4.4, 7.2</i>	<i>10.7</i>
<i>VSS Global Strain vs VIS Average Global Strain</i>	<i>-18.4 (4.1)</i>	<i>-0.1 (2.0)</i>	<i>0.67</i>	<i>-4.1, 3.9</i>	<i>7.7</i>
<i>VSS Global Strain vs VIS LoL Global Strain</i>	<i>-19.1 (4.2)</i>	<i>1.2 (2.0)</i>	<i>0.01</i>	<i>-2.9, 5.3</i>	<i>7.5</i>
<u>Regional Strain</u>					
<u>2 Chamber</u>					
<i>Apical Anterior Strain</i>	<i>-19.3 (7.3)</i>	<i>-3.2 (10.7)</i>	<i><.01</i>	<i>-24.6, 18.1</i>	<i>38.9</i>
<i>Apical Inferior Strain</i>	<i>-22.2 (6.4)</i>	<i>0.1 (8.0)</i>	<i>0.01</i>	<i>-15.9, 16.0</i>	<i>25.4</i>
<i>Basal Anterior Strain</i>	<i>-15.7 (5.8)</i>	<i>2.5 (6.0)</i>	<i>0.13</i>	<i>-9.5, 14.6</i>	<i>27.2</i>
<i>Basal Inferior Strain</i>	<i>-17.4 (6.4)</i>	<i>1.8 (7.9)</i>	<i>0.01</i>	<i>-14.0, 17.5</i>	<i>32.1</i>
<i>Mid Anterior Strain</i>	<i>-16.4 (5.8)</i>	<i>3.9 (5.7)</i>	<i><.01</i>	<i>-7.4, 15.3</i>	<i>24.5</i>
<i>Mid Inferior Strain</i>	<i>-20.5 (6.1)</i>	<i>-1.1 (6.5)</i>	<i>0.37</i>	<i>-14.0, 11.9</i>	<i>22.2</i>
<u>3 Chamber</u>					
<i>Apical Anterior Strain</i>	<i>-20.4 (7.3)</i>	<i>0.3 (8.9)</i>	<i>0.01</i>	<i>-17.5, 18.1</i>	<i>30.8</i>
<i>Apical Posterior Strain</i>	<i>-17.7 (8.3)</i>	<i>-1.9 (8.6)</i>	<i><.01</i>	<i>-19.0, 15.2</i>	<i>34.2</i>
<i>Basal Anteroseptal Strain</i>	<i>-16.5 (4.6)</i>	<i>1.5 (7.0)</i>	<i>0.58</i>	<i>-12.6, 15.6</i>	<i>30.1</i>
<i>Basal Posterior Strain</i>	<i>-20.5 (5.5)</i>	<i>2.7 (10.7)</i>	<i><.01</i>	<i>-18.7, 24.0</i>	<i>36.8</i>
<i>Mid Anteroseptal Strain</i>	<i>-17.8 (4.4)</i>	<i>2.2 (6.6)</i>	<i>0.01</i>	<i>-11.1, 15.5</i>	<i>26.3</i>
<i>Mid Posterior Strain</i>	<i>-20.0 (7.1)</i>	<i>-2.5 (8.2)</i>	<i>0.48</i>	<i>-19.0, 14.0</i>	<i>29.2</i>
<u>4 Chamber</u>					
<i>Apical Lateral Strain</i>	<i>-21.0 (7.3)</i>	<i>0.9 (8.1)</i>	<i>0.08</i>	<i>-15.4, 17.1</i>	<i>27.4</i>
<i>Apical Septal Strain</i>	<i>-18.5 (6.3)</i>	<i>-2.9 (9.1)</i>	<i><.01</i>	<i>-21.2, 15.4</i>	<i>35.0</i>
<i>Basal Lateral Strain</i>	<i>-18.4 (6.2)</i>	<i>3.6 (8.0)</i>	<i><.01</i>	<i>-12.4, 19.6</i>	<i>30.9</i>
<i>Basal Septal Strain</i>	<i>-15.9 (3.9)</i>	<i>1.6 (5.8)</i>	<i>0.11</i>	<i>-10.0, 13.1</i>	<i>25.7</i>
<i>Mid Lateral Strain</i>	<i>-20.9 (6.4)</i>	<i>-2.4 (6.7)</i>	<i>0.04</i>	<i>-15.9, 11.1</i>	<i>22.8</i>
<i>Mid Septal Strain</i>	<i>-17.2 (3.4)</i>	<i>2.9 (5.5)</i>	<i><.01</i>	<i>-8.0, 13.9</i>	<i>22.5</i>

VSS = vendor-specific software; VIS = vendor-independent software; LOA = limits of agreement; COV = coefficient of variation