

Database Name	Description	Protocol Support
V2-GSRD/TC/NC/F	<p>Database for Tc-99m Sestamibi or Tetrofosmin specific to uncorrected (NC) female patients.</p> <p>Reconstruction algorithm is specific to filtered backprojection (FBP).</p> <p>This database has perfusion and functional information; thus, it can be used for gated and ungated studies.</p>	<ul style="list-style-type: none"> Stress study in a dual isotope protocol. Stress or rest studies in a 2-day Tc-99m labeled radiotracer protocol. Stress or rest study in a 1-day rest/stress Tc-99m labeled radiotracer protocol. Stress or rest study in a 1-day stress/rest Tc-99m labeled radiotracer protocol. <p>WARNING</p> <p>This database should only be used for uncorrected FEMALE studies reconstructed with FBP from data acquired from RAO-to-LPO.</p>
V2-GSRD/TC/NC/M	<p>Database for Tc-99m Sestamibi or Tetrofosmin tracers specific to uncorrected (NC) male patients.</p> <p>Reconstruction algorithm is specific to filtered backprojection (FBP).</p> <p>This database has perfusion and functional information; thus, it can be used for gated and ungated studies.</p>	<ul style="list-style-type: none"> Stress study in a dual isotope protocol. Stress or rest studies in a 2-day Tc-99m labeled radiotracer protocol. Stress or rest study in a 1-day rest/stress Tc-99m labeled radiotracer protocol. Stress or rest study in a 1-day stress/rest Tc-99m labeled radiotracer protocol. <p>WARNING</p> <p>This database should only be used for uncorrected MALE studies reconstructed with FBP from data acquired from RAO-to-LPO.</p>
V-CCAM/TC/NC/F	<p>Database for Tc-99m Sestamibi or Tetrofosmin specific to uncorrected (NC) female patients acquired on the Siemens C.CAM imaging system.</p> <p>Reconstruction algorithm is specific to filtered backprojection (FBP).</p> <p>This database has perfusion and functional information; thus, it can be used for gated and ungated studies.</p>	<ul style="list-style-type: none"> Stress study in a dual isotope protocol. Stress or rest studies in a 2-day Tc-99m labeled radiotracer protocol. Stress or rest study in a 1-day rest/stress Tc-99m labeled radiotracer protocol. Stress or rest study in a 1-day stress/rest Tc-99m labeled radiotracer protocol. <p>WARNING</p> <p>This database should only be used for uncorrected FEMALE studies reconstructed with FBP from data acquired on the Siemens C.CAM imaging system.</p>
V-CCAM/TC/NC/M	<p>Database for Tc-99m Sestamibi or Tetrofosmin database specific to uncorrected (NC) male patients acquired on the Siemens C.CAM imaging system.</p> <p>Reconstruction algorithm is specific to filtered backprojection (FBP).</p> <p>This database has perfusion and functional information; thus, it can be used for gated and ungated studies.</p>	<ul style="list-style-type: none"> Stress study in a dual isotope protocol. Stress or rest studies in a 2-day Tc-99m labeled radiotracer protocol. Stress or rest study in a 1-day rest/stress Tc-99m labeled radiotracer protocol. Stress or rest study in a 1-day stress/rest Tc-99m labeled radiotracer protocol. <p>WARNING</p> <p>This database should only be used for uncorrected MALE studies reconstructed with FBP from data acquired on the Siemens C.CAM imaging system.</p>
V-SRD/TL/NC/F	<p>TI-201 database specific to uncorrected (NC) female patients.</p> <p>Reconstruction algorithm is specific to filtered backprojection (FBP).</p> <p>This database can only be used for analysis of the perfusion data.</p>	<ul style="list-style-type: none"> Rest study in a dual isotope protocol. Stress, rest, delay or redistribution study in a TI-201 only protocol. <p>WARNING</p> <p>This database should only be used for uncorrected FEMALE studies reconstructed with FBP from data acquired from RAO-to-LPO.</p>
V-SRD/TL/NC/M	<p>TI-201 database specific to uncorrected (NC) male patients.</p> <p>Reconstruction algorithm is specific to filtered backprojection (FBP).</p> <p>This database can only be used for analysis of the perfusion data.</p>	<ul style="list-style-type: none"> Rest study in a dual isotope protocol. Stress, rest, delay or redistribution study in a TI-201 only protocol. <p>WARNING</p> <p>This database should only be used for uncorrected MALE studies reconstructed with FBP from data acquired from RAO-to-LPO.</p>

Database Name	Description	Protocol Support
V-CCAM/TL/NC/F	<p>TI-201 database specific to uncorrected (NC) female patients acquired on the Siemens C.CAM imaging system.</p> <p>Reconstruction algorithm is specific to filtered backprojection (FBP).</p> <p>This database can only be used for analysis of the perfusion data.</p>	<ul style="list-style-type: none"> Rest study in a dual isotope protocol. Stress, rest, delay or redistribution study in a TI-201 only protocol. <p>WARNING</p> <p>This database should only be used for uncorrected FEMALE studies reconstructed with FBP from data acquired on the Siemens C.CAM imaging system.</p>
V-CCAM/TL/NC/M	<p>TI-201 database specific to uncorrected (NC) male patients acquired on the Siemens C.CAM imaging system.</p> <p>Reconstruction algorithm is specific to filtered backprojection (FBP).</p> <p>This database can only be used for analysis of the perfusion data.</p>	<ul style="list-style-type: none"> Rest study in a dual isotope protocol. Stress, rest, delay or redistribution study in a TI-201 only protocol. <p>WARNING</p> <p>This database should only be used for uncorrected MALE studies reconstructed with FBP from data acquired on the Siemens C.CAM imaging system.</p>
V-PSRD/TC/AC	<p>Database for Tc-99m Sestamibi or Tetrofosmin specific to Siemens PROFILE attenuation corrected (AC) studies.</p> <p>AC Reconstruction algorithm is specific to the Iter-W algorithm.</p> <p>This database can only be used for analysis of the perfusion data.</p>	<ul style="list-style-type: none"> Stress study in a dual isotope protocol. Stress or rest studies in a 2-day Tc-99m labeled radiotracer protocol. Stress or rest study in a 1-day rest/stress Tc-99m labeled radiotracer protocol. Stress or rest study in a 1-day stress/rest Tc-99m labeled radiotracer protocol. <p>WARNING</p> <p>In addition to the above, this database should only be used for PROFILE attenuation corrected images reconstructed with the Iter-W algorithm.</p>
V-PSRD/TL/AC	<p>TI-201 database specific to Siemens PROFILE attenuation corrected (AC) studies.</p> <p>AC Reconstruction algorithm is specific to the Iter-W algorithm.</p> <p>This database can only be used for analysis of the perfusion data.</p>	<ul style="list-style-type: none"> Rest study in a dual isotope protocol. Stress, rest, delay or redistribution study in a TI-201 only protocol. <p>WARNING</p> <p>In addition to the above, this database should only be used for PROFILE attenuation corrected images reconstructed with the Iter-W algorithm.</p>
V-Symbia/TC/CTAC,SC,RC	<p>Database for Tc-99m Sestamibi or Tetrofosmin tracers specific to Siemens Symbia CT based Attenuation+Scatter+Flash3D corrected studies.</p> <p>This database can only be used for analysis of the perfusion data.</p>	<ul style="list-style-type: none"> Stress study in a dual isotope protocol. Stress or rest studies in a 2-day Tc-99m labeled radiotracer protocol. Stress or rest study in a 1-day rest/stress Tc-99m labeled radiotracer protocol. Stress or rest study in a 1-day stress/rest Tc-99m labeled radiotracer protocol. <p>WARNING</p> <p>In addition to the above, this database should only be used for Symbia studies which have reconstructed incorporating CTAC, scatter, and Flash3D corrections.</p>
V-DIFF12/NC	<p>Comparison database for DIFF12 2-study comparison maps.</p> <p>NOTE</p> <p>This database was not constructed from patient data. It is spatially uniform with a mean of 0.0 and a standard deviation of 4.0. Using a threshold of 2.5σ (default) will provide a uniform defect threshold of 10%.</p>	<p>Any protocol with the knowledge of the NOTE to the left.</p>

Database Name	Description	Protocol Support
V-DIFF21/NC	<p>Comparison database for DIFF21 2-study comparison maps.</p> <p>NOTE</p> <p>This database was not constructed from patient data. It is spatially uniform with a mean of 0.0 and a standard deviation of 4.0. Using a threshold of 2.5σ (default) will provide a uniform defect threshold of 10%.</p>	Any protocol with the knowledge of the NOTE to the left.
V-REVER/NC V-REVER/AC V-REVER/CTAC	<p>Comparison database for Reversibility maps.</p> <p>NOTE</p> <p>These databases were not constructed from patient data. Each is spatially uniform with a mean of 0.0 and a standard deviation of 4.0. Using a threshold of 2.5σ (default) will provide a uniform defect threshold of 10%.</p>	Any protocol with the knowledge of the NOTE to the left.
V-FDG Viability	<p>Comparison database for FDG Viability maps.</p> <p>NOTE</p> <p>This database was not constructed from patient data. It is spatially uniform with a mean of 40.0 and a standard deviation of 4.0. Using a threshold of 2.5σ (default) will provide a uniform defect threshold of 50%.</p>	Any protocol with the knowledge of the NOTE to the left.