# Echo MRI HAROLD\_041\_001

# **Parameters and Metadata**

# Date of procedure HAROLD\_041\_001\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: true Is Annotated: false

Description: date\_of\_procedure

# Weight HAROLD\_041\_001\_002 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: true Is Annotated: false

Unit Measured: g

**Description:** weight

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#### Fat mass HAROLD\_041\_001\_003 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: true Is Annotated: false

Unit Measured: g				
Description: fat_mass				
Lean mass HAROLD_simpleParameter	041_001_004   v1.0			
Req. Analysis: false	Req. Upload: true	Is Annotated: false		
Unit Measured: g				
Description: lean_mass				
Free water HAROLD_041_001_005   v1.0				
simpleParameter				
Req. Analysis: false	Req. Upload: true	Is Annotated: false		
Description: free_water				
Total water HAROLD_	_041_001_006   v1.0			
simpleParameter				

Req. Analysis: false Req. Upload: true Is Annotated: false

**Description:** total\_water

### Percentage fat HAROLD\_041\_001\_701 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: true

**Description:** percentage\_fat

**Derivation:** archived('(HAROLD\_041\_001\_003/HAROLD\_041\_001\_002)\*100')

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### Percentage lean HAROLD\_041\_001\_702 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: true

**Description:** percentage\_lean

**Derivation:** archived('(HAROLD\_041\_001\_004/HAROLD\_041\_001\_002)\*100')

### Percentage free water HAROLD\_041\_001\_703 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: true

**Description:** percentage\_free\_water

Derivation: archived('(HAROL	_D_041_001_005/HAROLD_04	l1_001_002)*100')		
Percentage total water HAROLD_041_001_704   v1.0 simpleParameter				
Req. Analysis: false	Req. Upload: false	Is Annotated: true		
Description: percentage_total_water				
<b>Derivation:</b> archived('(HAROLD_041_001_006/HAROLD_041_001_002)*100')				