

# MeSH Tree Structures - 2008

## G6 - BIOCHEMICAL PHENOMENA, METABOLISM, AND NUTRITION

### Biochemical Phenomena, Metabolism, and Nutrition

<b>Biochemical Phenomena, Metabolism, and Nutrition</b>	G6		
<b>Biochemical Phenomena</b>	G6.184		
<b>Acid-Base Equilibrium</b>	G6.184.29	G7.621. H1.181.	G9.188.
<b>Aerobiosis</b>	G6.184.35		
<b>Allosteric Regulation</b>	G6.184.60		
<b>Amino Acid Substitution</b>	G6.184.65	E5.393.	
<b>Anaerobiosis</b>	G6.184.70		
<b>Base Composition</b>	G6.184.124		
<b>Binding Sites</b>	G6.184.154		
<b>Allosteric Site</b>	G6.184.154.147		
<b>Bay Region (Chemistry)</b>	G6.184.154.290		
<b>Binding, Competitive</b>	G6.184.154.309		
<b>Binding Sites, Antibody</b>	G6.184.154.408	D12.776. D12.776.	D12.776. G4.610.
<b>Body Composition</b>	G6.184.179		
<b>Body Fat Distribution</b>	G6.184.179.134	E1.370.	
<b>Adiposity</b>	G6.184.179.134.500	E1.370.	
<b>Body Fluid Compartments</b>	G6.184.179.180	A12.207.	
<b>Brain Chemistry</b>	G6.184.202		
<b>Calcification, Physiologic</b>	G6.184.227	G7.574.	G11.427.
<b>Tooth Calcification</b>	G6.184.227.710	G10.549.	
<b>Diffusion</b>	G6.184.307	H1.671.	
<b>Down-Regulation</b>	G6.184.320	G5.315. G12.307	G7.621.
<b>Energy Transfer</b>	G6.184.348	H1.181.	H1.671.
<b>Linear Energy Transfer</b>	G6.184.348.400	H1.181.	H1.671.
<b>Enzyme Activation</b>	G6.184.368	G6.535.	
<b>Enzyme Stability</b>	G6.184.378	E5.916.	
<b>Fermentation</b>	G6.184.405	G6.535.	
<b>Ion Transport</b>	G6.184.480	G6.535.	
<b>Calcium Signaling</b>	G6.184.480.100	G4.335. G6.535.	G6.184.
<b>Membrane Fluidity</b>	G6.184.568		
<b>Molecular Mimicry</b>	G6.184.589		
<b>Molecular Structure</b>	G6.184.603	H1.181.	
<b>Amino Acid Sequence</b>	G6.184.603.60	L1.453.	
<b>Amino Acid Motifs</b>	G6.184.603.60.40	G6.184.	
<b>F-Box Motifs</b>	G6.184.603.60.40.500	G6.184.	
<b>Extensins</b>	G6.184.603.60.270		
<b>Histone Code</b>	G6.184.603.60.360	G14.360	
<b>Immunoglobulin Variable Region</b>	G6.184.603.60.425	D12.776. D12.776. D12.776. D12.776. D12.776. D12.776.	D12.776. D12.776. D12.776. D12.776. D12.776.
<b>Complementarity Determining Regions</b>	G6.184.603.60.425.160	D12.776. D12.776. D12.776. D12.776.	D12.776. D12.776. D12.776. D12.776.
<b>Inteins</b>	G6.184.603.60.440		
<b>Peptide Library</b>	G6.184.603.60.620	D12.644.	G14.325.
<b>Protein Sorting Signals</b>	G6.184.603.60.670	D12.644.	
<b>Nuclear Export Signals</b>	G6.184.603.60.670.600	D12.644.	
<b>Nuclear Localization Signals</b>	G6.184.603.60.670.610	D12.644.	
<b>Repetitive Sequences, Amino Acid</b>	G6.184.603.60.720		
<b>Ankyrin Repeat</b>	G6.184.603.60.720.30	G6.184.	G6.184.
<b>Base Sequence</b>	G6.184.603.80	G14.80	L1.453.
<b>AT Rich Sequence</b>	G6.184.603.80.40	G14.80.	
<b>GC Rich Sequence</b>	G6.184.603.80.380	G14.80.	

## G6 - BIOCHEMICAL PHENOMENA, METABOLISM, AND NUTRITION

### Biochemical Phenomena, Metabolism, and Nutrition

#### Biochemical Phenomena

##### Molecular Structure

##### Base Sequence

##### GC Rich Sequence

##### CpG Islands

##### CpG Islands

##### Matrix Attachment Regions

##### Regulatory Sequences, Nucleic Acid

##### Enhancer Elements (Genetics)

##### E-Box Elements

##### HIV Enhancer

##### Response Elements

##### Serum Response Element

##### Vitamin D Response Element

##### Insulator Elements

##### Locus Control Region

##### Operator Regions (Genetics)

##### Promoter Regions (Genetics)

##### Response Elements

##### Serum Response Element

##### Vitamin D Response Element

##### TATA Box

##### Regulatory Sequences, Ribonucleic Acid

##### RNA 3' Polyadenylation Signals

##### RNA Splice Sites

##### RNA 5' Terminal Oligopyrimidine Sequence

##### Silencer Elements, Transcriptional

##### Terminator Regions (Genetics)

##### Repetitive Sequences, Nucleic Acid

##### Interspersed Repetitive Sequences

##### DNA Transposable Elements

##### Integrans

##### Genomic Islands

##### Retroelements

##### Endogenous Retroviruses

##### Genes, Intracisternal A-Particle

##### Long Interspersed Nucleotide Elements

##### Short Interspersed Nucleotide Elements

##### Alu Elements

##### Tandem Repeat Sequences

##### DNA Repeat Expansion

G6.184.603.80.380.160	G14.80.	G14.340.
G6.184.603.80.534	G14.80.	
G6.184.603.80.689	G14.80.	
G6.184.603.80.689.330	G14.80.	G14.340.
G6.184.603.80.689.330.240	G14.80.	G14.340.
G6.184.603.80.689.330.400	G6.184.	G14.80.
	G14.80.	G14.340.
G6.184.603.80.689.330.700	G6.184.	G14.80.
	G14.80.	G14.340.
	G14.340.	
G6.184.603.80.689.330.700.800	G6.184.	G14.80.
	G14.80.	G14.340.
	G14.340.	
G6.184.603.80.689.330.700.920	G6.184.	G14.80.
	G14.80.	G14.340.
	G14.340.	
G6.184.603.80.689.390	G14.80.	G14.340.
G6.184.603.80.689.450	G14.80.	G14.340.
G6.184.603.80.689.650	G14.80.	G14.340.
	G14.340.	
G6.184.603.80.689.675	G14.80.	G14.340.
G6.184.603.80.689.675.700	G6.184.	G14.80.
	G14.80.	G14.340.
	G14.340.	
G6.184.603.80.689.675.700.800	G6.184.	G14.80.
	G14.80.	G14.340.
	G14.340.	
G6.184.603.80.689.675.700.920	G6.184.	G14.80.
	G14.80.	G14.340.
	G14.340.	
G6.184.603.80.689.675.850	G14.80.	G14.340.
G6.184.603.80.689.687	D13.444.	G14.80.
G6.184.603.80.689.687.249	G14.80.	G14.340.
G6.184.603.80.689.687.490	D13.444.	G14.80.
	G14.340.	
G6.184.603.80.689.687.500	G14.80.	G14.340.
G6.184.603.80.689.755	G14.80.	G14.340.
G6.184.603.80.689.810	G14.80.	G14.340.
G6.184.603.80.708	G14.80.	
G6.184.603.80.708.330	G14.80.	G14.340.
G6.184.603.80.708.330.200	D13.444.	G14.80.
	G14.340.	
G6.184.603.80.708.330.200.500		
G6.184.603.80.708.330.330	G14.80.	G14.340.
G6.184.603.80.708.330.800	D13.444.	G14.80.
	G14.340.	
G6.184.603.80.708.330.800.175	B4.820.	B4.909.
	B4.909.	G14.80.
	G14.340.	
G6.184.603.80.708.330.800.200	G14.80.	G14.340.
	G14.340.	
G6.184.603.80.708.330.800.400	G14.80.	G14.340.
G6.184.603.80.708.330.800.800	G14.80.	G14.340.
G6.184.603.80.708.330.800.800.50	G14.80.	G14.340.
G6.184.603.80.708.800	G14.80.	G14.340.
G6.184.603.80.708.800.140	G5.600.	G13.920.
	G14.80.	G14.340.
	G14.340.	

## G6 - BIOCHEMICAL PHENOMENA, METABOLISM, AND NUTRITION

### Biochemical Phenomena, Metabolism, and Nutrition

#### Biochemical Phenomena

##### Molecular Structure

##### Base Sequence

##### Repetitive Sequences, Nucleic Acid

##### Tandem Repeat Sequences

##### Trinucleotide Repeat Expansion

G6.184.603.80.708.800.140.865

G5.600.  
G13.920.  
G14.80.  
G14.340.

G6.184.  
G14.80.  
G14.340.

##### DNA, Satellite

G6.184.603.80.708.800.150

D13.444.  
G14.340.

G14.80.  
G14.340.

##### Microsatellite Repeats

G6.184.603.80.708.800.500

G14.80.

G14.340.

##### Dinucleotide Repeats

G6.184.603.80.708.800.500.150

G14.80.

G14.340.

##### Trinucleotide Repeats

G6.184.603.80.708.800.500.850

G14.80.

G14.340.

##### Trinucleotide Repeat Expansion

G6.184.603.80.708.800.500.850.200

G5.600.  
G13.920.  
G14.80.  
G14.340.

G6.184.  
G14.80.  
G14.340.

##### Minisatellite Repeats

G6.184.603.80.708.800.550

G14.80.

G14.340.

##### Terminal Repeat Sequences

G6.184.603.80.708.850

G14.80.

##### HIV Long Terminal Repeat

G6.184.603.80.708.850.400

G14.80.

##### HIV Enhancer

G6.184.603.80.708.850.400.400

G6.184.  
G14.80.

G14.80.  
G14.340.

##### Carbohydrate Sequence

G6.184.603.160

L1.453.

##### Conserved Sequence

G6.184.603.580

##### Consensus Sequence

G6.184.603.580.175

##### Molecular Conformation

G6.184.603.790

H1.181.

##### Carbohydrate Conformation

G6.184.603.790.235

##### Nucleic Acid Conformation

G6.184.603.790.486

##### Base Pairing

G6.184.603.790.486.100

##### DNA, Cruciform

G6.184.603.790.486.325

D13.444.

##### G-Quadruplexes

G6.184.603.790.486.550

##### Protein Conformation

G6.184.603.790.709

##### Protein Structure, Quaternary

G6.184.603.790.709.550

##### Protein Structure, Secondary

G6.184.603.790.709.600

##### Amino Acid Motifs

G6.184.603.790.709.600.40

G6.184.

##### Ankyrin Repeat

G6.184.603.790.709.600.40.30

G6.184.

G6.184.

##### AT-Hook Motifs

G6.184.603.790.709.600.40.50

##### Cystine Knot Motifs

G6.184.603.790.709.600.40.127

##### F-Box Motifs

G6.184.603.790.709.600.40.205

G6.184.

##### Helix-Loop-Helix Motifs

G6.184.603.790.709.600.40.360

##### EF Hand Motifs

G6.184.603.790.709.600.40.360.240

##### Helix-Turn-Helix Motifs

G6.184.603.790.709.600.40.360.360

##### Leucine Zippers

G6.184.603.790.709.600.40.520

##### Zinc Fingers

G6.184.603.790.709.600.40.985

##### RING Finger Domains

G6.184.603.790.709.600.40.985.500

G6.184.

##### Protein Structure, Tertiary

G6.184.603.790.709.610

##### Catalytic Domain

G6.184.603.790.709.610.189

##### HMG-Box Domains

G6.184.603.790.709.610.380

##### Kringles

G6.184.603.790.709.610.480

##### Protein Interaction Domains and Motifs

G6.184.603.790.709.610.640

##### Ankyrin Repeat

G6.184.603.790.709.610.640.80

G6.184.

G6.184.

##### PDZ Domains

G6.184.603.790.709.610.640.500

##### RING Finger Domains

G6.184.603.790.709.610.640.625

G6.184.

##### src Homology Domains

G6.184.603.790.709.610.640.750

##### Structural Homology, Protein

G6.184.603.790.709.805

G6.184.

G13.820

##### Nitrogen Fixation

G6.184.620

G4.185.  
G6.535.

G4.185.

##### Nitrosation

G6.184.627

G6.535.

H1.181.

##### Nucleic Acid Denaturation

G6.184.642

E5.393.

G5.720

##### Nucleic Acid Hybridization

G6.184.661

E5.393.

##### Nucleic Acid Renaturation

G6.184.680

##### Osmosis

G6.184.708

H1.181.

##### Protein Binding

G6.184.775

G6.535.

## G6 - BIOCHEMICAL PHENOMENA, METABOLISM, AND NUTRITION

### Biochemical Phenomena, Metabolism, and Nutrition

#### Biochemical Phenomena

##### Protein Denaturation

Protein Denaturation	G6.184.794		
Protein Folding	G6.184.797		
Protein Hybridization	G6.184.800		
Protein Renaturation	G6.184.807		
RNA Stability	G6.184.835		
Sequence Homology	G6.184.842	G13.810	
Sequence Homology, Amino Acid	G6.184.842.200	G13.810.	
Structural Homology, Protein	G6.184.842.200.820	G6.184.	G13.820
Sequence Homology, Nucleic Acid	G6.184.842.550	G13.810.	
Synteny	G6.184.842.550.830	G13.810.	
Signal Transduction	G6.184.850	G4.335.	G4.335.
Ion Channel Gating	G6.184.850.400	G4.335.	G4.335.
MAP Kinase Signaling System	G6.184.850.560	G4.335.	
Mechanotransduction, Cellular	G6.184.850.580	G4.335.	G4.335.
Phototransduction	G6.184.850.700	G4.335.	G11.697.
Second Messenger Systems	G6.184.850.800	G4.335.	
Calcium Signaling	G6.184.850.800.100	G4.335.	G6.184.
		G6.535.	
Synaptic Transmission	G6.184.850.850	G4.335.	G11.561.
Structure-Activity Relationship	G6.184.872	G12.869	H1.158.
		H1.181.	H1.181.
Quantitative Structure-Activity Relationship	G6.184.872.500	G12.869.	H1.158.
		H1.181.	H1.181.
Substrate Specificity	G6.184.880		
Up-Regulation	G6.184.910	G5.315.	G7.621.
		G12.955	
Virus Replication	G6.184.941	G4.185.	
Virus Assembly	G6.184.941.950	G4.185.	
Water-Electrolyte Balance	G6.184.963	G7.621.	
Kallikrein-Kinin System	G6.184.963.500	G6.535.	G6.535.
		G7.621.	G9.330.
Water Loss, Insensible	G6.184.963.750	G7.621.	
Metabolism	G6.535		
Absorption	G6.535.23	H1.181.	
Intestinal Absorption	G6.535.23.500	G6.696.	G10.261.
Skin Absorption	G6.535.23.750	G10.795.	
Acylation	G6.535.95	H1.181.	
Acetylation	G6.535.95.52	H1.181.	
Aminoacylation	G6.535.95.55	G6.535.	H1.181.
Transfer RNA Aminoacylation	G6.535.95.55.860	G6.535.	G6.535.
		H1.181.	
Alkylation	G6.535.130	H1.181.	
Methylation	G6.535.130.538	H1.181.	
DNA Methylation	G6.535.130.538.161	G5.190	H1.181.
Amination	G6.535.140	H1.181.	
Autotrophic Processes	G6.535.153		
Chemoautotrophic Growth	G6.535.153.314		
Nitrogen Fixation	G6.535.153.630	G4.185.	G4.185.
		G6.184.	
Biological Transport	G6.535.166		
Biological Transport, Active	G6.535.166.310		
Active Transport, Cell Nucleus	G6.535.166.310.100	G6.535.	
Capillary Permeability	G6.535.166.330	G9.330.	
Cell Membrane Permeability	G6.535.166.335	G4.335.	
Cytoplasmic Streaming	G6.535.166.355	G4.335.	
Axonal Transport	G6.535.166.355.40	G4.335.	G11.561.
Ion Transport	G6.535.166.500	G6.184.	
Calcium Signaling	G6.535.166.500.100	G4.335.	G6.184.
		G6.184.	

## G6 - BIOCHEMICAL PHENOMENA, METABOLISM, AND NUTRITION

### Biochemical Phenomena, Metabolism, and Nutrition

#### Metabolism

##### Biological Transport

##### Protein Transport

##### Protein Transport

##### Active Transport, Cell Nucleus

##### Respiratory Transport

##### Pulmonary Gas Exchange

##### RNA Transport

##### Biotinylation

##### Carbohydrate Metabolism

##### Fermentation

##### Gluconeogenesis

##### Glycogenolysis

##### Glycolysis

##### Pentose Phosphate Pathway

##### Photosynthesis

##### Photophosphorylation

##### Cyclization

##### Dealkylation

##### Deamination

##### Decarboxylation

##### Dimerization

##### Electron Transport

##### Energy Metabolism

##### Basal Metabolism

##### Citric Acid Cycle

##### Glycolysis

##### Oxidation-Reduction

##### Electron Transport

##### Lipid Peroxidation

##### Oxidative Phosphorylation

##### Pentose Phosphate Pathway

##### Photophosphorylation

##### Proton-Motive Force

##### Membrane Potential, Mitochondrial

##### Substrate Cycling

##### Enterohepatic Circulation

##### Enzyme Activation

##### Esterification

##### Glycosylation

##### Halogenation

##### Heterotrophic Processes

##### Hydrogenation

##### Hydrolysis

##### Hydroxylation

##### Kallikrein-Kinin System

##### Lipid Metabolism

##### Lipogenesis

##### Lipolysis

##### Lipid Mobilization

##### Lipoylation

##### Metabolic Networks and Pathways

##### Biosynthetic Pathways

##### Citric Acid Cycle

##### Electron Transport

##### Glycolysis

	G6.535.166.700		
	G6.535.166.700.100	G6.535.	
	G6.535.166.775	G9.772.	
	G6.535.166.775.602	E1.370.	G9.772.
	G6.535.166.850		
	G6.535.233	E5.393.	
	G6.535.256		
	G6.535.256.249	G6.184.	
	G6.535.256.500		
	G6.535.256.625		
	G6.535.256.750	G6.535.	G6.535.
	G6.535.256.875	G6.535.	G6.535.
	G6.535.256.937	G4.742.	G6.535.
		G6.535.	H1.181.
	G6.535.256.937.700	G6.535.	G6.535.
		G6.535.	H1.181.
	G6.535.300	H1.181.	
	G6.535.305	H1.181.	
	G6.535.310	H1.181.	
	G6.535.315	H1.181.	
	G6.535.320	H1.181.	
	G6.535.327	G6.535.	G6.535.
	G6.535.335		
	G6.535.335.154	E1.370.	
	G6.535.335.342	G6.535.	
	G6.535.335.436	G6.535.	G6.535.
	G6.535.335.531	H1.181.	
	G6.535.335.531.403	G6.535.	G6.535.
	G6.535.335.531.587		
	G6.535.335.631	G6.535.	
	G6.535.335.693	G6.535.	G6.535.
	G6.535.335.724	G6.535.	G6.535.
		G6.535.	H1.181.
	G6.535.335.770	H1.181.	
	G6.535.335.770.500	G7.453.	
	G6.535.335.835	G6.535.	
	G6.535.355		
	G6.535.368	G6.184.	
	G6.535.475	H1.181.	
	G6.535.488	H1.181.	
	G6.535.496		
	G6.535.505		
	G6.535.506	H1.181.	
	G6.535.508	H1.181.	
	G6.535.510	H1.181.	
	G6.535.516	G6.184.	G6.535.
		G7.621.	G9.330.
	G6.535.520		
	G6.535.520.249		
	G6.535.520.500		
	G6.535.520.500.500		
	G6.535.536		
	G6.535.553		
	G6.535.553.100		
	G6.535.553.170	G6.535.	
	G6.535.553.350	G6.535.	G6.535.
	G6.535.553.360	G6.535.	G6.535.

## G6 - BIOCHEMICAL PHENOMENA, METABOLISM, AND NUTRITION

### Biochemical Phenomena, Metabolism, and Nutrition

#### Metabolism

##### Metabolic Networks and Pathways

##### Kallikrein-Kinin System

Kallikrein-Kinin System	G6.535.553.485	G6.184. G7.621.	G6.535. G9.330.
Pentose Phosphate Pathway	G6.535.553.695	G6.535.	G6.535.
Photosynthesis	G6.535.553.700	G4.742. G6.535.	G6.535. H1.181.
Nitrosation	G6.535.655	G6.184.	H1.181.
Oxidative Stress	G6.535.710		
Protein Carbonylation	G6.535.710.690		
Oxygen Consumption	G6.535.723		
Anaerobic Threshold	G6.535.723.110	G11.427.	
Respiratory Burst	G6.535.723.620	G4.335.	
Peptide Biosynthesis	G6.535.770		
Aminoacylation	G6.535.770.50	G6.535.	H1.181.
Transfer RNA Aminoacylation	G6.535.770.50.860	G6.535. H1.181.	G6.535.
Peptide Biosynthesis, Nucleic Acid-Independent	G6.535.770.333		
Protein Biosynthesis	G6.535.770.871	G5.310.	
Frameshifting, Ribosomal	G6.535.770.871.200	G5.315.	
Peptide Chain Elongation, Translational	G6.535.770.871.640		
Peptide Chain Initiation, Translational	G6.535.770.871.650		
Peptide Chain Termination, Translational	G6.535.770.871.720		
Protein Modification, Translational	G6.535.770.871.790	G5.315.	
Protein Processing, Post-Translational	G6.535.770.871.790.600	G5.315.	
Protein Prenylation	G6.535.770.871.790.600.400	G5.315.	G6.535.
Protein Splicing	G6.535.770.871.790.600.700	G5.315.	
Transfer RNA Aminoacylation	G6.535.770.871.850	G6.535. H1.181.	G6.535.
Pharmacokinetics	G6.535.780	G12.91.	
Area Under Curve	G6.535.780.101	E5.318. G12.91.	G3.850.
Biological Availability	G6.535.780.151	G12.91.	
Biotransformation	G6.535.780.202	G12.91.	
Metabolic Clearance Rate	G6.535.780.202.640	E1.450.	G12.91.
Metabolic Detoxication, Drug	G6.535.780.202.820	G12.91.	
Metabolic Detoxication, Phase I	G6.535.780.202.820.500	G12.91.	
Metabolic Detoxication, Phase II	G6.535.780.202.820.750	G12.91.	
Therapeutic Equivalency	G6.535.780.559	G12.91.	
Tissue Distribution	G6.535.780.917	G6.535.	G12.91.
Phosphorylation	G6.535.790	H1.181.	
Oxidative Phosphorylation	G6.535.790.550	G6.535.	
Photophosphorylation	G6.535.790.605	G6.535. G6.535.	G6.535. H1.181.
Phototrophic Processes	G6.535.795		
Photosynthesis	G6.535.795.700	G4.742. G6.535.	G6.535. H1.181.
Photophosphorylation	G6.535.795.700.700	G6.535. G6.535.	G6.535. H1.181.
Prenylation	G6.535.797		
Protein Prenylation	G6.535.797.500	G5.315.	G6.535.
Protein Binding	G6.535.800	G6.184.	
Renin-Angiotensin System	G6.535.805	G9.330.	
RNA Processing, Post-Transcriptional	G6.535.839	G5.315.	
RNA 3' End Processing	G6.535.839.225	G5.315.	
Polyadenylation	G6.535.839.225.710	G5.315.	
RNA Editing	G6.535.839.250	G5.315.	
RNA Splicing	G6.535.839.700	G5.315.	
Alternative Splicing	G6.535.839.700.100	G5.315.	
Trans-Splicing	G6.535.839.700.750	G5.315.	
Secretory Rate	G6.535.873		

## G6 - BIOCHEMICAL PHENOMENA, METABOLISM, AND NUTRITION

### Biochemical Phenomena, Metabolism, and Nutrition

#### Metabolism

##### Substrate Cycling

Substrate Cycling	G6.535.890	G6.535.	
Tissue Distribution	G6.535.910	G6.535.	G12.91.
Ubiquitination	G6.535.955		
Nutrition Physiology	G6.696		
Animal Nutrition Physiology	G6.696.161		
Child Nutrition Physiology	G6.696.220		
Adolescent Nutrition Physiology	G6.696.220.60		
Infant Nutrition Physiology	G6.696.220.500		
Bottle Feeding	G6.696.220.500.249	E2.421.	G6.696.
Breast Feeding	G6.696.220.500.500	G6.696.	
Weaning	G6.696.220.500.750	G6.696.	
Maternal Nutrition Physiology	G6.696.566		
Prenatal Nutrition Physiology	G6.696.566.624	G8.520.	
Nutrition Phenomena	G6.696.648		
Diet	G6.696.648.240	E5.272	
Diabetic Diet	G6.696.648.240.240	E2.642.	
Diet, Carbohydrate-Restricted	G6.696.648.240.245	E2.642.	
Diet Fads	G6.696.648.240.250	E2.642.	13.287
Diet, Fat-Restricted	G6.696.648.240.260	E2.642.	
Diet, Mediterranean	G6.696.648.240.270	E2.642.	
Diet, Protein-Restricted	G6.696.648.240.280	E2.642.	
Diet, Reducing	G6.696.648.240.285	E2.642.	
Diet, Sodium-Restricted	G6.696.648.240.290	E2.642.	
Diet, Vegetarian	G6.696.648.240.300	E2.642.	
Diet, Macrobiotic	G6.696.648.240.300.500	E2.642.	
Energy Intake	G6.696.648.240.340		
Caloric Restriction	G6.696.648.240.340.150	E2.642.	
Hunger	G6.696.648.390	F1.658.	G10.261.
Appetite	G6.696.648.390.70	F2.830.	G10.261.
Appetite Regulation	G6.696.648.390.70.290	G6.696.	G10.261.
Nutritional Requirements	G6.696.648.620		
Nutritional Status	G6.696.648.810	N1.224.	
Nutritive Value	G6.696.648.905		
Glycemic Index	G6.696.648.905.500		
Nutrition Processes	G6.696.692		
Appetite Regulation	G6.696.692.80	G6.696.	G10.261.
Bottle Feeding	G6.696.692.150	E2.421.	G6.696.
Breast Feeding	G6.696.692.185	G6.696.	
Digestion	G6.696.692.220	G10.261.	
Salivation	G6.696.692.220.800	G10.261.	
Eating	G6.696.692.260	G10.261.	
Drinking	G6.696.692.260.249	G10.261.	
Mastication	G6.696.692.260.500	G10.261.	
Intestinal Absorption	G6.696.692.431	G6.535.	G10.261.
Weaning	G6.696.692.762	G6.696.	