| Summary of November Update to 2017 MeSH: | | | |
|--|---|---|--|
| Field Changed | Old value | New Value | |
| Annotation | IM GEN only; NIM with specific organ / pathol (IM) + dis / pathol (IM); coord with specific technique if appropriate | usually NIM with specific organ / pathol + disease /diag and/or /pathol | |
| Annotation | usually NIM with specific organ / pathol (IM) + disease / pathol (IM) | usually NIM with specific organ / pathol + disease /diag and/or /pathol | |
| Annotation | usually NIM with specific organ/pathol (IM) + disease/pathol (IM) | usually NIM with specific organ / pathol + disease /diag and/or /pathol | |
| Annotation | usually NIM with specific organ/ pathol (IM) + disease /pathol (IM) | usually NIM with specific organ / pathol + disease /diag and/or /pathol | |
| Annotation | never IM; coord NIM with BASE SEQUENCE or AMINO ACID SEQUENCE or CARBOHYDRATE SEQUENCE (NIM) + specific nucleotide or protein or carbohydrate (IM); index here only for base sequences containing 50 or more nucleotides (exclude primers or probes), only for amino acid sequences containing 15 or more amino acids, only for carbohydrate sequences containing 3 or more carbohydrate units (trisaccharides); DF: MOL SEQ DATA | index only for general articles on sequence data; 1988-2016 indexed for articles which contained databank accession numbers for sequences deposited in a molecular sequence databank such as Genbank or for the presence of molecular sequences in the text | |
| | Annotation Annotation Annotation | Annotation IM GEN only; NIM with specific organ / pathol (IM) + dis / pathol (IM); coord with specific technique if appropriate Annotation usually NIM with specific organ / pathol (IM) + disease / pathol (IM) Annotation usually NIM with specific organ/pathol (IM) + disease/pathol (IM) Annotation usually NIM with specific organ/pathol (IM) + disease / pathol (IM) Annotation never IM; coord NIM with BASE SEQUENCE or AMINO ACID SEQUENCE or CARBOHYDRATE SEQUENCE (NIM) + specific nucleotide or protein or carbohydrate (IM); index here only for base sequences containing 50 or more nucleotides (exclude primers or probes), only for amino acid sequences containing 15 or more amino acids, only for carbohydrate sequences containing 3 or more carbohydrate units | |

| Iridium Radioisotopes | Annotation | Ir-182-190, 192, 194-198; / blood-csf-urine / Ir-182-190, 192, 194-198; Ir-182-190, 192, 194-198; for | |
|-----------------------|------------|---|--|
| | | metab permitted but not for tracer studies (routine use as tracer, use no qualifiers | |
| | | = / diag use) | |