3328	Amide A: N-H and O-H stretching vibrations of polysaccharides, proteins
3015	Olefin =CH stretching: unsaturated lipids, cholesterol esters
2960	CH ₃ asymmetric stretching: mainly lipids with the little contribution from carbohydrates, proteins, nucleic acids.
2920	CH2 asymmetric stretching: mainly lipids
2875	CH ₃ symmetric stretching: protein side chains, lipids
2850	CH2 symmetric stretching: mainly lipids

Supplementary Table - 1 Wavenumber (cm=1)

1720-1745

1710-1690

1705-1690

1654

1630-1640

1610 1578

1540-1550

1530

1467

1455

1370-1400

1330-1200

1230-1244

1090-1080

1060, 1050

996

965

CH2 symmetric stretching: mainly lipids C=O stretching vibrations of lipids (triglycerides and cholesterol esters) C=O asymmetric stretching: RNA and purine base C=O asymmetric stretching vibrations: RNA. DNA

Assignment

bending vibrations: proteins a-helix Amide I: (80%) and C-N (10%) stretching, N-H (10%) bending vibrations: proteins B-structure

Amide I: C=O (80%) and C-N (10%) stretching, N-H (10%) C4-C5 and C=N stretching in imidazole ring of DNA, RNA 1515 Aromatic tyrosine rine Amide II: N-H (60%) bending and C-N (40%) stretching vibrations: proteins α-helix Amide II: N-H (60%) bending and C-N (40%) stretching vibrations:

proteins B-structure CH2 bending vibrations: lipids and proteins CH3 bending and CH2 seissoring vibrations: lipids and proteins

COO- symmetric stretching and CH3 bending vibrations: lipids. Proteins Amide III: proteins PO3- antisymmetric stretching vibrations; RNA, DNA and

phospholipids PO2- symmetric stretching vibrations; RNA, DNA C-O stretching vibrations: deoxyribose/ribose DNA, RNA RNA stretch and bend ring of uracil PO41- symmetric stretch (DNA) and deoxyribose-phosphate