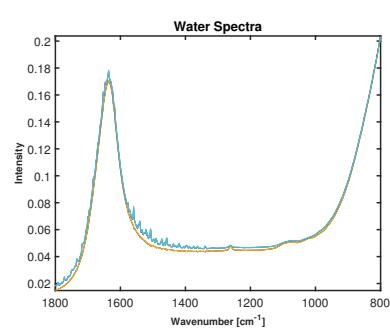
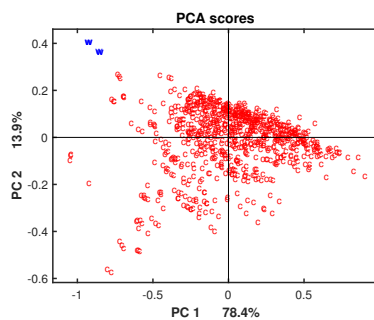


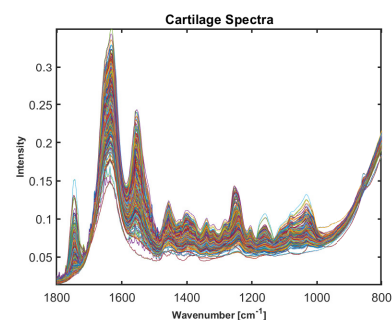
(a) Analyte-poor Spectra (b) Analyte-rich Spectra
Figure S1. Visual presentation of broadband spectrum after using MSC preclassification approach (Collagen as reference spectrum), (a) Water/Analyte-poor Spectra (b) Cartilage/Analyte-rich Spectra.



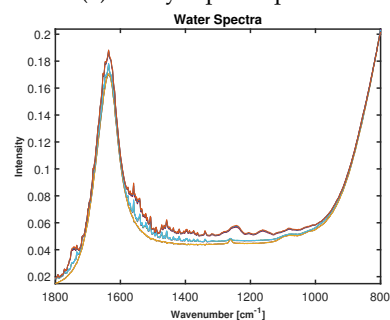
(a) Analyte-poor Spectra



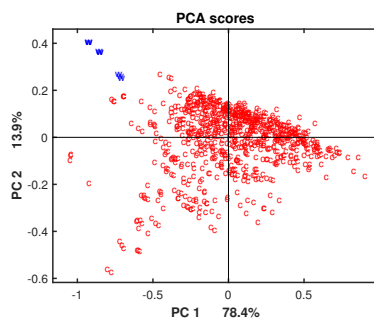
(b) PCA



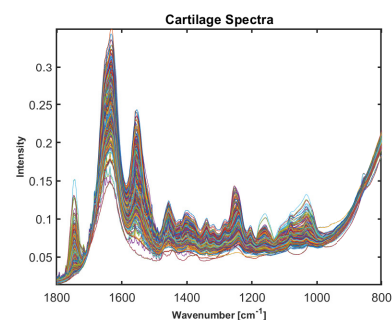
(c) Analyte-rich Spectra



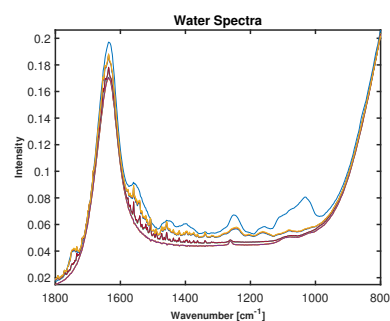
(d) Analyte-poor Spectra



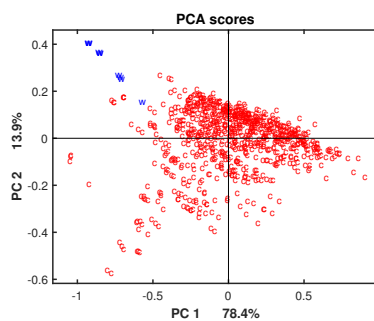
(e) PCA



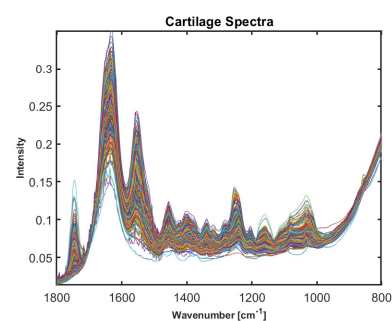
(f) Analyte-rich Spectra



(g) Analyte-poor Spectra



(h) PCA



(i) Analyte-rich Spectra

Figure S2. Visual presentation of Human 12 (set 2) broadband spectra after using MSC preclassification approach, (a,d,g) Water/Analyte-poor Spectra (c,f,i) Cartilage/Analyte-rich Spectra and (b,e,h) are PCA scores obtained by different *RMSE* thresholds, 0.1 for (b), 0.2 for (c) and 0.22 for (h). c and w in PCA scores correspond to cartilage and water spectrum.

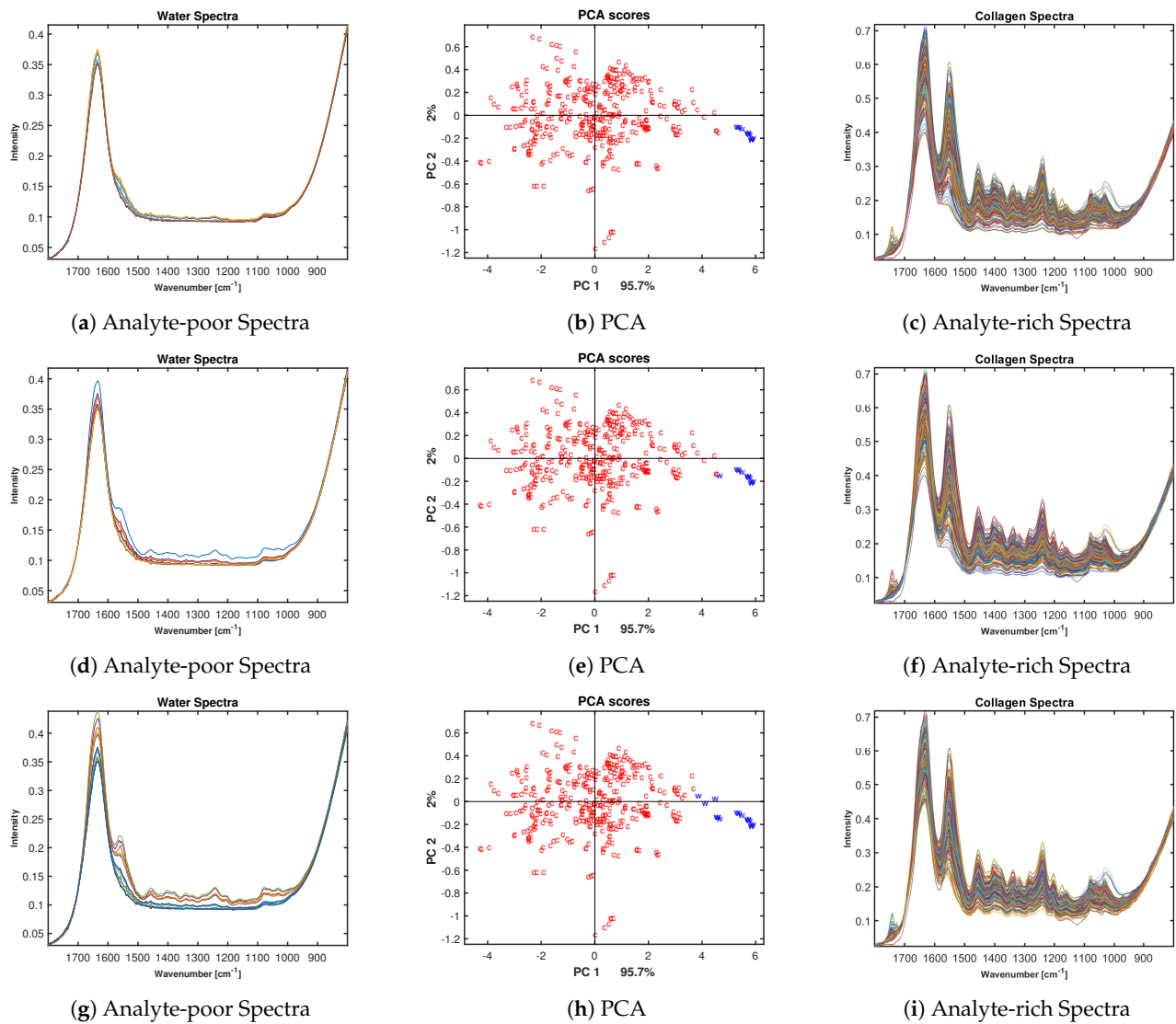


Figure S3. Visual presentation of Bovine (set 3) broadband spectra after using MSC preclassification approach, (a,d,g) Water/Analyte-poor Spectra (c,f,i) Cartilage/Analyte-rich Spectra and (b,e,h) are PCA scores obtained by different RMSE thresholds, 0.5 for (b), 0.6 for (c) and 0.12 for (h). c and w in PCA scores correspond to cartilage and water spectrum.