ICC analysis

The result of ICC analysis for Classic/Common-type descriptor is below:

```
##
                            type
                                    ICC
                                            F df1 df2
                                                               p lower bound
                                                                     0.04046
## Single_raters_absolute
                            ICC1 0.1466 3.062
                                                8 99 0.0040162
## Single random raters
                            ICC2 0.1587 3.839
                                                8 88 0.0006504
                                                                     0.05515
## Single_fixed_raters
                            ICC3 0.1913 3.839
                                                8 88 0.0006504
                                                                     0.06809
## Average_raters_absolute ICC1k 0.6734 3.062
                                                8 99 0.0040162
                                                                     0.33600
## Average_random_raters
                           ICC2k 0.6937 3.839
                                                8
                                                   88 0.0006504
                                                                     0.41193
## Average_fixed_raters
                           ICC3k 0.7395 3.839
                                                8 88 0.0006504
                                                                     0.46716
##
                           upper bound
## Single_raters_absolute
                                0.4033
## Single_random_raters
                                0.4097
## Single_fixed_raters
                                0.4653
## Average_raters_absolute
                                0.8902
## Average_random_raters
                                0.8928
## Average_fixed_raters
                                0.9126
```

The result of fitted random effects models is as follows:

```
## Linear mixed model fit by REML ['lmerMod']
## Formula: values ~ 1 + (1 | id) + (1 | items)
##
      Data: x.df
## REML criterion at convergence: 36.75
## Random effects:
## Groups
                         Std.Dev.
             Name
## items
             (Intercept) 0.167
## id
             (Intercept) 0.161
## Residual
                         0.332
## Number of obs: 36, groups: items, 12; id, 9
## Fixed Effects:
## (Intercept)
##
         0.812
##
            variance Percent
## ID
             0.02607 0.1587
## Items
             0.02795 0.1702
## Residual 0.11021 0.6711
## Total
            0.16423 1.0000
```

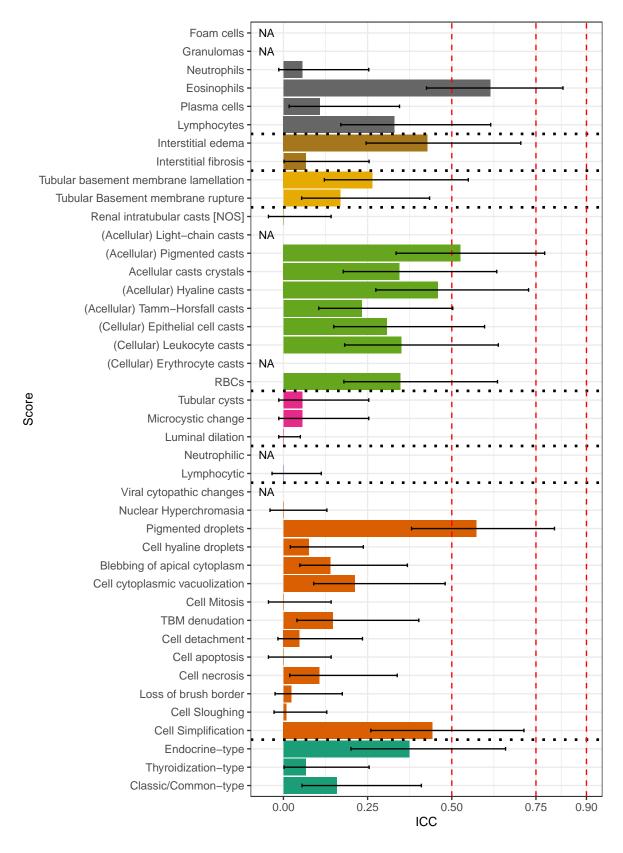


Figure 1: Estimated ICC of presence/abscence scoring of individual descriptors at the sample level with 95% CI. A descriptor is counted as present if a reviewer called it present in any grid square, and absent if they said no for all grid squares. Marked NA when the features were never present in any of 9 samples.

Gwet AC

 $Description\ of\ function:\ https://www.rdocumentation.org/packages/irrCAC/versions/1.0/topics/gwet.ac1.$

```
gwet.ac1.raw(score_df)
```

```
## $est
     coeff.name
                  pa pe coeff.val coeff.se conf.int p.value
##
## 1
           AC1 0.737 0.29
                             0.6297
                                      0.1962 (0.177,1) 0.01244 unweighted
##
## $weights
        [,1] [,2]
##
## [1,]
          1
## [2,]
          0
##
## $categories
## [1] 0 1
```

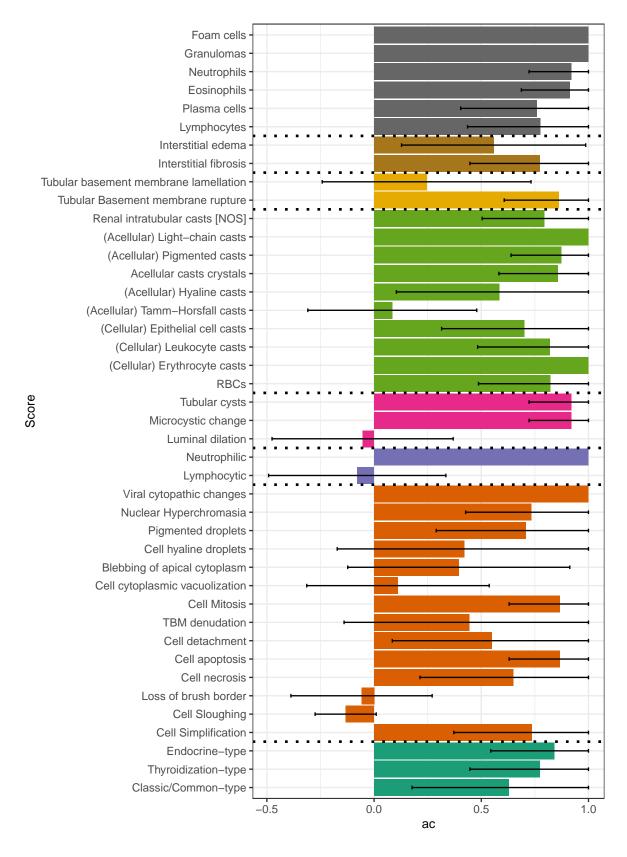


Figure 2: Estimated Gwet AC1 of individual descriptors with 95% CI. No CI was presented when there is no variation.

Supplementary material

Scoring

In the above analyses, presence scoring of each descriptor is as follows. A descriptor is counted as present if a reviewer called it present in any grid square, and absent if they said no for all grid squares.

Cell Sloughing

Domain: Tubular Cell Injury (other than atrophy)



Presence(1) or Absence(0) Score

Loss of brush border

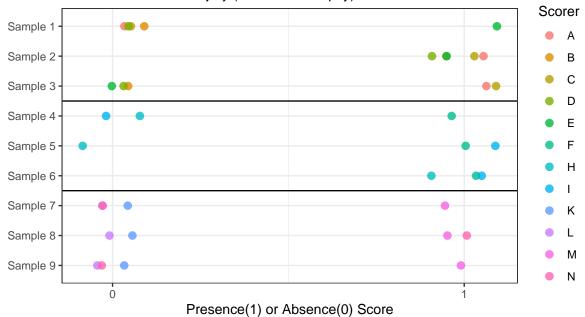
Domain: Tubular Cell Injury (other than atrophy)



Presence(1) or Absence(0) Score

Cell cytoplasmic vacuolization

Domain: Tubular Cell Injury (other than atrophy)



Blebbing of apical cytoplasm

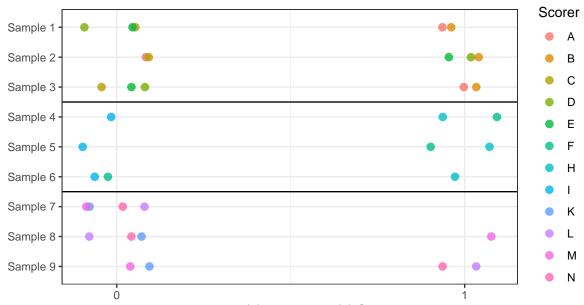
Domain: Tubular Cell Injury (other than atrophy)



Presence(1) or Absence(0) Score

Lymphocytic

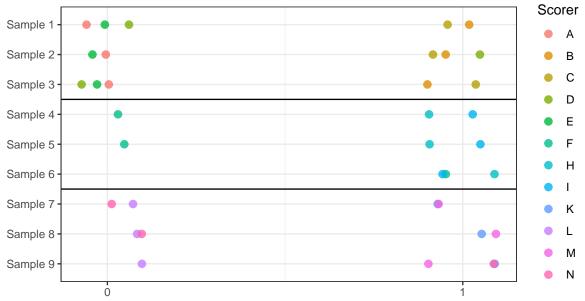




Presence(1) or Absence(0) Score

Luminal dilation

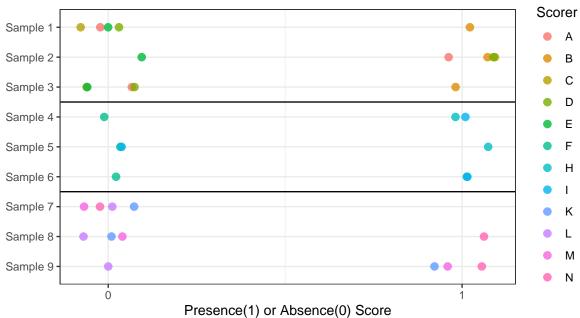
Domain: Abnormal Tubular Luminal Morphology



Presence(1) or Absence(0) Score

(Acellular) Tamm-Horsfall casts

Domain: Intratubular Casts



Tubular basement membrane lamellation

Domain: Tubular basement membrane Morphology



Presence(1) or Absence(0) Score