

ML_Core/ Utils/ SequenceInField

[Go Up](#)

DESCRIPTIONS

MACRO SequenceInField

SequenceInField
<code>(infile,infield,seq,wi_name='wi')</code>

Given a file which is sorted by the work item identifier and INFIELD (and possibly other values), add sequence numbers within the range of each infield. Slightly elaborate code is to avoid having to partition the data to one value of infield per node and to work with very large numbers of records where a global count project would be inappropriate. This is useful for assigning rank positions with the groupings.

PARAMETER infile the input file, any type

PARAMETER infield field name of grouping field

PARAMETER seq name of the field to receive the sequence number

PARAMETER wi_name work item field name, default is wi

RETURN a file of the same type with sequence numbers applied
