PROBLEM B: PEPTIDES

Input file: standard input

Output file: standard output

Problem

In a laboratory it has been decided to study the peptides with respect to aminoacids which build them. All aminoacids have been assigned one character codes. Thus a sample aminoacids sequence of a peptide may look like this: FFAFTOKA

The aim of the study is to find among obtained peptides those which:

- a) are built with the same aminoacids e.g. FFAFTOKA and OKATFA
- b) are built with the same aminoacids and each aminoacid occur the same number of times in both peptides. The last example does not fulfil this condition because aminoacid F occurs three times in the first peptide and only once in the second one.

Your task is to write a program which searches input data for aminoacids fulfilling conditions a) or b).

Input

Any number of tests, each one ending with symbol 0. First line of test data contains the test name. Consecutive ones contain peptide data: peptide identifier and a symbol ":" followed by aminoacids sequence.

Output

Name of the test and below groups of peptides fulfilling conditions a) or b) (their identifiers). If for two peptides both conditions were fulfilled then their identifiers should be written. If only condition a) were fulfilled then identifier should be followed by symbol "*"

Groups should be listed in the same order as they appeared in input data.

Do not repeat groups what means that the output:

PT2 PT7 and later PT7 PT2 is wrong.

Assumptions

Number of peptides in single test does not exceed 1000, number of aminoacids in a peptide does not exceed 200.

Example

Input file

test2 test1 PT1:XYXYXYXYXYXIKL Man:FPFHJWWER PT2:UJKLCTCTC Gorilla:FPFJHEWRW PT3:VICDAWR Pig:FPFJHER Mouse:FFPFJHRE PT4:KXXXXXXYYYYYLI PT5:RRWADIVC PT6:XYXYXYXYXYXIKL PT7:UJKLCTCTC PT8:VICDAWR PT9:KXXXXXXYYYYYLI PT10:RWADIVC 0

Output file

test1 test2
PT1 PT4 PT6 PT9 Man Gorilla
PT2 PT7 Pig Mouse*
PT3 PT5* PT8 PT10
PT4 PT6 PT9
PT5 PT8* PT10*
PT6 PT9
PT8 PT10