Total Points: 30/30

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/*
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Due Date: August 8 2016
* /
/*
Exercise 5.1. Suppose that you would like to invest $1,000 each year at
a bank. The investment earns 5% annual interest, compounded
monthly (that means the interest for each month will be 0.05/12).
Write a program using an explicit loop (or loops) to calculate your
balance for each month if you are investing for 2 years (that means
you deposit $1,000 at the beginning of each year).
* /
data ex 1;
     do year = 1 to 2;
           capital + 1000;
           do month = 1 to 12;
                 interest = capital*(0.05/12);
                 capital+interest;
                 output;
            end;
      end;
run;
/*
Exercise 6.1. Consider the following data set, PROB6_1.SAS7BDAT:
ID G1 G2 G3 S1 S2 S3
1 1 A A C 3 4 9
2 2 A B F 3 7 4
3 3 A C B 5 8 9
Transform PROB6_1 to multiple observations per subject by using
array processing like below:
ID TIME GRADE SCORE
1 1 1 A 3
2 1 2 A 4
3 1 3 C 9
4 2 1 A 3
5 2 2 B 7
6 2 3 F 4
7 3 1 A 5
8 3 2 C 8
9 3 3 B 9
* /
data ex_2 (drop=G1-G3 & S1-S3);
      set PROB6_1.SAS7BDAT;
      array g[3];
      do time = 1 to 3:
           grade = g[time];
            if not missing(grade) then output;
      end;
      array s[3];
      do time =1 to 3;
           score = s[time];
            if not missing(score) then output;
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end;
run;
/*
Problem 1
You are given the SAS data set SPEED (speed.sas), created by running the
program below. Create a new
data set SPEED2 from SPEED, with some new variables. The new variables
LX1 - LX5 are the natural
\log of the variable X1 - X5, and variables SY1 - SY3 are the square roots
of the variables Y1 - Y3. Use
arrays to create the new variables.
data speed;
input X1-X5 Y1-Y3;
datalines;
1 2 3 4 5 6 7 8
11 22 33 44 55 66 77 88
* /
data SPEED2 (drop=X1-X5 Y1-Y3);
     set SPEED;
     array speedo X1 X2 X3 X4 X5;
     do over speedo;
           LX = log10(speedo);
           if not missing(LX) then output;
     end;
     array squared Y1 Y2 Y3;
     do over squared;
           SY = sqrt(squared);
           if not missing(SY) then output;
     end;
run;
/*
Problem 2
You will work with the dna.sas
```

le that reads 15 DNA sequences (See below). The length of each sequence is 60 characters. Based on these DNA sequences, create 60 variables, D1 - D60. D1 will hold the DNA at the $\,$

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rst position, D2 will hold the DNA at the second position, and so on. You
must use array processing
to complete this problem. Hint: use the SUBSTR function.
data dna;
length dna $ 60;
input dna $;
datalines;
TGGAAGGGCTAATTTGGTCCCAAAAAAGACAAGAGATCCTTGATCTGTGGATCTACCACA
TGATTGGCAGAACTACACACCAGGGCCAGGGATCAGATATCCACTGACCTTTGGATGGTG
CTTCAAGTTAGTACCAGTTGAACCAGAGCAAGTAGAAGAGGCCAAATAAGGAGAGAAAA
GTGGAAGTTTGACAGCCTCCTAGCATTTCGTCACATGGCCCGAGAGCTGCATCCGGAGTA
\tt CTACAAAGACTGCTGACATCGAGCTTTCTACAAGGGACTTTCCGCTGGGGACTTTCCAGG
GAGGTGTGGCCTGGGCGGACTGGGGAGTGCCAGATGCTACATATAAGCAGC
TGCTTTTTGCCTGTACTGGGTCTCTCTGGTTAGACCAGATCTGAGCCTGGGAGCTCTCTG
GCTAACTAGGGAACCCACTGCTTAAGCCTCAATAAAGCTTGCCTTGAGTGCTCAAAGTAG
TGTGTGCCCGTCTGTTGTGTGACTCTGGTAACTAGAGATCCCTCAGACCCTTTTAGTCAG
TGTGGAAAATCTCTAGCAGTGGCGCCCGAACAGGGACTTGAAAGCGAAAGTAAAGCCAGA
GGAGATCTCTCGACGCAGGACTCGGCTTGCTGAAGCGCCACGGCAAGAGGCGAGGGGCG
CGAGAGCGTCGGTATTAAGCGGGGGAGAATTAGATAAATGGGAAAAAATTCGGTTAAGGC
CAGGGGGAAAGAACATATAAACTAAAACATATAGTATGGGCAAGCAGGGAGCTAGAAC
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/
/*/
data Problem_2;
    do until (last.dna);
        set DNA;
        by dna;
        array D[60] D1-D60;
        do i = 1 to 60;
            SUBSTR(dna, 1, 15);
        end;
end;
run;
```

