Accompanying Paper For

"Detecting similarities of rational plane curves using complex differential invariants"

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1. DETECTING SIMILARITIES/SYMMETRIES

All implementations constructed in computer algebra system MAPLE [1].

1.1. The Main Algorithm: The Maple Procedure CSim

CSim is the main algorithm that computes the similarities/symmetries of the input curves. The procedure returns a list consisting of the number of similarities/symmetries and computation time. We present the whole procedure below.

(Download the Maple Worksheet)

```
> CSim := \operatorname{proc}(x1, y1, x2, y2)
  local tm, st, x11, x12, x21, x22, y11, y12, y21, y22, x13, x23, y13, y23, delta1, delta2, R11, R12,
      RJ1, RJ2, ImI1, ImI2, ImJ1, ImJ2, Kp, Kq, K1p, K1q, EQPO, EQPR, EQPO1, EQPR1,
      EQPO2, S1, S2, TRFPO := Array(1..0), TRFPO1, FSPOS1, MBSPO := Array(1..0),
      MBSPO1, FSPRS1, FSPO, FSPR, MBPO, MBSPR := Array(1..0), MBSPR1, MBPR, APO,
      APR, BPO, BPR, FSPOS := Array(1..0), FSPRS := Array(1..0), TRFPR := Array(1..0),
      TRFPR1, j, APO1, APO2, APR1, APR2, BPO1, BPO2, BPR1, BPR2, i, tm1, C1, C2, D1, D2,
      k, x21f, x21ff, y21f, y21ff, po, pr,
  uses VectorCalculus;
   st := time():
   x11 \coloneqq diff(x1, t);
  x12 := diff(x11, t);
  x13 := diff(x12, t);
  x21 := diff(x2, t);
  x22 \coloneqq diff(x21, t);
  x23 \coloneqq diff(x22, t);
  y11 := diff(y1, t);
  y12 := diff(y11, t);
  y13 := diff(y12, t);
  y21 := diff(y2, t);
  y22 := diff(y21, t);
  y23 \coloneqq diff(y22, t);
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$$\begin{aligned} delta1 &\coloneqq x11^2 + y11^2; \\ delta2 &\coloneqq x21^2 + y21^2; \\ RI1 &\coloneqq normal \left(\frac{x11 \cdot x12 + y11 \cdot y12}{delta1} \right); \\ RI2 &\coloneqq normal \left(\frac{x11 \cdot x13 + y11 \cdot y13}{delta1} \right); \\ ImI1 &\coloneqq normal \left(\frac{y11 \cdot x13 - x11 \cdot y12}{delta1} \right); \\ ImI2 &\coloneqq normal \left(\frac{y11 \cdot x13 - x11 \cdot y13}{delta1} \right); \\ RJ1 &\coloneqq normal \left(\frac{x21 \cdot x22 + y21 \cdot y22}{delta2} \right); \\ RJ2 &\coloneqq normal \left(\frac{x21 \cdot x23 + y21 \cdot y23}{delta2} \right); \\ ImJ1 &\coloneqq normal \left(\frac{y21 \cdot x23 - x21 \cdot y23}{delta2} \right); \\ ImJ2 &\coloneqq normal \left(\frac{y21 \cdot x23 - x21 \cdot y23}{delta2} \right); \\ Kp &\coloneqq normal \left(\frac{3 \cdot RI1 \cdot ImI1 - ImI2}{ImI1^2} \right); \\ Kq &\coloneqq normal \left(\frac{3 \cdot RJ1 \cdot ImJ1 - ImJ2}{ImJ1^2} \right); \end{aligned}$$

$$K1p := normal\left(\frac{3 \cdot RII^2 - 2 \cdot RI2}{ImII^2}\right);$$

$$K1q := normal\left(\frac{3 \cdot RJI^2 - 2 \cdot RJ2}{ImJI^2}\right);$$

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 $EQPO1 := numer(Kp) \cdot eval(denom(Kq), t = s) - denom(Kp) \cdot eval(numer(Kq), t = s);$ $EQPR1 := numer(Kp) \cdot eval(denom(Kq), t = s) + denom(Kp) \cdot eval(numer(Kq), t = s);$ $EQPO2 := numer(Klp) \cdot eval(denom(Klq), t = s) - denom(Klp) \cdot eval(numer(Klq), t = s);$ EQPO := evala(Gcd(EQPO1, EQPO2)); EQPR := evala(Gcd(EQPR1, EQPO2)); tml := time() - st :if type(EQPO, constant) = true and type(EQPR, constant) = true then return tml,
"There is no similarity"; end if; FSPO := evala(AFactors(EQPO));for i from 1 to nops(FSPO[2]) do
if degree(FSPO[2][i][1], t) = 1 and degree(FSPO[2][i][1], s) = 1 then ArrayTools[Append](FSPOS, FSPO[2][i][1]);end if;

FSPOS1 := convert(FSPOS, list);

for *i* from 1 to *nops*(*FSPOS1*) do MBPO := normal(eval(s, isolate(convert(FSPOS1[i], radical), s)));if $simplify(coeff(numer(MBPO), t, 1) \cdot coeff(denom(MBPO), t, 0) - coeff(numer(MBPO), t, t)$ $0) \cdot coeff(denom(MBPO), t, 1)) \neq 0$ then $ArrayTools[Append] \Big(MBSPO, \frac{coeff(numer(MBPO), t, 1) \cdot t + coeff(numer(MBPO), t, 0)}{coeff(denom(MBPO), t, 1) \cdot t + coeff(denom(MBPO), t, 0)} \Big) \Big)$ end if: end do: MBSPO1 := convert(MBSPO, list);if $nops(MBSPO1) \neq 0$ then for *i* from 1 to nops(MBSPO1) do po := diff(MBSPO1[i], t);x21f := normal(eval(x21, t = MBSPO1[i]));y21f := normal(eval(y21, t = MBSPO1[i])); $APO1 := \frac{po \cdot (x21f \cdot x11 + y21f \cdot y11)}{x11 + y21f \cdot y11}.$ delta 1 $APO2 := \frac{po \cdot (y2lf \cdot x11 - x2lf \cdot y11)}{deltal};$ for j from 0 to 4 do if simplify(eval(denom(MBSPO1[i]), t=j)) $\neq 0$ and simplify(eval(denom(x21f), t=j)) $\neq 0$ and simplify(eval(denom(y21f), t=j)) $\neq 0$ and simplify(eval(denom(x11), t=j)) $\neq 0$ and simplify(eval(denom(y11), t=j)) $\neq 0$ and simplify(eval(numer(x11), t=j)) $\neq 0$ and simplify(eval(numer(y11), t=j)) $\neq 0$ then D1 := [normal(eval(APO1, t=j)),normal(eval(APO2, t=j)), j; break; end if: end do : $ArrayTools[Append](TRFPO, [MBSPO1[i], factor(D1[1] + I \cdot D1[2]), factor(eval(eval(x2 + I))))$ $y_2, t = MBSPO1[i]) - (D1[1] + I \cdot D1[2]) \cdot (x1 + I \cdot y1), t = D1[3]))]);$ end do: TRFPO1 := convert(TRFPO, list);FSPR := evala(AFactors(EQPR));for *i* from 1 to *nops*(*FSPR*[2]) do if degree(FSPR[2][i][1], t) = 1 and degree(FSPR[2][i][1], s) = 1 then ArrayTools[Append](FSPRS, FSPR[2][i][1]); end if: end do: FSPRS1 := convert(FSPRS, list);for i from 1 to nops(FSPRS1) do MBPR := normal(eval(s, isolate(convert(FSPRS1[i], radical), s)));if $simplify(coeff(numer(MBPR), t, 1) \cdot coeff(denom(MBPR), t, 0) - coeff(numer(MBPR), t, 0)$ $\cdot coeff(denom(MBPR), t, 1)) \neq 0$ then $ArrayTools[Append] \bigg(MBSPR, \frac{coeff(numer(MBPR), t, 1) \cdot t + coeff(numer(MBPR), t, 0)}{coeff(denom(MBPR), t, 1) \cdot t + coeff(denom(MBPR), t, 0)} \bigg);$ end if: end do: MBSPR1 := convert(MBSPR, list);

```
MBSPR1 := convert(MBSPR, list);
if nops(MBSPR1) \neq 0 then
for i from 1 to nops(MBSPR1) do
pr := diff(MBSPR1[i], t);
x21ff := normal(eval(x21, t = MBSPR1[i]));
y21ff := normal(eval(y21, t = MBSPR1[i]));
APRI := \frac{pr \cdot (x21 ff \cdot x11 - y21 ff \cdot y11)}{delta1};
APR2 := \frac{pr \cdot (y21ff \cdot x11 + x21ff \cdot y11)}{y}
if simplify(eval(denom(MBSPR1[i]), t = D1[3])) \neq 0 and simplify(eval(denom(x21ff), t
     =DI[3]) \neq 0 and simplify(eval(denom(y21ff), t=DI[3])) \neq 0 then CI :=
    [normal(eval(APR1, t = D1[3])), normal(eval(APR2, t = D1[3])), D1[3]];
else
for j from DI[3] + 1 to DI[3] + 5 do
if simplify(eval(denom(MBSPR1[i]), t=j)) \neq 0 and simplify(eval(denom(x21ff), t=j)) \neq 0
    and simplify(eval(denom(y21ff), t=j)) \neq 0 and simplify(eval(denom(x11), t=j)) \neq 0
    and simplify(eval(denom(y11), t=j)) \neq 0 and simplify(eval(numer(x11), t=j)) \neq 0
    and simplify(eval(numer(y11), t=j)) \neq 0 then C1 := [normal(eval(APR1, t=j)),
    normal(eval(APR2, t=j)), j; break;
end if;
end do :
end if;
ArrayTools[Append](TRFPR, [MBSPR1[i], factor(C1[1] + I \cdot C1[2]), factor(eval(eval(x2 + I))))
    y_2, t = MBSPR1[i]) - (CI[1] + I \cdot CI[2]) \cdot (xI - I \cdot yI), t = CI[3]))]);
end do:
TRFPR1 := convert(TRFPR, list);
end if:
end if:
tm := time() - st:
if nops(MBSPO1) = 0 and nops(MBSPR1) = 0 then return tm, "There is no similarity";
```

```
else
if nops(MBSPO1) = 0 then TRFPO1 := [];
end if;
if nops(MBSPR1) = 0 then TRFPR1 := [];
end if;
return tm, TRFPO1, TRFPR1;
end if;
return tm, TRFPO1, TRFPR1;
end if;
```

2. TESTS AND EXAMPLES

2.1. Examples and Codes Table 2

Timings in Table 2 are generated by the following MAPLE worksheets. CPU times in this Table are average time in detecting symmetries for well-known curves in Table 1.

For rose curves, (Download Maple Worksheeet).

For other miscallenaus curves, (Download Maple Worksheet).

Together with them, the following Maple Worksheets are of the algorithms in [2, 3], which we used to compare timings with ours, respectively.

(Download Maple Worksheet)

(Download Maple Worksheet)

2.2. Codes Table 3 and Table 4

To generate similar curves which have timings in Table 3 we apply the following complex numbers a, b and Möbius transformation $\varphi(t)$ to a rondom parametrization w of degree n and bitsize τ .

 $a = 3 + 2i, b = 1 - 2i, \varphi(t) = t + 1.$

So, using the equation $\mathbf{z} = a\mathbf{w}(\varphi) + b$, we run $CSim(\mathbf{z}, \mathbf{w})$ to compute similarities. The following Maple worksheet gives how to generate similar random curves.

(Download Maple Worksheet)

Finally, the following Maple worksheet gives how to generate non-similar random curves.

(Download Maple Worksheet)

REFERENCES

[1] MapleTM, 2021. Maplesoft, a division of Waterloo Maple Inc. Waterloo, Ontario.

[2] Bizzarri M., Lavicka M., Vrsek J. (2020), *Computing projective equivalences of special algebraic varieties*, Journal of Computational and Applied Mathematics Vol. 367.

[3] Hauer M., Jüttler B. (2018), *Projective and affine symmetries and equivalences of rational curves in arbitrary dimension*, Journal of Symbolic Computation Vol. 87, pp. 68–86.