User Guide for uc_algorithm

Calculating Unimodular Completions for Hyperregular Polynomial Matrices with Entries in $\frac{d}{dt}$ [6, 7]

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1 Introduction

Unimodular matrices are a topic related to differential flatness in the context of nonlinear control.

Given a dynamical systems in implicit form

$$\mathbf{0} = \mathbf{F}(\mathbf{x}, \dot{\mathbf{x}}), \quad \mathbf{x}(t) \in \mathbb{R}^n$$
 (1)

with m < n homogeneous equations, the so called $tangent\ system$ is determined with

$$\mathbf{0} = \left(\frac{\partial \mathbf{F}}{\partial \dot{\mathbf{x}}} \frac{\mathrm{d}}{\mathrm{d}t} + \frac{\partial \mathbf{F}}{\partial \mathbf{x}}\right) \bigg|_{\mathbf{F} = \mathbf{0}} \mathbf{v} = \mathbf{P}\left(\frac{\mathrm{d}}{\mathrm{d}t}\right) \mathbf{v}$$
(2)

and $\boldsymbol{v}(t) \in \mathbb{R}^n$, $\boldsymbol{P}_1 \frac{\mathrm{d}}{\mathrm{d}t} + \boldsymbol{P}_0 =: \boldsymbol{P}\left(\frac{\mathrm{d}}{\mathrm{d}t}\right) \in \mathcal{M}^{m \times n}[\frac{\mathrm{d}}{\mathrm{d}t}]$, where $\mathcal{M}^{m \times n}[\frac{\mathrm{d}}{\mathrm{d}t}]$ denotes the set of $(m \times n)$ -matrices with entries in $\frac{\mathrm{d}}{\mathrm{d}t}$. The matrix $\boldsymbol{P}\left(\frac{\mathrm{d}}{\mathrm{d}t}\right)$ is assumed to be hyperregular, such that there always exists a completion $\boldsymbol{Q} \in \mathcal{M}^{(n-m) \times n}$ with

$$\begin{pmatrix} \mathbf{P} \begin{pmatrix} \frac{\mathrm{d}}{\mathrm{d}t} \end{pmatrix} \\ \mathbf{Q} \end{pmatrix} \in \mathcal{U}^n[\frac{\mathrm{d}}{\mathrm{d}t}], \tag{3}$$

where $\mathcal{U}^n\left[\frac{\mathrm{d}}{\mathrm{d}t}\right]$ denotes matrices of $\mathcal{M}^{n\times n}\left[\frac{\mathrm{d}}{\mathrm{d}t}\right]$ that possess a polynomial inverse in $\mathcal{M}^{n\times n}\left[\frac{\mathrm{d}}{\mathrm{d}t}\right]$ and will be called *unimodular*. The matrix \boldsymbol{Q} is called *unimodular completion of* $\boldsymbol{P}\left(\frac{\mathrm{d}}{\mathrm{d}t}\right)$ and provides information about the flatness property of the system (1).

The algorithm at hand performs both steps (2) and (3) for a specified system (1) and determines a matrix Q. In addition, this guide outlines how to interpret this result.

2 Installation

2.1 Prerequisities

The algorithm is implemented in Python 2.7.11, so make sure it is installed on your system. To check your Python version from the command line type

```
$ python --version
Python 2.7.11
```

Additionally, the algorithm is based on the following packages that need to be up to date:

- sympy
- numpy
- symbtools
- pycartan

It is recommended to install these using the Python package index PyPI (see [1] for instructions):

```
$ pip install sympy numpy symbtools pycartan
```

2.2 Installing the algorithm

There are two ways of installing the algorithm. The quickest way is to clone the git repository:

```
$ git clone https://github.com/klim-/franke_algorithm.git
```

3 Template example

Given an implicit system of the form (1), the following template code shows how it can be prepared for the algorithm:

```
# -*- coding: utf-8 -*-
    import sympy as sp
import symbtools as st
 2
 3
    from sympy import sin, cos, tan
 4
     # Number of state variables
 6
    vec_x = st.symb_vector('x1:%i' % (n+1))
 9
    vec_xdot = st.time_deriv(vec_x, vec_x)
st.make_global(vec_x)
10
11
    st.make_global(vec_xdot)
12
13
     # Additional symbols
14
    g, k1, k2 = sp.symbols("g, k1, k2")
15
16
     # Time-dependent symbols
17
     diff_symbols = sp.Matrix([k1, k2])
18
20
    # Nonlinear system (state space representation)
    # 0 = F_eq(x, xdot)
F_eq = sp.Matrix([
22
               [ xdot1 - x4 ],
[ xdot2 - x5 ],
[ xdot3 - x6 ],
[ g*sin(x1) + xdot4*x3 ]])
23
24
25
26
27
    # Container carrying additional information
28
    # (will be stored in pickle-file)
data = st.Container()
29
30
    data.F_eq = F_eq
data.diff_symbols = diff_symbols
31
```

The example file should be located within the example folder.

4 Usage of the algorithm

It is recommended to use the IPython Qt Console by running the bash command

```
$ ipython qtconsole
```

This will render the output of the algorithm in LATEX. Inside the Qt Console run

```
>>> run franke_algorithm example/my_example.py
```

to start the script. Alternatively, the command from a standard shell is

```
$ python franke_algorithm example/my_example.py
```

The algorithm is iterative and the intermediate steps are not unique. This degree of freedom may well have an impact on the usability of the resulting matrix Q or the script itself such that the user can choose between a manual and an automatic mode. However, we believe the automatic mode is utilizing this freedom fairly well and the bottlenecks are to be found elsewhere.

4.1 Example output

The following system is an academic example taken from [5]

Running the algorithm results in the following output:

```
$ franke_algorithm.py examples/franke_ex1.py
x =
Matrix([
[x1],
[x2],
[x3],
[x4],
[x5]])

xdot =
Matrix([
[xdot1],
[xdot2],
[xdot3],
[xdot4],
[xdot5]])

0 = F(x,xdot) =
Matrix([
[-x3*x4 + xdot1],
[ -x4 + xdot2],
[ -x5 + xdot3]])

Enter "auto" for automatic mode, or "manual" for manual mode:
```

Hitting *Enter* is parsed as auto by default and gives:

```
P10_roc [5 x 2] =
Matrix([
[0, 0],
[0, 0],
[0, 0],
[1, 0],
[0, 1]])
P10_rpinv [5 x 3] =
Matrix([
[1, 0, 0],
[0, 1, 0],
[0, 0, 1],
[0, 0, 0],
[0, 0, 0]])
P10_dot [3 x 5] = Matrix([ [0, 0, 0, 0, 0, 0], [0, 0, 0, 0, 0], [0, 0, 0, 0, 0]])
A0 [3 x 3] =
Matrix([
[0, 0, -x4],
[0, 0, 0],
[0, 0, 0]])
B0 [3 x 2] = Matrix([
[-x3, 0],
[-1, 0],
[ 0, -1]])
B0_loc [1 x 3] = Matrix([[-1, x3, 0]])
P11 [1 x 3] =
Matrix([[-1, x3, 0]])
P01 [1 x 3] = Matrix([[0, 0, x4]])
P11_roc [3 x 2] = Matrix([
 [x3, 0],
[1, 0],
[0, 1]])
P11\_rpinv [3 x 1] =
Matrix([
[-1],
[ 0],
[ 0]])
P11_dot [1 x 3] = Matrix([[0, xdot3, 0]])
A1 [1 x 1] = Matrix([[0]])
```

```
B1 [1 x 2] = Matrix([[-xdot3, x4]])
--- Sonderfall 4.7 ----
B1_tilde [1 x 1] =
Matrix([[-xdot3]])
P11_{tilde_{roc}} [3 x 1] =
Matrix([
[x3],
[1],
[0]])
Z1 [3 x_1] =
Matrix([
[x3*x4/xdot3],
  x4/xdot3],
1]])
Z1_lpinv [1 x 3] =
Matrix([[0, 0, 1]])
Algorithmus am Ende
Q-matrix =
Matrix([
[-1, x3, 0, 0, 0],
[ 0, 0, 1, 0, 0]])
w[0] = (-1)dx1 + (x3)dx2
w[1] = (1)dx3
Data saved to examples/franke_ex1.pcl
```

5 Interpreting the results, checking for integrability

As indicated, the resulting matrix Q may help to decide whether the nonlinear system is flat and if so in determining a flat output. The details about this process are stated in [7]. Helpful for this analysis are the toolboxes pycartan and symbtools. The results of the algorithm are stored in a pcl-file and can be parsed from a fresh Python-shell. For this, type

```
>>> import symbtools as st
>>> data = st.pickle_full_load("/path/to/franke_ex1.pcl")
```

```
>>> data.F_eq
Matrix([
    [-x3*x4 + xdot1],
    [ -x4 + xdot2],
    [ -x5 + xdot3]])
>>> data.Q
Matrix([
    [-1, x3, 0, 0, 0],
    [ 0, 0, 1, 0, 0]])
```

In order to study the vector 1-form $\boldsymbol{\omega} = \boldsymbol{Q} \mathrm{d} \boldsymbol{x}$, a basis vector needs to be generated.

```
>>> basis_vec = data.vec_x
>>> basis_vec
Matrix([
[x1],
[x2],
[x3],
[x4],
[x5]])
```

The vector 1-form can now be generated using the package pycartan:

```
>>> import pycartan as pc
>>> omega = pc.VectorDifferentialForm(1, basis_vec, coeff=data.Q)
>>> omega
Matrix([
[-1, x3, 0, 0, 0],
[ 0,  0, 1, 0, 0]])"dX"
```

The resulting vector 1-form omega can now be unpacked into ordinary 1-forms:

```
>>> omega1, omega2 = omega_tilde4.unpack()
>>> omega1
(-1)dx1 + (x3)dx2
>>> omega2
(1)dx3
```

Checking a 1-form for integrability can be achieved by computing its exterior derivative:

```
>>> omega1.d
(-1)dx2^dx3
>>> omega2.d
(0)dx1^dx2
```

Since the exterior derivative of the 1-form omega2 is zero, it can be integrated which yields one of two components of a flat output y = (y1, y2):

```
>>> y2 = omega2.integrate()
>>> y2
x3
```

For the second component y1, it can be shown that the frobenius theorem is fulfilled [3]:

```
>>> omega1.d^omega2
(0)dx1^dx2^dx3^dx4
```

Thus, the system is flat and there exists a factor μ (x_1, x_2, x_3) such that $d\tilde{\omega}_1 = 0$ with $\tilde{\omega}_1 = \omega_1 + \mu \omega_2$. In order to determine μ , the exterior derivative of $\tilde{\omega}_1$ can be calculated which results in $d\tilde{\omega}_1 = dx_3 \wedge dx_2 + d\mu \wedge dx_3 \stackrel{!}{=} 0$. It can easily be seen that $d\mu = dx_2$ and therefore $\mu = x_2$. This can be verified:

```
>>> st.makeGlobal(data.vec_x) # load elements from vec_x into namespace
>>> mu = x2
>>> omega1_tilde = omega1 + mu*omega2
>>> omega1_tilde.d
(0)dx1^dx2
```

To determine the remaining component y1 of a flat output y, we can integrate omega1_tilde as before:

```
>>> y1 = omega1_tilde.integrate()
>>> y1
-x1 + x2*x3
```

A flat output of this example system is $\mathbf{y} = (-x_1 + x_2 x_3, x_3)$.

6 Implementation details

The algorithm described in [4] makes use of pseudo inverses and orthogonal complements of rectangular matrices. Since unimodular completions of hyperregular matrices are not unique, there are degrees of freedom in the algorithm at hand. Additionally, the iterative character makes it makes the symbolic computation of intermediate steps slow if these freedoms are not chosen well. This section describes these underlying heuristics.

6.1 Pseudoinverses

Definition 6.1. Given a matrix $P \in \mathcal{M}^{m \times n}$ with rank P = m < n, the matrix $P^{+R} \in \mathcal{M}^{n \times m}$ is called right pseudo inverse, if $PP^{+R} = I_m$ is fulfilled.

Definition 6.2. Given a matrix $P \in \mathcal{M}^{m \times n}$ with rank P = n < m, the matrix $P^{+L} \in \mathcal{M}^{n \times m}$ is called left pseudo inverse, if $P^{+L}P = I_n$ is fulfilled.

As indicated, the goal is to choose pseudo inverses as *simple* as possible such that further iterations can be computed efficiently. While MOORE-PENROSE pseudo inverses are easy to determine [2], they lead to matrices with complicated entries in general and due to the iterative nature of the algorithm, the entries grow rapidly. Instead, pseudo inverses with preferably many 0 or 1 entries have proven favorable for further computations.

Let P be matrix in $\mathcal{M}^{m \times n}$ with rank P = m < n. By right multiplying with a permutation matrix $V_{\pi} \in \mathbb{R}^{n}$, P can be sorted column-wise such that

$$\tilde{\boldsymbol{P}} = \boldsymbol{P} \boldsymbol{V}_{\pi} = (\boldsymbol{A}, \boldsymbol{B}), \quad \boldsymbol{A} \in \mathcal{M}^{m \times m}, \boldsymbol{B} \in \mathcal{M}^{m \times (n-m)}, \operatorname{rank} \boldsymbol{A} = m.$$
 (4)

A right pseudo inverse of \tilde{P} can now be specified with

$$\tilde{\boldsymbol{P}}^{+R} = \begin{pmatrix} \boldsymbol{A}^{-1} \\ \mathbf{0}_{(n-m)\times m} \end{pmatrix}. \tag{5}$$

This guarantees at least $m \cdot (n-m)$ zeros in $\tilde{\boldsymbol{P}}^{+R}$. With $\boldsymbol{P}\boldsymbol{V}_{\pi}\tilde{\boldsymbol{P}}^{+R} = \boldsymbol{I}_m$ this leads to a right pseudo inverse for \boldsymbol{P} :

$$\mathbf{P}^{+R} = \mathbf{V}_{\pi} \tilde{\mathbf{P}}^{+R} = \mathbf{V}_{\pi} \begin{pmatrix} \mathbf{A}^{-1} \\ \mathbf{0}_{(n-m)\times m} \end{pmatrix}. \tag{6}$$

The computation of the left pseudo inverse can be done with equation (6) as well:

Let T be a matrix in $\mathcal{M}^{m \times n}$ with rank T = n < m. Transposing the equation $T^{+L}T = I_n$ gives

$$\boldsymbol{I}_{n} = \left(\boldsymbol{T}^{+L}\boldsymbol{T}\right)^{\top} = \boldsymbol{T}^{\top} \left(\boldsymbol{T}^{+L}\right)^{\top} \equiv \boldsymbol{T}^{\top} \left(\boldsymbol{T}^{\top}\right)^{+R}.$$
 (7)

Transposing this equation again yields

$$\mathbf{T}^{+L} = \left(\left(\mathbf{T}^{\top} \right)^{+R} \right)^{\top}. \tag{8}$$

The heuristic for these computations is in V_{π} and is about a good choice of simple and linearly independent columns. For that, we need to specify simple. Criterias for that are preferably many zeros, preferably few mathematical operations or preferably few variables. In order to choose the criteria, the variable pinv_optimization in core/algebra.py needs to be set accordingly.

6.2 Orthogonal complements

The computation of orthogonal complements is not unique, either. Again, we are interested in *simple* complements such that further computations are efficient.

Definition 6.3. The right ortho complement $\mathbf{P}^{\perp R}$ of a matrix $\mathbf{P} \in \mathbb{M}^{m \times n}$ with rank $\mathbf{P} = m < n$ is defined by $\mathbf{P}\mathbf{P}^{\perp R} = \mathbf{0}_{m \times (n-m)}$, where $\mathbf{P}^{\perp R} \in \mathbb{M}^{n \times (n-m)}$ and rank $\mathbf{P}^{\perp R} = n - m$ holds.

Definition 6.4. The left ortho complement $\mathbf{P}^{\perp L}$ of a matrix $\mathbf{P} \in \mathcal{M}^{m \times n}$ with rank $\mathbf{P} = n < m$ is defined by $\mathbf{P}^{\perp L} \mathbf{P} = \mathbf{0}_{(m-n) \times n}$, where $\mathbf{P}^{\perp L} \in \mathcal{M}^{(m-n) \times m}$ and rank $\mathbf{P}^{\perp L} = m - n$ holds.

The preceding heuristic can be used for the computation of orthogonal complements, too:

Let P be a matrix in $\mathbb{M}^{m \times n}$ with rank P = m < n. The matrix $K := (P^{+R}, P^{\perp R}) \in \mathbb{M}^{n \times n}$ is regular and there exists an inverse $K^{-1} \in \mathbb{M}^{n \times n}$ which can be arrange into two hyper-columns Δ_1 and Δ_2 such that

$$I_n = K^{-1}K = \begin{pmatrix} \mathbf{\Delta}_1 \\ \mathbf{\Delta}_2 \end{pmatrix} \begin{pmatrix} \mathbf{P}^{+R} & \mathbf{P}^{\perp R} \end{pmatrix} = \begin{pmatrix} \mathbf{\Delta}_1 \mathbf{P}^{+R} & \mathbf{\Delta}_1 \mathbf{P}^{\perp R} \\ \mathbf{\Delta}_2 \mathbf{P}^{+R} & \mathbf{\Delta}_2 \mathbf{P}^{\perp R} \end{pmatrix}. \tag{9}$$

It is obvious that $\Delta_1 = P$. As explained in the previous section, P can be sorted column-wise such that $\tilde{P} = PV_{\pi} = (A, B)$ with $A \in \mathcal{M}^{m \times m}$ and rank A = m. The matrix \tilde{P} can then be completed such that

$$\tilde{\boldsymbol{K}}^{-1} := \boldsymbol{K} \boldsymbol{V}_{\pi} = \begin{pmatrix} \boldsymbol{A} & \boldsymbol{B} \\ \boldsymbol{0}_{(n-m) \times m} & \boldsymbol{I}_{n-m} \end{pmatrix}. \tag{10}$$

Due to equation (5), this results in

$$I_{n} = \tilde{K}^{-1}\tilde{K} = \begin{pmatrix} A & B \\ \mathbf{0}_{(n-m)\times m} & I_{n-m} \end{pmatrix} \begin{pmatrix} A^{-1} & C \\ \mathbf{0}_{(n-m)\times m} & D \end{pmatrix}$$
(11a)
$$= \begin{pmatrix} I_{m} & AC + BD \\ \mathbf{0}_{(n-m)\times m} & D \end{pmatrix}.$$
(11b)

$$= \begin{pmatrix} I_m & AC + BD \\ \mathbf{0}_{(n-m)\times m} & D \end{pmatrix}. \tag{11b}$$

It follows that $D = I_{n-m}$ and thus $C = -A^{-1}B$. From

$$\tilde{\boldsymbol{K}} = \left(\tilde{\boldsymbol{P}}^{+R}, \tilde{\boldsymbol{P}}^{\perp R}\right) = \begin{pmatrix} \boldsymbol{A}^{-1} & -\boldsymbol{A}^{-1}\boldsymbol{B} \\ \boldsymbol{0}_{(n-m)\times m} & \boldsymbol{I}_{n-m} \end{pmatrix}$$
(12)

follows

$$(I_m, \mathbf{0}_{m \times (n-m)}) = PK = PV_{\pi} \begin{pmatrix} A^{-1} & -A^{-1}B \\ \mathbf{0}_{(n-m) \times m} & I_{n-m} \end{pmatrix}.$$
(13)

A right orthogonal complement can be computed as

$$\mathbf{P}^{\perp R} = \mathbf{V}_{\pi} \begin{pmatrix} -\mathbf{A}^{-1} \mathbf{B} \\ \mathbf{I}_{n-m} \end{pmatrix}. \tag{14}$$

The computation of a left orthogonal complement can deduced from that:

Let T be a matrix in $\mathcal{M}^{m \times n}$ with rank T = n < m. Transposing the equation $T^{\perp L}T = \mathbf{0}_{(m-n)\times n}$ leads to

$$\mathbf{0}_{n \times (m-n)} = \left(\mathbf{T}^{\perp L} \mathbf{T} \right)^{\top} = \mathbf{T}^{\top} \left(\mathbf{T}^{\perp L} \right)^{\top} \equiv \mathbf{T}^{\top} \left(\mathbf{T}^{\top} \right)^{\perp R}. \tag{15}$$

Transposing this equation again yields

$$\boldsymbol{T}^{\perp L} = \left(\left(\boldsymbol{T}^{\top} \right)^{\perp R} \right)^{\top}. \tag{16}$$

6.3Heuristic for pseudo inverses and orthogonal complements

The heuristics for computing preferably simple pseudo inverses and orthogonal complements of $P \in \mathcal{M}^{m \times n}$ with rank P = m < n is in the function reshape_matrix_columns(P) in

core/algebra.py. This function works as follows:

- 1. Remove zero columns of \boldsymbol{P}
- 2. Sort columns by *complexity* specified by pinv_optimization (free symbols or count operations)
- 3. Sort columns by number of zeros
- 4. Pick the first m linear independent columns and add to a matrix A
- 5. Compare **A** and **P** and calculate permutation matrix V_{π} from that
- 6. Use V_{π} to calculate B
- 7. Verify that $PV_{\pi} = (A, B)$
- 8. Return $\boldsymbol{A}, \boldsymbol{B}, \boldsymbol{V}_{\pi}$

References

- [1] Python package index pypi. https://pypi.python.org/pypi, February 2016.
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