

Reaction and Flow Invariants in Chemical Reaction Systems with Inlet and Outlet Streams

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The concept of reaction invariants is extended to include flow invariants of chemical reaction systems with inlet and outlet streams. A transformation to normal form is introduced that allows the separation of: (i) the evolution of the reactions, (ii) the influence of inlet streams, and (iii) the behavior of the reaction and flow invariants. Resulting from this transformation to normal form, model reduction, construction of the linearizing feedback and properties such as state accessibility, state reconstruction, and feedback linearizability are analyzed.

Keywords: Chemical reaction systems, Reaction invariant, Flow invariant, Model reduction, State accessibility, State reconstruction, Feedback linearizability.

1 Introduction

Models of chemical reaction systems include information regarding both the chemical reactions (stoichiometry and kinetics) and the operation mode of the reactor. For the analysis of such models, it is important to distinguish between the states that intrinsically vary with time and those which do not. A transformation has been proposed by Waller and Mäkilä (1981) to separate the reaction variant (same dimension as the number of reactions) from the reaction invariant states. For reaction systems with neither inlet nor outlet streams, the complete concentration vector can then be reconstructed by integrating only the reaction variants.

However, for considerations of safety and productivity, the reactors used in the production of specialty chemicals are often of the semi-batch type (systems with inlets), or run continuously (systems with inlet and outlets). The aim here is to extend the concept of invariance to chemical reaction systems with inlet and outlet streams. This requires the concept of invariants to include flow invariants. A transformation to normal form is proposed herein that performs a three level decomposition of the states into: (i) the evolution of the reactions, (ii) the influence of inlet streams, and (iii) the behavior of the reaction and flow invariants. No kinetic information (except for the assumption of independent kinetics) is necessary for the analysis.

The proposed transformation to normal form helps determine the minimal dimensionality of the model (i.e., the minimal number of differential equations) and analyze control-relevant properties

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such as state accessibility and feedback linearizability. The important results regarding state accessibility proposed by Bastin and Lévine (1993) can be readily derived from the transformation to normal form. Under certain conditions, the system can be proven to be feedback linearizable, and the linearizing control can be obtained easily.

For simplicity, reaction systems with constant temperature, density, and inlet molar concentrations are considered first. However, isothermal operation is sometimes difficult to achieve or simply undesirable because it is non-optimal. Furthermore, as there is a considerable effort to reduce the amount of solvent used in liquid-phase homogeneous reaction systems, the assumption of constant density is often not valid. Also, the inlet molar concentrations might vary due to variability in the raw materials or in the preprocessing steps. Thus, the assumptions of constant temperature, density, and inlet molar concentrations are relaxed, and indications for constructing the transformation to normal form are provided. Semi-batch reaction systems and continuous stirred-tank reaction systems (CSTRs) are considered as special cases.

The paper is organized as follows: Section 2 describes the basic model used in this study. Section 3 develops the transformation to normal form, the implications of which are discussed in Section 4. Extensions to the basic model are provided in Section 5 and special cases are considered in Section 6. A simulated example is presented in Section 7, and Section 8 concludes the paper.

2 Model of a homogeneous, constant-density, isothermal reaction system

Consider a homogeneous, constant-density, isothermal, semi-batch or continuous stirred-tank chemical reaction system comprising S species and R independent reactions, *independent reactions* being those which have both independent stoichiometries *and* independent kinetics (Amrhein *et al.*, 1996). The material balance equations and the continuity equation are described by:

$$\begin{aligned} \dot{\mathbf{n}} &= V \mathbf{K} \mathbf{r}_n(\mathbf{n}, V) + \mathbf{C}_{in} \mathbf{q}_{in} - (q_{out}/V)\mathbf{n}, & \mathbf{n}(0) &= \mathbf{n}_0, \\ \dot{V} &= (\mathbf{1}_p^T \mathbf{q}_{in}) - q_{out}, & V(0) &= V_0, \end{aligned} \quad (1)$$

where \mathbf{n} is the S -dimensional vector of the number of moles, \mathbf{q}_{in} the p -dimensional inlet volumetric flowrate vector, q_{out} the outlet volumetric flowrate expressed as the sum of q outlet volumetric flowrates, V the reactor volume, \mathbf{K} the $S \times R$ stoichiometric matrix, \mathbf{r}_n the R -dimensional reaction rate vector, which is a function of \mathbf{n} and V , $\mathbf{C}_{in} = [\mathbf{c}_{in}^1, \dots, \mathbf{c}_{in}^p]$, \mathbf{c}_{in}^i the molar concentrations of the i th inlet stream, \mathbf{n}_0 the initial number of moles, V_0 the initial volume, and $\mathbf{1}_p$ a p -dimensional vector with all elements being 1. To completely describe (1), the R -dimensional vector function $\mathbf{r}_n(\mathbf{n}, V)$ must be specified.

The terms $V\mathbf{K}\mathbf{r}_n(\mathbf{n}, V)$, $\mathbf{C}_{in}\mathbf{q}_{in}$, and $(q_{out}/V)\mathbf{n}$ represent the effect of the reactions, the inlet streams, and outlet stream on the number of moles, respectively. The molar concentrations are given by $\mathbf{c} = \mathbf{n}/V$. A model similar to (1) can be derived for bioreactions by expressing the material balances and the continuity equation in mass units. \mathbf{K} then represents the matrix of yield coefficients.

It is assumed that the time profiles of the inlet flowrates \mathbf{q}_{in} are linearly independent and the inlet molar concentrations \mathbf{c}_{in}^i ($i = 1, \dots, p$) are constant. Finally, note that the outlet flowrate q_{out} does not necessarily depend on \mathbf{q}_{in} .

3 Transformation to normal form

3.1 Reaction invariants in the absence of inlets and outlets

The aim here is the extension of the concept of reaction invariants to obtain a normal form for chemical reaction systems. As a first step towards understanding reaction invariants, a chemical reaction system with neither inlet nor outlet streams is considered:

$$\begin{aligned}\dot{\mathbf{n}} &= V \mathbf{K} \mathbf{r}_n(\mathbf{n}, V), & \mathbf{n}(0) &= \mathbf{n}_0, \\ \dot{V} &= 0, & V(0) &= V_0.\end{aligned}\tag{2}$$

Though there are $(S + 1)$ differential equations in (2), it will be shown below that the reaction system has intrinsically only R states or directions in which \mathbf{n} evolves with time since there are only R independent reactions. The other $(S - R)$ directions are unaffected by the reactions and are termed the *invariants of the reaction system* (Waller and Mäkilä, 1981).

The state space will be separated into: (i) an R -dimensional vector space which evolves due to the R independent reactions (the *reaction variants*), and (ii) the $(S - R)$ -dimensional reaction invariants. The ideas of Moore–Penrose pseudo-inverse (denoted by superscript $+$) and completion of space will be used for this purpose. Let $\mathbf{P} \in \mathbb{R}^{S \times (S-R)}$ be a matrix with orthonormal columns which satisfies: (i) $\text{rank}([\mathbf{K} \ \mathbf{P}]) = S$, and (ii) $\mathbf{P}^T \mathbf{K} = \mathbf{0}_{(S-R) \times R}$. Then, the *linear* transformation which separates the R -dimensional reaction variants, \mathbf{z}_1 , and the $(S - R)$ -dimensional reaction invariants, \mathbf{z}_2 , is given by

$$\begin{bmatrix} \mathbf{z}_1 \\ \mathbf{z}_2 \end{bmatrix} = \begin{bmatrix} \mathbf{K}^+ \mathbf{n} \\ \mathbf{P}^T \mathbf{n} \end{bmatrix}.\tag{3}$$

Substituting (3) in (2) and using $\mathbf{r}_z(\mathbf{z}, V)$ for the reaction rates in the new coordinates gives:

$$\begin{aligned}\dot{\mathbf{z}}_1 &= V \mathbf{r}_z(\mathbf{z}, V), & \mathbf{z}_1(0) &= \mathbf{K}^+ \mathbf{n}_0, \\ \dot{\mathbf{z}}_2 &= \mathbf{0}_{S-R}, & \mathbf{z}_2(0) &= \mathbf{P}^T \mathbf{n}_0, \\ \dot{V} &= 0, & V(0) &= V_0.\end{aligned}\tag{4}$$

Remark 1 (Construction of \mathbf{P}) Note that the columns of \mathbf{P} span the null space of \mathbf{K}^T . Therefore, one way to construct \mathbf{P} is by Singular Value Decomposition (SVD) of \mathbf{K} (Golub and Van Loan, 1983). \mathbf{P} is the matrix of the right singular vectors corresponding to the zero singular values of \mathbf{K} .

3.2 Reaction invariants in the presence of inlets and outlets

Even when inlets and outlets are present, the reactions affect only an R -dimensional space. The reaction invariants can be separated using the transformation (3). Applying (3) to (1) gives,

$$\begin{aligned}\dot{\mathbf{z}}_1 &= V \mathbf{r}_z(\mathbf{z}, V) + \mathbf{K}^+ \mathbf{C}_{in} \mathbf{q}_{in} - (q_{out}/V) \mathbf{z}_1, & \mathbf{z}_1(0) &= \mathbf{K}^+ \mathbf{n}_0, \\ \dot{\mathbf{z}}_2 &= \mathbf{P}^T \mathbf{C}_{in} \mathbf{q}_{in} - (q_{out}/V) \mathbf{z}_2, & \mathbf{z}_2(0) &= \mathbf{P}^T \mathbf{n}_0, \\ \dot{V} &= (\mathbf{1}_p^T \mathbf{q}_{in}) - q_{out}, & V(0) &= V_0.\end{aligned}\tag{5}$$

Note that, though \mathbf{z}_2 and V are reaction invariants, they do not remain constant. Thus, the evolution of \mathbf{n} is no longer restricted to an R -dimensional space as previously; it evolves in a larger space. A similar conclusion has been drawn by Fjeld *et al.* (1974).

In the absence of inlets and outlets, both isolating the manifold in which \mathbf{n} evolves and spotting the reaction invariants is the same problem. This is because that the variation in \mathbf{n} is caused only through the R independent reactions. Alternatively, in the presence of inlets and outlets, the variation in \mathbf{n} and V is not only due to reactions but also due to flows (inlets and outlets). Thus, isolating the manifold in which \mathbf{n} and V evolve means finding and eliminating states which are invariant with respect to both reactions and flows.

3.3 Reaction and flow invariants

The key aim of this paper is to provide a three-part decomposition of the system equations (1). The first part is that which is affected by the R independent reactions. Among the reaction invariants, the second part picks those states which evolve with the flows. The third part consists of those states which remain constant, i.e., reaction *and* flow invariants. Note that the transformation to normal form that will be proposed to this effect is *nonlinear*.

The dimension of the reaction variant space is R . As will be seen below, the dimension of the manifold in which the transformed states evolve is given by $\sigma + 1$ where $\sigma \equiv \text{rank} \left(\begin{bmatrix} \mathbf{K} & \mathbf{C}_{in} \end{bmatrix} \right)$. This means that the dimension increases with every inlet whose molar concentrations are independent of \mathbf{K} and the remaining inlet molar concentrations. First, it is assumed that $\sigma = R + p$. It will be shown that the dimension of the reaction variant space is R , that of the reaction invariant and flow variant space is $p + 1$, and that of the reaction and flow invariant space is $S - \sigma$. Next, the transformation to normal form will be developed, and it will then be generalized to the case of $\sigma < R + p$.

In comparison with (3), the completion of the state space involving \mathbf{n} and V takes a three-level structure. Let $\mathbf{L} \in \mathbb{R}^{S \times (\sigma - R)}$ and $\mathbf{N} \in \mathbb{R}^{S \times (S - \sigma)}$ be matrices with orthonormal columns which satisfy the following conditions: (i) $\text{rank} \left(\begin{bmatrix} \mathbf{K} & \mathbf{L} & \mathbf{N} \end{bmatrix} \right) = S$, (ii) \mathbf{K} , \mathbf{L} and \mathbf{N} are mutually orthogonal, (iii) $\mathbf{N}^T \mathbf{C}_{in} = \mathbf{0}_{S - \sigma \times p}$, (iv) $\mathbf{L}^T \mathbf{C}_{in}$ is invertible.

For the application of the three-part decomposition, the transformation to normal form (3) has to be changed as follows: (i) instead of \mathbf{P} , an $S \times (\sigma - R)$ matrix, \mathbf{M} , has to be chosen which renders $\mathbf{M}^T \mathbf{C}_{in} = \mathbf{I}_{(\sigma - R)}$, achieved by choosing $\mathbf{M} \equiv \mathbf{L} (\mathbf{C}_{in}^T \mathbf{L})^{-1}$; (ii) $\mathbf{K}^+ \mathbf{C}_{in} \mathbf{q}_{in}$ has to be eliminated. For this, an additional projection matrix $(\mathbf{I}_S - \mathbf{C}_{in} \mathbf{M}^T)$ is formulated which, by construction of \mathbf{M} , satisfies $(\mathbf{I}_S - \mathbf{C}_{in} \mathbf{M}^T) \mathbf{K} = \mathbf{K}$. With these modifications, the transformation to normal form reads:

$$\begin{bmatrix} \mathbf{z}_1 \\ \mathbf{z}_2 \\ \mathbf{z}_3 \end{bmatrix} = \begin{bmatrix} \mathbf{K}^+ (\mathbf{I}_S - \mathbf{C}_{in} \mathbf{M}^T) \mathbf{n} \\ \mathbf{M}^T \mathbf{n} \\ \mathbf{N}^T \mathbf{n} \end{bmatrix} \quad (6)$$

and

$$\begin{aligned}
\dot{\mathbf{z}}_1 &= V \mathbf{r}_z(\mathbf{z}, V) - (q_{out}/V) \mathbf{z}_1, & \mathbf{z}_1(0) &= \mathbf{K}^+ (\mathbf{I}_S - \mathbf{C}_{in} \mathbf{M}^T) \mathbf{n}_0, \\
\dot{\mathbf{z}}_2 &= \mathbf{q}_{in} - (q_{out}/V) \mathbf{z}_2, & \mathbf{z}_2(0) &= \mathbf{M}^T \mathbf{n}_0, \\
\dot{\mathbf{z}}_3 &= -(q_{out}/V) \mathbf{z}_3, & \mathbf{z}_3(0) &= \mathbf{N}^T \mathbf{n}_0, \\
\dot{V} &= (\mathbf{1}_p^T \mathbf{q}_{in}) - (q_{out}/V) V, & V(0) &= V_0,
\end{aligned} \tag{7}$$

where \mathbf{z}_1 , \mathbf{z}_2 , and \mathbf{z}_3 are vectors of dimension R , p , and $(S - \sigma)$, respectively. For the elimination of the term (q_{out}/V) , a *nonlinear* transformation can be used:

$$\begin{bmatrix} \mathbf{x} \\ \nu \end{bmatrix} = \begin{bmatrix} \mathbf{z}/(V - \mathbf{1}_p^T \mathbf{z}_2) \\ V - \mathbf{1}_p^T \mathbf{z}_2 \end{bmatrix}, \tag{8}$$

where \mathbf{x} and ν are the transformed states of dimension $(R + S + p - \sigma)$ and 1, respectively. By introducing a non-zero scaling factor η , the following theorem is proposed which formulates the so-called *normal form* of (1).

Theorem 2 *Let $\sigma = \text{rank}([\mathbf{K} \ \mathbf{C}_{in}]) = (R + p)$. Then, a diffeomorphism $\mathcal{T} : [\frac{\mathbf{n}}{V}] \leftrightarrow [\frac{\mathbf{x}}{\nu}]$ exists which transforms model (1) into:*

$$\begin{aligned}
\dot{\mathbf{x}}_1 &= h(\mathbf{x}_2) \mathbf{r}(\mathbf{x}), & \mathbf{x}_1(0) &= g_0 \mathbf{K}^+ (\mathbf{I}_S - \mathbf{C}_{in} \mathbf{M}^T) \mathbf{n}_0, \\
\dot{\mathbf{x}}_2 &= \mathbf{q}_{in}/\nu, & \mathbf{x}_2(0) &= g_0 \mathbf{M}^T \mathbf{n}_0, \\
\dot{\mathbf{x}}_3 &= \mathbf{0}_{S-\sigma}, & \mathbf{x}_3(0) &= g_0 \mathbf{N}^T \mathbf{n}_0, \\
\dot{\nu} &= -q_{out}/h(\mathbf{x}_2), & \nu(0) &= 1/g_0,
\end{aligned} \tag{9}$$

where

$$\begin{aligned}
h(\mathbf{x}_2) &= \eta + \mathbf{1}_p^T \mathbf{x}_2, \\
g_0 &= \eta/(V_0 - \mathbf{1}_p^T \mathbf{M}^T \mathbf{n}_0),
\end{aligned} \tag{10}$$

and \mathbf{x}_1 , \mathbf{x}_2 , and \mathbf{x}_3 are vectors of dimension R , p , and $(S - \sigma)$, respectively, η a non-zero arbitrary constant, \mathbf{r} the R -dimensional reaction rate vector expressed in terms of \mathbf{x} , $\mathbf{M} = \mathbf{L}(\mathbf{C}_{in}^T \mathbf{L})^{-1}$, $\mathbf{L} \in \mathbb{R}^{S \times (\sigma - R)}$, and $\mathbf{N} \in \mathbb{R}^{S \times (S - \sigma)}$ matrices with orthonormal columns which satisfy: (i) $\text{rank}([\mathbf{K} \ \mathbf{L} \ \mathbf{N}]) = S$, (ii) \mathbf{K} , \mathbf{L} and \mathbf{N} are mutually orthogonal, (iii) $\mathbf{N}^T \mathbf{C}_{in} = \mathbf{0}_{S - \sigma \times p}$, (iv) $\mathbf{L}^T \mathbf{C}_{in}$ is invertible.

The transformation to normal form is one-to-one and can be written as follows using $g(\mathbf{n}, V) = \eta/(V - \mathbf{1}_p^T \mathbf{M}^T \mathbf{n})$:

$$\begin{bmatrix} \mathbf{n} \\ V \end{bmatrix} \rightarrow \begin{bmatrix} \mathbf{x} \\ \nu \end{bmatrix} : \begin{bmatrix} \mathbf{x}_1 \\ \mathbf{x}_2 \\ \mathbf{x}_3 \\ \nu \end{bmatrix} = \begin{bmatrix} g(\mathbf{n}, V) \mathbf{K}^+ (\mathbf{I}_S - \mathbf{C}_{in} \mathbf{M}^T) \mathbf{n} \\ g(\mathbf{n}, V) \mathbf{M}^T \mathbf{n} \\ g(\mathbf{n}, V) \mathbf{N}^T \mathbf{n} \\ 1/g(\mathbf{n}, V) \end{bmatrix} \tag{11}$$

$$\begin{bmatrix} \mathbf{x} \\ \nu \end{bmatrix} \rightarrow \begin{bmatrix} \mathbf{n} \\ V \end{bmatrix} : \begin{bmatrix} \mathbf{n} \\ V \end{bmatrix} = \begin{bmatrix} \nu (\mathbf{K} \mathbf{x}_1 + \mathbf{C}_{in} \mathbf{x}_2 + \mathbf{N} \mathbf{x}_3) \\ \nu h(\mathbf{x}_2) \end{bmatrix}. \tag{12}$$

(see Appendix A for proof)

Table 1

Summary of the transformation to normal form.

Original space	Reaction variants	Reaction invariants and flow variants (Inlets)	Reaction and flow invariants	Reaction invariants and flow variants (Outlet)			
	\mathbf{x}_1	\mathbf{x}_2	\mathbf{x}_3	ν			
	$\dot{\mathbf{x}}_1 = h(\mathbf{x}_2) \mathbf{r}(\mathbf{x})$	$\dot{\mathbf{x}}_2 = \mathbf{q}_{in}/\nu$	$\dot{\mathbf{x}}_3 = \mathbf{0}$	$\dot{\nu} = -q_{out}/h(\mathbf{x}_2)$			
$\mathbf{n} =$	$(\mathbf{K} \mathbf{x}_1$	$+$	$\mathbf{C}_{in} \mathbf{x}_2$	$+$	$\mathbf{N} \mathbf{x}_3)$	\cdot	ν
$V =$	$h(\mathbf{x}_2)$				\cdot	ν	
$\mathbf{c} =$	$\frac{(\mathbf{K} \mathbf{x}_1 + \mathbf{C}_{in} \mathbf{x}_2 + \mathbf{N} \mathbf{x}_3)}{h(\mathbf{x}_2)}$						
\mathbb{R}^{S+1}	\mathbb{R}^R	\mathbb{R}^p	$\mathbb{R}^{(S-\sigma)}$	\mathbb{R}^1			

The matrices \mathbf{N} and \mathbf{L} required in Theorem 2 can be constructed similarly as in Remark 1 by noting that the columns of \mathbf{N} and \mathbf{L} span the null spaces of $[\mathbf{K}, \mathbf{C}_{in}]^T$ and $[\mathbf{K}, \mathbf{N}]^T$, respectively.

Theorem 2 can be easily extended to the case where $\sigma < R + p$. For this, the matrix \mathbf{M} is redefined using the pseudo-inverse of $(\mathbf{C}_{in}^T \mathbf{L})$ instead of the inverse such that $\mathbf{M} := \mathbf{L} (\mathbf{C}_{in}^T \mathbf{L})^+$. It will be shown in Subsection 3.4 that the transformation to normal form in this case is no longer one-to-one.

Corollary 3 *For $\sigma < R + p$, the transformed model (9) represents (1), where $\mathbf{M} := \mathbf{L} (\mathbf{C}_{in}^T \mathbf{L})^+$. The relation $[\mathbf{x}] \rightarrow [\mathbf{v}]$ is defined by (12) (see Appendix B for proof).*

3.4 Discussion

The results obtained in Theorem 2 and Corollary 3 are summarized in Table 1. Though a rigorous interpretation of the results is difficult due to the nonlinear term in the transformation to normal form, an intuitive meaning for the various parts of \mathbf{x} can be given. \mathbf{x}_1 describes the R -dimensional reaction variants, while \mathbf{x}_2 captures the p -dimensional flow part. \mathbf{x}_3 are the $(S - \sigma)$ -dimensional reaction and flow invariants with ν being related to the outflow.

The third row depicts the differential equations which explain why the interpretation of the states outlined above is possible. The fourth and fifth rows summarize the reconstruction of the original states \mathbf{n} and V . With this transformation to normal form, \mathbf{c} (sixth row) and \mathbf{n} are decomposed along the directions determined by the columns of: (i) the stoichiometric matrix \mathbf{K} , (ii) the inlet concentration matrix \mathbf{C}_{in} , and (iii) the reaction and flow invariant matrix \mathbf{N} . Note that, in contrast to \mathbf{n} , \mathbf{c} is not directly influenced by the outflow through ν .

In the last row, the dimensions of the various spaces are given. The transformed system is of

dimension $(S + R + p - \sigma + 1)$, whereas the original system has dimension $(S + 1)$. Thus, if $\sigma < (R + p)$, there is an increase in dimension as a result of the transformation to normal form. This is why the transformation to normal form is not one-to-one when $\sigma < (R + p)$. An example is now given as illustration.

Example 4 Consider a reaction system with two independent reactions ($R = 2$): $A \rightarrow B$, $B + C \rightarrow D$. When A and B are fed as two different inlets ($p = 2$), then

$$[\mathbf{K} \ \mathbf{C}_{in}] = \begin{bmatrix} -1 & 0 & c_{in,A} & 0 \\ 1 & -1 & 0 & c_{in,B} \\ 0 & -1 & 0 & 0 \\ 0 & 1 & 0 & 0 \end{bmatrix}$$

and $\sigma = 3 < 4 = R + p$. This is due to the fact that feeding A along with the reaction $A \rightarrow B$ is equivalent to feeding B . This redundancy is reflected in the loss of rank in $[\mathbf{K}, \mathbf{C}_{in}]$.

Owing to redundancy, a clear separation of the influence of reactions and flows is no longer possible. One way to handle this is to explain the same effect by more than one state, thus, artificially increasing the dimension of the state space (see Table 1).

Note that \mathbf{x}_2 and ν form the flow variant space. However, since ν enters nonlinearly in the reconstruction of \mathbf{n} and V , it is given a special symbol. Owing to the redundancy described above, amongst the $(p + 1)$ states present in $[\mathbf{x}_\nu^2]$ only $(\sigma - R + 1)$ states constitute the reaction invariant and flow variant space.

4 Implications of the transformation to normal form

The transformation to normal form for chemical reaction systems with *independent inlet and outlet streams* proposed in the previous section clearly shows the evolution of \mathbf{n} in three different spaces: the stoichiometric space (\mathbf{K}), the inlet space (\mathbf{C}_{in}), and the invariant space (\mathbf{N}). Such a separation has immediate implications for model reduction, state accessibility, state reconstruction, feedback linearizability, and the construction of the linearizing feedback.

4.1 Model reduction

From (9), it is clear that only $(R + p + 1)$ differential equations need to be integrated to compute the trajectories. In addition, a set of $(S - \sigma)$ constants must be computed from the initial conditions. In the case of $S > (R + p)$, analysis and design of controllers, observers and optimizers can be achieved with the reduced model of $(R + p + 1)$ states if the initial conditions \mathbf{n}_0 and V_0 are assumed to be known.

4.2 State accessibility

It is obvious from the transformation to normal form in Theorem 2 that: (i) \mathbf{x}_3 is inaccessible, and (ii) \mathbf{x}_2 and ν are accessible from the inlet and outlet streams. Moreover, a part of \mathbf{x}_1 can be inaccessible depending on the kinetics. Thus, the dimension of the inaccessible part is at least $(S - \sigma)$ which leads to the following corollary.

Corollary 5 *The maximum dimension of the accessible part is $(\sigma+1)$, where $\sigma = \text{rank} \left(\begin{bmatrix} \mathbf{K} & \mathbf{C}_{in} \end{bmatrix} \right)$.*

Therefore, if full state accessibility is required, the dimension of \mathbf{x}_3 should be zero (i.e., $\sigma = S$) implying that the number of inlet streams should be at least $(S - R)$ (i.e., $p \geq S - R$).

Corollary 6 *$\sigma = S$ is a necessary condition for full state accessibility.*

These results have been reported previously by Bastin and Lévine (1993). However, deriving them from the transformation to normal form in Theorem 2 provides an alternate proof.

Another interesting aspect of the transformation to normal form is that the dynamics of the inaccessible part can be transformed to $\dot{\mathbf{x}}_3 = \mathbf{0}_{S-\sigma}$, which is marginally stable.

4.3 State reconstruction

The reconstruction of \mathbf{n} as in (12) can be used to reconstruct the complete state information from a subset of measured molar concentrations. The key idea is that \mathbf{x}_2 , \mathbf{x}_3 and ν can be calculated at any point in time from the initial conditions and the flows without knowledge of the kinetics since they are reaction invariants. Then, if the molar concentrations, \mathbf{c}_s , of $S_s \geq R$ species are measured, \mathbf{x}_1 can be calculated under certain minor assumptions. When \mathbf{x}_1 is available, the number of moles \mathbf{n} , and also, the molar concentration vector $\mathbf{c} = \mathbf{n}/V$ can be reconstructed from (12). The idea is similar to that of the asymptotic observer proposed by Bastin and Dochain (1990).

Proposition 7 *Let the molar concentrations of $S_s \geq R$ species be measured and \mathbf{K}_s be the $S_s \times R$ submatrix of \mathbf{K} corresponding to these S_s species. Given \mathbf{K} , \mathbf{C}_{in} , \mathbf{q}_{in} , q_{out} , \mathbf{c}_0 , and V_0 , if \mathbf{K}_s has a unique left pseudo-inverse, the molar concentrations, \mathbf{c}_t , of the remaining $(S - S_s)$ species can be reconstructed without knowledge of reaction kinetics using*

$$\mathbf{c}_t = \mathbf{K}_t \mathbf{K}_s^+ \mathbf{c}_s + \left(\mathbf{C}_{in,t} - \mathbf{K}_t \mathbf{K}_s^+ \mathbf{C}_{in,s} \right) \mathbf{x}_2 / h(\mathbf{x}_2) + \left(\mathbf{N}_t - \mathbf{K}_t \mathbf{K}_s^+ \mathbf{N}_s \right) \mathbf{x}_3 / h(\mathbf{x}_2), \quad (13)$$

where the subscript s represents a quantity corresponding to the measured S_s molar concentrations and t to that of $(S - S_s)$ unmeasured molar concentrations.

Let only an estimate of the initial molar concentrations of the S_t species be available. Then the estimation error $\mathbf{e} \equiv \hat{\mathbf{c}}_t - \mathbf{c}_t$, with $\hat{\mathbf{c}}_t$ being the estimated molar concentrations, is given by

$$\mathbf{e} = \frac{h(\mathbf{x}_2(0))}{h(\mathbf{x}_2)} \mathbf{e}(0). \quad (14)$$

If at least one of the inlet flowrates is always non-zero, then the estimation error asymptotically converges to zero. (see Appendix C for proof).

The asymptotic convergence can intuitively be explained from (14). The numerator term is constant while the denominator monotonically increases if at least one of the inlet flowrates is always non-zero. $h(\mathbf{x}_2)$ tends to infinity, thereby pushing the error to zero. Note that since the estimation errors of the S_t species are independent of each other, an initial molar concentration error of any species does not propagate on the molar concentration estimates of the remaining species.

Note that, if molar concentrations of fewer than R species are measured, then observer techniques relying on the knowledge of reaction kinetics should be used (Soroush, 1997).

4.4 Feedback linearizability

Consider a reduced system that does not include the inaccessible part \mathbf{x}_3 . The system is affine in the inputs and is given by:

$$\dot{\mathbf{x}} = \begin{bmatrix} \dot{\mathbf{x}}_1 \\ \dot{\mathbf{x}}_2 \\ \dot{\nu} \end{bmatrix} = \begin{bmatrix} \mathbf{f}_1(\mathbf{x}) \\ \mathbf{0}_p \\ 0 \end{bmatrix} + \begin{bmatrix} \mathbf{0}_{R \times m} \\ \hline (1/\nu)\mathbf{I}_p & \mathbf{0}_p \\ \hline \mathbf{0}_p^\top & -1/h(\mathbf{x}_2) \end{bmatrix} \mathbf{u} \equiv \mathbf{f}(\mathbf{x}) + \sum_{j=1}^m \mathbf{g}_j u_j, \quad \mathbf{u} = \begin{bmatrix} \mathbf{q}_{in} \\ q_{out} \end{bmatrix}, \quad (15)$$

where \mathbf{f} and \mathbf{g}_j are n -dimensional vector fields, \mathbf{f}_1 an R -dimensional vector field, and \mathbf{u} the m -dimensional input vector. m and n are the dimensions of \mathbf{u} and \mathbf{x} , respectively, with, in this case, $m = (p + 1)$ and $n = (R + m)$. Owing to the special structure of the vector fields, important results regarding the feedback linearizability of system (15) can be obtained. A system is feedback linearizable if a state feedback and a diffeomorphism exist which renders the closed-loop system linear (see Appendix D for conditions of feedback linearizability).

Proposition 8 *Given system (15) with $\sigma = (R + p)$ and $m \geq R$ inputs. If the $R \times m$ matrix $\mathbf{J} = [(1/\nu)\frac{\partial \mathbf{f}_1}{\partial \mathbf{x}_2}, (-1/h(\mathbf{x}_2))\frac{\partial \mathbf{f}_1}{\partial \nu}]$ is of rank R , then (15) is feedback linearizable. (see Appendix D for proof)*

An advantage of model (1) transformed into (15) is that the linearizing feedback can be calculated easily. With this feedback, the system is linear — no further diffeomorphism is required.

\mathbf{x}_1 and a part of \mathbf{x}_2 form the flat outputs (Fliess *et al.*, 1995), linearization of (15) to $\ddot{\mathbf{x}}_1 = \mathbf{v}_u$ and $\dot{\mathbf{x}}_{2,v} = \mathbf{v}_v$ with the new external inputs $\mathbf{v}_u \in \mathbb{R}^R$ and $\mathbf{v}_v \in \mathbb{R}^{m-R}$ can be achieved, where $\mathbf{x}_2 = \begin{bmatrix} \mathbf{x}_{2,u} \\ \mathbf{x}_{2,v} \end{bmatrix}$. Differentiating $\dot{\mathbf{x}}_1$ with respect to time,

$$\mathbf{v}_u = \ddot{\mathbf{x}}_1 = \frac{\partial \mathbf{f}_1}{\partial \mathbf{x}_1} \dot{\mathbf{x}}_1 + \mathbf{J} \mathbf{u}. \quad (16)$$

Partitioning $\mathbf{J} = [\mathbf{J}_s, \mathbf{J}_t]$ such that $\mathbf{J}_s \in \mathbb{R}^{R \times R}$ is invertible, the following feedback is computed from (16):

$$\mathbf{u} = \begin{bmatrix} -\mathbf{J}_s^{-1} \frac{\partial \mathbf{f}_1}{\partial \mathbf{x}_1} \dot{\mathbf{x}}_1 \\ \mathbf{0}_{m-R} \end{bmatrix} + \begin{bmatrix} \mathbf{J}_s^{-1} & -\mathbf{J}_s^{-1} \mathbf{J}_t \\ \mathbf{0}_{(m-R) \times R} & \mathbf{I}_{m-R} \end{bmatrix} \begin{bmatrix} \mathbf{v}_u \\ \mathbf{v}_v \end{bmatrix}. \quad (17)$$

Note that $(m - R)$ inputs are superfluous and, thus, not necessary to control the R reactions.

The assumption of full rank for \mathbf{J} is not very restrictive, especially when only independent reactions are considered. If only non-reacting species are added, it is clear that they will not affect the

reaction rates, and \mathbf{J} will lose rank. In contrast, if different reacting species in at least R inlets are fed to the reaction system, the assumption of full rank for \mathbf{J} can be met easily.

In the above proposition, only sufficient conditions for full-state feedback linearizability have been provided. An example illustrating the conservatism of these conditions is given next. Consider the semi-batch reaction system $A + B \rightarrow C$ and $A + C \rightarrow D$ with second-order kinetics and B being fed ($m = p = 1$, $R = 2$). Although the conditions of Proposition 8 are not verified, it can easily be verified that the system is feedback linearizable.

5 Extensions to other types of reaction systems

The transformation to normal form was stated in Section 3 for reaction systems with temperature, constant density, and inlet molar concentrations. Below, these results will be extended to cover practical situations commonly found in chemical reaction systems, in particular those dealing with varying temperature, density, and inlet molar concentrations.

5.1 Non-isothermal reaction systems

In a non-isothermal scenario, the temperature, T , should be modeled as an additional state. Under the assumptions that: (i) the pressure remains constant, (ii) the mixing enthalpies are negligible, and (iii) the temperatures, \mathbf{T}_{in} , of the p inlet streams remain constant, the heat balance equation is:

$$\frac{d}{dt}(V \rho c_p T) = V (-\Delta \mathbf{h}_R^T) \mathbf{r}_n(\mathbf{n}, V, T) + \mathbf{T}_{in}^T \mathbf{S}_{in} \Phi_{in} \mathbf{q}_{in} - (q_{out}/V) (V \rho s T) + Q_{ext}, \quad (18)$$

where $\Delta \mathbf{h}_R$ is the R -dimensional vector of reaction enthalpies, c_p the specific heat capacity, ρ the density of the mixture, \mathbf{S}_{in} and Φ_{in} p -dimensional diagonal matrices with elements being the specific heat capacities and densities of the inlet streams, respectively, and Q_{ext} the external heat energy such as external heating/cooling energy or heat dissipation. Using $T^* = \rho s T$, the model for non-isothermal constant-density reaction systems is given by:

$$\begin{aligned} \dot{\mathbf{n}} &= V \mathbf{K} \mathbf{r}_n(\mathbf{n}, V, T^*) + \mathbf{C}_{in} \mathbf{q}_{in} - (q_{out}/V) \mathbf{n}, & \mathbf{n}(0) &= \mathbf{n}_0, \\ \frac{d}{dt}(V T^*) &= V (-\Delta \mathbf{h}_R^T) \mathbf{r}_n(\mathbf{n}, V, T^*) + \mathbf{T}_{in}^{*,T} \mathbf{q}_{in} - (q_{out}/V) (V T^*) + Q_{ext}, & T^*(0) &= T_0^*, \\ \dot{V} &= (\mathbf{1}_p^T \mathbf{q}_{in}) - q_{out}, & V(0) &= V_0. \end{aligned} \quad (19)$$

Furthermore, it is assumed that \mathbf{S}_{in} and Φ_{in} are constant. For the formulation of the transformation of (19) into normal form, the heat balance equation is not transformed when the external energy is non-negligible ($Q_{ext} \neq 0$). However, the remaining equations are transformed using (9) since the transformation to normal form proposed in Theorem 2 is independent of reaction kinetics and temperature. Note that the dimension of the inaccessible part \mathbf{x}_3 does not decrease even when the temperature is used as an additional input for control.

When $Q_{ext} = 0$, then (VT^*) can be combined with \mathbf{n} leading to,

$$\frac{d}{dt} \begin{bmatrix} \mathbf{n} \\ (VT^*) \end{bmatrix} = V \begin{bmatrix} \mathbf{K} \\ -\Delta \mathbf{h}_R^T \end{bmatrix} \mathbf{r}_n(\mathbf{n}, V, T^*) + \begin{bmatrix} \mathbf{C}_{in} \\ \mathbf{T}_{in}^{*,T} \end{bmatrix} \mathbf{q}_{in} - (q_{out}/V) \begin{bmatrix} \mathbf{n} \\ (VT^*) \end{bmatrix}. \quad (20)$$

From (20) the state (VT^*) can be modeled as an additional species and, hence, $S_T \equiv S + 1$. If $\sigma = \text{rank}([\mathbf{K}, \mathbf{C}_{in}]) = R + p$, then $\text{rank}(\begin{bmatrix} \mathbf{K} & \mathbf{C}_{in} \\ -\Delta \mathbf{h}_R^T & \mathbf{T}_{in}^{*,T} \end{bmatrix}) = R + p$. Since $(S_T - \sigma) = (S - \sigma) + 1$, the number of reaction and flow invariants (inaccessible states) is increased by 1. This means that the additional state (VT^*) is inaccessible in the absence of external energy. Intuitively, the temperature cannot vary independently of the reactions and the flows without external heating since the evolution of the reactions is already captured by \mathbf{n} .

5.2 Varying-density reaction systems

In reaction systems with constant density ρ , the conservation equation for total mass, m , translates directly into a conservation equation for the volume V , as was the case in equation (1). This is due to the relationship $V = m/\rho$ with ρ being constant. However, in the case of varying density, the mass conservation leads to,

$$\frac{d}{dt}(\rho V) = \mathbf{1}_p^T \Phi_{in} \mathbf{q}_{in} - \rho q_{out}, \quad (\rho V)(0) = \rho_0 V_0. \quad (21)$$

In the case of varying density, $\dot{\mathbf{n}}$ in (1) can still be transformed as in Theorem 2 if h is redefined using the following algebra. Substituting $V = h\nu$ in (21) and using (9), $\frac{d}{dt}(\rho h) = \mathbf{1}_p^T \Phi_{in} (\mathbf{q}_{in}/\nu)$. After integrating both sides, the following redefinition of h results:

$$h(\mathbf{x}, \nu) := (\mathbf{1}_p^T \Phi_{in} \mathbf{x}_2 + \eta^*)/\rho(\mathbf{x}, \nu), \quad (22)$$

where η^* is an arbitrary integration constant. An important difference between the original definition (10) and the redefinition (22) is that, since ρ depends on both \mathbf{x} and ν , the redefined h depends not only on \mathbf{x}_2 but also on ν and the *entire* \mathbf{x} vector. Note that since $g = h(\mathbf{x}, \nu)/V$, it gets redefined as $g(\mathbf{n}, V) := \eta^*/(\rho V - \mathbf{1}_p^T \Phi_{in} \mathbf{M}^T \mathbf{n})$.

Since Φ_{in} is assumed to be constant and is usually available, availability of the inlet volumetric flowrates \mathbf{q}_{in} and the inlet mass flowrates $\mathbf{u}_{in} = \Phi_{in} \mathbf{q}_{in}$ is equivalent. However, since the varying density ρ is usually unknown, the outlet mass flowrate, $u_{out} = \rho q_{out}$, cannot be calculated from q_{out} and *vice versa*.

If u_{out} is available, then $\dot{\nu}$ in the transformed system (9) has to be replaced by:

$$\begin{aligned} \dot{\nu} &= -q_{out}/h(\mathbf{x}, \nu) = -q_{out} \rho / (\mathbf{1}_p^T \Phi_{in} \mathbf{x}_2 + \eta^*) \\ &= -u_{out} / (\mathbf{1}_p^T \Phi_{in} \mathbf{x}_2 + \eta^*). \end{aligned} \quad (23)$$

In this case, ν , and hence \mathbf{x}_2 (see 9), are independent of ρ which implies that ν and \mathbf{x}_2 are reaction invariants. Note that h for $\dot{\mathbf{x}}_1$ in (9) and for V in (12) is as defined in (22).

In summary, the structure of the transformed system (9) is retained for varying-density reaction systems except for the redefinition of h . With volumetric flowrates, the interpretation of the three-part decomposition discussed in Section 3.4 is not possible due to coupling through the density.

Alternatively, if the mass flowrates are available, the interpretation of the three-part decomposition is still valid since ν and \mathbf{x}_2 are independent of the density.

5.3 Varying inlet concentrations

In the case of varying inlet concentrations, the reaction system is modeled by:

$$\begin{aligned}\dot{\mathbf{n}} &= V \mathbf{K} \mathbf{r}(\mathbf{c}) + \mathbf{C}_{in}^* \mathbf{q}_{in}^* - (q_{out}/V) \mathbf{n}, & \mathbf{n}(0) &= \mathbf{n}_0 \\ \dot{V} &= \mathbf{1}_{p^*}^T \mathbf{q}_{in}^* - q_{out}, & V(0) &= V_0,\end{aligned}\quad (24)$$

where \mathbf{C}_{in}^* and \mathbf{q}_{in}^* are the varying inlet molar concentrations and flowrates of the p^* physical inlets, respectively.

The transformation to normal form is carried out in two steps. In the first step, (24), a system with p^* inlet streams of *varying* inlet molar concentrations, is transformed into (1), a system with p streams of *constant* inlet molar concentrations. The second step is the transformation of (1) to normal form, which has been discussed in earlier sections.

For the first step, let the $S \times K$ matrix \mathbf{F}_{in} , and the $p^* \times K$ matrix \mathbf{Q}_{in}^* be defined as:

$$\mathbf{F}_{in} \equiv [\mathbf{C}_{in}^*(0) \mathbf{q}_{in}^*(0), \dots, \mathbf{C}_{in}^*(K-1) \mathbf{q}_{in}^*(K-1)], \quad \mathbf{Q}_{in}^* \equiv [\mathbf{q}_{in}^*(0), \dots, \mathbf{q}_{in}^*(K-1)], \quad (25)$$

where K is the number of discretization points. Let $f \equiv \text{rank}(\mathbf{F}_{in})$. \mathbf{F}_{in} can be decomposed using, for example, SVD such that $\mathbf{F}_{in} = \bar{\mathbf{C}}_{in} \bar{\mathbf{Q}}_{in}$, where $\bar{\mathbf{C}}_{in}$ and $\bar{\mathbf{Q}}_{in}$ are the $S \times f$ matrix of constant inlet molar concentrations and the $f \times K$ matrix of corresponding inlet volumetric flowrates for K discretization points, respectively. Redefining $p := f + 1$, the $S \times p$ inlet molar concentration matrix \mathbf{C}_{in} and the $p \times K$ matrix of flowrates, \mathbf{Q}_{in} , are obtained:

$$\mathbf{C}_{in} := [\bar{\mathbf{C}}_{in} \ \mathbf{0}_S], \quad \mathbf{Q}_{in} \equiv \begin{bmatrix} \bar{\mathbf{Q}}_{in} \\ \mathbf{1}_{p^*}^T \mathbf{Q}_{in}^* - \mathbf{1}_f^T \bar{\mathbf{Q}}_{in} \end{bmatrix}. \quad (26)$$

The p -dimensional inlet volumetric flowrate vector $\mathbf{q}_{in}(k)$ at the k th discretization instant is the $(k+1)$ th column of \mathbf{Q}_{in} ($k = 0, \dots, K-1$). With this redefinition:

$$\mathbf{C}_{in}^*(k) \mathbf{q}_{in}^*(k) = \bar{\mathbf{C}}_{in} \bar{\mathbf{q}}_{in}(k) = \mathbf{C}_{in} \mathbf{q}_{in}(k), \quad \mathbf{1}_{p^*}^T \mathbf{q}_{in}^*(k) = \mathbf{1}_p^T \mathbf{q}_{in}(k). \quad (27)$$

Substituting (27) in (24) leads to (1). Note that the structure of the reaction system in normal form (9) is retained even for reaction systems with varying inlet concentrations except for redefinition of the inlet streams. Also, the interpretation of the three-part decomposition is valid.

6 Special cases

In the development of the transformation to normal form in Section 3, it was assumed that the inlets and outlets are independent. However, in certain important special cases, q_{out} is not independent, for example, for (i) semi-batch reaction systems where $q_{out} = 0$, and (ii) CSTRs, where $q_{out} = \mathbf{1}_p^T \mathbf{q}_{in}$. That this dependence leads to inaccessibility of the volume since the outflow loses its direct influence on the volume will be shown next.

Semi-batch reaction systems. Owing to the absence of an outlet stream, $\dot{\nu} = 0$ implying $\nu(t) = \nu_0$. Thus, $V = \nu_0 h(\mathbf{x}_2)$. The transformed set of equations is given below.

$$\dot{\mathbf{x}}_1 = h(\mathbf{x}_2) \mathbf{r}(\mathbf{x}), \quad \dot{\mathbf{x}}_2 = \mathbf{q}_{in}/\nu_0, \quad \dot{\mathbf{x}}_3 = \mathbf{0}_{S-\sigma}, \quad \dot{\nu} = 0. \quad (28)$$

Note that the transformation given by (11)–(12) is linear since $g(\mathbf{n}, V)$ is a constant with $g(\mathbf{n}, V) = 1/\nu_0$.

Continuous stirred-tank reaction systems. In a CSTR, $q_{out} = \mathbf{1}_p^T \mathbf{q}_{in}$, $\dot{V} = 0$ and $V(t) = V_0$. Since the nonlinear transformation (8) of V to ν is not required to get to the normal form, a similar transformation as in (28) can be used to obtain the transformed system:

$$\dot{\mathbf{x}}_1 = h(\mathbf{x}_2) \mathbf{r}(\mathbf{x}), \quad \dot{\mathbf{x}}_2 = h(\mathbf{x}_2) \mathbf{q}_{in}/V_0, \quad \dot{\mathbf{x}}_3 = \mathbf{0}_{S-\sigma}, \quad \dot{V} = 0. \quad (29)$$

As discussed in Section 3.4, the reaction invariant and flow variant space has a maximum dimension of $(\sigma - R + 1)$. In the two special cases shown above, the same quantity is limited to a maximum of p since there are only p independent flows. Combining these two limits, the dimension of the reaction invariant and flow variant space is determined as $\min(\sigma - R + 1, p)$.

The number of reaction invariant and flow variants $\min(\sigma - R + 1, p)$ equals p for either of the following two cases: (i) $\sigma = (R+p)$, and (ii) $\sigma = R+p-1$. This means that the number of accessible states is not reduced even when the molar concentrations of one inlet stream are dependent on the stoichiometry and the molar concentrations of the remaining inlet streams. Intuitively, this is due to the volume becoming accessible in the case of $\sigma = R + p - 1$ as discussed below.

Addition of solvent. The addition of solvent is considered as an example for $\sigma = R + p - 1$. The inlet molar concentration matrix has the form $\mathbf{C}_{in} := [\mathbf{c}_{in}^1, \dots, \mathbf{c}_{in}^{(p-1)}, \mathbf{0}_S]$. If it is assumed that $\text{rank}([\mathbf{K} \mathbf{c}_{in}^1 \dots \mathbf{c}_{in}^{(p-1)}]) = R+p-1$, then $\text{rank}([\mathbf{K} \mathbf{C}_{in}]) = R+p-1$. Intuitively, when the outlet is dependent on the inlets, the addition of solvent makes the volume accessible, since adding the solvent changes the volume, thereby diluting other species and affecting \mathbf{n} and V . Note that, for the tangent-linearized model with no outlet stream and $\text{rank}([\mathbf{K} \mathbf{c}_{in}^1 \dots \mathbf{c}_{in}^{(p-1)}]) = R+p-1$, local controllability of the volume by solvent addition has been proven by Dochain and Chen (1992). However, the work presented herein proposes *global* results for *nonlinear* chemical reaction systems.

7 Example

The concept of model reduction, state reconstruction, state accessibility and feedback linearizability will now be illustrated on a homogeneous, constant-density, non-isothermal semi-batch reaction system ($R = 3$ and $S = 7$), namely the ethanolysis of phthalyl chloride (A) (de Vallière, 1989). In two successive irreversible ethanolysis reactions, phthalyl chloride monoethyl ester (C), the desired product, and phthalic diethylester (E) are produced from ethanol (B). Both reactions also produce hydrochloric acid (D). It is assumed that B also reacts with D in a reversible side reaction to produce ethyl chloride (F) and water (G). The reaction system can be described by the following reaction scheme:

Table 2

Numerical values for the simulated reaction systems.

Parameter	Value	Unit
$\{\kappa_{i,0}\}$	[18 17 38 30]	$\text{M}^{-1} \text{h}^{-1}$
$\{E_i^*\}$	[150 200 350 400]	K



The molar concentration vector reads $\mathbf{c} = [c_A, c_B, c_C, c_D, c_E, c_F, c_G]^T$. Since a known temperature profile $T(t)$ is imposed (see Fig. 1a), the differential equation for T can be neglected. Hence, the model is of order 8. The reaction rates are assumed to follow the so-called *mass-action principle*:

$$r_1 = \kappa_1(T)c_Ac_B, \quad r_2 = \kappa_2(T)c_Bc_C, \quad r_3 = \kappa_3(T)c_Bc_D - \kappa_4(T)c_Fc_G$$

with κ_i following the Arrhenius law $\kappa_i(T) = \kappa_{i,0} \exp(-E_i^*/T)$. The numerical values of the parameters are given in Table 2.

Semi-batch operation with $p = 2$ inlet streams and no outlet will now be studied with \mathbf{C}_{in} chosen such that $\text{rank}(\left[\mathbf{K} \ \mathbf{C}_{in}\right]) = R+p$. In this way, the transformation to normal form (diffeomorphism) of Theorem 2 can be illustrated. Ethanol ($V_0 = 1.5$ l with 2 w-% water) in 80 w-% solvent (dimethyl sulfoxide) is initially placed in a vessel, i.e. $\mathbf{c}_0 = [0, 4.2, 0, 0, 0, 0, 0.22]^T$ M. The first inlet stream contains phthaloyl dichloride and the second an ethanol/water mixture, i.e. $\mathbf{c}_{in}^1 = [4.9, 0, 0, 0, 0, 0, 0]^T$ M and $\mathbf{c}_{in}^2 = [0, 20, 0, 0, 0, 0, 1.5]^T$ M. The inlet volumetric flowrate profiles are given in Fig. 1c, the volume profile in Fig. 1b, and the molar concentration profiles in Fig. 2.

Model (1) can be reduced to $(R+p) = 5$ differential equations as given in Theorem 2. With the choice $\eta \equiv V_0 - \mathbf{1}_p^T \mathbf{M}^T \mathbf{n}_0 = 1.200$ l, $\nu(t) = \nu_0 = 1$. From (28) and (12), the reduced model and the reconstructed molar concentrations \mathbf{c} read:

$$\begin{bmatrix} \dot{\mathbf{x}}_1 \\ \dot{\mathbf{x}}_2 \end{bmatrix} = \begin{bmatrix} h(\mathbf{x}_2) \mathbf{r}(\mathbf{x}) \\ \mathbf{q}_{in} \end{bmatrix}, \quad \mathbf{x}_1(0) = \begin{bmatrix} -0.034 \\ -0.023 \\ -0.065 \end{bmatrix}, \quad \mathbf{x}_2(0) = \begin{bmatrix} -0.007 \\ 0.209 \end{bmatrix}, \quad (30)$$

$$\mathbf{c} = (\mathbf{K} \mathbf{x}_1 + \mathbf{C}_{in} \mathbf{x}_2 + \mathbf{N} \mathbf{x}_3) / h(\mathbf{x}_2),$$

$$V = h(\mathbf{x}_2),$$

where

$$\mathbf{N} = \begin{bmatrix} 0 & -0.014 & 0.329 & -0.343 & 0.657 & -0.550 & 0.194 \\ 0 & 0.049 & 0.222 & -0.172 & 0.444 & 0.536 & -0.659 \end{bmatrix}^T$$

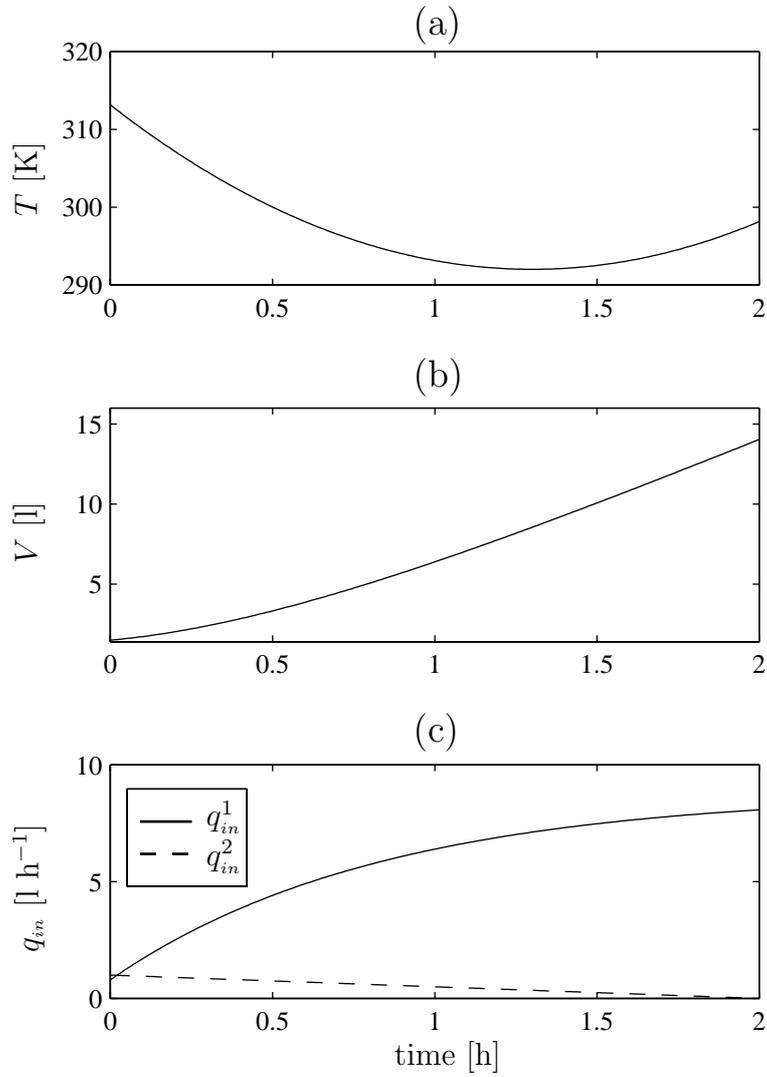


Fig. 1. Time profiles of: (a) the temperature T , (b) the volume V , and (c) the inlet volumetric flowrates q_{in} .

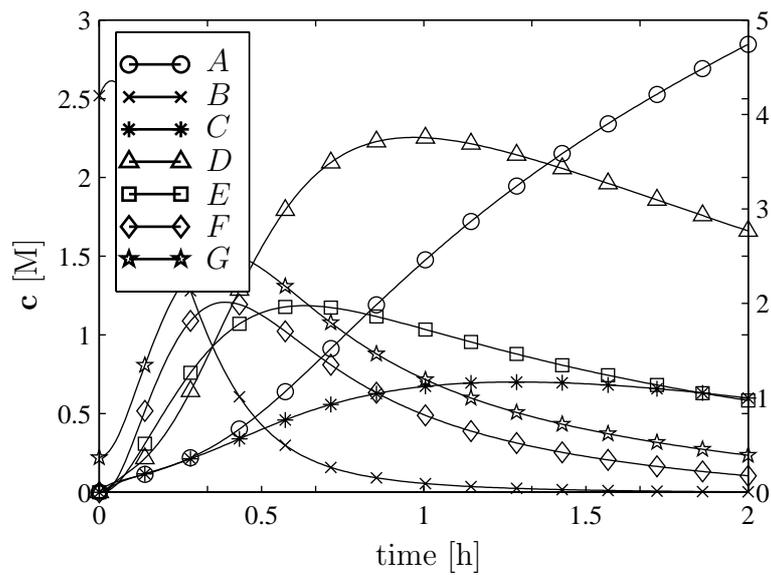


Fig. 2. Time profiles of the molar concentrations c (right axis for c_B , and left axis for the other species).

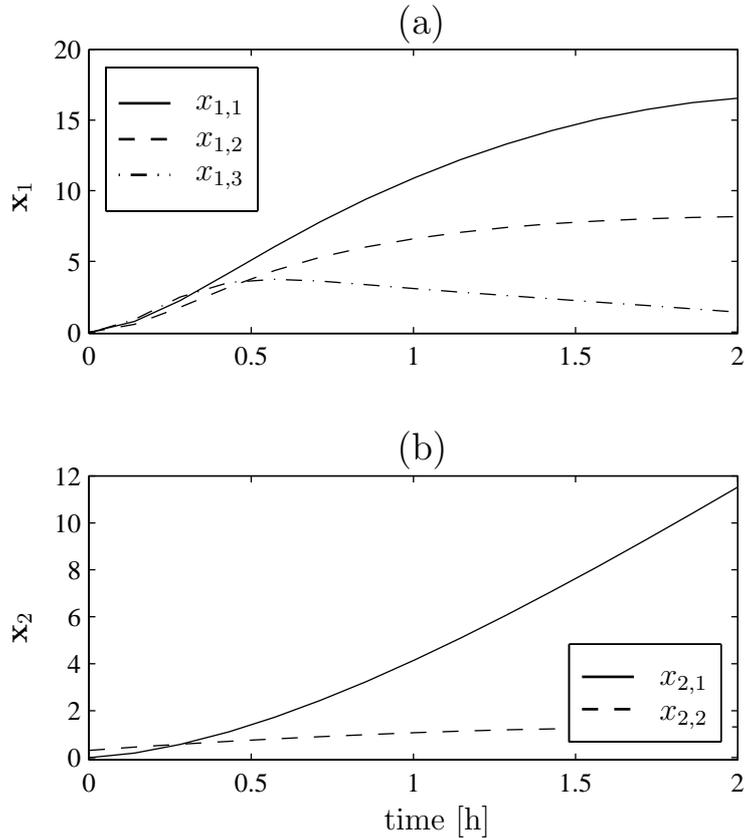


Fig. 3. Time profiles of: (a) \mathbf{x}_1 , and (b) \mathbf{x}_2 .

and $\mathbf{x}_3 = [-0.028, 0.094]^T$. The time profiles of \mathbf{x}_1 and \mathbf{x}_2 are shown in Fig. 3.

If the molar concentrations of $S_s = R = 3$ species are measured, e.g. $\mathbf{c}_s = [c_A, c_D, c_G]^T$, the molar concentrations of the remaining species can be estimated using (13) and (30) with

$$\mathbf{K}_s = \begin{bmatrix} -1 & 0 & 0 \\ 1 & 1 & -1 \\ 0 & 0 & 1 \end{bmatrix}.$$

From Corollary 6 it can be seen that $p = 2$ inlet streams are not sufficient to have full state accessibility since at least $(S - R) = 4$ inlet streams would be necessary. Furthermore, in the absence of an outlet stream ($q_{out} = 0$) — as is the case here — at least $p = R = 3$ inlet streams would be needed to guarantee feedback linearizability using Proposition 8.

8 Conclusions

A generic nonlinear transformation to normal form for reaction systems with inlet and outlet streams has been presented. It leads to a normal form where the reaction and flow invariants can be easily identified. Such a transformation was used to prove results on state accessibility. Implications of this transformation to normal form in the fields of model reduction, state reconstruction and

feedback linearizability were discussed. Using this transformation for feedback linearization has the advantage that the linearizing control can be obtained in a straightforward manner.

The results obtained for constant-density, isothermal reaction systems were extended to varying density and non-isothermal situations, and special cases such as semi-batch reaction systems and CSTRs were considered. Note that the results obtained are *global* in nature and deal with the *full-fledged nonlinear* system, also for extensions and special cases. In summary, major insight is gained by using the normal form for reaction systems which can help, in turn, to perform simulation, analysis, control, and optimization more efficiently.

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Appendices

A Proof of Theorem 2

The construction of the transformation to normal form was shown in the text. However, the reconstruction of the original states \mathbf{n} and V from \mathbf{x} and ν remains to be proven.

The relationship $\begin{bmatrix} \mathbf{z} \\ V \end{bmatrix} \rightarrow \begin{bmatrix} \mathbf{x} \\ \nu \end{bmatrix}$ is given in (8). The inverse can be obtained after a few algebraic manipulations as:

$$\begin{bmatrix} \mathbf{z} \\ V \end{bmatrix} = \begin{bmatrix} \mathbf{x} \nu \\ h(\mathbf{x}_2) \nu \end{bmatrix}. \quad (\text{A.1})$$

According to (6), the number of moles \mathbf{n} satisfies:

$$\mathbf{z} = \begin{bmatrix} \mathbf{K}^+ (\mathbf{I}_S - \mathbf{C}_{in} \mathbf{M}^T) \\ \mathbf{M}^T \\ \mathbf{N}^T \end{bmatrix} \mathbf{n}. \quad (\text{A.2})$$

Multiplying both sides of (A.2) by $\begin{bmatrix} \mathbf{K} & \mathbf{C}_{in} & \mathbf{N} \end{bmatrix}$ and noting that $\mathbf{K}\mathbf{K}^+ + \mathbf{L}\mathbf{L}^T + \mathbf{N}\mathbf{N}^T = \mathbf{I}_S$,

$$\begin{aligned} \begin{bmatrix} \mathbf{K} & \mathbf{C}_{in} & \mathbf{N} \end{bmatrix} \mathbf{z} &= \left(\mathbf{K}\mathbf{K}^+ (\mathbf{I}_S - \mathbf{C}_{in} \mathbf{M}^T) + \mathbf{C}_{in} \mathbf{M}^T + \mathbf{N}\mathbf{N}^T \right) \mathbf{n} \\ &= \left(\mathbf{K}\mathbf{K}^+ + (\mathbf{L}\mathbf{L}^T + \mathbf{N}\mathbf{N}^T) \mathbf{C}_{in} \mathbf{M}^T + \mathbf{N}\mathbf{N}^T \right) \mathbf{n} \\ &= \left(\mathbf{K}\mathbf{K}^+ + \mathbf{L}\mathbf{L}^T + \mathbf{N}\mathbf{N}^T \right) \mathbf{n} = \mathbf{n}. \end{aligned} \quad (\text{A.3})$$

Substituting $\mathbf{z} = \mathbf{x}\nu$ into (A.3) leads to $\mathbf{n} = \nu(\mathbf{K}\mathbf{x}_1 + \mathbf{C}_{in}\mathbf{x}_2 + \mathbf{N}\mathbf{x}_3)$ and, hence, the theorem follows. \square

B Proof of Corollary 3

The only difference between Corollary 3 and Theorem 2 is that $\mathbf{M} = \mathbf{L}(\mathbf{C}_{in}^T \mathbf{L})^+$. However, since $\mathbf{L}\mathbf{L}^T \mathbf{C}_{in} \mathbf{M}^T = \mathbf{L}(\mathbf{L}^T \mathbf{C}_{in})(\mathbf{L}^T \mathbf{C}_{in})^+ \mathbf{L}^T = \mathbf{L}\mathbf{L}^T$, the equation $\mathbf{K}\mathbf{K}^+ + \mathbf{L}\mathbf{L}^T + \mathbf{N}\mathbf{N}^T = \mathbf{I}_S$ in the proof of Theorem 2 still holds and, hence, the corollary follows. \square

C Proof of Proposition 7

Note that $\mathbf{c} = \mathbf{n}/V$. The measured and unmeasured molar concentrations induce the following partition in (12):

$$\frac{\mathbf{n}}{\nu} = h(\mathbf{x}_2) \mathbf{c} = h(\mathbf{x}_2) \begin{bmatrix} \mathbf{c}_s \\ \mathbf{c}_t \end{bmatrix} = \begin{bmatrix} \mathbf{K}_s \\ \mathbf{K}_t \end{bmatrix} \mathbf{x}_1 + \begin{bmatrix} \mathbf{C}_{in,s} \\ \mathbf{C}_{in,t} \end{bmatrix} \mathbf{x}_2 + \begin{bmatrix} \mathbf{N}_s \\ \mathbf{N}_t \end{bmatrix} \mathbf{x}_3. \quad (\text{C.1})$$

The states \mathbf{x}_2 , \mathbf{x}_3 , and ν can be reconstructed from the inlet and outlet streams, \mathbf{c}_0 , and V_0 without knowledge of the kinetics. Owing to the existence of \mathbf{K}_s^+ , by assumption \mathbf{x}_1 can be reconstructed from \mathbf{c}_s as:

$$\mathbf{x}_1 = \mathbf{K}_s^+ (h(\mathbf{x}_2) \mathbf{c}_s - \mathbf{C}_{in,s} \mathbf{x}_2 - \mathbf{N}_s \mathbf{x}_3). \quad (\text{C.2})$$

Having reconstructed \mathbf{x}_1 , (13) follows.

Adding and subtracting the initial conditions of \mathbf{x}_2 and noting that \mathbf{x}_3 stay at their initial value, the estimate can be written as

$$\begin{aligned} \hat{\mathbf{c}}_t &= \mathbf{K}_t \mathbf{K}_s^+ \mathbf{c}_s + \left(\mathbf{C}_{in,t} - \mathbf{K}_t \mathbf{K}_s^+ \mathbf{C}_{in,s} \right) (\hat{\mathbf{x}}_2 - \hat{\mathbf{x}}_2(0)) / \hat{h} \\ &\quad - \left(\mathbf{C}_{in,t} - \mathbf{K}_t \mathbf{K}_s^+ \mathbf{C}_{in,s} \right) \hat{\mathbf{x}}_2(0) / \hat{h} + \left(\mathbf{N}_t - \mathbf{K}_t \mathbf{K}_s^+ \mathbf{N}_s \right) \hat{\mathbf{x}}_3(0) / \hat{h}. \end{aligned} \quad (\text{C.3})$$

By assumption, there is no uncertainty in the initial volume, and material exchange terms. Using this fact and noting that $h_0 \nu_0 = \hat{h}_0 \hat{\nu}_0 = V_0$,

$$\hat{\nu} \nu_0 = \nu \hat{\nu}_0, \quad (\mathbf{x}_2 - \mathbf{x}_2(0)) \nu_0 = (\hat{\mathbf{x}}_2 - \hat{\mathbf{x}}_2(0)) \hat{\nu}_0, \quad h \nu_0 = \hat{h} \hat{\nu}_0$$

and, thus,

$$(\mathbf{x}_2 - \mathbf{x}_2(0)) / h = (\hat{\mathbf{x}}_2 - \hat{\mathbf{x}}_2(0)) / \hat{h}. \quad (\text{C.4})$$

This means that there is no error in the estimation of $(\mathbf{x}_2 - \mathbf{x}_2(0)) / h$. Using (C.4) and also the fact that \mathbf{c}_s is measured without uncertainty, the estimation error $\mathbf{e} \equiv \hat{\mathbf{c}}_t - \mathbf{c}_t$ becomes:

$$\mathbf{e} = \frac{-\left(\mathbf{C}_{in,t} - \mathbf{K}_t \mathbf{K}_s^+ \mathbf{C}_{in,s} \right) (\hat{\mathbf{x}}_2(0) \nu_0 / \hat{\nu}_0 - \mathbf{x}_2(0)) + \left(\mathbf{N}_t - \mathbf{K}_t \mathbf{K}_s^+ \mathbf{N}_s \right) (\hat{\mathbf{x}}_3(0) \nu_0 / \hat{\nu}_0 - \mathbf{x}_3(0))}{h}. \quad (\text{C.5})$$

Note that the numerator in (C.5) is constant, which implies $\frac{d}{dt}(h \mathbf{e}) = \mathbf{0}_{S_t}$. Thus, the numerator is given by $\mathbf{e}(0) h(\mathbf{x}_2(0))$, leading to (14). The error dynamics are given by

$$\dot{\mathbf{e}} = -d \mathbf{e}, \quad (\text{C.6})$$

where $d = (\mathbf{1}_p^\top \mathbf{q}_{in})/V$ is the so-called *dilution* term. If at least one of the inlet flowrates is non-zero for $t \rightarrow \infty$, then $d > 0$. Hence, the error dynamics (C.6) are stable, and \mathbf{e} converges asymptotically to $\mathbf{0}_{S_t}$. Consequently, the result proposed by Bastin and Dochain (1990) is obtained:

$$\lim_{t \rightarrow \infty} \hat{\mathbf{c}}_t = \mathbf{c}_t, \quad d > 0. \quad (\text{C.7})$$

□

D Proof of Proposition 8

A system is feedback linearizable iff

(FL1) the distributions $\Delta_i = \text{span}\{\mathbf{f}^k \mathbf{g}_j : 0 \leq k \leq i, 1 \leq j \leq m\}^2$ have constant rank everywhere and are closed under Lie-bracketing (involutive) for all $i \geq 0$, and

(FL2) $\text{rank}(\Delta_{n-1}) = n$.

Consider the distributions

$$\Delta_0 = \text{span}\{\mathbf{g}_j, j = 1, \dots, m\}, \quad \Delta_1 = \text{span}\{\mathbf{g}_j, [\mathbf{f}, \mathbf{g}_j], j = 1, \dots, m\}. \quad (\text{D.1})$$

It can be verified that arranging the vectors of the distribution Δ_1 leads to

$$\begin{bmatrix} \mathbf{g}_1 & \dots & \mathbf{g}_m & [\mathbf{f}, \mathbf{g}_1] & \dots & [\mathbf{f}, \mathbf{g}_m] \end{bmatrix} = \begin{bmatrix} \mathbf{0}_{R \times m} & -\mathbf{J} \\ \mathbf{I}_m & \mathbf{0}_{m \times m} \end{bmatrix}. \quad (\text{D.2})$$

Since by assumption \mathbf{J} is of rank R , it follows that $\text{rank}(\Delta_1) = (R + m)$ which is the maximum possible rank. Thus, Δ_1 is involutive. Furthermore, Δ_0 has rank m and is involutive since $[\mathbf{g}_i, \mathbf{g}_j] = \mathbf{0}, \forall i, j = 1, \dots, p$, and $[\mathbf{g}_i, \mathbf{g}_m] = (\mathbf{g}_m - \mathbf{g}_i)/(\nu h(\mathbf{x}_2)), \forall i = 1, \dots, m$. Thus, (15) is feedback linearizable. □

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² $[\cdot, \cdot]$ denotes the Lie-bracket and $[\mathbf{f}^k \mathbf{g}_j]$ the iterated Lie-bracket $[\mathbf{f}, \mathbf{f}^{(k-1)} \mathbf{g}_j]$.

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