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## **APPENDICES**

# APPENDIX A

## RELATIVE GAIN ARRAY

There are two basic methods used for calculating the RGA for a square, linear system, whose transfer function matrix  $\mathbf{G}(s)$  is available: (1) the “first principles” method, and (2) the matrix method.

### *Calculating RGA's from First Principles*

We will illustrate this procedure with the 2 x 2 system of Eqs. (3.1) and (3.2).

First, we observe that, by definition, the RGA is concerned with steady-state conditions, thus we only need the steady-state form of this model, which is:

$$y_1 = K_{11}m_1 + K_{12}m_2 \quad (\text{A.1})$$

$$y_2 = K_{21}m_1 + K_{22}m_2 \quad (\text{A.2})$$

To obtain  $\lambda_{11}$  from the definition in Eq. (3.6), we need to evaluate both of the indicated partial derivatives from Eqs. (A.1) and (A.2) as follows: Eqs. (A.1) and (A.2) represent steady-state, open-loop conditions; therefore, the numerator partial derivative in Eq. (3.6) is obtained from Eq. (A.1) by straightforward differentiation:

$$\left( \frac{\partial y_1}{\partial m_1} \right)_{\text{all loops open}} = K_{11} \quad (\text{A.3})$$

The second partial derivative calls for Loop 2 to be closed, so that in response to changes in  $m_1$ , the second control variable  $m_2$  can be used ultimately to restore  $y_2$  to its initial value of 0. Thus, to obtain the second partial derivative, we first find, from Eq. (A.2), the value  $m_2$  must take to keep  $y_2 = 0$  in the face of changes in  $m_1$ ; what effect this will have on  $y_1$  is deduced by substituting this value for  $m_2$  in Eq. (A.1).

Setting  $y_2 = 0$  and solving for  $m_2$  in Eq. (A.2) gives:

$$m_2 = -\frac{K_{21}}{K_{22}}m_1 \quad (\text{A.4})$$

and substituting this into Eq. (A.1) gives:

$$y_1 = K_{11}m_1 - \frac{K_{12}K_{21}}{K_{22}}m_1 \quad (\text{A.5})$$

Having thus “eliminated”  $m_2$  we may then differentiate, so that now:

$$\left( \frac{\partial y_1}{\partial m_1} \right)_{\text{loop 2 closed}} = K_{11} \left( 1 - \frac{K_{12}K_{21}}{K_{11}K_{22}} \right) \quad (\text{A.6})$$

and we obtain, finally, that:

$$\lambda_{11} = \frac{1}{1-\zeta} \quad (\text{A.7})$$

where

$$\zeta = \frac{K_{12}K_{21}}{K_{11}K_{22}} \quad (\text{A.8})$$

As an exercise, the reader is encouraged to follow the illustrated procedure and confirm that the other relative gains,  $\lambda_{12}$ ,  $\lambda_{21}$ , and  $\lambda_{22}$ , are given by:

$$\lambda_{12} = \lambda_{21} = \frac{-\zeta}{1-\zeta} \quad (\text{A.9})$$

and

$$\lambda_{22} = \lambda_{11} = \frac{1}{1-\zeta} \quad (\text{A.10})$$

Thus the RGA for the 2 x 2 system is given by:

$$\Lambda = \begin{bmatrix} \frac{1}{1-\zeta} & \frac{-\zeta}{1-\zeta} \\ \frac{-\zeta}{1-\zeta} & \frac{1}{1-\zeta} \end{bmatrix} \quad (\text{A.11})$$

Note carefully that the elements add up to unity across any row as well as down any column.

It is useful to observe that if we define:

$$\lambda = \lambda_{11} = \frac{1}{1-\zeta} \quad (\text{A.12})$$

for this 2 x 2 system, then the RGA may be written:

$$\Lambda = \begin{bmatrix} \lambda & 1-\lambda \\ 1-\lambda & \lambda \end{bmatrix} \quad (\text{A.13})$$

The immediate consequence is that the single element  $\lambda$  is sufficient to determine the RGA for a 2 x 2 system. Thus for 2 x 2 system,  $\lambda$  is often called the relative gain parameter.

If we realistically assess this first-principles method of obtaining the RGA, it will not be difficult to come to the conclusion that it will be too tedious to apply in cases where the system is of higher dimension than 2 x 2. The problem is not with the numerator partial derivatives – these are always easy to obtain. The problem is with the denominator partial derivatives: the procedure for their evaluation in higher dimensional systems will involve solving several systems of linear algebraic equations.

The fact that linear algebraic equations are quite conveniently solved by matrix methods should then alert us to the possibility of using matrix methods to simplify the tedium involved in calculating RGA's, especially for higher dimensional systems.

### *The Matrix Method for Calculating RGA's*

Let  $\mathbf{K}$  be the matrix of steady-state gains of the transfer function matrix  $\mathbf{G}(s)$ , i.e.:

$$\lim_{s \rightarrow 0} \mathbf{G}(s) = \mathbf{K} \quad (\text{A.14})$$

whose elements are  $K_{ij}$ ; further, let  $\mathbf{R}$  be the transpose of the inverse of this steady-state gain matrix, i.e.:

$$\mathbf{R} = (\mathbf{K}^{-1})^T$$

with elements  $r_{ij}$ . Then, it is possible to show that the elements of the RGA can be obtained from the elements of these two matrices according to:

$$\lambda_{ij} = K_{ij} r_{ij} \quad (\text{A.15})$$



It is important to note that the above equation indicates an element-by-element multiplication of the corresponding elements of the two matrices  $\mathbf{K}$  and  $\mathbf{R}$ ; it has nothing to do with the standard matrix product.

### Computing The RGA with MATLAB

If  $G$  is a constant matrix then the RGA can be computed using

$$\text{RGA} = G.*\text{pinv}(G.');$$

If  $G(j\omega)$  is a frequency-dependent matrix generated using the  $\mu$  toolbox, e.g.

$$G = \text{pck}(A,B,C,D); \text{omega}=\text{logspace}(-2,2,41); Gw=\text{frsp}(G,\text{omega});$$

Then the RGA as a function of frequency can be computed using

$$\text{RGA}_w = \text{veval}('.*',Gw,\text{vpinv}(\text{vtp}(Gw)));$$

## Appendix B

### Data of HDA Process for Simulation

Name	Fresh Hydrogen	Fresh Toluene	purge	stabilizer gas	benzene product	diphenyl product
Vapour Fraction	1.00	0.00	1.00	1.00	0.00	0.00
Temperature [F]	86.00	86.00	113.00	123.87	221.99	558.83
Pressure [psia]	605.00	635.00	476.80	150.00	30.00	31.00
Molar Flow [lbmole/hr]	490.38	289.00	483.40	19.73	270.81	6.28
Mass Flow [lb/hr]	1194.96	26628.69	5379.81	343.48	21154.42	968.31
Heat Flow [Btu/hr]	-4.45E+05	1.75E+06	-8.87E+06	-5.15E+05	7.09E+06	6.50E+05
Comp Mole Frac (Hydrogen)	0.9700	0.0000	0.4012	0.0902	0.0000	0.0000
Comp Mole Frac (Methane)	0.0300	0.0000	0.5875	0.8674	0.0000	0.0000
Comp Mole Frac (Benzene)	0.0000	0.0000	0.0101	0.0420	0.9997	0.0000
Comp Mole Frac (Toluene)	0.0000	1.0000	0.0012	0.0003	0.0003	0.0003
Comp Mole Frac (BiPhenyl)	0.0000	0.0000	0.0000	0.0000	0.0000	0.9997

### Data of HDA Process for Simulation (Continue)

Name	gas recycle	toluene recycle	furnance in	reactor in	reactor out	quench
Vapour Fraction	1.00	0.00	1.00	1.00	1.00	0.00
Temperature [F]	148.71	284.17	1106.00	1150.00	1230.97	113.83
Pressure [psia]	575.00	615.00	513.00	503.00	486.00	486.00
Molar Flow [lbmole/hr]	3519.20	84.67	4383.25	4383.25	4383.25	109.10
Mass Flow [lb/hr]	39119.40	7800.72	74743.78	74743.78	74742.70	8654.94
Heat Flow [Btu/hr]	-6.35E+07	1.18E+06	6.33E+05	3.76E+06	3.76E+06	1.88E+06
Comp Mole Frac (Hydrogen)	0.4020	0.0000	0.4313	0.4313	0.3668	0.0047
Comp Mole Frac (Methane)	0.5868	0.0000	0.4745	0.4745	0.5404	0.0449
Comp Mole Frac (Benzene)	0.0100	0.0006	0.0080	0.0080	0.0711	0.7124
Comp Mole Frac (Toluene)	0.0012	0.9993	0.0862	0.0862	0.0203	0.2215
Comp Mole Frac (BiPhenyl)	0.0000	0.0000	0.0000	0.0000	0.0014	0.0165

### Data of HDA Process for Simulation (Continue)

Name	fehe hot in	fehe hot out	sep gas out	stabilizer feed	stabilizer bottom	product bottom
Vapour Fraction	1.00	0.95	1.00	0.03	0	0
Temperature [F]	1149.90	228.56	113.00	114.75	374.7792337	292.5600107
Pressure [psia]	486.00	480.00	476.80	152.00	153	33
Molar Flow [lbmole/hr]	4492.35	4492.35	4002.60	381.25	361.5212442	90.70702121
Mass Flow [lb/hr]	83397.64	83397.64	44545.55	30244.74	29901.25883	8746.842286
Heat Flow [Btu/hr]	5.64E+06	-5.60E+07	-7.34E+07	6.56E+06	10784879.86	1660476.999
Comp Mole Frac (Hydrogen)	0.3580	0.3580	0.4012	0.0047	0.0000	0.0000
Comp Mole Frac (Methane)	0.5284	0.5284	0.5875	0.0449	0.0000	0.0000
Comp Mole Frac (Benzene)	0.0866	0.0866	0.0101	0.7124	0.7490	0.0006
Comp Mole Frac (Toluene)	0.0252	0.0252	0.0012	0.2215	0.2336	0.9302
Comp Mole Frac (BiPhenyl)	0.0018	0.0018	0.0000	0.0165	0.0174	0.0692

## Appendix C

### Parameter Tuning of Three Control Structures

Controller	Reference Control Structure		Control Structure 1		Control Structure 2	
	$K_c$	$T_i$	$K_c$	$T_i$	$K_c$	$T_i$
PCR	2	10	2	10	2	10
FC	0.225	0.0419	0.1	0.0125	0.1	0.0125
TCR	0.2	0.5	0.2	0.5	0.2	0.5
TCQ	0.105	0.383	0.105	0.383	0.0726	0.252
TCC	0.5	5	0.5	5	0.5	5
LC	2	-	2	-	2	-
CC	0.449	14.1	0.449	14.1	0.449	14.1
PC1	1	10	1	10	1	10
LC_C1	1	-	1	-	1	-
LC_R1	2	-	2	-	2	-
TC1	1	12	1	12	1	12
PC2	1	20	1	20	1	20
LC_C2	2	-	2	-	2	-
LC_R2	2	-	2	-	2	-
TC2	2	10	2	10	2	10
PC3	1	15	1	15	1	15
LC_C3	1	-	1	-	1	-
LC_R3	2.9	-	2.9	-	2.9	-
TC3	1	30	1	30	1	30

## VITA



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