Test Report: Stoichiometry Mass-Mass Program

Deemah Alomair

December 26, 2019

1 Revision History

Date	Version	Notes
26/12/2019	1.0	First version of the document

2 Symbols, Abbreviations and Acronyms

symbol	description
Τ	Test
SMMP	Stoichiometry Mass-Mass Program

Contents

1	Revision History						
2	Symbols, Abbreviations and Acronyms						
3	Functional Requirements Evaluation						
4	Nonfunctional Requirements Evaluation 4.1 Usability	1 1 1					
5	Comparison to Existing Implementation	2					
6	Unit Testing 6.1 Mass Input Test-id1 6.2 Reaction Input Test-id2 6.3 Reaction Output Test-id3, Mass Output Test-id4 6.4 Check Balance Test-id5 6.5 Molecular Weight Calculation Test-id6 6.6 Mole1 Calculation Test-id7 6.7 Mole Ratio Calculation Test-id8 6.8 Mole2 Calculation Test-id9 6.9 Mass Calculation Test-id10	2 3 5 5 6 7 7 8					
7	Changes Due to Testing	9					
8	Automated Testing	9					
9	Trace to Requirements	10					
10	Trace to Modules	10					
11	Code Coverage Metrics	11					
\mathbf{L}_{i}	ist of Tables						
	1 Reliability testing of SMMP as comparison to online balancer	9					

2	Traceability Matrix Showing the Connections Between unit
	test cases and functional requirements
3	Traceability Matrix Showing the Connections Between unit
	test cases and Nonfunctional requirements
4	Traceability Matrix Showing the Connections Between Mod-
	ules and Test Cases
- .	0 T1
List	of Figures
1	error massage if user enter non number mass value
2	error massage if user enter mass less than 1
3	error massage if user enter atom less than 1 after selecting an
	element
4	error massage if user enter non number atom value 4
5	final system output
6	balance reaction
7	non balance reaction
8	molecular weight value
9	Mole1 value
10	Mole Ratio value
11	Mole2 value
12	final mass value
13	coverage testing
10	COverage resulting \cdot , \cdot

This document report result from system test cases found in System Verification and Validation Plan document (1).

3 Functional Requirements Evaluation

Functional requirements are evaluated using system test cases from id1 to id10. All the details regarding functional requirements evaluation can be found in section 6. The traceability between system test cases and functional requirements found in table 2 of section 9.

4 Nonfunctional Requirements Evaluation

Nonfunctional Requirements evaluated in system test cases id11 - id12. All the details regarding Nonfunctional requirements evaluation can be found in this section. The traceability between system test cases and Nonfunctional requirements found in table 3 of section 9.

4.1 Usability

system test id11 measures the Usability of SMMP system. The survey measured the satisfaction level of potential user after using SMMP. How easy and understandable the system is?. It needs to be filled by any user and if satisfactory level is low then enhancement need to be taken into account. survey can be found in Appendix of (1).

4.2 Reliability

system test id12 measures the Reliability of SMMP system. 25 different unbalanced chemical reactions were tested and compared to answer of online balancer (2) as parallel testing. the aim was to get 100% of correct answers that includes right balance reaction and correct mass value. Below table shows the final result compared to the online balancer and the final percentage of correctness. The goal of Reliability had been accomplished. all tested reactions were balanced correctly as compared to the online balancer.

Unbalanced Chemical Reaction Online Balancer R		SMMP Result	correctness
$CH_4 + O_2 \rightarrow CO_2 + H_2O$	$CH_4 + O_2 \to CO_2 + H_2O$ $CH_4 + 2O_2 \to CO_2 + 2H_2O$		correct
$Fe_2O_3 + C \rightarrow Fe + CO_2$	$2Fe_2O_3 + 3C \rightarrow 4Fe + 3CO_2$	$2Fe_2O_3 + 3C \rightarrow 4Fe + 3CO_2$	correct
$N_2 + O_2 \rightarrow N_2 O_5$	$2N_2 + 5O_2 \rightarrow 2N_2O_5$	$2N_2 + 5O_2 \rightarrow 2N_2O_5$	correct
$CH_4 + Cl_2 \rightarrow CCl_4 + HCl$	$CH_4 + 4Cl_2 \rightarrow CCl_4 + 4HCl$	$4CH_4 + Cl_2 \rightarrow CCl_4 + 4HCl$	correct
$N_2 + H_2 \rightarrow NH_3$	$N_2 + 3H_2 \rightarrow 2NH_3$	$N_2 + 3H_2 \rightarrow 2NH_3$	correct
$Fe + H_2O \rightarrow Fe_3O_4 + H_2$	$3Fe + 4H_2O \rightarrow Fe_3O_4 + 4H_2$	$3Fe + 4H_2O \rightarrow Fe_3O_4 + 4H_2$	correct
$Xe + F_2 \rightarrow XeF_6$	$Xe + 3F_2 \rightarrow XeF_6$	$Xe + 3F_2 \rightarrow XeF_6$	correct
$\mathrm{Hg} + O_2 \to \mathrm{HgO}$	$2 \text{Hg} + O_2 \rightarrow 2 \text{HgO}$	$2 \text{Hg} + O_2 \rightarrow 2 \text{HgO}$	correct
$CaO + C \rightarrow CaC_2 + CO$	$CaO + 3C \rightarrow CaC_2 + CO$	$CaO + 3C \rightarrow CaC_2 + CO$	correct
$S_8 + F_2 \rightarrow SF_6$	$S_8 + 24F_2 \rightarrow 8SF_6$	$S_8 + 24F_2 \rightarrow 8SF_6$	correct
$Mg + N_2 \rightarrow Mg_3N_2$	$3Mg + N_2 \to Mg_3N_2$	$3Mg + N_2 \to Mg_3N_2$	correct
$BeF_2 + Mg \rightarrow MgF_2 + Be$ balance		balance	correct
$\operatorname{Zn} + \operatorname{HCl} \to \operatorname{Zn}Cl_2 + H_2$	$\operatorname{Zn} + 2\operatorname{HCl} \to \operatorname{Zn}Cl_2 + H_2$	$\operatorname{Zn} + 2\operatorname{HCl} \to \operatorname{Zn}Cl_2 + H_2$	correct
$SiC + Cl_2 \rightarrow SiCl_4 + C$ $SiC + 2Cl_2 \rightarrow SiCl_4 + C$		$SiC + 2Cl_2 \rightarrow SiCl_4 + C$	correct
$MnS + HCl \rightarrow H_2S + MnCl_2$	$MnS + 2HCl \rightarrow H_2S + MnCl_2$	$MnS + 2HCl \rightarrow H_2S + MnCl_2$	correct
$UF_4 + Mg \rightarrow MgF_2 + U$	$UF_4 + 2Mg \rightarrow 2MgF_2 + U$	$UF_4 + 2Mg \rightarrow 2MgF_2 + U$	correct
$S + N_2O \rightarrow SO_2 + N_2$	$S + 2N_2O \rightarrow SO_2 + 2N_2$	$S + 2N_2O \rightarrow SO_2 + 2N_2$	correct
$SiH_4 + O_2 \rightarrow SiO_2 + H_2O$	$\mathrm{Si}H_4 + 2O_2 \to \mathrm{Si}O_2 + 2H_2\mathrm{O}$	$SiH_4 + 2O_2 \rightarrow SiO_2 + 2H_2O$	correct
$TiCl_4 + Mg \rightarrow MgCl_2 + Ti$	$TiCl_4 + 2Mg \rightarrow 2MgCl_2 + Ti$	$TiCl_4 + 2Mg \rightarrow 2MgCl_2 + Ti$	correct
$Si + S_8 \rightarrow Si_2S_4$	$4Si + S_8 \rightarrow 2Si_2S_4$	$4\mathrm{Si} + S_8 \to 2Si_2S_4$	correct
$SiO_2 + HF \rightarrow SiF_4 + H_2O$	$SiO_2 + 4HF \rightarrow SiF_4 + 2H_2O$	$SiO_2 + 4HF \rightarrow SiF_4 + 2H_2O$	correct
$P_4 + O_2 \rightarrow P_2O_5$	$P_4 + 5O_2 \rightarrow 2P_2O_5$	$P_4 + 5O_2 \rightarrow 2P_2O_5$	correct
$Sb + O_2 \to Sb_4O_6$	$4Sb + 3O_2 \to Sb_4O_6$	$4Sb + 3O_2 \to Sb_4O_6$	correct
$UO_2 + HF \rightarrow UF_4 + H_2O$	$UO_2 + 4HF \rightarrow UF_4 + 2H_2O$	$UO_2 + 4HF \rightarrow UF_4 + 2H_2O$	correct
$Al + O_2 \rightarrow Al_2O_3$	$4Al + 3O_2 \rightarrow 2Al_2O_3$	$4Al + 3O_2 \rightarrow 2Al_2O_3$	correct

Table 1: Reliability testing of SMMP as comparison to online balancer.

5 Comparison to Existing Implementation

This is stand alone system. If there is any existing systems with same functionalities and goals the developer is not aware about them and no comparison had been made.

6 Unit Testing

6.1 Mass Input Test-id1

The goal of this test is to make sure user enter a positive number in mass value widget. the default mass value is 1. pictures below illustrate all possible entries by user that cause error massage which matches cases of id1, Table 1 in

(1). The tests had been passed correctly and system responded as intended.

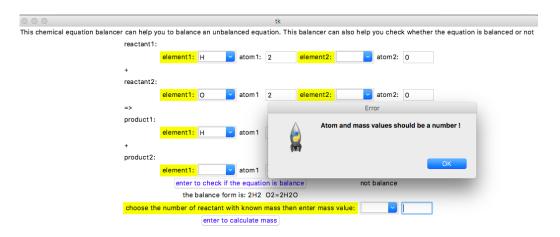


Figure 1: error massage if user enter non number mass value

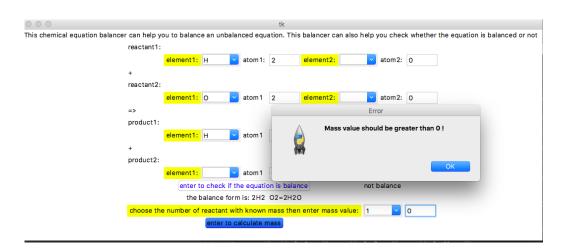


Figure 2: error massage if user enter mass less than 1

6.2 Reaction Input Test-id2

The goal of this test is to ensure that user had entered a correct chemical reaction format that satisfies the system requirements and enter a positive

number for atom value. More details of accepted reaction format can be found in Unit Verification and Validation Plan (3) and Unit Verification and Validation Test Report (4). Test case id2 testing entered atom value. pictures bellow show all possible entires of atom value and system responds to each case. The test passed with right system behavior.

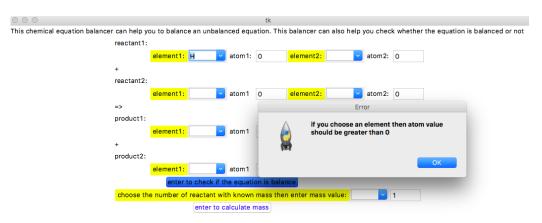


Figure 3: error massage if user enter atom less than 1 after selecting an element

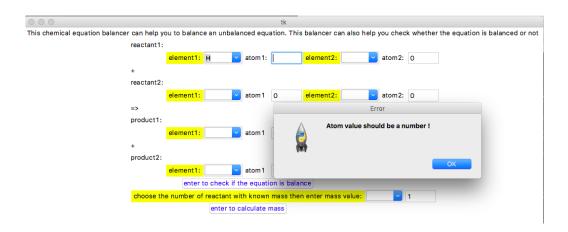


Figure 4: error massage if user enter non number atom value

6.3 Reaction Output Test-id3, Mass Output Test-id4

The goal of this test is to output the final result including the mass and balance reaction to GUI. This is the main goal of the system. Picture below shows how system will print out the final result to end user. This test is for test cases id3, id4 in (1). Test passed correctly and system responded as intended.

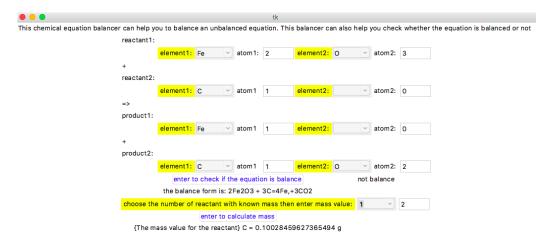


Figure 5: final system output

6.4 Check Balance Test-id5

The goal of this test is to check if the entered reaction is balance or not. Display "balance" if yes and "not balance" with new balanced form if not. Pictures below show how system respond to balance and non balance reactions. The test passed correctly and system responded as intended.

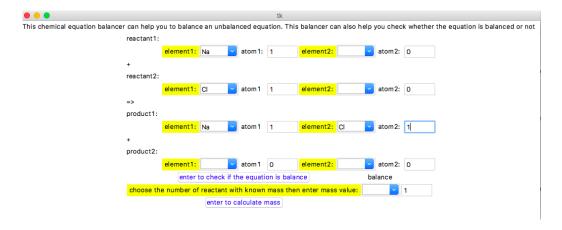


Figure 6: balance reaction

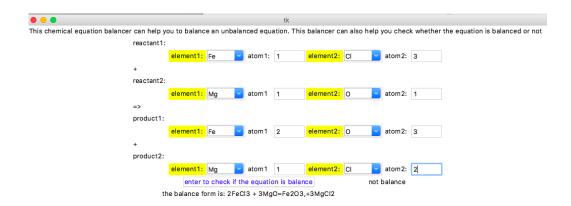


Figure 7: non balance reaction

6.5 Molecular Weight Calculation Test-id6

The goal of this test is to get molecular weight for a reactant. Picture below shows molecular weight is printed for reactant " Fe_2O_3 " as the test case id6 in (1) and the result is identical. Test passed correctly and system responded as intended.

```
Run: project ×

//Users/deemaalomair/PycharmProjects/CAS741/venv/bin/python /Users/deemaalomair/PycharmProjects/CAS741/project.py
Reactant dictionary: {'Fe': 2, '0': 3, 'C': 1}
Product dictionary: {'Fe': 4, '0': 6} 'C': 3}
Product1 dictionary: {'Fe': 4, '0': 6} 'C': 3}
Product1 dictionary: {'Fe': 4, 'C': 3, '0': 6} 'C': 3}

product1 dictionary: {'Fe': 4, 'C': 3, '0': 6} 'C': 3}
product1 dictionary: {'Fe': 4, 'C': 3, '0': 6} 'C': 3}
product1 dictionary: {'Fe': 4, 'C': 3, '0': 6} 'C': 3}
product1 dictionary: {'Fe': 4, 'C': 3, '0': 6} 'C': 3}
product1 dictionary: {'Fe': 4, 'C': 3, '0': 6} 'C': 3}
product1 dictionary: {'Fe': 4, 'C': 3, '0': 6} 'C': 3}
product1 dictionary: {'Fe': 4, 'C': 3, '0': 6} 'C': 3}
product1 dictionary: {'Fe': 4, 'C': 3, '0': 6} 'C': 3}
product1 dictionary: {'Fe': 4, 'C': 3, '0': 6} 'C': 3}
product1 dictionary: {'Fe': 4, 'C': 3, '0': 6} 'C': 3}
product1 dictionary: {'Fe': 4, 'C': 3, '0': 6} 'C': 3}
product1 dictionary: {'Fe': 4, 'C': 3, '0': 6} 'C': 3}
product1 dictionary: {'Fe': 4, 'C': 3, '0': 6} 'C': 3}
product1 dictionary: {'Fe': 4, 'C': 3, '0': 6} 'C': 3}
product1 dictionary: {'Fe': 4, 'C': 3, '0': 6} 'C': 3}
product1 dictionary: {'Fe': 4, 'C': 3, '0': 6} 'C': 3}
product1 dictionary: {'Fe': 4, 'C': 3, '0': 6} 'C': 3}
product1 dictionary: {'Fe': 4, 'C': 3, '0': 6} 'C': 3}
product1 dictionary: {'Fe': 4, 'C': 3, '0': 6} 'C': 3}
product1 dictionary: {'Fe': 4, 'C': 3, '0': 6} 'C': 3}
product1 dictionary: {'Fe': 4, 'C': 3, '0': 6} 'C': 3}
product1 dictionary: {'Fe': 4, 'C': 3, '0': 6} 'C': 3}
product1 dictionary: {'Fe': 4, 'C': 3, '0': 6} 'C': 3}
product1 dictionary: {'Fe': 4, 'C': 3, 'C': 3}
product1 dictionary: {'Fe': 4, '
```

Figure 8: molecular weight value

6.6 Mole1 Calculation Test-id7

The goal of this test is to get mole for reactant with known mass. we name this mole "Mole1". Picture below shows Mole1 value printed for reactant " Fe_2O_3 " as the test case id7 in (1) and the result is identical. Test passed correctly and system responded as intended.

```
Run: project ×

// Jers/deemaalomair/PycharmProjects/CAS741/venv/bin/python /Users/deemaalomair/PycharmProjects/CAS741/project.py
Reactant dictionary: {'Fe': 2, '0': 3, 'C': 1}
Product dictionary: {'Fe': 4, '0': 6, 'C': 3}
Reactant1 dictionary: {'Fe': 4, '0': 6, 'C': 3}
Product1 dictionary: {'Fe': 4, '0': 6, 'C': 3}
Coefficient1: 2
coefficient2: 3
coefficient3: 4
coefficient4: 3
element name: 0 atomic mass: 15.9994
element name: Fe atomic mass: 55.845
reactant1 mane: Featomic mass: 55.845
reactant2 mane: Featomic mass: 55.845
reactant3 mane: Featomic mass: 55.845
reactant4 mane: Featomic mass: 55.845
reactant5 mane: Featomic mass: 55.845
reactant6 mane: Featomic mass: 55.845
reactant7 mane: Featomic mass: 55.845
reactant6 mane: Featomic mass: 55.845
reactant6 mane: Featomic mass: 55.845
reactant7 mane: Featomic mass: 55.845
reactant3 mane: Featomic mass: 55.845
reactant3 mane: Featomic mass: 55.845
reactant4 dictionary: {'Fe': 4, 'C': 3, '0': 6}
reactant5 mane: Featomic mass: 55.845
reactant6 mane: Featomic mass: 55.845
reactant7 mane: Featomic mass: 55.845
reactant6 mane: Featomic mass: 55.845
reactant7 mane: Featomic mass: 55.845
reactant6 mane: Featomic mass: 55.845
reactant6 mane: Featomic mass: 55.845
reactant7 mane: Featomic mass: 55.845
reactant6 mane: Featomic mass: 55.845
reactant7 mane: Featomic mass: 55.845
```

Figure 9: Mole1 value

6.7 Mole Ratio Calculation Test-id8

The goal of this test is to get mole ratio between given reactants. Picture below shows Mole Ratio value for coefficient of reactant2/coefficient of reactant1 printed as the test case id8 in (1) and the result is identical. Test passed correctly and system responded as intended.

```
Run: project ×

// Josers/deemaalomair/PycharmProjects/CAS741/venv/bin/python /Users/deemaalomair/PycharmProjects/CAS741/project.py
Reactant dictionary: {'Fe': 2, '0': 3, 'C': 1}
Product dictionary: {'Fe': 4, '0': 6, 'C': 3}
Product dictionary: {'Fe': 4, 'C': 3, '0': 6}
coefficient1: 2
coefficient2: 3
coefficient3: 4
coefficient4: 3
element name: 0 atomic mass: 15.9994
element name: Fe atomic mass: 55.845
reactant name: Fe203 Molecular Weigh: 159.6882
Mole1: 0.012524406938020467
ratio 1.5
```

Figure 10: Mole Ratio value

6.8 Mole2 Calculation Test-id9

The goal of this test is to get mole for reactant with unknown mass. we name this mole "Mole2". Picture below shows Mole2 value printed as the test case id9 in (1) and the result is identical. Test passed correctly and system responded as intended.

```
Run: project ×

//Users/deemaalomair/PycharmProjects/CAS741/venv/bin/python /Users/deemaalomair/PycharmProjects/CAS741/project.py
Reactant dictionary: {'Fe': 2, '0': 3, 'C': 1}
Product dictionary: {'Fe': 4, '0': 6, 'C': 3}
Reactant1 dictionary: {'Fe': 4, 'C': 3, '0': 6}
coefficient1: 2
coefficient2: 3
coefficient3: 4
coefficient4: 3
element name: 0 atomic mass: 15.9994
element name: Fe atomic mass: 55.845
reactant name: Fe 203 Molecular Weigh: 159.6882
Mole1: 0.008349604625346977
```

Figure 11: Mole2 value

6.9 Mass Calculation Test-id10

The goal of this test is to get the final mass result for reactant with unknown mass. Picture below shows mass value printed as the test case id10 in (1) and the result is identical. Test passed correctly and system responded as intended.

```
Run:  project ×

// Jers/deemaalomair/PycharmProjects/CAS741/venv/bin/python /Users/deemaalomair/PycharmProjects/CAS741/project.py
Reactant dictionary: {'Fe': 2, '0': 3, 'C': 1}
Product dictionary: {'Fe': 4, '0': 6, 'C': 3}
Product dictionary: {'Fe': 4, 'C': 3, '0': 6}
coefficient1: 2
coefficient2: 3
coefficient3: 4
coefficient4: 3
element name: Pe atomic mass: 15.994
element name: Fe atomic mass: 55.845
reactant name: Fe203 Molecular Weigh: 159.6882
Mole1: 0.008349604625346977
ratio 1.5
mol2: 0.008349604625346977
reactant name: C10 Molecular Weigh: 12.0107
Final Mass: 0.10028459627365494
```

Figure 12: final mass value

7 Changes Due to Testing

No changes are necessary to the first stage of implementation due to these test results.

8 Automated Testing

- unit testing preformed for id1, id2 are all automated testing. they are done using pre-install ""unittest" package used by python. if assertion is not met then error massage will arise directly.
- coverage testing was preformed using coverage package without any error. blew picture shows coverage testing.

```
(venv) deema-2:CA5741 deemaalomair$ coverage run project.py
Rectant dictionary: ("H: 2, "0': 2)
Product dictionary: ("H: 4, "0': 2)
Rectant1 dictionary: ("H: 4, "0': 2)
Product1 dictionary: ("H: 4, "0': 2)
coefficient1: 2
coefficient3: 2
coefficient3: 2
coefficient3: 2
coefficient3: 2
coefficient4: 4
reactant name: Hatonic mass: 1.00794
reactant name: Hatonic mass: 1.00794
reactant name: HOW blocular Weigh: 2.01588
Molta: 0.92225496778025
ratio 0.5
molt2: 1.98042599394005
reactant name: COB Molecular Weigh: 31.9988
Final Mass: 63.49346191241541

■ Terminal ◆ PrimonComose ▶ 4:Rhm ◆ 5:Debog : IE 5:DODO

2 Eventog
```

Figure 13: coverage testing

9 Trace to Requirements

	id1	id2	id3	id4	id5	id6	id7	id8	id9	id10
R1(input)	X	X								
R2(output)			X	X						
R3(calculation)						X	X	X	X	X
R4(VerifyInputOutput)	X	X	X	X						

Table 2: Traceability Matrix Showing the Connections Between unit test cases and functional requirements

	id11	id12				
NF1		X				
NF2	X					

Table 3: Traceability Matrix Showing the Connections Between unit test cases and Nonfunctional requirements

10 Trace to Modules

A complete description of modules is found in the MG (5).

	id1	id2	id3	id4	id5	id6	id7	id8	id9	id10
Input Module		X								
Atomic Mass Module						X				
Balancing Chemical Reaction Module			X		X					
Mass Calculation Module						X	X	X	X	X
GUI Module			X	X						

Table 4: Traceability Matrix Showing the Connections Between Modules and Test Cases

11 Code Coverage Metrics

coverage test was done for the whole system and covers all functional requirements. It covers test cases from id1 to id10.

References

- [1] https://github.com/deemaalomair1/CAS741project/blob/master/docs/VnVPlan/SystVnVPlan/SystVnVPlan.pdf
- [2] http://www.endmemo.com/chem/balancer.php
- [3] https://github.com/deemaalomair1/CAS741project/blob/master/docs/VnVPlan/UnitVnVPlan/UnitVnVPlan.pdf
- [4] https://github.com/deemaalomair1/CAS741project/tree/master/docs/VnVReport
- [5] https://github.com/deemaalomair1/CAS741project/tree/master/docs/Design