

## **BCH714: Computational Systems Biology Minicourse 2024:**

### **The Program:**

<b>28/03 Thursday</b> <b>09:00</b> Peer grading of Assessment I JMR & DvN <b>11:00</b> Research lect. DvN <b>14:00</b> Introduction to Practical	<b>08/04 Monday</b> <b>9:00</b> Group 1 (First set of experiments) Group 2 (work on seminar) <b>14:00</b> Group 2 (First set of experiments) Group 1 (work on seminar)	<b>09/04 Tuesday</b> <b>9:00</b> Group 2 (Second set of experiments) Group 2 (work on seminar) <b>13:30</b> Group 1 (Second set of experiments) Group 1 (work on seminar)
<b>10/04 Wednesday</b> <b>9:00</b> Data analysis JMR <b>11:00</b> Redo experiments DvN	<b>11/04 Thursday</b> <b>09:00</b> Research lect. JMR <b>11:00</b> Data analysis JMR Redo experiments DvN	<b>12/04 Friday</b> <b>09:00</b> Data analysis (whole day) JMR
<b>15/04 Monday</b> <b>10:30</b> Data analysis (whole day) JMR	<b>16/04 Tuesday</b> <b>09:30</b> Research lect. JLS Finish final report <b>Hand in final report (16/04/2024) at 23:59</b>	

### **GROUPS:**

#### IA

- James Clarke
- Sophie de Villiers
- Elsa Pretorius

#### IB

- Courtney Campbell
- Amy Josephus
- Amogelang Mataboge

#### IC

- Haley Lewendal
- Suleiman Sungay
- Dom Beaumont

#### ID

- Brett Arnolds
- Faith Lekhu

#### 2A

- Theo Willemse
- Katinka Louw
- Cayley Sampson

#### 2B

- Conli Titus
- Cameron Bunch
- Marenique Smit

#### 2C

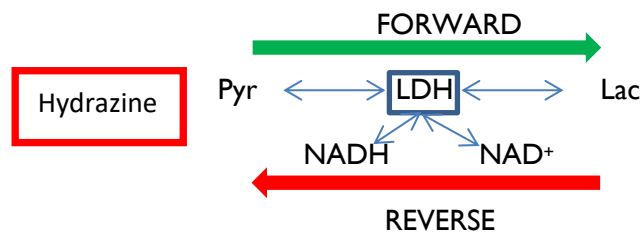
- Le Roux Strydom
- Chrisilie Henning

#### 2D

- Elna Kruger
- Kurauwone Marongwe

## ABOUT THE PRAC:

### (1) Characterise the enzyme:



### (2) The experiments:

#### Experiment 1: Rate as a function of substrate in the presence of low F16BP [0.5 $\mu$ M]

##### GROUP A

- Reverse direction
- Varying [ NAD<sup>+</sup>]
- Same [Lac] + [F16BP] + [CFE] + [Hydrazine]

##### GROUP B

- Reverse direction
- Varying [ Lac]
- Same [NAD<sup>+</sup>] + [F16BP] + [CFE] + [Hydrazine]

##### GROUP C

- Forward direction
- Varying [NADH]
- Same [Pyr] + [F16BP] + [CFE]

##### GROUP D

- Forward direction
- Varying [Pyr]
- Same [NADH] + [F16BP] + [CFE]

#### Experiment 2: Rate as a function of substrate in the presence of high F16BP [10 mM]

##### GROUP A

- Reverse direction
- Varying [NAD<sup>+</sup>]
- Same [Lac] + [F16BP] + [CFE] + [Hydrazine]

##### GROUP B

- Reverse direction
- Varying [Lac]
- Same [NAD<sup>+</sup>] + [F16BP] + [CFE] + [Hydrazine]

##### GROUP C

- Forward direction
- Varying [NADH]
- Same [Pyr] + [CFE] + [F16BP]

##### GROUP D

- Forward direction
- Varying [Pyr]
- Same [NADH] + [CFE] + [F16BP]

### (3) An enzyme assay:

- a- Pipette metabolites into the cuvettes, the reaction is initiated by the addition of the cell free extract (cfe), (which contains the enzyme of interest: LDH).
- b- Once the cell extract is added, measure the absorbance readings for the reaction over a period of 3mins.
- c- The types of data we expect to see- Forward reaction: decreasing Abs; Reverse reaction: increasing Abs

### (4) What needs to be done:

Create a pipetting schema so that we investigate 10 different concentrations around the Km values. Take the following into consideration:

Km		Stock concentrations	
Km(NADH)	= 0.08mM	[NADH]	= 5mM
Km(NAD <sup>+</sup> )	= 2.4mM	[NAD]	= 100mM
Km(Pyr)	= 1.5mM	[Pyr]	= 50mM
Km(Lac)	= 100mM	[Lac]	= 5M
		[F16BP]	= 0.1 mM, 100mM

#### b - Volumes:

- The total cuvette volume is 1 ml.
- If you are doing reverse direction experiments (only) 25  $\mu$ l of the 1 ml is the Hydrazine.
- 100  $\mu$ L of the total volume is cfe.
- The other volumes are of the substrate and co-factor which you calculated in the pipetting schema.
- Lastly, the remaining volume to make up this 1 ml volume is done by adding buffer to the cocktail mixture.

#### c - The ten substrate concentrations:

- The range you are going to investigate is between Km/10 and 10Km and the fixed substrates are usually at saturating condition i.e. 10 x Km. Use Excel to create an incremental pipetting schema.