# Proteomics data for androgen receptor splice variant-7 in circulating tumour cell project, led by Therese Becker

#### **Description**

The dataset contains the raw data for samples 7 and 8, proteins digested from a gel band excised from proteins extracted from PC3 cells. It also contains the ProteinLynx Global Server (Waters) search results of the proteins identified.

Expand/Collapse all

- Why?

# Why was the data collected and the research project that who it was produced by?

Was there a Research Data Management Plan (RDMP)? If so, identify the research project. Search by name to locate or edit RDMP.

#### **Project name**

Choice of antibody is critical for specific and sensitive detection of androgen receptor splice variant-7 in circulating tumor cells.

#### **Funding source**

Ingham Institute of Applied Medical Research

**Grant number(s)** 

FoR codes

**SEO** codes

# What is the data about?

#### **Data title**

Proteomics data for androgen receptor splice variant-7 in circulating tumour cell project, led by Therese Becker

#### **Description**

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#### Data type

Dataset

#### **Keywords**

androgen receptor deprivation therapy androgen receptor splice variant-7 antibody specificity, biomarker castrate resistant prostate cancer circulating tumor cells liquid biopsy

#### Data record size

11.8 GB

## Who was involved?

#### Principal investigator

Title Name Email ORCID

Dr Therese Becker T.Becker@westernsydney.edu.au 0000-0002-5636-9902

Data manager

Title Name Email ORCID

Dr David Harman d.harman@westernsydney.edu.au0000-0003-4376-832X

**Collaborators** 

Title Name Email ORCID

Associat Kieran Scott Kieran. Scott @westernsydney.edu. @000-0002-

Professor 6012-4107

Tanzila Khan 19153083@student.westernsydne 0.0001-a002-

7957-5944

John Lock john.lock@unsw.edu.au

Yafeng Ma yafeng.ma@unsw.edu.au

ProfessoPaul de Souza p.desouza@westernsydney.edu.a@000-0002-

7380-1170

Dr Wei Chua W.Chua@westernsydney.edu.au

Bavanthi Balakrishn Bava

ar

Bavanthi.Balakrishnar@helth.nsw.gov.au

**Principal Supervisor (students only)** 

Title Name Email ORCID

#### - When?

# When should your data be reviewed for disposal?

#### Applicable minimum retention period

7 years (contract/consultancy related)

Justification for extended retention period

Select the earliest possible disposal date of the data

#### - Where?

# Where is the data?

Туре	Data location	Notes
URL	https://rds.westernsydney.edu.au/Schools/SSH/2020/Becker_T/	

## Software and equipment used

Waters Synapt QTof Mass Spectrometer, nanoAcquity UPLC, Waters MassLynx software, and Waters PLGS software. Data also extracted into Excel format.