

# Molecular characterisation of drug-tolerant persister cells to overcome chemotherapy induced relapse in breast cancer

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## Background

The majority of primary **breast cancers** respond to **chemotherapeutic intervention**. However, years or even decades later many women suffer from **relapse**, which **gradually lose drug sensitivity** until ultimately **resistance develops**.

To prevent these, we aim to **investigate the few cancer cells**, which **withstand initial chemotherapy**, without being resistant a priori. These cells are called "**drug tolerant persister cells**" (DTPs).

## Hypothesis

Chemotherapeutic drug treatment bequeaths a rare surviving **quiescent persister cell population**, which represents a **lurking reservoir of surviving cells**. These cells eventually give rise to **relapsing tumours** and potentially constitute the **causative factor for drug resistance**.

## Aims

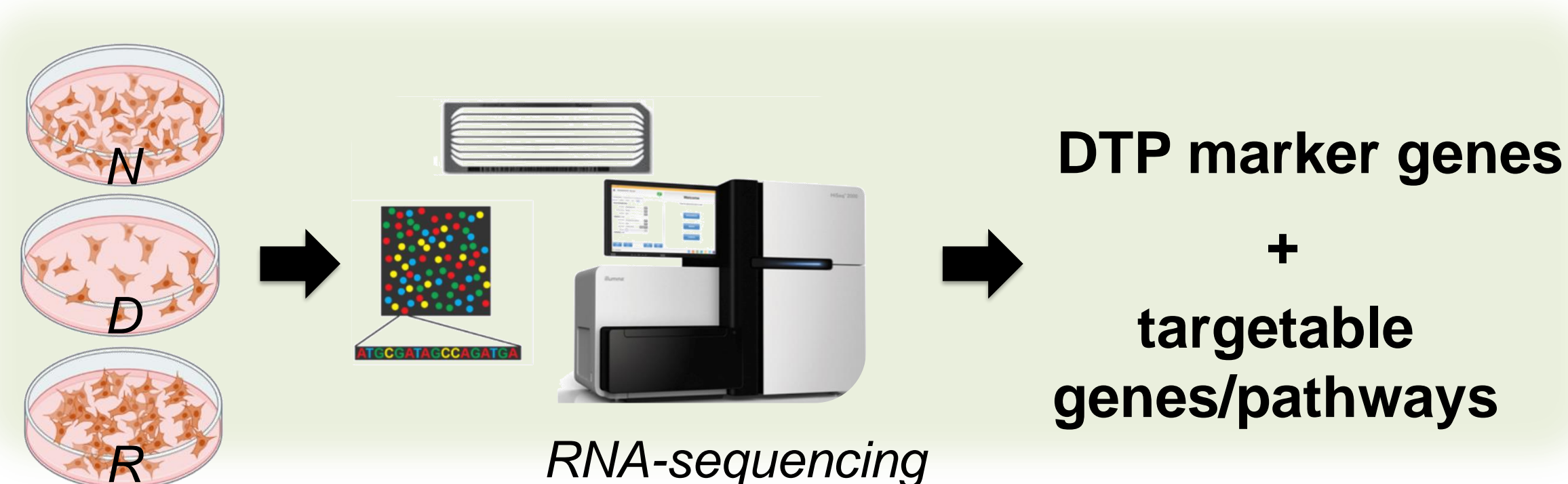
**AIM 1:** *In vivo* identification and tracking of DTPs



**AIM 2:** Morphological characterisation of DTPs



**AIM 3:** Molecular and functional characterisation of DTPs

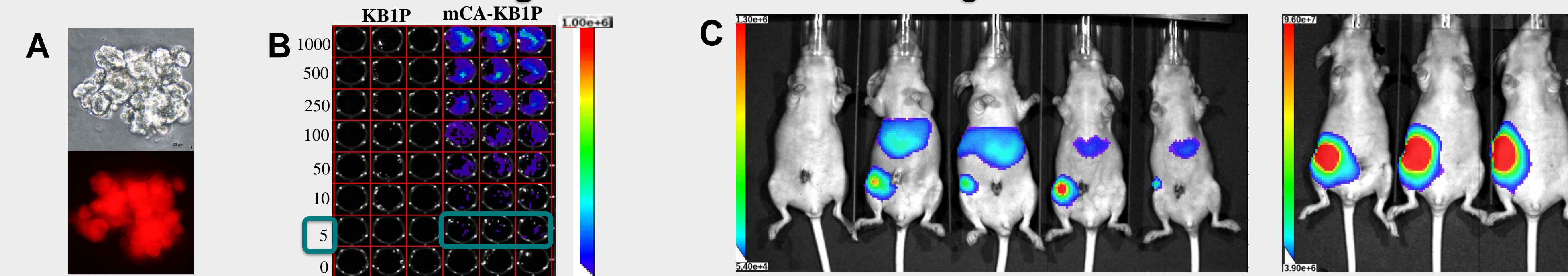


KB1P...breast cancer organoid, mCA...mCherry/Akaluc plasmid, BLI...biolum. imaging, N= naive, D= DTP, R= repopulated

## Results

**AIM 1:**

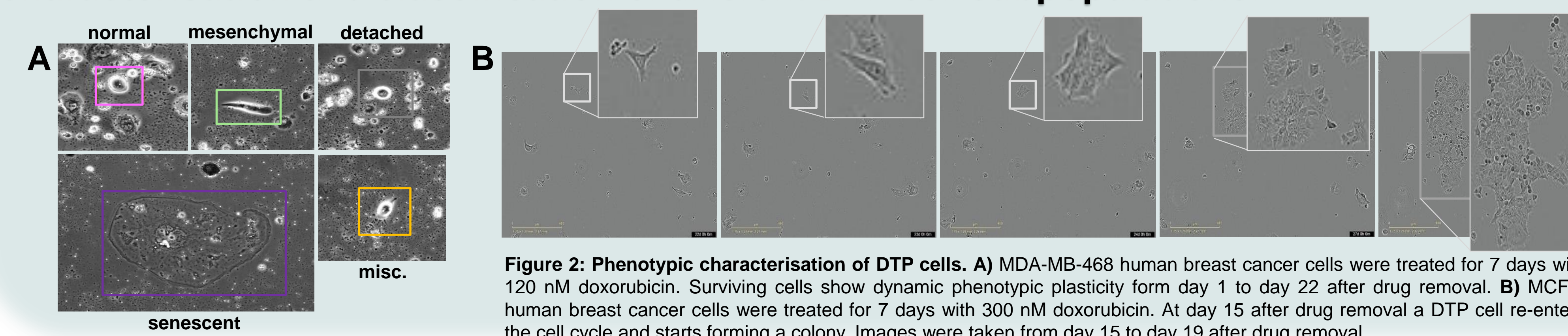
The successful generation of a **polycistronic lentiviral construct** expressing the **AkaLuc luciferase** and the **fluorescent reporter protein mCherry (mCA)** allowed the transduction of **murine breast cancer organoids (KB1P)** resulting in the **mCA-KB1P** model for the **non-invasive intravital detection and tracking of rare DTP cells** in living mice.



**Figure 1:** Establishment of a non-invasive *in vivo* imaging tool. **A)** Murine breast cancer organoids are successfully genetically modified by lentiviral transduction to overexpress an mCherry-AkaLuc reporter. Bottom panel shows KB1P organoids strongly expressing the fluorescent reporter, mCherry. **B)** *In vitro* AkaBLI detects ~ 5 cells in a plate based titration assay. **C)** Engrafted mCA-KB1P organoids retain their BLI signal. Left panel shows a control KB1P-transplanted mouse and four mCA-KB1P4 transplanted mice 48 hrs after mammary fat pad transplantation of  $5 \times 10^4$  cells. The liver signal is an unspecific signal of the substrate. Right panel shows three mCA-KB1P-transplanted mice at day 44 after transplantation.

**AIM 2:**

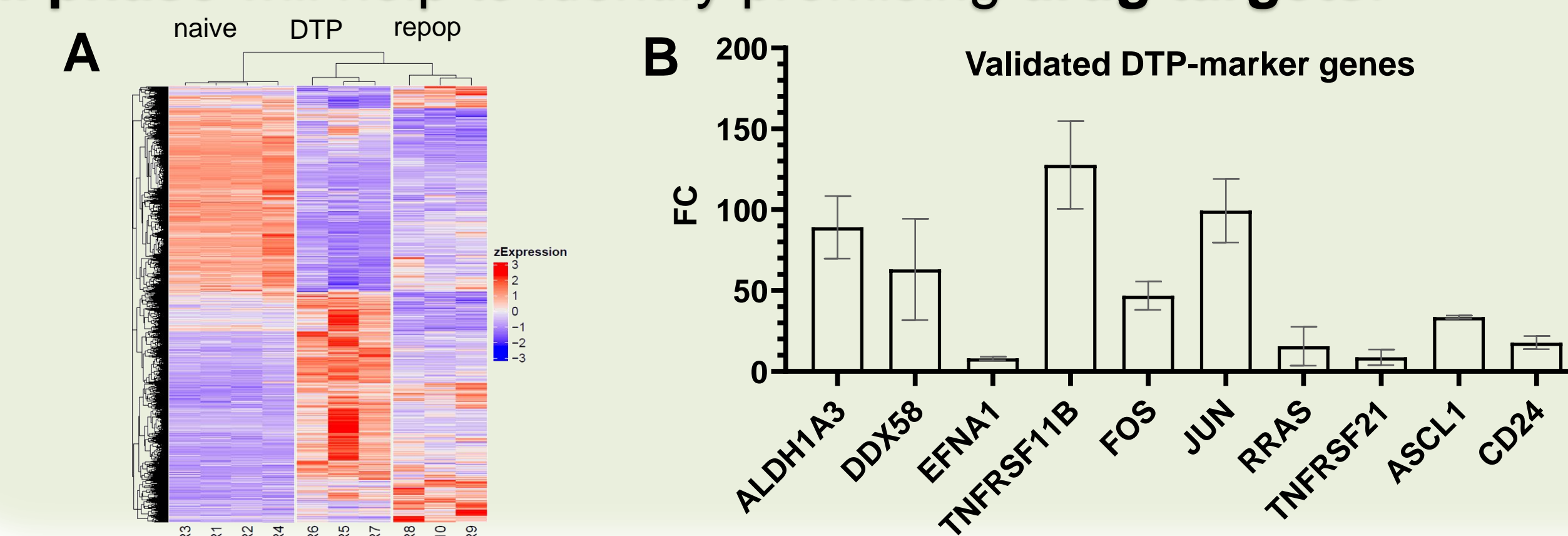
**Live cell imaging pilot studies of the DTP cell phase** showed **dynamic changes of the motility and morphology** of DTPs and recorded their **reawakening after a dormancy period of 15 days**. We generated a cell line with fluorescently labelled nuclei and cytoplasm, which enables *in vitro* DTP tracking and cell size determination of DTP cells. This will allow the **morphological characterisation and classification of different DTP cell subpopulations**.



**Figure 2:** Phenotypic characterisation of DTP cells. **A)** MDA-MB-468 human breast cancer cells were treated for 7 days with 120 nM doxorubicin. Surviving cells show dynamic phenotypic plasticity from day 1 to day 22 after drug removal. **B)** MCF-7 human breast cancer cells were treated for 7 days with 300 nM doxorubicin. At day 15 after drug removal a DTP cell re-enters the cell cycle and starts forming a colony. Images were taken from day 15 to day 19 after drug removal.

**AIM 3:**

**RNA-sequencing of drug-naïve, DTP and repopulated cells** shows the distinctness of the three populations, suggests **subpopulations within the DTP cell pool** and reveals a **transcriptional DTP signature**. We validated selected **DTP marker genes** which will be used to investigate the fate of the DTP cells both *in vitro* and *in vivo*. The determination of potential **key regulators of the DTP cell phase** will help to identify promising **drug targets**.



**Figure 3:** Molecular characterisation of DTP cells. **A)** RNAseq reveals distinct transcriptional profiles for DTP and repopulated cells. Up-regulated genes are depicted in red and down-regulated genes are shown in blue. Noteworthy, subpopulations within the DTP cell pool can be determined. **B)** Selected *in vitro* validated DTP-marker genes show substantial upregulation in the DTP cell phase compared to the untreated, naïve cells. Fold change (FC) values of the DTP cell phase are shown for each gene.

## Conclusions

Summarising, we have established a **non-invasive *in vivo* imaging tool**, which will be used to **identify and track DTP cells** in living animals.

Moreover, our **live cell imaging** approach revealed the plasticity of the surviving cells after chemotherapy. This will allow **sub-classification of DTP cells** to identify cells with the potential of re-entering the cell cycle and thereby giving rise to proliferating colonies.

The determination of a **DTP cell specific transcriptional signature** will help to identify **potential drug targets** and will contribute to **unveil the Achilles' heel** of the few surviving DTP cells to eventually **overcome chemotherapy induced relapse in breast cancer**.

## Literature cited

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## Acknowledgments

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## Further sources

[https://sydney-informatics-hub.github.io/training-RNAseq-slides/01\\_IntroductionToRNASeq/assets/tracy2.png](https://sydney-informatics-hub.github.io/training-RNAseq-slides/01_IntroductionToRNASeq/assets/tracy2.png)  
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