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• Full Name: Alexander Swarbrick

 Current Position & Affiliation: Strategic Program Co-Lead, Targeting the tumour

ecosystems and Lab Head, Garvan Institute of Medical

Research

• Country: Australia

## Educational Background:

1995: 1st class Honours, Molecular & Cellular Biology, UNSW Sydney

2003: Doctorate of Philosophy, UNSW Sydney

## • Professional Experience:

2023: NHMRC Leadership Fellowship (Level 2)

2023-present Professor, University of New South Wales, Sydney

2022-present Strategic Program Co-Lead: Targeting the tumour ecosystems

Petre Foundation Chair of Breast Cancer Research 2020-present

2019: NHMRC Senior Research Fellowship

Petre Foundation Chair of Breast Cancer Research 2017-2022:

Associate Professor, University of New South Wales, Sydney 2016-2023

2013: NHMRC Career Development Fellowship II

2011: NSW Premier's award for Outstanding Research Fellow,

Cancer Institute NSW

National Breast Cancer Foundation Fellowship 2011:

2011-present Member Scientific Faculty, Garvan Institute of Medical Research 2009: CINSW Early Career Fellowship and NBCF Early Career Fellowship 2009-2016 Conjoint Senior Lecturer, University of New South Wales, Sydney

2008-present Lab Head, Cancer Research Program, Garvan Institute of Medical Research

2007: Sydney University Medal for Excellence in Medical Research Postdoctoral Fellow, Garvan Institute of Medical Research 2007

2005: CJ Martin Fellowship from the NHMRC

Postdoctoral Fellow, University of California, San Francisco Francisco 2003-2007

## Professional Organizations:

- Inaugural senior section editor for Cancer Research Communications, a new journal from the American Association for Cancer Research 2021-current
- Associate Editor for Breast Cancer Research in the areas of cellular and molecular pathogenesis. 2020-2021
- Member of grant review committee for Cancer Australia 2021
- External grant assessor for the Welcome Leap Program 2021
- External assessor for ~ 7 applications/year for NHMRC project grants (2008-current), as well as other bodies including Cancer Institute NSW, Cancer Research UK, Sparks UK and the UK MRC.
- GRP (Oncology) for the NHMRC in 2011, 2014, 2017.

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- Grant review committee for the Victorian Cancer Agency's startup fund, 2013
- National Breast Cancer Grant Committee, 2012, 2013, 2017, 2019
- Sydney Catalyst Grant review panel, 2016, 2017, 2019
- Regular reviewer for journals including Nature (5x in 2021), Nature Genetics, Nature Communications, Cell Reports, Nature Biotechnology, Breast Cancer Research, Cancer Research, Oncogene, Cell Genomics, e-Life, Nucleic acids Research.

## • Main Scientific Publications (2019-Present):

- Chaudagar K et al., (2023). Reversal of lactate and PD-1-mediated macrophage 1. immunosuppression controls growth of PTEN/p53-deficient prostate cancer. Clin Cancer Res doi: 10.1158/1078-0432.CCR-22-3350.
- Masle-Farquhar E et al., (2022). STAT3 gain-of-function mutations connect leukemia with 2. autoimmune disease by pathological NKG2D(hi) CD8(+) T cell dysregulation and accumulation. Immunity 55: 2386-2404.
- Baldwin LA et al., (2022). DNA barcoding reveals ongoing immunoediting of clonal cancer 3. populations during metastatic progression and immunotherapy response. Nat Commun 13: 6539.
- Deng N et al., (2022). Deep whole genome sequencing identifies recurrent genomic alterations 4. in commonly used breast cancer cell lines and patient-derived xenograft models. Breast Cancer Res 24: 63.
- Papanicolaou M et al., (2022). Temporal profiling of the breast tumour microenvironment 5. reveals collagen XII as a driver of metastasis. Nat Commun 13: 4587.
- Roden D and Swarbrick A. (2022) Mapping the cancer cell states conserved across solid tumors. 6. Nat Genet 54: 1066-1067.
- 7. Crist SB et al., (2022). Unchecked oxidative stress in skeletal muscle prevents outgrowth of disseminated tumour cells. Nat Cell Biol 24: 538-553.
- Holliday H et al., (2022). miR-99b-5p, miR-380-3p, and miR-485-3p are novel 8. chemosensitizing miRNAs in high-risk neuroblastoma. Mol Ther 30:1 119-1134.
- Meyer B et al., (2021). Identification of DNA methylation biomarkers with potential to predict 9. response to neoadjuvant chemotherapy in triple-negative breast cancer. Clin Epigenetics 13: 226.
- Andersson A et al., (2021). Spatial deconvolution of HER2-positive breast cancer delineates 10. tumor-associated cell type interactions. Nat Commun 12: 6012.
- Bergholtz H et al., (2021). Best Practices for Spatial Profiling for Breast Cancer Research with 11. the GeoMx((R)) Digital Spatial Profiler. Cancers (Basel) 13: 4456.
- Wu SZ et al., (2021). A single-cell and spatially resolved atlas of human breast cancers. Nat 12. Genet 53: 1334-1347.
- Chew NJ et al., (2021). Evaluation of FGFR targeting in breast cancer through interrogation of 13. patient-derived models. Breast Cancer Res 23: 82.
- Mulder K et al., (2021). Cross-tissue single-cell landscape of human monocytes and 14. macrophages in health and disease. Immunity 54: 1883-1900.
- 15. Wu SZ and Swarbrick A (2021). Single-cell advances in stromal-leukocyte interactions in cancer. Immunol Rev 302: 286-298.
- Holliday H, Khoury A and Swarbrick A. Chromatin immunoprecipitation of transcription 16. factors and histone modifications in Comma-Dbeta mammary epithelial cells. STAR Protoc 2: 100514.
- Wu SZ et al., (2021). Cryopreservation of human cancers conserves tumour heterogeneity for 17. single-cell multi-omics analysis. Genome Med 13: 81.



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- 18. Holliday H et al., (2021). Inhibitor of Differentiation 4 (ID4) represses mammary myoepithelial differentiation via inhibition of HEB. iScience 24: 102072.
- 19. Hickey TE et al., (2021). The androgen receptor is a tumor suppressor in estrogen receptorpositive breast cancer. Nat Med 27: 310-320.
- 20. McEvoy CR et al., (2021). A MXI1-NUTM1 fusion protein with MYC-like activity suggests a novel oncogenic mechanism in a subset of NUTM1-rearranged tumors. Lab Invest. 101: 26-
- 21. Millar EK et al., (2020). Tumour Stroma Ratio Assessment Using Digital Image Analysis Predicts Survival in Triple Negative and Luminal Breast Cancer. Cancers (Basel) 12: 3749.
- 22. Thankamony AP et al., (2020). Targeting the Id1-Kif11 Axis in Triple-Negative Breast Cancer Using Combination Therapy. Biomolecules 10: 1295.
- Harvey K et al., (2020). Stromal cell diversity associated with immune evasion in human triple-23. negative breast cancer. EMBO J 39: e104063.
- 24. Portman N et al., (2020). MDM2 inhibition in combination with endocrine therapy and CDK4/6 inhibition for the treatment of ER-positive breast cancer. Breast Cancer Res 22: 87.
- 25. Teo WS et al., (2020). Id Proteins Promote a Cancer Stem Cell Phenotype in Mouse Models of Triple Negative Breast Cancer via Negative Regulation of Robo1. Front Cell Dev Biol 8: 552.
- 26. Baker LA et al., (2020). Proteogenomic analysis of Inhibitor of Differentiation 4 (ID4) in basallike breast cancer. Breast Cancer Res 22: 63.
- 27. Owen KL et al., (2020). Prostate cancer cell-intrinsic interferon signaling regulates dormancy and metastatic outgrowth in bone. EMBO Rep 21: e50162.
- Piggin CL et al., (2020). ELF5 modulates the estrogen receptor cistrome in breast cancer. PLoS 28. Genet 16: e1008531.
- 29. Hoque M et al., (2020). Annexin A6 improves anti-migratory and anti-invasive properties of tyrosine kinase inhibitors in EGFR overexpressing human squamous epithelial cells. FEBS J 287: 2961-2978.
- 30. McCabe MJ et al., (2019). Development and validation of a targeted gene sequencing panel for application to disparate cancers. Sci Rep 9: 17052.
- Brockwell NK et al., (2019). Tumor inherent interferon regulators as biomarkers of long-term 31. chemotherapeutic response in TNBC. NPJ Precis Oncol 3: 21.
- 32. Singh M et al., (2019). High-throughput targeted long-read single cell sequencing reveals the clonal and transcriptional landscape of lymphocytes. Nat Commun 10: 3120.
- Khoo WH et al., (2019). A niche-dependent myeloid transcriptome signature defines dormant 33. myeloma cells. Blood 134: 30-43.
- Chia K et al., (2019). Non-canonical AR activity facilitates endocrine resistance in breast cancer. 34. Endocr Relat Cancer 26: 251-264.